

Codebook ▾

Codes for Missing Data	
Code / Value	Label
UNK	Unknown
NASK	Not asked
NAVU	Not available

### Data Dictionary Codebook

^ Collapse all instruments

#	Variable / Field Name	Field Label <i>Field Note</i>	Field Attributes (Field Type, Validation, Choices, Calculations, etc.)								
Instrument: <b>Demographics</b> (demographics) <span style="float: right;">^ Collapse</span>											
1	recid	Record ID	text								
2	ikdr_regid	IKDR registry ID <i>sites should leave this blank. data coordinator centre use only.</i>	text Field Annotation: @PLACEHOLDER = "DCC use only"								
3	siteid	Site IKDR ID (site-assigned patient ID) <i>Please assign a patient ID for tracking purposes at your own site. Do not include any identifiable information e.g., names. MRN.</i>	text, Required								
4	patguid	Patient GUID	text								
5	ds_substudy	Patient enrolled in the IKDR - data science sub-study(PreVAIL Kids study)	yesno, Required <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table>	1	Yes	0	No				
1	Yes										
0	No										
6	ds_quest Show the field ONLY if: [ds_substudy] = '1'	Is the patient participating in the environmental epidemiology questionnaire?	radio, Required <table border="1"> <tr><td>0</td><td>No, not approached</td></tr> <tr><td>1</td><td>No, patient refused</td></tr> <tr><td>2</td><td>Yes</td></tr> </table>	0	No, not approached	1	No, patient refused	2	Yes		
0	No, not approached										
1	No, patient refused										
2	Yes										
7	ds_questid Show the field ONLY if: [ds_quest] = '2'	Patient unique access code for environmental epidemiology questionnaire	text, Required								
8	group	Group(variable to be decommissioned following data migration) <i>KD + CAA is applicable before Jan01, 2020Acute disease diagnosed on or after Jan 01, 2020</i>	radio <table border="1"> <tr><td>1</td><td>KD + CAA</td></tr> <tr><td>2</td><td>Acute disease</td></tr> </table> Field Annotation: @HIDDEN @READONLY	1	KD + CAA	2	Acute disease				
1	KD + CAA										
2	Acute disease										
9	groupenrol	Enrollment group <i>KD CAA group is only applicable before Jan01, 2020MIS-C, Covid &amp; KD acute diagnosed on or after Jan 01, 2020</i>	radio, Required <table border="1"> <tr><td>1</td><td>MIS-C</td></tr> <tr><td>2</td><td>COVID</td></tr> <tr><td>3</td><td>KD acute</td></tr> <tr><td>4</td><td>KD CAA only</td></tr> </table>	1	MIS-C	2	COVID	3	KD acute	4	KD CAA only
1	MIS-C										
2	COVID										
3	KD acute										
4	KD CAA only										
10	miscptid Show the field ONLY if: [ds_substudy] = '1' and ([groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3')	Section Header: <i>DEMOGRAPHICS</i> ID of the qualified MIS-C patient <i>Leave blank if this is the qualified MIS-C patientSites should include an ID for both the KD acute and COVID patients that are associated with the index MIS-C patient</i>	text								
11	consent_stat	Consent status	radio <table border="1"> <tr><td>1</td><td>Consent acquired</td></tr> <tr><td>2</td><td>Waiver of consent</td></tr> </table>	1	Consent acquired	2	Waiver of consent				
1	Consent acquired										
2	Waiver of consent										
12	inf_stat Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Section Header: <i>SECTION: SARS-COV2- INFECTION STATUS</i> Infection status	radio, Required <table border="1"> <tr><td>1</td><td>Positive</td></tr> <tr><td>2</td><td>Negative (but with symptoms)</td></tr> <tr><td>3</td><td>Test not done (but with symptoms)</td></tr> <tr><td>4</td><td>Negative/asymptomatic</td></tr> </table>	1	Positive	2	Negative (but with symptoms)	3	Test not done (but with symptoms)	4	Negative/asymptomatic
1	Positive										
2	Negative (but with symptoms)										
3	Test not done (but with symptoms)										
4	Negative/asymptomatic										

13	inf_positive Show the field ONLY if: [inf_stat] = '1'	If positive, check the tests that are positive (check all that apply)	checkbox, Required <table border="1"> <tr> <td>1</td> <td>inf_positive__1</td> <td>Nasopharyngeal swab PCR positive</td> </tr> <tr> <td>2</td> <td>inf_positive__2</td> <td>Bronchoalveolar lavage PCR positive</td> </tr> <tr> <td>3</td> <td>inf_positive__3</td> <td>Stool PCR positive</td> </tr> <tr> <td>4</td> <td>inf_positive__4</td> <td>Blood PCR positive</td> </tr> <tr> <td>5</td> <td>inf_positive__5</td> <td>Serology: Ig G positive</td> </tr> <tr> <td>6</td> <td>inf_positive__6</td> <td>Serology: Ig M positive</td> </tr> <tr> <td>7</td> <td>inf_positive__7</td> <td>Serology: total antibodies (IgG &amp; IgM combined) positive</td> </tr> </table>	1	inf_positive__1	Nasopharyngeal swab PCR positive	2	inf_positive__2	Bronchoalveolar lavage PCR positive	3	inf_positive__3	Stool PCR positive	4	inf_positive__4	Blood PCR positive	5	inf_positive__5	Serology: Ig G positive	6	inf_positive__6	Serology: Ig M positive	7	inf_positive__7	Serology: total antibodies (IgG & IgM combined) positive
1	inf_positive__1	Nasopharyngeal swab PCR positive																						
2	inf_positive__2	Bronchoalveolar lavage PCR positive																						
3	inf_positive__3	Stool PCR positive																						
4	inf_positive__4	Blood PCR positive																						
5	inf_positive__5	Serology: Ig G positive																						
6	inf_positive__6	Serology: Ig M positive																						
7	inf_positive__7	Serology: total antibodies (IgG & IgM combined) positive																						
14	inf_positivedt Show the field ONLY if: [inf_stat] = '1'	Date of first positive Covid test(this infection event) <i>if patient becomes Covid positive &gt;1 time, please contact us (DCC) on how to enroll the patient in the database.</i>	text (date_ymd), Required Field Annotation: @HIDEBUTTON																					
15	inf_negative Show the field ONLY if: [inf_stat] = '2'	If negative, check all that apply	checkbox, Required <table border="1"> <tr> <td>1</td> <td>inf_negative__1</td> <td>PCR is negative</td> </tr> <tr> <td>2</td> <td>inf_negative__2</td> <td>Serology (Ig G and/or Ig M) is negative</td> </tr> </table>	1	inf_negative__1	PCR is negative	2	inf_negative__2	Serology (Ig G and/or Ig M) is negative															
1	inf_negative__1	PCR is negative																						
2	inf_negative__2	Serology (Ig G and/or Ig M) is negative																						
16	inf_contact6wk Show the field ONLY if: [group] = '2' or [groupenrol] = '1' or [groupenrol] = '2'	Contact within 6 weeks of onset of symptoms	yesno <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>No</td> </tr> </table>	1	Yes	0	No																	
1	Yes																							
0	No																							
17	inf_contacttyp Show the field ONLY if: [inf_contact6wk] = '1'	If yes, check all that apply	checkbox, Required <table border="1"> <tr> <td>1</td> <td>inf_contacttyp__1</td> <td>Close contact with an individual with laboratory-confirmed SARS-COV2 infection</td> </tr> <tr> <td>2</td> <td>inf_contacttyp__2</td> <td>Close contact with an individual with illness clinically compatible with COVID-19 disease who had close contact with an individual with laboratory-confirmed SARS-COV2 infection</td> </tr> <tr> <td>3</td> <td>inf_contacttyp__3</td> <td>Travel to or residence in an area with sustained, ongoing community transmission of SARS-COV2</td> </tr> </table>	1	inf_contacttyp__1	Close contact with an individual with laboratory-confirmed SARS-COV2 infection	2	inf_contacttyp__2	Close contact with an individual with illness clinically compatible with COVID-19 disease who had close contact with an individual with laboratory-confirmed SARS-COV2 infection	3	inf_contacttyp__3	Travel to or residence in an area with sustained, ongoing community transmission of SARS-COV2												
1	inf_contacttyp__1	Close contact with an individual with laboratory-confirmed SARS-COV2 infection																						
2	inf_contacttyp__2	Close contact with an individual with illness clinically compatible with COVID-19 disease who had close contact with an individual with laboratory-confirmed SARS-COV2 infection																						
3	inf_contacttyp__3	Travel to or residence in an area with sustained, ongoing community transmission of SARS-COV2																						
18	covidvac Show the field ONLY if: [group] = '2' or [groupenrol] = '1' or [groupenrol] = '2'	Section Header: SECTION: COVID VACCINATION COVID vaccination?	yesno <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>No</td> </tr> </table>	1	Yes	0	No																	
1	Yes																							
0	No																							
19	covidvactyp Show the field ONLY if: [covidvac] = '1'	Vaccine provider <i>if different providers/brands were used for the doses, please select "other" and then type the brands/providers</i>	radio <table border="1"> <tr><td>1</td><td>Pfizer/BioNTech</td></tr> <tr><td>2</td><td>Moderna</td></tr> <tr><td>3</td><td>AstraZeneca</td></tr> <tr><td>4</td><td>Sinovac</td></tr> <tr><td>5</td><td>Sinopharm</td></tr> <tr><td>6</td><td>Sputnik V</td></tr> <tr><td>7</td><td>Johnson &amp; Johnson</td></tr> <tr><td>8</td><td>Novavax</td></tr> <tr><td>9</td><td>Other</td></tr> </table>	1	Pfizer/BioNTech	2	Moderna	3	AstraZeneca	4	Sinovac	5	Sinopharm	6	Sputnik V	7	Johnson & Johnson	8	Novavax	9	Other			
1	Pfizer/BioNTech																							
2	Moderna																							
3	AstraZeneca																							
4	Sinovac																							
5	Sinopharm																							
6	Sputnik V																							
7	Johnson & Johnson																							
8	Novavax																							
9	Other																							
20	covidvactytx Show the field ONLY if: [covidvactyp]= '9'	if other vaccine provider(s), please specify	text																					
21	covidvacdt1 Show the field ONLY if: [covidvac] = '1'	Date of COVID vaccination dose 1 <i>Estimate date if no documentation available</i>	text (date_ymd) Field Annotation: @HIDEBUTTON																					
22	covidvacstat2 Show the field ONLY if: [covidvac] = '1'	Status of COVID vaccination dose 2	radio <table border="1"> <tr><td>1</td><td>Administered</td></tr> <tr><td>2</td><td>Pending</td></tr> <tr><td>3</td><td>Not needed</td></tr> <tr><td>4</td><td>Contraindicated</td></tr> </table>	1	Administered	2	Pending	3	Not needed	4	Contraindicated													
1	Administered																							
2	Pending																							
3	Not needed																							
4	Contraindicated																							

23	covidvacdt2 Show the field ONLY if: [covidvacstat2]='1'	if administered, date of COVID vaccination dose 2 <i>Estimate date if no documentation available</i>	text (date_ymd) Field Annotation: @HIDEBUTTON												
24	brk0201 Show the field ONLY if: [covidvacdt1] <> " and [covidvacdt2] <> " and [covidvacdt1] >= [covidvacdt2]	DATA ERRORVaccine dose 2 must be later than dose 1. Please correct the dates.	descriptive												
25	sex	Section Header: SECTION: DEMOGRAPHICS Sex	radio, Identifier <table border="1"> <tr><td>1</td><td>Male</td></tr> <tr><td>2</td><td>Female</td></tr> </table>	1	Male	2	Female								
1	Male														
2	Female														
26	agey	Age at diagnosis (years) <i>e.g., for age 10 years and 11 months &gt; put "10"</i>	text (integer, Min: 0, Max: 100) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'Put the "year" value of age'												
27	agem	Age at diagnosis (months) <i>e.g., for age 10 years and 11 months &gt; put "11"</i>	text (integer, Min: 0, Max: 12) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'Put the "month" value of age'												
28	age	Age in years (calculated field) <i>e.g., for age 10 years &amp; 11 months &gt; this field displays "10.92"</i>	calc Calculation: round(((agey]+if([agem]=0, .00, if([agem]=1, .08, if([agem]=2, 0.14, if([agem]=3, 0.25, if([agem]=4, .33, if([agem]=5, 0.42, if([agem]=6, 0.50, if([agem]=7, 0.58, if([agem]=8, .67, if([agem]=9, 0.75, if([agem]=10, 0.83, if([agem]=11, .92, if([agem]=12, 1.00, "NaN")))))))))))))/3) Custom alignment: RH												
29	location	Location (country)	dropdown (autocomplete) <table border="1"> <tr><td>2</td><td>Canada</td></tr> <tr><td>4</td><td>Chile</td></tr> <tr><td>5</td><td>Italy</td></tr> <tr><td>6</td><td>Mexico</td></tr> <tr><td>1</td><td>USA</td></tr> <tr><td>3</td><td>Other</td></tr> </table>	2	Canada	4	Chile	5	Italy	6	Mexico	1	USA	3	Other
2	Canada														
4	Chile														
5	Italy														
6	Mexico														
1	USA														
3	Other														
30	location_us Show the field ONLY if: [location] = '1'	Zip code (5 digits): US	text (integer), Identifier												
31	location_canada Show the field ONLY if: [location] = '2'	Postal code (first 3 digits): Canada	text, Identifier												
32	location_othertx Show the field ONLY if: [location] = '3'	Other country: please specify	text												
33	location_citytx Show the field ONLY if: [location] = '4' and [location] = '5' and [location] = '6' and [location] = '3'	City of residence	text												
34	ethnicity	Section Header: SECTION: ETHNICITY - BIOLOGIC MOTHER Ethnicity - availability	radio <table border="1"> <tr><td>1</td><td>Both parents</td></tr> <tr><td>2</td><td>Patient only</td></tr> <tr><td>3</td><td>No/Unknown</td></tr> </table>	1	Both parents	2	Patient only	3	No/Unknown						
1	Both parents														
2	Patient only														
3	No/Unknown														

35	ethnicity_m Show the field ONLY if: [ethnicity] = '1' or [ethnicity] = '2'	Ethnicity of biologic mother	checkbox, Identifier <table border="1" data-bbox="1040 107 1524 590"> <tr> <td>1</td> <td>ethnicity_m__1</td> <td>African/African American/Caribbean (Black)/Black</td> </tr> <tr> <td>2</td> <td>ethnicity_m__2</td> <td>Arabic/Middle Eastern/West Asian</td> </tr> <tr> <td>3</td> <td>ethnicity_m__3</td> <td>Caucasian/European/White</td> </tr> <tr> <td>4</td> <td>ethnicity_m__4</td> <td>Eastern Asian</td> </tr> <tr> <td>5</td> <td>ethnicity_m__5</td> <td>Indigenous people of North/South America</td> </tr> <tr> <td>6</td> <td>ethnicity_m__6</td> <td>Latin American/Hispanic/Caribbean (non-Black)</td> </tr> <tr> <td>7</td> <td>ethnicity_m__7</td> <td>South Asian</td> </tr> <tr> <td>8</td> <td>ethnicity_m__8</td> <td>Southeast Asian</td> </tr> <tr> <td>9</td> <td>ethnicity_m__9</td> <td>Other (do not use Canadian/American)</td> </tr> <tr> <td>10</td> <td>ethnicity_m__10</td> <td>Unknown</td> </tr> </table> Field Annotation: @NONEOFTHEABOVE='10'	1	ethnicity_m__1	African/African American/Caribbean (Black)/Black	2	ethnicity_m__2	Arabic/Middle Eastern/West Asian	3	ethnicity_m__3	Caucasian/European/White	4	ethnicity_m__4	Eastern Asian	5	ethnicity_m__5	Indigenous people of North/South America	6	ethnicity_m__6	Latin American/Hispanic/Caribbean (non-Black)	7	ethnicity_m__7	South Asian	8	ethnicity_m__8	Southeast Asian	9	ethnicity_m__9	Other (do not use Canadian/American)	10	ethnicity_m__10	Unknown
1	ethnicity_m__1	African/African American/Caribbean (Black)/Black																															
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9	ethnicity_m__9	Other (do not use Canadian/American)																															
10	ethnicity_m__10	Unknown																															
36	ethnicity_m1 Show the field ONLY if: [ethnicity_m(1)] = '1'	if African/African American/Caribbean (Black)/Black, please select <i>Wikipedia link: list of regions in Africa</i>	radio, Identifier <table border="1" data-bbox="1040 674 1247 961"> <tr><td>1.1</td><td>African-American</td></tr> <tr><td>1.2</td><td>Black Caribbean</td></tr> <tr><td>1.3</td><td>Northern Africa</td></tr> <tr><td>1.4</td><td>Southern Africa</td></tr> <tr><td>1.5</td><td>Western Africa</td></tr> <tr><td>1.6</td><td>Eastern Africa</td></tr> <tr><td>1.7</td><td>Central Africa</td></tr> <tr><td>1.10</td><td>Unknown</td></tr> </table>	1.1	African-American	1.2	Black Caribbean	1.3	Northern Africa	1.4	Southern Africa	1.5	Western Africa	1.6	Eastern Africa	1.7	Central Africa	1.10	Unknown														
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37	ethnicity_m2 Show the field ONLY if: [ethnicity_m(2)] = '1'	if Arabic/Middle Eastern/West Asian, please select	radio, Identifier <table border="1" data-bbox="1040 999 1346 1182"> <tr><td>2.1</td><td>Israeli</td></tr> <tr><td>2.2</td><td>Persian/Arabic/Iranian/Saudis</td></tr> <tr><td>2.3</td><td>Turks</td></tr> <tr><td>2.4</td><td>Western/Central Asian</td></tr> <tr><td>2.10</td><td>Unknown</td></tr> </table>	2.1	Israeli	2.2	Persian/Arabic/Iranian/Saudis	2.3	Turks	2.4	Western/Central Asian	2.10	Unknown																				
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38	ethnicity_m3 Show the field ONLY if: [ethnicity_m(3)] = '1'	if Caucasian/European/White, please select	radio, Identifier <table border="1" data-bbox="1040 1220 1471 1472"> <tr><td>3.1</td><td>British/Celtic (England, Scotland, Ireland)</td></tr> <tr><td>3.2</td><td>Germanic (Germany/Austria/Netherlands)</td></tr> <tr><td>3.3</td><td>Greek/Albanian (Greece, Albania, Macedonia)</td></tr> <tr><td>3.4</td><td>Northern (Scandinavia)</td></tr> <tr><td>3.5</td><td>Slavic (Poland, Ukraine, Russia)</td></tr> <tr><td>3.6</td><td>Southern (France, Spain, Portugal, Italy)</td></tr> <tr><td>3.10</td><td>Unknown</td></tr> </table>	3.1	British/Celtic (England, Scotland, Ireland)	3.2	Germanic (Germany/Austria/Netherlands)	3.3	Greek/Albanian (Greece, Albania, Macedonia)	3.4	Northern (Scandinavia)	3.5	Slavic (Poland, Ukraine, Russia)	3.6	Southern (France, Spain, Portugal, Italy)	3.10	Unknown																
3.1	British/Celtic (England, Scotland, Ireland)																																
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3.10	Unknown																																
39	ethnicity_m4 Show the field ONLY if: [ethnicity_m(4)] = '1'	if Eastern Asian, please select	radio, Identifier <table border="1" data-bbox="1040 1509 1261 1692"> <tr><td>4.1</td><td>Chinese/Taiwanese</td></tr> <tr><td>4.2</td><td>Japanese</td></tr> <tr><td>4.3</td><td>Korean</td></tr> <tr><td>4.4</td><td>Mongolian</td></tr> <tr><td>4.10</td><td>Unknown</td></tr> </table>	4.1	Chinese/Taiwanese	4.2	Japanese	4.3	Korean	4.4	Mongolian	4.10	Unknown																				
4.1	Chinese/Taiwanese																																
4.2	Japanese																																
4.3	Korean																																
4.4	Mongolian																																
4.10	Unknown																																
40	ethnicity_m5 Show the field ONLY if: [ethnicity_m(5)] = '1'	if Indigenous people of North/South America, please select	radio, Identifier <table border="1" data-bbox="1040 1730 1382 1908"> <tr><td>5.1</td><td>Inuit</td></tr> <tr><td>5.2</td><td>Native North American/Metis</td></tr> <tr><td>5.3</td><td>Native South American/Caribbean</td></tr> <tr><td>5.4</td><td>Pacific Islander</td></tr> <tr><td>5.10</td><td>Unknown</td></tr> </table>	5.1	Inuit	5.2	Native North American/Metis	5.3	Native South American/Caribbean	5.4	Pacific Islander	5.10	Unknown																				
5.1	Inuit																																
5.2	Native North American/Metis																																
5.3	Native South American/Caribbean																																
5.4	Pacific Islander																																
5.10	Unknown																																

41	ethnicity_m6 Show the field ONLY if: [ethnicity_m(6)] = '1'	if Latin American/Hispanic/Caribbean (non-Black), please select	radio, Identifier <table border="1"> <tr><td>6.1</td><td>Brazilian</td></tr> <tr><td>6.2</td><td>Caribbean</td></tr> <tr><td>6.3</td><td>Mexican/central American</td></tr> <tr><td>6.4</td><td>South American (other than Brazilian)</td></tr> <tr><td>6.10</td><td>Unknown</td></tr> </table>	6.1	Brazilian	6.2	Caribbean	6.3	Mexican/central American	6.4	South American (other than Brazilian)	6.10	Unknown																				
6.1	Brazilian																																
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6.10	Unknown																																
42	ethnicity_m7 Show the field ONLY if: [ethnicity_m(7)] = '1'	if South Asian, please select	radio, Identifier <table border="1"> <tr><td>7.1</td><td>Indian/Bengali</td></tr> <tr><td>7.2</td><td>Nepalese/Bhutanese</td></tr> <tr><td>7.3</td><td>Pakistani/Afghani</td></tr> <tr><td>7.4</td><td>Sri Lankan/Maldives</td></tr> <tr><td>7.10</td><td>Unknown</td></tr> </table>	7.1	Indian/Bengali	7.2	Nepalese/Bhutanese	7.3	Pakistani/Afghani	7.4	Sri Lankan/Maldives	7.10	Unknown																				
7.1	Indian/Bengali																																
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7.4	Sri Lankan/Maldives																																
7.10	Unknown																																
43	ethnicity_m8 Show the field ONLY if: [ethnicity_m(8)] = '1'	if Southeast Asian, please select	radio, Identifier <table border="1"> <tr><td>8.1</td><td>Filipino</td></tr> <tr><td>8.2</td><td>Indonesian/Malaysian/Singaporean</td></tr> <tr><td>8.3</td><td>Thai/Cambodian/Vietnamese/Burmese/Laotian</td></tr> <tr><td>8.10</td><td>Unknown</td></tr> </table>	8.1	Filipino	8.2	Indonesian/Malaysian/Singaporean	8.3	Thai/Cambodian/Vietnamese/Burmese/Laotian	8.10	Unknown																						
8.1	Filipino																																
8.2	Indonesian/Malaysian/Singaporean																																
8.3	Thai/Cambodian/Vietnamese/Burmese/Laotian																																
8.10	Unknown																																
44	ethnicity_mtx Show the field ONLY if: [ethnicity_m(9)] = '1'	if other ethnicity, please specify	notes, Identifier																														
45	ethnicity_f Show the field ONLY if: [ethnicity] = '1'	Section Header: SECTION: ETHNICITY - BIOLOGIC FATHER Ethnicity of biologic father	checkbox, Identifier <table border="1"> <tr><td>1</td><td>ethnicity_f__1</td><td>African/African American/Caribbean (Black)/Black</td></tr> <tr><td>2</td><td>ethnicity_f__2</td><td>Arabic/Middle Eastern/West Asian</td></tr> <tr><td>3</td><td>ethnicity_f__3</td><td>Caucasian/European/White</td></tr> <tr><td>4</td><td>ethnicity_f__4</td><td>Eastern Asian</td></tr> <tr><td>5</td><td>ethnicity_f__5</td><td>Indigenous people of North/South America</td></tr> <tr><td>6</td><td>ethnicity_f__6</td><td>Latin American/Hispanic/Caribbean (non-Black)</td></tr> <tr><td>7</td><td>ethnicity_f__7</td><td>South Asian</td></tr> <tr><td>8</td><td>ethnicity_f__8</td><td>Southeast Asian</td></tr> <tr><td>9</td><td>ethnicity_f__9</td><td>Other (do not use Canadian/American)</td></tr> <tr><td>10</td><td>ethnicity_f__10</td><td>Unknown</td></tr> </table> Field Annotation: @NONEOFTHEABOVE='10'	1	ethnicity_f__1	African/African American/Caribbean (Black)/Black	2	ethnicity_f__2	Arabic/Middle Eastern/West Asian	3	ethnicity_f__3	Caucasian/European/White	4	ethnicity_f__4	Eastern Asian	5	ethnicity_f__5	Indigenous people of North/South America	6	ethnicity_f__6	Latin American/Hispanic/Caribbean (non-Black)	7	ethnicity_f__7	South Asian	8	ethnicity_f__8	Southeast Asian	9	ethnicity_f__9	Other (do not use Canadian/American)	10	ethnicity_f__10	Unknown
1	ethnicity_f__1	African/African American/Caribbean (Black)/Black																															
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9	ethnicity_f__9	Other (do not use Canadian/American)																															
10	ethnicity_f__10	Unknown																															
46	ethnicity_f1 Show the field ONLY if: [ethnicity_f(1)] = '1'	if African/African American/Caribbean (Black)/Black, please select <i>Wikipedia link: list of regions in Africa</i>	radio, Identifier <table border="1"> <tr><td>1.1</td><td>African-American</td></tr> <tr><td>1.2</td><td>Black Caribbean</td></tr> <tr><td>1.3</td><td>Northern Africa</td></tr> <tr><td>1.4</td><td>Southern Africa</td></tr> <tr><td>1.5</td><td>Western Africa</td></tr> <tr><td>1.6</td><td>Eastern Africa</td></tr> <tr><td>1.7</td><td>Central Africa</td></tr> <tr><td>1.10</td><td>Unknown</td></tr> </table>	1.1	African-American	1.2	Black Caribbean	1.3	Northern Africa	1.4	Southern Africa	1.5	Western Africa	1.6	Eastern Africa	1.7	Central Africa	1.10	Unknown														
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1.7	Central Africa																																
1.10	Unknown																																
47	ethnicity_f2 Show the field ONLY if: [ethnicity_f(2)] = '1'	if Arabic/Middle Eastern/West Asian, please select	radio, Identifier <table border="1"> <tr><td>2.1</td><td>Israeli</td></tr> <tr><td>2.2</td><td>Persian/Arabic/Iranian/Saudis</td></tr> <tr><td>2.3</td><td>Turks</td></tr> <tr><td>2.4</td><td>Western/Central Asian</td></tr> <tr><td>2.10</td><td>Unknown</td></tr> </table>	2.1	Israeli	2.2	Persian/Arabic/Iranian/Saudis	2.3	Turks	2.4	Western/Central Asian	2.10	Unknown																				
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2.4	Western/Central Asian																																
2.10	Unknown																																

48	ethnicity_f3 Show the field ONLY if: [ethnicity_f(3)] = '1'	if Caucasian/European/White, please select	radio, Identifier 3.1 British/Celtic (England, Scotland, Ireland) 3.2 Germanic (Germany/Austria/Netherlands) 3.3 Greek/Albanian (Greece, Albania, Macedonia) 3.4 Northern (Scandinavia) 3.5 Slavic (Poland, Ukraine, Russia) 3.6 Southern (France, Spain, Portugal, Italy) 3.10 Unknown
49	ethnicity_f4 Show the field ONLY if: [ethnicity_f(4)] = '1'	if Eastern Asian, please select	radio, Identifier 4.1 Chinese/Taiwanese 4.2 Japanese 4.3 Korean 4.4 Mongolian 4.10 Unknown
50	ethnicity_f5 Show the field ONLY if: [ethnicity_f(5)] = '1'	if Indigenous people of North/South America, please select	radio, Identifier 5.1 Inuit 5.2 Native North American/Metis 5.3 Native South American/Caribbean 5.4 Pacific Islander 5.10 Unknown
51	ethnicity_f6 Show the field ONLY if: [ethnicity_f(6)] = '1'	if Latin American/Hispanic/Caribbean (non-Black), please select	radio, Identifier 6.1 Brazilian 6.2 Caribbean 6.3 Mexican/central American 6.4 South American (other than Brazilian) 6.10 Unknown
52	ethnicity_f7 Show the field ONLY if: [ethnicity_f(7)] = '1'	if South Asian, please select	radio, Identifier 7.1 Indian/Bengali 7.2 Nepalese/Bhutanese 7.3 Pakistani/Afghani 7.4 Sri Lankan/Maldives 7.10 Unknown
53	ethnicity_f8 Show the field ONLY if: [ethnicity_f(8)] = '1'	if Southeast Asian, please select	radio, Identifier 8.1 Filipino 8.2 Indonesian/Malaysian/Singaporean 8.3 Thai/Cambodian/Vietnamese/Burmese/Laotian 8.10 Unknown
54	ethnicity_ftx Show the field ONLY if: [ethnicity_f(9)] = '1'	if other ethnicity, please specify	notes, Identifier
55	demographics_complete	Section Header: <i>Form Status</i> Complete?	dropdown 0 Incomplete 1 Unverified 2 Complete
Instrument: <b>Biological Sample</b> (biological_sample) <a href="#">^ Collapse</a>			
56	brk_0101 Show the field ONLY if: [groupenrol] = '4'	Section Header: <i>BIOLOGICAL SAMPLE/SPECIMEN DATA</i> THIS SECTION IS NOT APPLICABLE for KD patients (prior to Jan 01, 2020)	descriptive
57	sample Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Biological sample collected	radio 1 Yes 0 No

58	<p>ds_samplotyp</p> <p>Show the field ONLY if: [ds_substudy] = '1' and [sample] = '1'</p>	<p>If yes, biological sample collection</p>	<p>radio</p> <table border="1"> <tr> <td>1</td> <td>Biobanked</td> </tr> <tr> <td>2</td> <td>Prospective</td> </tr> </table>	1	Biobanked	2	Prospective																		
1	Biobanked																								
2	Prospective																								
59	<p>samplestudy</p> <p>Show the field ONLY if: [sample] = '1'</p>	<p>if yes, study name(s) <i>e.g., IKDR - data science sub-study (PreVAIL Kids)</i></p>	<p>notes</p>																						
60	<p>brk_02011</p> <p>Show the field ONLY if: ([groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3') and [ds_substudy] = '0'</p>	<p>Section Header: <i>SECTION: SAMPLE DETAILS</i></p> <p>THIS SECTION IS ONLY APPLICABLE for patients enrolled in PreVAILKids study</p>	<p>descriptive</p>																						
61	<p>ds_shipdt</p> <p>Show the field ONLY if: [ds_substudy] = '1' and ([sample] = '1' or [sample] = '2')</p>	<p>Date samples shipped to JHU</p>	<p>text (date_ymd)</p> <p>Field Annotation: @HIDEBUTTON</p>																						
62	<p>ds_receivedt</p> <p>Show the field ONLY if: [ds_substudy] = '1' and ([sample] = '1' or [sample] = '2')</p>	<p>Date samples received at JHU</p>	<p>text (date_ymd)</p> <p>Field Annotation: @HIDEBUTTON</p>																						
63	<p>ds_shipconcern</p> <p>Show the field ONLY if: [ds_substudy] = '1' and ([sample] = '1' or [sample] = '2')</p>	<p>Shipping concerns</p>	<p>yesno</p> <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>No</td> </tr> </table>	1	Yes	0	No																		
1	Yes																								
0	No																								
64	<p>ds_shipconcerntx</p> <p>Show the field ONLY if: [ds_shipconcern] = '1'</p>	<p>Describe shipping concerns</p>	<p>notes</p>																						
65	<p>ds_aliquiplasma</p> <p>Show the field ONLY if: [ds_substudy] = '1' and ([sample] = '1' or [sample] = '2')</p>	<p>Number of aliquots acquired - Plasma <i>0 if no sample acquired</i></p>	<p>dropdown</p> <table border="1"> <tr> <td>0</td> <td>No sample acquired</td> </tr> <tr> <td>1</td> <td>01 aliquot</td> </tr> <tr> <td>2</td> <td>02 aliquots</td> </tr> <tr> <td>3</td> <td>03 aliquots</td> </tr> <tr> <td>4</td> <td>04 aliquots</td> </tr> <tr> <td>5</td> <td>05 aliquots</td> </tr> <tr> <td>6</td> <td>06 aliquots</td> </tr> <tr> <td>7</td> <td>07 aliquots</td> </tr> <tr> <td>8</td> <td>08 aliquots</td> </tr> <tr> <td>9</td> <td>09 aliquots</td> </tr> <tr> <td>10</td> <td>10 aliquots</td> </tr> </table> <p>Custom alignment: RH</p>	0	No sample acquired	1	01 aliquot	2	02 aliquots	3	03 aliquots	4	04 aliquots	5	05 aliquots	6	06 aliquots	7	07 aliquots	8	08 aliquots	9	09 aliquots	10	10 aliquots
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9	09 aliquots																								
10	10 aliquots																								
66	<p>ds_aliquserum</p> <p>Show the field ONLY if: [ds_substudy] = '1' and ([sample] = '1' or [sample] = '2')</p>	<p>Number of aliquots acquired - Serum <i>0 if no sample acquired</i></p>	<p>dropdown</p> <table border="1"> <tr> <td>0</td> <td>No sample acquired</td> </tr> <tr> <td>1</td> <td>01 aliquot</td> </tr> <tr> <td>2</td> <td>02 aliquots</td> </tr> <tr> <td>3</td> <td>03 aliquots</td> </tr> <tr> <td>4</td> <td>04 aliquots</td> </tr> <tr> <td>5</td> <td>05 aliquots</td> </tr> <tr> <td>6</td> <td>06 aliquots</td> </tr> <tr> <td>7</td> <td>07 aliquots</td> </tr> <tr> <td>8</td> <td>08 aliquots</td> </tr> <tr> <td>9</td> <td>09 aliquots</td> </tr> <tr> <td>10</td> <td>10 aliquots</td> </tr> </table>	0	No sample acquired	1	01 aliquot	2	02 aliquots	3	03 aliquots	4	04 aliquots	5	05 aliquots	6	06 aliquots	7	07 aliquots	8	08 aliquots	9	09 aliquots	10	10 aliquots
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8	08 aliquots																								
9	09 aliquots																								
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67	<p>ds_aliqubuffy</p> <p>Show the field ONLY if: [ds_substudy] = '1' and ([sample] = '1' or [sample] = '2')</p>	<p>Frozen blood buffy coat obtained for DNA extraction?</p>	<p>yesno</p> <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>No</td> </tr> </table>	1	Yes	0	No																		
1	Yes																								
0	No																								

68	ds_plasma_obtdt Show the field ONLY if: [ds_aliquplasma] >= 1	Section Header: SECTION: SAMPLE LOG - PLASMAonly applicable for PreVAiLKids study Date aliquots obtained for sample	text (date_ymd) Field Annotation: @HIDEBUTTON										
69	ds_brkplasma Show the field ONLY if: [ds_aliquplasma] >= 1	INSTRUCTIONS: Date aliquot obtained should be at the sample level, unless it's a leftover sample. [ds_aliquplasma] have been obtained for this patient. Enter each aliquot info in the log below. For each aliquot label, follow format "patientID_sampletype_aliquotnumber" e.g., "123-12_plasma_01" If aliquot volume remaining, please log a new aliquot with a revised sample number (suffix: _R1, _R2, etc.)	descriptive										
70	ds_plasma_label_01 Show the field ONLY if: [ds_aliquplasma] >= 1	Aliquot label #01	text Field Annotation: @PLACEHOLDER = 'Aliquot 01'										
71	ds_plasma_usedt_01 Show the field ONLY if: [ds_aliquplasma] >= 1	Date aliquot used #01	text (date_ymd) Field Annotation: @HIDEBUTTON										
72	ds_plasma_stat_01 Show the field ONLY if: [ds_aliquplasma] >= 1	Aliquot status #01	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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4	Leftover 2nd cycle												
73	ds_plasma_study_01 Show the field ONLY if: [ds_plasma_stat_01] = '2'	PI/study name #01	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
74	ds_plasma_vol_01 Show the field ONLY if: [ds_aliquplasma] >= 1	Volume (ul) #01	text (integer)										
75	ds_plasma_rem_01 Show the field ONLY if: [ds_aliquplasma] >= 1	Volume remaining #01	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_01__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_01__1	Yes							
1	ds_plasma_rem_01__1	Yes											
76	ds_plasma_label_01r1 Show the field ONLY if: [ds_plasma_rem_01(1)] = '1'	Aliquot label #01R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
77	ds_plasma_usedt_01r1 Show the field ONLY if: [ds_plasma_rem_01(1)] = '1'	Date aliquot used #01R1	text (date_ymd) Field Annotation: @HIDEBUTTON										
78	ds_plasma_stat_01r1 Show the field ONLY if: [ds_plasma_rem_01(1)] = '1'	Aliquot status #01R1	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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79	ds_plasma_study_01r1 Show the field ONLY if: [ds_plasma_stat_01r1] = '2'	PI/study name #01R1	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
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80	ds_plasma_vol_01r1 Show the field ONLY if: [ds_plasma_rem_01(1)] = '1'	Volume (ul) #01R1	text (integer)										
81	ds_plasma_rem_01r1 Show the field ONLY if: [ds_plasma_rem_01(1)] = '1'	Volume remaining #01R1	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_01r1__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_01r1__1	Yes							
1	ds_plasma_rem_01r1__1	Yes											



82	ds_plasma_label_01r2 Show the field ONLY if: [ds_plasma_rem_01(1)] = '1'	Aliquot label #01r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
83	ds_plasma_usedt_01r2 Show the field ONLY if: [ds_plasma_rem_01(1)] = '1'	Date aliquot used #01r2	text (date_ymd) Field Annotation: @HIDEBUTTON										
84	ds_plasma_stat_01r2 Show the field ONLY if: [ds_plasma_rem_01(1)] = '1'	Aliquot status #01r2	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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4	Leftover 2nd cycle												
85	ds_plasma_study_01r2 Show the field ONLY if: [ds_plasma_stat_01r2] = '2'	PI/study name #01r2	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
86	ds_plasma_vol_01r2 Show the field ONLY if: [ds_plasma_rem_01(1)] = '1'	Volume (ul) #01r2	text (integer)										
87	ds_plasma_rem_01r2 Show the field ONLY if: [ds_plasma_rem_01(1)] = '1'	Volume remaining #01r2	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_01r2__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_01r2__1	Yes							
1	ds_plasma_rem_01r2__1	Yes											
88	ds_plasma_label_01r3 Show the field ONLY if: [ds_plasma_rem_01(1)] = '1'	Aliquot label #01r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'										
89	ds_plasma_usedt_01r3 Show the field ONLY if: [ds_plasma_rem_01(1)] = '1'	Date aliquot used #01r3	text (date_ymd) Field Annotation: @HIDEBUTTON										
90	ds_plasma_stat_01r3 Show the field ONLY if: [ds_plasma_rem_01(1)] = '1'	Aliquot status #01r3	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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4	Leftover 2nd cycle												
91	ds_plasma_study_01r3 Show the field ONLY if: [ds_plasma_stat_01r3] = '2'	PI/study name #01r3	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
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92	ds_plasma_vol_01r3 Show the field ONLY if: [ds_plasma_rem_01(1)] = '1'	Volume (ul) #01r3	text (integer)										
93	ds_plasma_rem_01r3 Show the field ONLY if: [ds_plasma_rem_01(1)] = '1'	Volume remaining #01r3	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_01r3__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_01r3__1	Yes							
1	ds_plasma_rem_01r3__1	Yes											
94	ds_plasma_label_02 Show the field ONLY if: [ds_aliquiplasma] >= 2	Aliquot label #02	text Field Annotation: @PLACEHOLDER = 'Aliquot 02'										
95	ds_plasma_usedt_02 Show the field ONLY if: [ds_aliquiplasma] >= 2	Date aliquot used #02	text (date_ymd) Field Annotation: @HIDEBUTTON										

96	ds_plasma_stat_02 Show the field ONLY if: [ds_aliquiplasma] >= 2	Aliquot status #02	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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4	Leftover 2nd cycle												
97	ds_plasma_study_02 Show the field ONLY if: [ds_plasma_stat_02] = '2'	PI/study name #02	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
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98	ds_plasma_vol_02 Show the field ONLY if: [ds_aliquiplasma] >= 2	Volume (ul) #02	text (integer)										
99	ds_plasma_rem_02 Show the field ONLY if: [ds_aliquiplasma] >= 2	Volume remaining #02	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_02__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_02__1	Yes							
1	ds_plasma_rem_02__1	Yes											
100	ds_plasma_label_02r1 Show the field ONLY if: [ds_plasma_rem_02(1)] = '1'	Aliquot label #02R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
101	ds_plasma_usedt_02r1 Show the field ONLY if: [ds_plasma_rem_02(1)] = '1'	Date aliquot used #02R1	text (date_ymd) Field Annotation: @HIDEBUTTON										
102	ds_plasma_stat_02r1 Show the field ONLY if: [ds_plasma_rem_02(1)] = '1'	Aliquot status #02R1	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
103	ds_plasma_study_02r1 Show the field ONLY if: [ds_plasma_stat_02r1] = '2'	PI/study name #02R1	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
104	ds_plasma_vol_02r1 Show the field ONLY if: [ds_plasma_rem_02(1)] = '1'	Volume (ul) #02R1	text (integer)										
105	ds_plasma_rem_02r1 Show the field ONLY if: [ds_plasma_rem_02(1)] = '1'	Volume remaining #02R1	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_02r1__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_02r1__1	Yes							
1	ds_plasma_rem_02r1__1	Yes											
106	ds_plasma_label_02r2 Show the field ONLY if: [ds_plasma_rem_02(1)] = '1'	Aliquot label #02r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
107	ds_plasma_usedt_02r2 Show the field ONLY if: [ds_plasma_rem_02(1)] = '1'	Date aliquot used #02r2	text (date_ymd) Field Annotation: @HIDEBUTTON										
108	ds_plasma_stat_02r2 Show the field ONLY if: [ds_plasma_rem_02(1)] = '1'	Aliquot status #02r2	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												

109	ds_plasma_study_02r2 Show the field ONLY if: [ds_plasma_stat_02r2] = '2'	PI/study name #02r2	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
110	ds_plasma_vol_02r2 Show the field ONLY if: [ds_plasma_rem_02(1)] = '1'	Volume (ul) #02r2	text (integer)										
111	ds_plasma_rem_02r2 Show the field ONLY if: [ds_plasma_rem_02(1)] = '1'	Volume remaining #02r2	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_02r2__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_02r2__1	Yes							
1	ds_plasma_rem_02r2__1	Yes											
112	ds_plasma_label_02r3 Show the field ONLY if: [ds_plasma_rem_02(1)] = '1'	Aliquot label #02r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'										
113	ds_plasma_usedt_02r3 Show the field ONLY if: [ds_plasma_rem_02(1)] = '1'	Date aliquot used #02r3	text (date_ymd) Field Annotation: @HIDEBUTTON										
114	ds_plasma_stat_02r3 Show the field ONLY if: [ds_plasma_rem_02(1)] = '1'	Aliquot status #02r3	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
115	ds_plasma_study_02r3 Show the field ONLY if: [ds_plasma_stat_02r3] = '2'	PI/study name #02r3	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
116	ds_plasma_vol_02r3 Show the field ONLY if: [ds_plasma_rem_02(1)] = '1'	Volume (ul) #02r3	text (integer)										
117	ds_plasma_rem_02r3 Show the field ONLY if: [ds_plasma_rem_02(1)] = '1'	Volume remaining #02r3	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_02r3__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_02r3__1	Yes							
1	ds_plasma_rem_02r3__1	Yes											
118	ds_plasma_label_03 Show the field ONLY if: [ds_aliquiplasma] >= 3	Aliquot label #03	text Field Annotation: @PLACEHOLDER = 'Aliquot 03'										
119	ds_plasma_usedt_03 Show the field ONLY if: [ds_aliquiplasma] >= 3	Date aliquot used #03	text (date_ymd) Field Annotation: @HIDEBUTTON										
120	ds_plasma_stat_03 Show the field ONLY if: [ds_aliquiplasma] >= 3	Aliquot status #03	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
121	ds_plasma_study_03 Show the field ONLY if: [ds_plasma_stat_03] = '2'	PI/study name #03	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												

122	ds_plasma_vol_03 Show the field ONLY if: [ds_aliquiplasma] >= 3	Volume (ul) #03	text (integer)										
123	ds_plasma_rem_03 Show the field ONLY if: [ds_aliquiplasma] >= 3	Volume remaining #03	checkbox <input type="checkbox"/> ds_plasma_rem_03__1 <input type="checkbox"/> Yes										
124	ds_plasma_label_03r1 Show the field ONLY if: [ds_plasma_rem_03(1)] = '1'	Aliquot label #03R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
125	ds_plasma_usedt_03r1 Show the field ONLY if: [ds_plasma_rem_03(1)] = '1'	Date aliquot used #03R1	text (date_ymd) Field Annotation: @HIDEBUTTON										
126	ds_plasma_stat_03r1 Show the field ONLY if: [ds_plasma_rem_03(1)] = '1'	Aliquot status #03R1	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
127	ds_plasma_study_03r1 Show the field ONLY if: [ds_plasma_stat_03r1] = '2'	PI/study name #03R1	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
128	ds_plasma_vol_03r1 Show the field ONLY if: [ds_plasma_rem_03(1)] = '1'	Volume (ul) #03R1	text (integer)										
129	ds_plasma_rem_03r1 Show the field ONLY if: [ds_plasma_rem_03(1)] = '1'	Volume remaining #03R1	checkbox <input type="checkbox"/> ds_plasma_rem_03r1__1 <input type="checkbox"/> Yes										
130	ds_plasma_label_03r2 Show the field ONLY if: [ds_plasma_rem_03(1)] = '1'	Aliquot label #03r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
131	ds_plasma_usedt_03r2 Show the field ONLY if: [ds_plasma_rem_03(1)] = '1'	Date aliquot used #03r2	text (date_ymd) Field Annotation: @HIDEBUTTON										
132	ds_plasma_stat_03r2 Show the field ONLY if: [ds_plasma_rem_03(1)] = '1'	Aliquot status #03r2	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
133	ds_plasma_study_03r2 Show the field ONLY if: [ds_plasma_stat_03r2] = '2'	PI/study name #03r2	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
134	ds_plasma_vol_03r2 Show the field ONLY if: [ds_plasma_rem_03(1)] = '1'	Volume (ul) #03r2	text (integer)										
135	ds_plasma_rem_03r2 Show the field ONLY if: [ds_plasma_rem_03(1)] = '1'	Volume remaining #03r2	checkbox <input type="checkbox"/> ds_plasma_rem_03r2__1 <input type="checkbox"/> Yes										

136	ds_plasma_label_03r3 Show the field ONLY if: [ds_plasma_rem_03(1)] = '1'	Aliquot label #03r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'										
137	ds_plasma_usedt_03r3 Show the field ONLY if: [ds_plasma_rem_03(1)] = '1'	Date aliquot used #03r3	text (date_ymd) Field Annotation: @HIDEBUTTON										
138	ds_plasma_stat_03r3 Show the field ONLY if: [ds_plasma_rem_03(1)] = '1'	Aliquot status #03r3	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
139	ds_plasma_study_03r3 Show the field ONLY if: [ds_plasma_stat_03r3] = '2'	PI/study name #03r3	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
140	ds_plasma_vol_03r3 Show the field ONLY if: [ds_plasma_rem_03(1)] = '1'	Volume (ul) #03r3	text (integer)										
141	ds_plasma_rem_03r3 Show the field ONLY if: [ds_plasma_rem_03(1)] = '1'	Volume remaining #03r3	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_03r3__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_03r3__1	Yes							
1	ds_plasma_rem_03r3__1	Yes											
142	ds_plasma_label_04 Show the field ONLY if: [ds_aliquiplasma] >= 4	Aliquot label #04	text Field Annotation: @PLACEHOLDER = 'Aliquot 04'										
143	ds_plasma_usedt_04 Show the field ONLY if: [ds_aliquiplasma] >= 4	Date aliquot used #04	text (date_ymd) Field Annotation: @HIDEBUTTON										
144	ds_plasma_stat_04 Show the field ONLY if: [ds_aliquiplasma] >= 4	Aliquot status #04	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
145	ds_plasma_study_04 Show the field ONLY if: [ds_plasma_stat_04] = '2'	PI/study name #04	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
146	ds_plasma_vol_04 Show the field ONLY if: [ds_aliquiplasma] >= 4	Volume (ul) #04	text (integer)										
147	ds_plasma_rem_04 Show the field ONLY if: [ds_aliquiplasma] >= 4	Volume remaining #04	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_04__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_04__1	Yes							
1	ds_plasma_rem_04__1	Yes											
148	ds_plasma_label_04r1 Show the field ONLY if: [ds_plasma_rem_04(1)] = '1'	Aliquot label #04R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
149	ds_plasma_usedt_04r1 Show the field ONLY if: [ds_plasma_rem_04(1)] = '1'	Date aliquot used #04R1	text (date_ymd) Field Annotation: @HIDEBUTTON										

150	ds_plasma_stat_04r1 Show the field ONLY if: [ds_plasma_rem_04(1)] = '1'	Aliquot status #04R1	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
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2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
151	ds_plasma_study_04r1 Show the field ONLY if: [ds_plasma_stat_04r1] = '2'	PI/study name #04R1	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
152	ds_plasma_vol_04r1 Show the field ONLY if: [ds_plasma_rem_04(1)] = '1'	Volume (ul) #04R1	text (integer)										
153	ds_plasma_rem_04r1 Show the field ONLY if: [ds_plasma_rem_04(1)] = '1'	Volume remaining #04R1	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_04r1__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_04r1__1	Yes							
1	ds_plasma_rem_04r1__1	Yes											
154	ds_plasma_label_04r2 Show the field ONLY if: [ds_plasma_rem_04(1)] = '1'	Aliquot label #04r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
155	ds_plasma_usedt_04r2 Show the field ONLY if: [ds_plasma_rem_04(1)] = '1'	Date aliquot used #04r2	text (date_ymd) Field Annotation: @HIDEBUTTON										
156	ds_plasma_stat_04r2 Show the field ONLY if: [ds_plasma_rem_04(1)] = '1'	Aliquot status #04r2	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
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3	Leftover 1st cycle												
4	Leftover 2nd cycle												
157	ds_plasma_study_04r2 Show the field ONLY if: [ds_plasma_stat_04r2] = '2'	PI/study name #04r2	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
158	ds_plasma_vol_04r2 Show the field ONLY if: [ds_plasma_rem_04(1)] = '1'	Volume (ul) #04r2	text (integer)										
159	ds_plasma_rem_04r2 Show the field ONLY if: [ds_plasma_rem_04(1)] = '1'	Volume remaining #04r2	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_04r2__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_04r2__1	Yes							
1	ds_plasma_rem_04r2__1	Yes											
160	ds_plasma_label_04r3 Show the field ONLY if: [ds_plasma_rem_04(1)] = '1'	Aliquot label #04r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'										
161	ds_plasma_usedt_04r3 Show the field ONLY if: [ds_plasma_rem_04(1)] = '1'	Date aliquot used #04r3	text (date_ymd) Field Annotation: @HIDEBUTTON										
162	ds_plasma_stat_04r3 Show the field ONLY if: [ds_plasma_rem_04(1)] = '1'	Aliquot status #04r3	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												

163	ds_plasma_study_04r3 Show the field ONLY if: [ds_plasma_stat_04r3] = '2'	PI/study name #04r3	dropdown 1 NIH DASH repository 2 Other  Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"
164	ds_plasma_vol_04r3 Show the field ONLY if: [ds_plasma_rem_04(1)] = '1'	Volume (ul) #04r3	text (integer)
165	ds_plasma_rem_04r3 Show the field ONLY if: [ds_plasma_rem_04(1)] = '1'	Volume remaining #04r3	checkbox 1 ds_plasma_rem_04r3__1 Yes
166	ds_plasma_label_05 Show the field ONLY if: [ds_aliquiplasma] >= 5	Aliquot label #05	text Field Annotation: @PLACEHOLDER = 'Aliquot 05'
167	ds_plasma_usedt_05 Show the field ONLY if: [ds_aliquiplasma] >= 5	Date aliquot used #05	text (date_ymd) Field Annotation: @HIDEBUTTON
168	ds_plasma_stat_05 Show the field ONLY if: [ds_aliquiplasma] >= 5	Aliquot status #05	dropdown 0 Available 1 Used for primary study 2 Used for ancillary study 3 Leftover 1st cycle 4 Leftover 2nd cycle  Custom alignment: RH
169	ds_plasma_study_05 Show the field ONLY if: [ds_plasma_stat_05] = '2'	PI/study name #05	dropdown 1 NIH DASH repository 2 Other  Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"
170	ds_plasma_vol_05 Show the field ONLY if: [ds_aliquiplasma] >= 5	Volume (ul) #05	text (integer)
171	ds_plasma_rem_05 Show the field ONLY if: [ds_aliquiplasma] >= 5	Volume remaining #05	checkbox 1 ds_plasma_rem_05__1 Yes
172	ds_plasma_label_05r1 Show the field ONLY if: [ds_plasma_rem_05(1)] = '1'	Aliquot label #05R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'
173	ds_plasma_usedt_05r1 Show the field ONLY if: [ds_plasma_rem_05(1)] = '1'	Date aliquot used #05R1	text (date_ymd) Field Annotation: @HIDEBUTTON
174	ds_plasma_stat_05r1 Show the field ONLY if: [ds_plasma_rem_05(1)] = '1'	Aliquot status #05R1	dropdown 0 Available 1 Used for primary study 2 Used for ancillary study 3 Leftover 1st cycle 4 Leftover 2nd cycle  Custom alignment: RH
175	ds_plasma_study_05r1 Show the field ONLY if: [ds_plasma_stat_05r1] = '2'	PI/study name #05R1	dropdown 1 NIH DASH repository 2 Other  Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"

176	ds_plasma_vol_05r1 Show the field ONLY if: [ds_plasma_rem_05(1)] = '1'	Volume (ul) #05R1	text (integer)										
177	ds_plasma_rem_05r1 Show the field ONLY if: [ds_plasma_rem_05(1)] = '1'	Volume remaining #05R1	checkbox <table border="1"><tr><td>1</td><td>ds_plasma_rem_05r1__1</td><td>Yes</td></tr></table>	1	ds_plasma_rem_05r1__1	Yes							
1	ds_plasma_rem_05r1__1	Yes											
178	ds_plasma_label_05r2 Show the field ONLY if: [ds_plasma_rem_05(1)] = '1'	Aliquot label #05r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
179	ds_plasma_usedt_05r2 Show the field ONLY if: [ds_plasma_rem_05(1)] = '1'	Date aliquot used #05r2	text (date_ymd) Field Annotation: @HIDEBUTTON										
180	ds_plasma_stat_05r2 Show the field ONLY if: [ds_plasma_rem_05(1)] = '1'	Aliquot status #05r2	dropdown <table border="1"><tr><td>0</td><td>Available</td></tr><tr><td>1</td><td>Used for primary study</td></tr><tr><td>2</td><td>Used for ancillary study</td></tr><tr><td>3</td><td>Leftover 1st cycle</td></tr><tr><td>4</td><td>Leftover 2nd cycle</td></tr></table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
181	ds_plasma_study_05r2 Show the field ONLY if: [ds_plasma_stat_05r2] = '2'	PI/study name #05r2	dropdown <table border="1"><tr><td>1</td><td>NIH DASH repository</td></tr><tr><td>2</td><td>Other</td></tr></table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
182	ds_plasma_vol_05r2 Show the field ONLY if: [ds_plasma_rem_05(1)] = '1'	Volume (ul) #05r2	text (integer)										
183	ds_plasma_rem_05r2 Show the field ONLY if: [ds_plasma_rem_05(1)] = '1'	Volume remaining #05r2	checkbox <table border="1"><tr><td>1</td><td>ds_plasma_rem_05r2__1</td><td>Yes</td></tr></table>	1	ds_plasma_rem_05r2__1	Yes							
1	ds_plasma_rem_05r2__1	Yes											
184	ds_plasma_label_05r3 Show the field ONLY if: [ds_plasma_rem_05(1)] = '1'	Aliquot label #05r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'										
185	ds_plasma_usedt_05r3 Show the field ONLY if: [ds_plasma_rem_05(1)] = '1'	Date aliquot used #05r3	text (date_ymd) Field Annotation: @HIDEBUTTON										
186	ds_plasma_stat_05r3 Show the field ONLY if: [ds_plasma_rem_05(1)] = '1'	Aliquot status #05r3	dropdown <table border="1"><tr><td>0</td><td>Available</td></tr><tr><td>1</td><td>Used for primary study</td></tr><tr><td>2</td><td>Used for ancillary study</td></tr><tr><td>3</td><td>Leftover 1st cycle</td></tr><tr><td>4</td><td>Leftover 2nd cycle</td></tr></table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
187	ds_plasma_study_05r3 Show the field ONLY if: [ds_plasma_stat_05r3] = '2'	PI/study name #05r3	dropdown <table border="1"><tr><td>1</td><td>NIH DASH repository</td></tr><tr><td>2</td><td>Other</td></tr></table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
188	ds_plasma_vol_05r3 Show the field ONLY if: [ds_plasma_rem_05(1)] = '1'	Volume (ul) #05r3	text (integer)										
189	ds_plasma_rem_05r3 Show the field ONLY if: [ds_plasma_rem_05(1)] = '1'	Volume remaining #05r3	checkbox <table border="1"><tr><td>1</td><td>ds_plasma_rem_05r3__1</td><td>Yes</td></tr></table>	1	ds_plasma_rem_05r3__1	Yes							
1	ds_plasma_rem_05r3__1	Yes											



190	ds_plasma_label_06 Show the field ONLY if: [ds_aliquiplasma] >= 6	Aliquot label #06	text Field Annotation: @PLACEHOLDER = 'Aliquot 06'										
191	ds_plasma_usedt_06 Show the field ONLY if: [ds_aliquiplasma] >= 6	Date aliquot used #06	text (date_ymd) Field Annotation: @HIDEBUTTON										
192	ds_plasma_stat_06 Show the field ONLY if: [ds_aliquiplasma] >= 6	Aliquot status #06	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
193	ds_plasma_study_06 Show the field ONLY if: [ds_plasma_stat_06] = '2'	PI/study name #06	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
194	ds_plasma_vol_06 Show the field ONLY if: [ds_aliquiplasma] >= 6	Volume (ul) #06	text (integer)										
195	ds_plasma_rem_06 Show the field ONLY if: [ds_aliquiplasma] >= 6	Volume remaining #06	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_06__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_06__1	Yes							
1	ds_plasma_rem_06__1	Yes											
196	ds_plasma_label_06r1 Show the field ONLY if: [ds_plasma_rem_06(1)] = '1'	Aliquot label #06R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
197	ds_plasma_usedt_06r1 Show the field ONLY if: [ds_plasma_rem_06(1)] = '1'	Date aliquot used #06R1	text (date_ymd) Field Annotation: @HIDEBUTTON										
198	ds_plasma_stat_06r1 Show the field ONLY if: [ds_plasma_rem_06(1)] = '1'	Aliquot status #06R1	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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4	Leftover 2nd cycle												
199	ds_plasma_study_06r1 Show the field ONLY if: [ds_plasma_stat_06r1] = '2'	PI/study name #06R1	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
200	ds_plasma_vol_06r1 Show the field ONLY if: [ds_plasma_rem_06(1)] = '1'	Volume (ul) #06R1	text (integer)										
201	ds_plasma_rem_06r1 Show the field ONLY if: [ds_plasma_rem_06(1)] = '1'	Volume remaining #06R1	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_06r1__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_06r1__1	Yes							
1	ds_plasma_rem_06r1__1	Yes											
202	ds_plasma_label_06r2 Show the field ONLY if: [ds_plasma_rem_06(1)] = '1'	Aliquot label #06r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
203	ds_plasma_usedt_06r2 Show the field ONLY if: [ds_plasma_rem_06(1)] = '1'	Date aliquot used #06r2	text (date_ymd) Field Annotation: @HIDEBUTTON										

204	ds_plasma_stat_06r2 Show the field ONLY if: [ds_plasma_rem_06(1)] = '1'	Aliquot status #06r2	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
205	ds_plasma_study_06r2 Show the field ONLY if: [ds_plasma_stat_06r2] = '2'	PI/study name #06r2	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> <p>Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"</p>	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
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206	ds_plasma_vol_06r2 Show the field ONLY if: [ds_plasma_rem_06(1)] = '1'	Volume (ul) #06r2	text (integer)										
207	ds_plasma_rem_06r2 Show the field ONLY if: [ds_plasma_rem_06(1)] = '1'	Volume remaining #06r2	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_06r2__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_06r2__1	Yes							
1	ds_plasma_rem_06r2__1	Yes											
208	ds_plasma_label_06r3 Show the field ONLY if: [ds_plasma_rem_06(1)] = '1'	Aliquot label #06r3	text Field Annotation: @PLACEHOLDER = "suffix "_R3"										
209	ds_plasma_usedt_06r3 Show the field ONLY if: [ds_plasma_rem_06(1)] = '1'	Date aliquot used #06r3	text (date_ymd) Field Annotation: @HIDEBUTTON										
210	ds_plasma_stat_06r3 Show the field ONLY if: [ds_plasma_rem_06(1)] = '1'	Aliquot status #06r3	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
211	ds_plasma_study_06r3 Show the field ONLY if: [ds_plasma_stat_06r3] = '2'	PI/study name #06r3	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> <p>Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"</p>	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
212	ds_plasma_vol_06r3 Show the field ONLY if: [ds_plasma_rem_06(1)] = '1'	Volume (ul) #06r3	text (integer)										
213	ds_plasma_rem_06r3 Show the field ONLY if: [ds_plasma_rem_06(1)] = '1'	Volume remaining #06r3	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_06r3__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_06r3__1	Yes							
1	ds_plasma_rem_06r3__1	Yes											
214	ds_plasma_label_07 Show the field ONLY if: [ds_aliquiplasma] >= 7	Aliquot label #07	text Field Annotation: @PLACEHOLDER = 'Aliquot 07'										
215	ds_plasma_usedt_07 Show the field ONLY if: [ds_aliquiplasma] >= 7	Date aliquot used #07	text (date_ymd) Field Annotation: @HIDEBUTTON										
216	ds_plasma_stat_07 Show the field ONLY if: [ds_aliquiplasma] >= 7	Aliquot status #07	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												

217	ds_plasma_study_07 Show the field ONLY if: [ds_plasma_stat_07] = '2'	PI/study name #07	dropdown 1 NIH DASH repository 2 Other  Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"
218	ds_plasma_vol_07 Show the field ONLY if: [ds_aliquiplasma] >= 7	Volume (ul) #07	text (integer)
219	ds_plasma_rem_07 Show the field ONLY if: [ds_aliquiplasma] >= 7	Volume remaining #07	checkbox 1 ds_plasma_rem_07__1 Yes
220	ds_plasma_label_07r1 Show the field ONLY if: [ds_plasma_rem_07(1)] = '1'	Aliquot label #07R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'
221	ds_plasma_usedt_07r1 Show the field ONLY if: [ds_plasma_rem_07(1)] = '1'	Date aliquot used #07R1	text (date_ymd) Field Annotation: @HIDEBUTTON
222	ds_plasma_stat_07r1 Show the field ONLY if: [ds_plasma_rem_07(1)] = '1'	Aliquot status #07R1	dropdown 0 Available 1 Used for primary study 2 Used for ancillary study 3 Leftover 1st cycle 4 Leftover 2nd cycle  Custom alignment: RH
223	ds_plasma_study_07r1 Show the field ONLY if: [ds_plasma_stat_07r1] = '2'	PI/study name #07R1	dropdown 1 NIH DASH repository 2 Other  Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"
224	ds_plasma_vol_07r1 Show the field ONLY if: [ds_plasma_rem_07(1)] = '1'	Volume (ul) #07R1	text (integer)
225	ds_plasma_rem_07r1 Show the field ONLY if: [ds_plasma_rem_07(1)] = '1'	Volume remaining #07R1	checkbox 1 ds_plasma_rem_07r1__1 Yes
226	ds_plasma_label_07r2 Show the field ONLY if: [ds_plasma_rem_07(1)] = '1'	Aliquot label #07r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'
227	ds_plasma_usedt_07r2 Show the field ONLY if: [ds_plasma_rem_07(1)] = '1'	Date aliquot used #07r2	text (date_ymd) Field Annotation: @HIDEBUTTON
228	ds_plasma_stat_07r2 Show the field ONLY if: [ds_plasma_rem_07(1)] = '1'	Aliquot status #07r2	dropdown 0 Available 1 Used for primary study 2 Used for ancillary study 3 Leftover 1st cycle 4 Leftover 2nd cycle  Custom alignment: RH
229	ds_plasma_study_07r2 Show the field ONLY if: [ds_plasma_stat_07r2] = '2'	PI/study name #07r2	dropdown 1 NIH DASH repository 2 Other  Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"

230	ds_plasma_vol_07r2 Show the field ONLY if: [ds_plasma_rem_07(1)] = '1'	Volume (ul) #07r2	text (integer)
231	ds_plasma_rem_07r2 Show the field ONLY if: [ds_plasma_rem_07(1)] = '1'	Volume remaining #07r2	checkbox 1 ds_plasma_rem_07r2__1 Yes
232	ds_plasma_label_07r3 Show the field ONLY if: [ds_plasma_rem_07(1)] = '1'	Aliquot label #07r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'
233	ds_plasma_usedt_07r3 Show the field ONLY if: [ds_plasma_rem_07(1)] = '1'	Date aliquot used #07r3	text (date_ymd) Field Annotation: @HIDEBUTTON
234	ds_plasma_stat_07r3 Show the field ONLY if: [ds_plasma_rem_07(1)] = '1'	Aliquot status #07r3	dropdown 0 Available 1 Used for primary study 2 Used for ancillary study 3 Leftover 1st cycle 4 Leftover 2nd cycle Custom alignment: RH
235	ds_plasma_study_07r3 Show the field ONLY if: [ds_plasma_stat_07r3] = '2'	PI/study name #07r3	dropdown 1 NIH DASH repository 2 Other Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"
236	ds_plasma_vol_07r3 Show the field ONLY if: [ds_plasma_rem_07(1)] = '1'	Volume (ul) #07r3	text (integer)
237	ds_plasma_rem_07r3 Show the field ONLY if: [ds_plasma_rem_07(1)] = '1'	Volume remaining #07r3	checkbox 1 ds_plasma_rem_07r3__1 Yes
238	ds_plasma_label_08 Show the field ONLY if: [ds_aliquiplasma] >= 8	Aliquot label #08	text Field Annotation: @PLACEHOLDER = 'Aliquot 08'
239	ds_plasma_usedt_08 Show the field ONLY if: [ds_aliquiplasma] >= 8	Date aliquot used #08	text (date_ymd) Field Annotation: @HIDEBUTTON
240	ds_plasma_stat_08 Show the field ONLY if: [ds_aliquiplasma] >= 8	Aliquot status #08	dropdown 0 Available 1 Used for primary study 2 Used for ancillary study 3 Leftover 1st cycle 4 Leftover 2nd cycle Custom alignment: RH
241	ds_plasma_study_08 Show the field ONLY if: [ds_plasma_stat_08] = '2'	PI/study name #08	dropdown 1 NIH DASH repository 2 Other Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"
242	ds_plasma_vol_08 Show the field ONLY if: [ds_aliquiplasma] >= 8	Volume (ul) #08	text (integer)
243	ds_plasma_rem_08 Show the field ONLY if: [ds_aliquiplasma] >= 8	Volume remaining #08	checkbox 1 ds_plasma_rem_08__1 Yes

244	ds_plasma_label_08r1 Show the field ONLY if: [ds_plasma_rem_08(1)] = '1'	Aliquot label #08R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
245	ds_plasma_usedt_08r1 Show the field ONLY if: [ds_plasma_rem_08(1)] = '1'	Date aliquot used #08R1	text (date_ymd) Field Annotation: @HIDEBUTTON										
246	ds_plasma_stat_08r1 Show the field ONLY if: [ds_plasma_rem_08(1)] = '1'	Aliquot status #08R1	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
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247	ds_plasma_study_08r1 Show the field ONLY if: [ds_plasma_stat_08r1] = '2'	PI/study name #08R1	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
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248	ds_plasma_vol_08r1 Show the field ONLY if: [ds_plasma_rem_08(1)] = '1'	Volume (ul) #08R1	text (integer)										
249	ds_plasma_rem_08r1 Show the field ONLY if: [ds_plasma_rem_08(1)] = '1'	Volume remaining #08R1	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_08r1__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_08r1__1	Yes							
1	ds_plasma_rem_08r1__1	Yes											
250	ds_plasma_label_08r2 Show the field ONLY if: [ds_plasma_rem_08(1)] = '1'	Aliquot label #08r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
251	ds_plasma_usedt_08r2 Show the field ONLY if: [ds_plasma_rem_08(1)] = '1'	Date aliquot used #08r2	text (date_ymd) Field Annotation: @HIDEBUTTON										
252	ds_plasma_stat_08r2 Show the field ONLY if: [ds_plasma_rem_08(1)] = '1'	Aliquot status #08r2	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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253	ds_plasma_study_08r2 Show the field ONLY if: [ds_plasma_stat_08r2] = '2'	PI/study name #08r2	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
254	ds_plasma_vol_08r2 Show the field ONLY if: [ds_plasma_rem_08(1)] = '1'	Volume (ul) #08r2	text (integer)										
255	ds_plasma_rem_08r2 Show the field ONLY if: [ds_plasma_rem_08(1)] = '1'	Volume remaining #08r2	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_08r2__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_08r2__1	Yes							
1	ds_plasma_rem_08r2__1	Yes											
256	ds_plasma_label_08r3 Show the field ONLY if: [ds_plasma_rem_08(1)] = '1'	Aliquot label #08r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'										
257	ds_plasma_usedt_08r3 Show the field ONLY if: [ds_plasma_rem_08(1)] = '1'	Date aliquot used #08r3	text (date_ymd) Field Annotation: @HIDEBUTTON										

258	ds_plasma_stat_08r3 Show the field ONLY if: [ds_plasma_rem_08(1)] = '1'	Aliquot status #08r3	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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3	Leftover 1st cycle												
4	Leftover 2nd cycle												
259	ds_plasma_study_08r3 Show the field ONLY if: [ds_plasma_stat_08r3] = '2'	PI/study name #08r3	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> <p>Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"</p>	1	NIH DASH repository	2	Other						
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2	Other												
260	ds_plasma_vol_08r3 Show the field ONLY if: [ds_plasma_rem_08(1)] = '1'	Volume (ul) #08r3	text (integer)										
261	ds_plasma_rem_08r3 Show the field ONLY if: [ds_plasma_rem_08(1)] = '1'	Volume remaining #08r3	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_08r3__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_08r3__1	Yes							
1	ds_plasma_rem_08r3__1	Yes											
262	ds_plasma_label_09 Show the field ONLY if: [ds_aliquiplasma] >= 9	Aliquot label #09	text Field Annotation: @PLACEHOLDER = 'Aliquot 09'										
263	ds_plasma_usedt_09 Show the field ONLY if: [ds_aliquiplasma] >= 9	Date aliquot used #09	text (date_ymd) Field Annotation: @HIDEBUTTON										
264	ds_plasma_stat_09 Show the field ONLY if: [ds_aliquiplasma] >= 9	Aliquot status #09	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
265	ds_plasma_study_09 Show the field ONLY if: [ds_plasma_stat_09] = '2'	PI/study name #09	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> <p>Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"</p>	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
266	ds_plasma_vol_09 Show the field ONLY if: [ds_aliquiplasma] >= 9	Volume (ul) #09	text (integer)										
267	ds_plasma_rem_09 Show the field ONLY if: [ds_aliquiplasma] >= 9	Volume remaining #09	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_09__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_09__1	Yes							
1	ds_plasma_rem_09__1	Yes											
268	ds_plasma_label_09r1 Show the field ONLY if: [ds_plasma_rem_09(1)] = '1'	Aliquot label #09R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
269	ds_plasma_usedt_09r1 Show the field ONLY if: [ds_plasma_rem_09(1)] = '1'	Date aliquot used #09R1	text (date_ymd) Field Annotation: @HIDEBUTTON										
270	ds_plasma_stat_09r1 Show the field ONLY if: [ds_plasma_rem_09(1)] = '1'	Aliquot status #09R1	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												

271	ds_plasma_study_09r1 Show the field ONLY if: [ds_plasma_stat_09r1] = '2'	PI/study name #09R1	dropdown 1 NIH DASH repository 2 Other  Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"
272	ds_plasma_vol_09r1 Show the field ONLY if: [ds_plasma_rem_09(1)] = '1'	Volume (ul) #09R1	text (integer)
273	ds_plasma_rem_09r1 Show the field ONLY if: [ds_plasma_rem_09(1)] = '1'	Volume remaining #09R1	checkbox 1 ds_plasma_rem_09r1__1 Yes
274	ds_plasma_label_09r2 Show the field ONLY if: [ds_plasma_rem_09(1)] = '1'	Aliquot label #09r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'
275	ds_plasma_usedt_09r2 Show the field ONLY if: [ds_plasma_rem_09(1)] = '1'	Date aliquot used #09r2	text (date_ymd) Field Annotation: @HIDEBUTTON
276	ds_plasma_stat_09r2 Show the field ONLY if: [ds_plasma_rem_09(1)] = '1'	Aliquot status #09r2	dropdown 0 Available 1 Used for primary study 2 Used for ancillary study 3 Leftover 1st cycle 4 Leftover 2nd cycle  Custom alignment: RH
277	ds_plasma_study_09r2 Show the field ONLY if: [ds_plasma_stat_09r2] = '2'	PI/study name #09r2	dropdown 1 NIH DASH repository 2 Other  Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"
278	ds_plasma_vol_09r2 Show the field ONLY if: [ds_plasma_rem_09(1)] = '1'	Volume (ul) #09r2	text (integer)
279	ds_plasma_rem_09r2 Show the field ONLY if: [ds_plasma_rem_09(1)] = '1'	Volume remaining #09r2	checkbox 1 ds_plasma_rem_09r2__1 Yes
280	ds_plasma_label_09r3 Show the field ONLY if: [ds_plasma_rem_09(1)] = '1'	Aliquot label #09r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'
281	ds_plasma_usedt_09r3 Show the field ONLY if: [ds_plasma_rem_09(1)] = '1'	Date aliquot used #09r3	text (date_ymd) Field Annotation: @HIDEBUTTON
282	ds_plasma_stat_09r3 Show the field ONLY if: [ds_plasma_rem_09(1)] = '1'	Aliquot status #09r3	dropdown 0 Available 1 Used for primary study 2 Used for ancillary study 3 Leftover 1st cycle 4 Leftover 2nd cycle  Custom alignment: RH
283	ds_plasma_study_09r3 Show the field ONLY if: [ds_plasma_stat_09r3] = '2'	PI/study name #09r3	dropdown 1 NIH DASH repository 2 Other  Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"

284	ds_plasma_vol_09r3 Show the field ONLY if: [ds_plasma_rem_09(1)] = '1'	Volume (ul) #09r3	text (integer)										
285	ds_plasma_rem_09r3 Show the field ONLY if: [ds_plasma_rem_09(1)] = '1'	Volume remaining #09r3	checkbox <table border="1"><tr><td>1</td><td>ds_plasma_rem_09r3__1</td><td>Yes</td></tr></table>	1	ds_plasma_rem_09r3__1	Yes							
1	ds_plasma_rem_09r3__1	Yes											
286	ds_plasma_label_10 Show the field ONLY if: [ds_aliquiplasma] >= 10	Aliquot label #10	text Field Annotation: @PLACEHOLDER = 'Aliquot 10'										
287	ds_plasma_usedt_10 Show the field ONLY if: [ds_aliquiplasma] >= 10	Date aliquot used #10	text (date_ymd) Field Annotation: @HIDEBUTTON										
288	ds_plasma_stat_10 Show the field ONLY if: [ds_aliquiplasma] >= 10	Aliquot status #10	dropdown <table border="1"><tr><td>0</td><td>Available</td></tr><tr><td>1</td><td>Used for primary study</td></tr><tr><td>2</td><td>Used for ancillary study</td></tr><tr><td>3</td><td>Leftover 1st cycle</td></tr><tr><td>4</td><td>Leftover 2nd cycle</td></tr></table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
289	ds_plasma_study_10 Show the field ONLY if: [ds_plasma_stat_10] = '2'	PI/study name #10	dropdown <table border="1"><tr><td>1</td><td>NIH DASH repository</td></tr><tr><td>2</td><td>Other</td></tr></table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
290	ds_plasma_vol_10 Show the field ONLY if: [ds_aliquiplasma] >= 10	Volume (ul) #10	text (integer)										
291	ds_plasma_rem_10 Show the field ONLY if: [ds_aliquiplasma] >= 10	Volume remaining #10	checkbox <table border="1"><tr><td>1</td><td>ds_plasma_rem_10__1</td><td>Yes</td></tr></table>	1	ds_plasma_rem_10__1	Yes							
1	ds_plasma_rem_10__1	Yes											
292	ds_plasma_label_10r1 Show the field ONLY if: [ds_plasma_rem_10(1)] = '1'	Aliquot label #10R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
293	ds_plasma_usedt_10r1 Show the field ONLY if: [ds_plasma_rem_10(1)] = '1'	Date aliquot used #10R1	text (date_ymd) Field Annotation: @HIDEBUTTON										
294	ds_plasma_stat_10r1 Show the field ONLY if: [ds_plasma_rem_10(1)] = '1'	Aliquot status #10R1	dropdown <table border="1"><tr><td>0</td><td>Available</td></tr><tr><td>1</td><td>Used for primary study</td></tr><tr><td>2</td><td>Used for ancillary study</td></tr><tr><td>3</td><td>Leftover 1st cycle</td></tr><tr><td>4</td><td>Leftover 2nd cycle</td></tr></table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
295	ds_plasma_study_10r1 Show the field ONLY if: [ds_plasma_stat_10r1] = '2'	PI/study name #10R1	dropdown <table border="1"><tr><td>1</td><td>NIH DASH repository</td></tr><tr><td>2</td><td>Other</td></tr></table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
296	ds_plasma_vol_10r1 Show the field ONLY if: [ds_plasma_rem_10(1)] = '1'	Volume (ul) #10R1	text (integer)										
297	ds_plasma_rem_10r1 Show the field ONLY if: [ds_plasma_rem_10(1)] = '1'	Volume remaining #10R1	checkbox <table border="1"><tr><td>1</td><td>ds_plasma_rem_10r1__1</td><td>Yes</td></tr></table>	1	ds_plasma_rem_10r1__1	Yes							
1	ds_plasma_rem_10r1__1	Yes											



298	ds_plasma_label_10r2 Show the field ONLY if: [ds_plasma_rem_10(1)] = '1'	Aliquot label #10r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
299	ds_plasma_usedt_10r2 Show the field ONLY if: [ds_plasma_rem_10(1)] = '1'	Date aliquot used #10r2	text (date_ymd) Field Annotation: @HIDEBUTTON										
300	ds_plasma_stat_10r2 Show the field ONLY if: [ds_plasma_rem_10(1)] = '1'	Aliquot status #10r2	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
301	ds_plasma_study_10r2 Show the field ONLY if: [ds_plasma_stat_10r2] = '2'	PI/study name #10r2	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
302	ds_plasma_vol_10r2 Show the field ONLY if: [ds_plasma_rem_10(1)] = '1'	Volume (ul) #10r2	text (integer)										
303	ds_plasma_rem_10r2 Show the field ONLY if: [ds_plasma_rem_10(1)] = '1'	Volume remaining #10r2	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_10r2__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_10r2__1	Yes							
1	ds_plasma_rem_10r2__1	Yes											
304	ds_plasma_label_10r3 Show the field ONLY if: [ds_plasma_rem_10(1)] = '1'	Aliquot label #10r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'										
305	ds_plasma_usedt_10r3 Show the field ONLY if: [ds_plasma_rem_10(1)] = '1'	Date aliquot used #10r3	text (date_ymd) Field Annotation: @HIDEBUTTON										
306	ds_plasma_stat_10r3 Show the field ONLY if: [ds_plasma_rem_10(1)] = '1'	Aliquot status #10r3	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
307	ds_plasma_study_10r3 Show the field ONLY if: [ds_plasma_stat_10r3] = '2'	PI/study name #10r3	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
308	ds_plasma_vol_10r3 Show the field ONLY if: [ds_plasma_rem_10(1)] = '1'	Volume (ul) #10r3	text (integer)										
309	ds_plasma_rem_10r3 Show the field ONLY if: [ds_plasma_rem_10(1)] = '1'	Volume remaining #10r3	checkbox <table border="1"> <tr><td>1</td><td>ds_plasma_rem_10r3__1</td><td>Yes</td></tr> </table>	1	ds_plasma_rem_10r3__1	Yes							
1	ds_plasma_rem_10r3__1	Yes											
310	emb_plasma0200 Show the field ONLY if: [ds_aliquiplasma] >= 1	Aliquot## Aliquot label Date aliquotused Aliquot status(study name if ancillary study) Volume(uL) Volume remaining	descriptive										
311	emb_plasma0201 Show the field ONLY if: [ds_aliquiplasma] >= 1	01 {ds_plasma_label_01} {ds_plasma_usedt_01} {ds_plasma_stat_01} {ds_plasma_study_01} {ds_plasma_vol_01} {ds_plasma_rem_01}	descriptive										

312	emb_plasma0201r1 Show the field ONLY if: [ds_plasma_rem_01(1)] = '1'	01 {ds_plasma_label_01r1} {ds_plasma_usedt_01r1} {ds_plasma_stat_01r1}{ds_plasma_study_01r1} {ds_plasma_vol_01r1} {ds_plasma_rem_01r1}	descriptive
313	emb_plasma0201r2 Show the field ONLY if: [ds_plasma_rem_01r1(1)] = '1'	01 {ds_plasma_label_01r2} {ds_plasma_usedt_01r2} {ds_plasma_stat_01r2}{ds_plasma_study_01r2} {ds_plasma_vol_01r2} {ds_plasma_rem_01r2}	descriptive
314	emb_plasma0201r3 Show the field ONLY if: [ds_plasma_rem_01r2(1)] = '1'	01 {ds_plasma_label_01r3} {ds_plasma_usedt_01r3} {ds_plasma_stat_01r3}{ds_plasma_study_01r3} {ds_plasma_vol_01r3} {ds_plasma_rem_01r3}	descriptive
315	emb_plasma0202 Show the field ONLY if: [ds_aliquiplasma] >= 2	02 {ds_plasma_label_02} {ds_plasma_usedt_02} {ds_plasma_stat_02} {ds_plasma_study_02} {ds_plasma_vol_02} {ds_plasma_rem_02}	descriptive
316	emb_plasma0202r1 Show the field ONLY if: [ds_plasma_rem_02(1)] = '1'	02 {ds_plasma_label_02r1} {ds_plasma_usedt_02r1} {ds_plasma_stat_02r1}{ds_plasma_study_02r1} {ds_plasma_vol_02r1} {ds_plasma_rem_02r1}	descriptive
317	emb_plasma0202r2 Show the field ONLY if: [ds_plasma_rem_02r1(1)] = '1'	02 {ds_plasma_label_02r2} {ds_plasma_usedt_02r2} {ds_plasma_stat_02r2}{ds_plasma_study_02r2} {ds_plasma_vol_02r2} {ds_plasma_rem_02r2}	descriptive
318	emb_plasma0202r3 Show the field ONLY if: [ds_plasma_rem_02r2(1)] = '1'	02 {ds_plasma_label_02r3} {ds_plasma_usedt_02r3} {ds_plasma_stat_02r3}{ds_plasma_study_02r3} {ds_plasma_vol_02r3} {ds_plasma_rem_02r3}	descriptive
319	emb_plasma0203 Show the field ONLY if: [ds_aliquiplasma] >= 3	03 {ds_plasma_label_03} {ds_plasma_usedt_03} {ds_plasma_stat_03} {ds_plasma_study_03} {ds_plasma_vol_03} {ds_plasma_rem_03}	descriptive
320	emb_plasma0203r1 Show the field ONLY if: [ds_plasma_rem_03(1)] = '1'	03 {ds_plasma_label_03r1} {ds_plasma_usedt_03r1} {ds_plasma_stat_03r1}{ds_plasma_study_03r1} {ds_plasma_vol_03r1} {ds_plasma_rem_03r1}	descriptive
321	emb_plasma0203r2 Show the field ONLY if: [ds_plasma_rem_03r1(1)] = '1'	03 {ds_plasma_label_03r2} {ds_plasma_usedt_03r2} {ds_plasma_stat_03r2}{ds_plasma_study_03r2} {ds_plasma_vol_03r2} {ds_plasma_rem_03r2}	descriptive
322	emb_plasma0203r3 Show the field ONLY if: [ds_plasma_rem_03r2(1)] = '1'	03 {ds_plasma_label_03r3} {ds_plasma_usedt_03r3} {ds_plasma_stat_03r3}{ds_plasma_study_03r3} {ds_plasma_vol_03r3} {ds_plasma_rem_03r3}	descriptive
323	emb_plasma0204 Show the field ONLY if: [ds_aliquiplasma] >= 4	04 {ds_plasma_label_04} {ds_plasma_usedt_04} {ds_plasma_stat_04} {ds_plasma_study_04} {ds_plasma_vol_04} {ds_plasma_rem_04}	descriptive
324	emb_plasma0204r1 Show the field ONLY if: [ds_plasma_rem_04(1)] = '1'	04 {ds_plasma_label_04r1} {ds_plasma_usedt_04r1} {ds_plasma_stat_04r1}{ds_plasma_study_04r1} {ds_plasma_vol_04r1} {ds_plasma_rem_04r1}	descriptive
325	emb_plasma0204r2 Show the field ONLY if: [ds_plasma_rem_04r1(1)] = '1'	04 {ds_plasma_label_04r2} {ds_plasma_usedt_04r2} {ds_plasma_stat_04r2}{ds_plasma_study_04r2} {ds_plasma_vol_04r2} {ds_plasma_rem_04r2}	descriptive
326	emb_plasma0204r3 Show the field ONLY if: [ds_plasma_rem_04r2(1)] = '1'	04 {ds_plasma_label_04r3} {ds_plasma_usedt_04r3} {ds_plasma_stat_04r3}{ds_plasma_study_04r3} {ds_plasma_vol_04r3} {ds_plasma_rem_04r3}	descriptive
327	emb_plasma0205 Show the field ONLY if: [ds_aliquiplasma] >= 5	05 {ds_plasma_label_05} {ds_plasma_usedt_05} {ds_plasma_stat_05} {ds_plasma_study_05} {ds_plasma_vol_05} {ds_plasma_rem_05}	descriptive
328	emb_plasma0205r1 Show the field ONLY if: [ds_plasma_rem_05(1)] = '1'	05 {ds_plasma_label_05r1} {ds_plasma_usedt_05r1} {ds_plasma_stat_05r1}{ds_plasma_study_05r1} {ds_plasma_vol_05r1} {ds_plasma_rem_05r1}	descriptive
329	emb_plasma0205r2 Show the field ONLY if: [ds_plasma_rem_05r1(1)] = '1'	05 {ds_plasma_label_05r2} {ds_plasma_usedt_05r2} {ds_plasma_stat_05r2}{ds_plasma_study_05r2} {ds_plasma_vol_05r2} {ds_plasma_rem_05r2}	descriptive
330	emb_plasma0205r3 Show the field ONLY if: [ds_plasma_rem_05r2(1)] = '1'	05 {ds_plasma_label_05r3} {ds_plasma_usedt_05r3} {ds_plasma_stat_05r3}{ds_plasma_study_05r3} {ds_plasma_vol_05r3} {ds_plasma_rem_05r3}	descriptive

331	emb_plasma0206 Show the field ONLY if: [ds_aliquiplasma] >= 6	06 {ds_plasma_label_06} {ds_plasma_usedt_06} {ds_plasma_stat_06} {ds_plasma_study_06} {ds_plasma_vol_06} {ds_plasma_rem_06}	descriptive
332	emb_plasma0206r1 Show the field ONLY if: [ds_plasma_rem_06(1)] = '1'	06 {ds_plasma_label_06r1} {ds_plasma_usedt_06r1} {ds_plasma_stat_06r1}{ds_plasma_study_06r1} {ds_plasma_vol_06r1} {ds_plasma_rem_06r1}	descriptive
333	emb_plasma0206r2 Show the field ONLY if: [ds_plasma_rem_06r1(1)] = '1'	06 {ds_plasma_label_06r2} {ds_plasma_usedt_06r2} {ds_plasma_stat_06r2}{ds_plasma_study_06r2} {ds_plasma_vol_06r2} {ds_plasma_rem_06r2}	descriptive
334	emb_plasma0206r3 Show the field ONLY if: [ds_plasma_rem_06r2(1)] = '1'	06 {ds_plasma_label_06r3} {ds_plasma_usedt_06r3} {ds_plasma_stat_06r3}{ds_plasma_study_06r3} {ds_plasma_vol_06r3} {ds_plasma_rem_06r3}	descriptive
335	emb_plasma0207 Show the field ONLY if: [ds_aliquiplasma] >= 7	07 {ds_plasma_label_07} {ds_plasma_usedt_07} {ds_plasma_stat_07} {ds_plasma_study_07} {ds_plasma_vol_07} {ds_plasma_rem_07}	descriptive
336	emb_plasma0207r1 Show the field ONLY if: [ds_plasma_rem_07(1)] = '1'	07 {ds_plasma_label_07r1} {ds_plasma_usedt_07r1} {ds_plasma_stat_07r1}{ds_plasma_study_07r1} {ds_plasma_vol_07r1} {ds_plasma_rem_07r1}	descriptive
337	emb_plasma0207r2 Show the field ONLY if: [ds_plasma_rem_07r1(1)] = '1'	07 {ds_plasma_label_07r2} {ds_plasma_usedt_07r2} {ds_plasma_stat_07r2}{ds_plasma_study_07r2} {ds_plasma_vol_07r2} {ds_plasma_rem_07r2}	descriptive
338	emb_plasma0207r3 Show the field ONLY if: [ds_plasma_rem_07r2(1)] = '1'	07 {ds_plasma_label_07r3} {ds_plasma_usedt_07r3} {ds_plasma_stat_07r3}{ds_plasma_study_07r3} {ds_plasma_vol_07r3} {ds_plasma_rem_07r3}	descriptive
339	emb_plasma0208 Show the field ONLY if: [ds_aliquiplasma] >= 8	08 {ds_plasma_label_08} {ds_plasma_usedt_08} {ds_plasma_stat_08} {ds_plasma_study_08} {ds_plasma_vol_08} {ds_plasma_rem_08}	descriptive
340	emb_plasma0208r1 Show the field ONLY if: [ds_plasma_rem_08(1)] = '1'	08 {ds_plasma_label_08r1} {ds_plasma_usedt_08r1} {ds_plasma_stat_08r1}{ds_plasma_study_08r1} {ds_plasma_vol_08r1} {ds_plasma_rem_08r1}	descriptive
341	emb_plasma0208r2 Show the field ONLY if: [ds_plasma_rem_08r1(1)] = '1'	08 {ds_plasma_label_08r2} {ds_plasma_usedt_08r2} {ds_plasma_stat_08r2}{ds_plasma_study_08r2} {ds_plasma_vol_08r2} {ds_plasma_rem_08r2}	descriptive
342	emb_plasma0208r3 Show the field ONLY if: [ds_plasma_rem_08r2(1)] = '1'	08 {ds_plasma_label_08r3} {ds_plasma_usedt_08r3} {ds_plasma_stat_08r3}{ds_plasma_study_08r3} {ds_plasma_vol_08r3} {ds_plasma_rem_08r3}	descriptive
343	emb_plasma0209 Show the field ONLY if: [ds_aliquiplasma] >= 9	09 {ds_plasma_label_09} {ds_plasma_usedt_09} {ds_plasma_stat_09} {ds_plasma_study_09} {ds_plasma_vol_09} {ds_plasma_rem_09}	descriptive
344	emb_plasma0209r1 Show the field ONLY if: [ds_plasma_rem_09(1)] = '1'	09 {ds_plasma_label_09r1} {ds_plasma_usedt_09r1} {ds_plasma_stat_09r1}{ds_plasma_study_09r1} {ds_plasma_vol_09r1} {ds_plasma_rem_09r1}	descriptive
345	emb_plasma0209r2 Show the field ONLY if: [ds_plasma_rem_09r1(1)] = '1'	09 {ds_plasma_label_09r2} {ds_plasma_usedt_09r2} {ds_plasma_stat_09r2}{ds_plasma_study_09r2} {ds_plasma_vol_09r2} {ds_plasma_rem_09r2}	descriptive
346	emb_plasma0209r3 Show the field ONLY if: [ds_plasma_rem_09r2(1)] = '1'	09 {ds_plasma_label_09r3} {ds_plasma_usedt_09r3} {ds_plasma_stat_09r3}{ds_plasma_study_09r3} {ds_plasma_vol_09r3} {ds_plasma_rem_09r3}	descriptive
347	emb_plasma0210 Show the field ONLY if: [ds_aliquiplasma] >= 10	10 {ds_plasma_label_10} {ds_plasma_usedt_10} {ds_plasma_stat_10} {ds_plasma_study_10} {ds_plasma_vol_10} {ds_plasma_rem_10}	descriptive
348	emb_plasma0210r1 Show the field ONLY if: [ds_plasma_rem_10(1)] = '1'	10 {ds_plasma_label_10r1} {ds_plasma_usedt_10r1} {ds_plasma_stat_10r1}{ds_plasma_study_10r1} {ds_plasma_vol_10r1} {ds_plasma_rem_10r1}	descriptive
349	emb_plasma0210r2 Show the field ONLY if: [ds_plasma_rem_10r1(1)] = '1'	10 {ds_plasma_label_10r2} {ds_plasma_usedt_10r2} {ds_plasma_stat_10r2}{ds_plasma_study_10r2} {ds_plasma_vol_10r2} {ds_plasma_rem_10r2}	descriptive

350	emb_plasma0210r3 Show the field ONLY if: [ds_plasma_rem_10r2(1)] = '1'	10 {ds_plasma_label_10r3} {ds_plasma_usedt_10r3} {ds_plasma_stat_10r3}{ds_plasma_study_10r3} {ds_plasma_vol_10r3} {ds_plasma_rem_10r3}	descriptive										
351	ds_serum_obtdt Show the field ONLY if: [ds_aliqu serum] >= 1	Section Header: SECTION: SAMPLE LOG - serumonly applicable for PreVAIL Kids substudy Date aliquots obtained for sample	text (date_ymd) Field Annotation: @HIDEBUTTON										
352	ds_brkserum Show the field ONLY if: [ds_aliqu serum] >= 1	INSTRUCTIONS: Date aliquot obtained should be at the sample level, unless it's a leftover sample. [ds_aliqu serum] have been obtained for this patient. Enter each aliquot info in the log below. For each aliquot label, follow format "patientID_sampletype_aliquotnumber" e.g., "123-12_serum_01" If aliquot volume remaining, please log a new aliquot with a revised sample number (suffix: _R1, _R2, etc.)	descriptive										
353	ds_serum_label_01 Show the field ONLY if: [ds_aliqu serum] >= 1	Aliquot label #01	text Field Annotation: @PLACEHOLDER = 'Aliquot 01'										
354	ds_serum_usedt_01 Show the field ONLY if: [ds_aliqu serum] >= 1	Date aliquot used #01	text (date_ymd) Field Annotation: @HIDEBUTTON										
355	ds_serum_stat_01 Show the field ONLY if: [ds_aliqu serum] >= 1	Aliquot status #01	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
356	ds_serum_study_01 Show the field ONLY if: [ds_serum_stat_01] = '2'	PI/study name #01	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
357	ds_serum_vol_01 Show the field ONLY if: [ds_aliqu serum] >= 1	Volume (ul) #01	text (integer)										
358	ds_serum_rem_01 Show the field ONLY if: [ds_aliqu serum] >= 1	Volume remaining #01	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_01__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_01__1	Yes							
1	ds_serum_rem_01__1	Yes											
359	ds_serum_label_01r1 Show the field ONLY if: [ds_serum_rem_01(1)] = '1'	Aliquot label #01R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
360	ds_serum_usedt_01r1 Show the field ONLY if: [ds_serum_rem_01(1)] = '1'	Date aliquot used #01R1	text (date_ymd) Field Annotation: @HIDEBUTTON										
361	ds_serum_stat_01r1 Show the field ONLY if: [ds_serum_rem_01(1)] = '1'	Aliquot status #01R1	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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3	Leftover 1st cycle												
4	Leftover 2nd cycle												
362	ds_serum_study_01r1 Show the field ONLY if: [ds_serum_stat_01r1] = '2'	PI/study name #01R1	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
363	ds_serum_vol_01r1 Show the field ONLY if: [ds_serum_rem_01(1)] = '1'	Volume (ul) #01R1	text (integer)										

364	ds_serum_rem_01r1 Show the field ONLY if: [ds_serum_rem_01(1)] = '1'	Volume remaining #01R1	checkbox <table border="1"><tr><td>1</td><td>ds_serum_rem_01r1__1</td><td>Yes</td></tr></table>	1	ds_serum_rem_01r1__1	Yes							
1	ds_serum_rem_01r1__1	Yes											
365	ds_serum_label_01r2 Show the field ONLY if: [ds_serum_rem_01(1)] = '1'	Aliquot label #01r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
366	ds_serum_usedt_01r2 Show the field ONLY if: [ds_serum_rem_01(1)] = '1'	Date aliquot used #01r2	text (date_ymd) Field Annotation: @HIDEBUTTON										
367	ds_serum_stat_01r2 Show the field ONLY if: [ds_serum_rem_01(1)] = '1'	Aliquot status #01r2	dropdown <table border="1"><tr><td>0</td><td>Available</td></tr><tr><td>1</td><td>Used for primary study</td></tr><tr><td>2</td><td>Used for ancillary study</td></tr><tr><td>3</td><td>Leftover 1st cycle</td></tr><tr><td>4</td><td>Leftover 2nd cycle</td></tr></table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
368	ds_serum_study_01r2 Show the field ONLY if: [ds_serum_stat_01r2] = '2'	PI/study name #01r2	dropdown <table border="1"><tr><td>1</td><td>NIH DASH repository</td></tr><tr><td>2</td><td>Other</td></tr></table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
369	ds_serum_vol_01r2 Show the field ONLY if: [ds_serum_rem_01(1)] = '1'	Volume (ul) #01r2	text (integer)										
370	ds_serum_rem_01r2 Show the field ONLY if: [ds_serum_rem_01(1)] = '1'	Volume remaining #01r2	checkbox <table border="1"><tr><td>1</td><td>ds_serum_rem_01r2__1</td><td>Yes</td></tr></table>	1	ds_serum_rem_01r2__1	Yes							
1	ds_serum_rem_01r2__1	Yes											
371	ds_serum_label_01r3 Show the field ONLY if: [ds_serum_rem_01(1)] = '1'	Aliquot label #01r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'										
372	ds_serum_usedt_01r3 Show the field ONLY if: [ds_serum_rem_01(1)] = '1'	Date aliquot used #01r3	text (date_ymd) Field Annotation: @HIDEBUTTON										
373	ds_serum_stat_01r3 Show the field ONLY if: [ds_serum_rem_01(1)] = '1'	Aliquot status #01r3	dropdown <table border="1"><tr><td>0</td><td>Available</td></tr><tr><td>1</td><td>Used for primary study</td></tr><tr><td>2</td><td>Used for ancillary study</td></tr><tr><td>3</td><td>Leftover 1st cycle</td></tr><tr><td>4</td><td>Leftover 2nd cycle</td></tr></table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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374	ds_serum_study_01r3 Show the field ONLY if: [ds_serum_stat_01r3] = '2'	PI/study name #01r3	dropdown <table border="1"><tr><td>1</td><td>NIH DASH repository</td></tr><tr><td>2</td><td>Other</td></tr></table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
375	ds_serum_vol_01r3 Show the field ONLY if: [ds_serum_rem_01(1)] = '1'	Volume (ul) #01r3	text (integer)										
376	ds_serum_rem_01r3 Show the field ONLY if: [ds_serum_rem_01(1)] = '1'	Volume remaining #01r3	checkbox <table border="1"><tr><td>1</td><td>ds_serum_rem_01r3__1</td><td>Yes</td></tr></table>	1	ds_serum_rem_01r3__1	Yes							
1	ds_serum_rem_01r3__1	Yes											
377	ds_serum_label_02 Show the field ONLY if: [ds_aliquoserum] >= 2	Aliquot label #02	text Field Annotation: @PLACEHOLDER = 'Aliquot 02'										

378	ds_serum_usedt_02 Show the field ONLY if: [ds_aliqserum] >= 2	Date aliquot used #02	text (date_ymd) Field Annotation: @HIDEBUTTON										
379	ds_serum_stat_02 Show the field ONLY if: [ds_aliqserum] >= 2	Aliquot status #02	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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3	Leftover 1st cycle												
4	Leftover 2nd cycle												
380	ds_serum_study_02 Show the field ONLY if: [ds_serum_stat_02] = '2'	PI/study name #02	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
381	ds_serum_vol_02 Show the field ONLY if: [ds_aliqserum] >= 2	Volume (ul) #02	text (integer)										
382	ds_serum_rem_02 Show the field ONLY if: [ds_aliqserum] >= 2	Volume remaining #02	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_02__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_02__1	Yes							
1	ds_serum_rem_02__1	Yes											
383	ds_serum_label_02r1 Show the field ONLY if: [ds_serum_rem_02(1)] = '1'	Aliquot label #02R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
384	ds_serum_usedt_02r1 Show the field ONLY if: [ds_serum_rem_02(1)] = '1'	Date aliquot used #02R1	text (date_ymd) Field Annotation: @HIDEBUTTON										
385	ds_serum_stat_02r1 Show the field ONLY if: [ds_serum_rem_02(1)] = '1'	Aliquot status #02R1	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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386	ds_serum_study_02r1 Show the field ONLY if: [ds_serum_stat_02r1] = '2'	PI/study name #02R1	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
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387	ds_serum_vol_02r1 Show the field ONLY if: [ds_serum_rem_02(1)] = '1'	Volume (ul) #02R1	text (integer)										
388	ds_serum_rem_02r1 Show the field ONLY if: [ds_serum_rem_02(1)] = '1'	Volume remaining #02R1	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_02r1__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_02r1__1	Yes							
1	ds_serum_rem_02r1__1	Yes											
389	ds_serum_label_02r2 Show the field ONLY if: [ds_serum_rem_02(1)] = '1'	Aliquot label #02r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
390	ds_serum_usedt_02r2 Show the field ONLY if: [ds_serum_rem_02(1)] = '1'	Date aliquot used #02r2	text (date_ymd) Field Annotation: @HIDEBUTTON										

391	ds_serum_stat_02r2 Show the field ONLY if: [ds_serum_rem_02(1)] = '1'	Aliquot status #02r2	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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4	Leftover 2nd cycle												
392	ds_serum_study_02r2 Show the field ONLY if: [ds_serum_stat_02r2] = '2'	PI/study name #02r2	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> <p>Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"</p>	1	NIH DASH repository	2	Other						
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393	ds_serum_vol_02r2 Show the field ONLY if: [ds_serum_rem_02(1)] = '1'	Volume (ul) #02r2	text (integer)										
394	ds_serum_rem_02r2 Show the field ONLY if: [ds_serum_rem_02(1)] = '1'	Volume remaining #02r2	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_02r2__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_02r2__1	Yes							
1	ds_serum_rem_02r2__1	Yes											
395	ds_serum_label_02r3 Show the field ONLY if: [ds_serum_rem_02(1)] = '1'	Aliquot label #02r3	text Field Annotation: @PLACEHOLDER = "suffix "_R3"										
396	ds_serum_usedt_02r3 Show the field ONLY if: [ds_serum_rem_02(1)] = '1'	Date aliquot used #02r3	text (date_ymd) Field Annotation: @HIDEBUTTON										
397	ds_serum_stat_02r3 Show the field ONLY if: [ds_serum_rem_02(1)] = '1'	Aliquot status #02r3	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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398	ds_serum_study_02r3 Show the field ONLY if: [ds_serum_stat_02r3] = '2'	PI/study name #02r3	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> <p>Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"</p>	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
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399	ds_serum_vol_02r3 Show the field ONLY if: [ds_serum_rem_02(1)] = '1'	Volume (ul) #02r3	text (integer)										
400	ds_serum_rem_02r3 Show the field ONLY if: [ds_serum_rem_02(1)] = '1'	Volume remaining #02r3	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_02r3__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_02r3__1	Yes							
1	ds_serum_rem_02r3__1	Yes											
401	ds_serum_label_03 Show the field ONLY if: [ds_aliqserum] >= 3	Aliquot label #03	text Field Annotation: @PLACEHOLDER = 'Aliquot 03'										
402	ds_serum_usedt_03 Show the field ONLY if: [ds_aliqserum] >= 3	Date aliquot used #03	text (date_ymd) Field Annotation: @HIDEBUTTON										
403	ds_serum_stat_03 Show the field ONLY if: [ds_aliqserum] >= 3	Aliquot status #03	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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404	ds_serum_study_03 Show the field ONLY if: [ds_serum_stat_03] = '2'	PI/study name #03	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
405	ds_serum_vol_03 Show the field ONLY if: [ds_aliqu serum] >= 3	Volume (ul) #03	text (integer)										
406	ds_serum_rem_03 Show the field ONLY if: [ds_aliqu serum] >= 3	Volume remaining #03	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_03__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_03__1	Yes							
1	ds_serum_rem_03__1	Yes											
407	ds_serum_label_03r1 Show the field ONLY if: [ds_serum_rem_03(1)] = '1'	Aliquot label #03R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
408	ds_serum_usedt_03r1 Show the field ONLY if: [ds_serum_rem_03(1)] = '1'	Date aliquot used #03R1	text (date_ymd) Field Annotation: @HIDEBUTTON										
409	ds_serum_stat_03r1 Show the field ONLY if: [ds_serum_rem_03(1)] = '1'	Aliquot status #03R1	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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410	ds_serum_study_03r1 Show the field ONLY if: [ds_serum_stat_03r1] = '2'	PI/study name #03R1	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
411	ds_serum_vol_03r1 Show the field ONLY if: [ds_serum_rem_03(1)] = '1'	Volume (ul) #03R1	text (integer)										
412	ds_serum_rem_03r1 Show the field ONLY if: [ds_serum_rem_03(1)] = '1'	Volume remaining #03R1	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_03r1__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_03r1__1	Yes							
1	ds_serum_rem_03r1__1	Yes											
413	ds_serum_label_03r2 Show the field ONLY if: [ds_serum_rem_03(1)] = '1'	Aliquot label #03r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
414	ds_serum_usedt_03r2 Show the field ONLY if: [ds_serum_rem_03(1)] = '1'	Date aliquot used #03r2	text (date_ymd) Field Annotation: @HIDEBUTTON										
415	ds_serum_stat_03r2 Show the field ONLY if: [ds_serum_rem_03(1)] = '1'	Aliquot status #03r2	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
416	ds_serum_study_03r2 Show the field ONLY if: [ds_serum_stat_03r2] = '2'	PI/study name #03r2	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
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417	ds_serum_vol_03r2 Show the field ONLY if: [ds_serum_rem_03(1)] = '1'	Volume (ul) #03r2	text (integer)
418	ds_serum_rem_03r2 Show the field ONLY if: [ds_serum_rem_03(1)] = '1'	Volume remaining #03r2	checkbox 1 ds_serum_rem_03r2__1 Yes
419	ds_serum_label_03r3 Show the field ONLY if: [ds_serum_rem_03(1)] = '1'	Aliquot label #03r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'
420	ds_serum_usedt_03r3 Show the field ONLY if: [ds_serum_rem_03(1)] = '1'	Date aliquot used #03r3	text (date_ymd) Field Annotation: @HIDEBUTTON
421	ds_serum_stat_03r3 Show the field ONLY if: [ds_serum_rem_03(1)] = '1'	Aliquot status #03r3	dropdown 0 Available 1 Used for primary study 2 Used for ancillary study 3 Leftover 1st cycle 4 Leftover 2nd cycle Custom alignment: RH
422	ds_serum_study_03r3 Show the field ONLY if: [ds_serum_stat_03r3] = '2'	PI/study name #03r3	dropdown 1 NIH DASH repository 2 Other Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"
423	ds_serum_vol_03r3 Show the field ONLY if: [ds_serum_rem_03(1)] = '1'	Volume (ul) #03r3	text (integer)
424	ds_serum_rem_03r3 Show the field ONLY if: [ds_serum_rem_03(1)] = '1'	Volume remaining #03r3	checkbox 1 ds_serum_rem_03r3__1 Yes
425	ds_serum_label_04 Show the field ONLY if: [ds_aliqserum] >= 4	Aliquot label #04	text Field Annotation: @PLACEHOLDER = 'Aliquot 04'
426	ds_serum_usedt_04 Show the field ONLY if: [ds_aliqserum] >= 4	Date aliquot used #04	text (date_ymd) Field Annotation: @HIDEBUTTON
427	ds_serum_stat_04 Show the field ONLY if: [ds_aliqserum] >= 4	Aliquot status #04	dropdown 0 Available 1 Used for primary study 2 Used for ancillary study 3 Leftover 1st cycle 4 Leftover 2nd cycle Custom alignment: RH
428	ds_serum_study_04 Show the field ONLY if: [ds_serum_stat_04] = '2'	PI/study name #04	dropdown 1 NIH DASH repository 2 Other Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"
429	ds_serum_vol_04 Show the field ONLY if: [ds_aliqserum] >= 4	Volume (ul) #04	text (integer)
430	ds_serum_rem_04 Show the field ONLY if: [ds_aliqserum] >= 4	Volume remaining #04	checkbox 1 ds_serum_rem_04__1 Yes

431	ds_serum_label_04r1 Show the field ONLY if: [ds_serum_rem_04(1)] = '1'	Aliquot label #04R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
432	ds_serum_usedt_04r1 Show the field ONLY if: [ds_serum_rem_04(1)] = '1'	Date aliquot used #04R1	text (date_ymd) Field Annotation: @HIDEBUTTON										
433	ds_serum_stat_04r1 Show the field ONLY if: [ds_serum_rem_04(1)] = '1'	Aliquot status #04R1	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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434	ds_serum_study_04r1 Show the field ONLY if: [ds_serum_stat_04r1] = '2'	PI/study name #04R1	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
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435	ds_serum_vol_04r1 Show the field ONLY if: [ds_serum_rem_04(1)] = '1'	Volume (ul) #04R1	text (integer)										
436	ds_serum_rem_04r1 Show the field ONLY if: [ds_serum_rem_04(1)] = '1'	Volume remaining #04R1	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_04r1__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_04r1__1	Yes							
1	ds_serum_rem_04r1__1	Yes											
437	ds_serum_label_04r2 Show the field ONLY if: [ds_serum_rem_04(1)] = '1'	Aliquot label #04r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
438	ds_serum_usedt_04r2 Show the field ONLY if: [ds_serum_rem_04(1)] = '1'	Date aliquot used #04r2	text (date_ymd) Field Annotation: @HIDEBUTTON										
439	ds_serum_stat_04r2 Show the field ONLY if: [ds_serum_rem_04(1)] = '1'	Aliquot status #04r2	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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440	ds_serum_study_04r2 Show the field ONLY if: [ds_serum_stat_04r2] = '2'	PI/study name #04r2	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
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441	ds_serum_vol_04r2 Show the field ONLY if: [ds_serum_rem_04(1)] = '1'	Volume (ul) #04r2	text (integer)										
442	ds_serum_rem_04r2 Show the field ONLY if: [ds_serum_rem_04(1)] = '1'	Volume remaining #04r2	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_04r2__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_04r2__1	Yes							
1	ds_serum_rem_04r2__1	Yes											
443	ds_serum_label_04r3 Show the field ONLY if: [ds_serum_rem_04(1)] = '1'	Aliquot label #04r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'										
444	ds_serum_usedt_04r3 Show the field ONLY if: [ds_serum_rem_04(1)] = '1'	Date aliquot used #04r3	text (date_ymd) Field Annotation: @HIDEBUTTON										

445	ds_serum_stat_04r3 Show the field ONLY if: [ds_serum_rem_04(1)] = '1'	Aliquot status #04r3	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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446	ds_serum_study_04r3 Show the field ONLY if: [ds_serum_stat_04r3] = '2'	PI/study name #04r3	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> <p>Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"</p>	1	NIH DASH repository	2	Other						
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447	ds_serum_vol_04r3 Show the field ONLY if: [ds_serum_rem_04(1)] = '1'	Volume (ul) #04r3	text (integer)										
448	ds_serum_rem_04r3 Show the field ONLY if: [ds_serum_rem_04(1)] = '1'	Volume remaining #04r3	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_04r3__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_04r3__1	Yes							
1	ds_serum_rem_04r3__1	Yes											
449	ds_serum_label_05 Show the field ONLY if: [ds_aliqu serum] >= 5	Aliquot label #05	text Field Annotation: @PLACEHOLDER = 'Aliquot 05'										
450	ds_serum_usedt_05 Show the field ONLY if: [ds_aliqu serum] >= 5	Date aliquot used #05	text (date_ymd) Field Annotation: @HIDEBUTTON										
451	ds_serum_stat_05 Show the field ONLY if: [ds_aliqu serum] >= 5	Aliquot status #05	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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3	Leftover 1st cycle												
4	Leftover 2nd cycle												
452	ds_serum_study_05 Show the field ONLY if: [ds_serum_stat_05] = '2'	PI/study name #05	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> <p>Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"</p>	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
453	ds_serum_vol_05 Show the field ONLY if: [ds_aliqu serum] >= 5	Volume (ul) #05	text (integer)										
454	ds_serum_rem_05 Show the field ONLY if: [ds_aliqu serum] >= 5	Volume remaining #05	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_05__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_05__1	Yes							
1	ds_serum_rem_05__1	Yes											
455	ds_serum_label_05r1 Show the field ONLY if: [ds_serum_rem_05(1)] = '1'	Aliquot label #05R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
456	ds_serum_usedt_05r1 Show the field ONLY if: [ds_serum_rem_05(1)] = '1'	Date aliquot used #05R1	text (date_ymd) Field Annotation: @HIDEBUTTON										
457	ds_serum_stat_05r1 Show the field ONLY if: [ds_serum_rem_05(1)] = '1'	Aliquot status #05R1	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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458	ds_serum_study_05r1 Show the field ONLY if: [ds_serum_stat_05r1] = '2'	PI/study name #05R1	dropdown 1 NIH DASH repository 2 Other  Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"
459	ds_serum_vol_05r1 Show the field ONLY if: [ds_serum_rem_05(1)] = '1'	Volume (ul) #05R1	text (integer)
460	ds_serum_rem_05r1 Show the field ONLY if: [ds_serum_rem_05(1)] = '1'	Volume remaining #05R1	checkbox 1 ds_serum_rem_05r1__1 Yes
461	ds_serum_label_05r2 Show the field ONLY if: [ds_serum_rem_05(1)] = '1'	Aliquot label #05r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'
462	ds_serum_usedt_05r2 Show the field ONLY if: [ds_serum_rem_05(1)] = '1'	Date aliquot used #05r2	text (date_ymd) Field Annotation: @HIDEBUTTON
463	ds_serum_stat_05r2 Show the field ONLY if: [ds_serum_rem_05(1)] = '1'	Aliquot status #05r2	dropdown 0 Available 1 Used for primary study 2 Used for ancillary study 3 Leftover 1st cycle 4 Leftover 2nd cycle  Custom alignment: RH
464	ds_serum_study_05r2 Show the field ONLY if: [ds_serum_stat_05r2] = '2'	PI/study name #05r2	dropdown 1 NIH DASH repository 2 Other  Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"
465	ds_serum_vol_05r2 Show the field ONLY if: [ds_serum_rem_05(1)] = '1'	Volume (ul) #05r2	text (integer)
466	ds_serum_rem_05r2 Show the field ONLY if: [ds_serum_rem_05(1)] = '1'	Volume remaining #05r2	checkbox 1 ds_serum_rem_05r2__1 Yes
467	ds_serum_label_05r3 Show the field ONLY if: [ds_serum_rem_05(1)] = '1'	Aliquot label #05r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'
468	ds_serum_usedt_05r3 Show the field ONLY if: [ds_serum_rem_05(1)] = '1'	Date aliquot used #05r3	text (date_ymd) Field Annotation: @HIDEBUTTON
469	ds_serum_stat_05r3 Show the field ONLY if: [ds_serum_rem_05(1)] = '1'	Aliquot status #05r3	dropdown 0 Available 1 Used for primary study 2 Used for ancillary study 3 Leftover 1st cycle 4 Leftover 2nd cycle  Custom alignment: RH
470	ds_serum_study_05r3 Show the field ONLY if: [ds_serum_stat_05r3] = '2'	PI/study name #05r3	dropdown 1 NIH DASH repository 2 Other  Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"

471	ds_serum_vol_05r3 Show the field ONLY if: [ds_serum_rem_05(1)] = '1'	Volume (ul) #05r3	text (integer)										
472	ds_serum_rem_05r3 Show the field ONLY if: [ds_serum_rem_05(1)] = '1'	Volume remaining #05r3	checkbox <table border="1"><tr><td>1</td><td>ds_serum_rem_05r3__1</td><td>Yes</td></tr></table>	1	ds_serum_rem_05r3__1	Yes							
1	ds_serum_rem_05r3__1	Yes											
473	ds_serum_label_06 Show the field ONLY if: [ds_aliqu serum] >= 6	Aliquot label #06	text Field Annotation: @PLACEHOLDER = 'Aliquot 06'										
474	ds_serum_usedt_06 Show the field ONLY if: [ds_aliqu serum] >= 6	Date aliquot used #06	text (date_ymd) Field Annotation: @HIDEBUTTON										
475	ds_serum_stat_06 Show the field ONLY if: [ds_aliqu serum] >= 6	Aliquot status #06	dropdown <table border="1"><tr><td>0</td><td>Available</td></tr><tr><td>1</td><td>Used for primary study</td></tr><tr><td>2</td><td>Used for ancillary study</td></tr><tr><td>3</td><td>Leftover 1st cycle</td></tr><tr><td>4</td><td>Leftover 2nd cycle</td></tr></table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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476	ds_serum_study_06 Show the field ONLY if: [ds_serum_stat_06] = '2'	PI/study name #06	dropdown <table border="1"><tr><td>1</td><td>NIH DASH repository</td></tr><tr><td>2</td><td>Other</td></tr></table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
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477	ds_serum_vol_06 Show the field ONLY if: [ds_aliqu serum] >= 6	Volume (ul) #06	text (integer)										
478	ds_serum_rem_06 Show the field ONLY if: [ds_aliqu serum] >= 6	Volume remaining #06	checkbox <table border="1"><tr><td>1</td><td>ds_serum_rem_06__1</td><td>Yes</td></tr></table>	1	ds_serum_rem_06__1	Yes							
1	ds_serum_rem_06__1	Yes											
479	ds_serum_label_06r1 Show the field ONLY if: [ds_serum_rem_06(1)] = '1'	Aliquot label #06R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
480	ds_serum_usedt_06r1 Show the field ONLY if: [ds_serum_rem_06(1)] = '1'	Date aliquot used #06R1	text (date_ymd) Field Annotation: @HIDEBUTTON										
481	ds_serum_stat_06r1 Show the field ONLY if: [ds_serum_rem_06(1)] = '1'	Aliquot status #06R1	dropdown <table border="1"><tr><td>0</td><td>Available</td></tr><tr><td>1</td><td>Used for primary study</td></tr><tr><td>2</td><td>Used for ancillary study</td></tr><tr><td>3</td><td>Leftover 1st cycle</td></tr><tr><td>4</td><td>Leftover 2nd cycle</td></tr></table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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482	ds_serum_study_06r1 Show the field ONLY if: [ds_serum_stat_06r1] = '2'	PI/study name #06R1	dropdown <table border="1"><tr><td>1</td><td>NIH DASH repository</td></tr><tr><td>2</td><td>Other</td></tr></table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
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483	ds_serum_vol_06r1 Show the field ONLY if: [ds_serum_rem_06(1)] = '1'	Volume (ul) #06R1	text (integer)										
484	ds_serum_rem_06r1 Show the field ONLY if: [ds_serum_rem_06(1)] = '1'	Volume remaining #06R1	checkbox <table border="1"><tr><td>1</td><td>ds_serum_rem_06r1__1</td><td>Yes</td></tr></table>	1	ds_serum_rem_06r1__1	Yes							
1	ds_serum_rem_06r1__1	Yes											

485	ds_serum_label_06r2 Show the field ONLY if: [ds_serum_rem_06(1)] = '1'	Aliquot label #06r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
486	ds_serum_usedt_06r2 Show the field ONLY if: [ds_serum_rem_06(1)] = '1'	Date aliquot used #06r2	text (date_ymd) Field Annotation: @HIDEBUTTON										
487	ds_serum_stat_06r2 Show the field ONLY if: [ds_serum_rem_06(1)] = '1'	Aliquot status #06r2	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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488	ds_serum_study_06r2 Show the field ONLY if: [ds_serum_stat_06r2] = '2'	PI/study name #06r2	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
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489	ds_serum_vol_06r2 Show the field ONLY if: [ds_serum_rem_06(1)] = '1'	Volume (ul) #06r2	text (integer)										
490	ds_serum_rem_06r2 Show the field ONLY if: [ds_serum_rem_06(1)] = '1'	Volume remaining #06r2	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_06r2__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_06r2__1	Yes							
1	ds_serum_rem_06r2__1	Yes											
491	ds_serum_label_06r3 Show the field ONLY if: [ds_serum_rem_06(1)] = '1'	Aliquot label #06r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'										
492	ds_serum_usedt_06r3 Show the field ONLY if: [ds_serum_rem_06(1)] = '1'	Date aliquot used #06r3	text (date_ymd) Field Annotation: @HIDEBUTTON										
493	ds_serum_stat_06r3 Show the field ONLY if: [ds_serum_rem_06(1)] = '1'	Aliquot status #06r3	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
494	ds_serum_study_06r3 Show the field ONLY if: [ds_serum_stat_06r3] = '2'	PI/study name #06r3	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
495	ds_serum_vol_06r3 Show the field ONLY if: [ds_serum_rem_06(1)] = '1'	Volume (ul) #06r3	text (integer)										
496	ds_serum_rem_06r3 Show the field ONLY if: [ds_serum_rem_06(1)] = '1'	Volume remaining #06r3	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_06r3__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_06r3__1	Yes							
1	ds_serum_rem_06r3__1	Yes											
497	ds_serum_label_07 Show the field ONLY if: [ds_aliqu serum] >= 7	Aliquot label #07	text Field Annotation: @PLACEHOLDER = 'Aliquot 07'										
498	ds_serum_usedt_07 Show the field ONLY if: [ds_aliqu serum] >= 7	Date aliquot used #07	text (date_ymd) Field Annotation: @HIDEBUTTON										

499	ds_serum_stat_07 Show the field ONLY if: [ds_aliqu serum] >= 7	Aliquot status #07	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
500	ds_serum_study_07 Show the field ONLY if: [ds_serum_stat_07] = '2'	PI/study name #07	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> <p>Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"</p>	1	NIH DASH repository	2	Other						
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2	Other												
501	ds_serum_vol_07 Show the field ONLY if: [ds_aliqu serum] >= 7	Volume (ul) #07	text (integer)										
502	ds_serum_rem_07 Show the field ONLY if: [ds_aliqu serum] >= 7	Volume remaining #07	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_07__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_07__1	Yes							
1	ds_serum_rem_07__1	Yes											
503	ds_serum_label_07r1 Show the field ONLY if: [ds_serum_rem_07(1)] = '1'	Aliquot label #07R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
504	ds_serum_usedt_07r1 Show the field ONLY if: [ds_serum_rem_07(1)] = '1'	Date aliquot used #07R1	text (date_ymd) Field Annotation: @HIDEBUTTON										
505	ds_serum_stat_07r1 Show the field ONLY if: [ds_serum_rem_07(1)] = '1'	Aliquot status #07R1	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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506	ds_serum_study_07r1 Show the field ONLY if: [ds_serum_stat_07r1] = '2'	PI/study name #07R1	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> <p>Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"</p>	1	NIH DASH repository	2	Other						
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507	ds_serum_vol_07r1 Show the field ONLY if: [ds_serum_rem_07(1)] = '1'	Volume (ul) #07R1	text (integer)										
508	ds_serum_rem_07r1 Show the field ONLY if: [ds_serum_rem_07(1)] = '1'	Volume remaining #07R1	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_07r1__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_07r1__1	Yes							
1	ds_serum_rem_07r1__1	Yes											
509	ds_serum_label_07r2 Show the field ONLY if: [ds_serum_rem_07(1)] = '1'	Aliquot label #07r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
510	ds_serum_usedt_07r2 Show the field ONLY if: [ds_serum_rem_07(1)] = '1'	Date aliquot used #07r2	text (date_ymd) Field Annotation: @HIDEBUTTON										
511	ds_serum_stat_07r2 Show the field ONLY if: [ds_serum_rem_07(1)] = '1'	Aliquot status #07r2	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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3	Leftover 1st cycle												
4	Leftover 2nd cycle												

512	ds_serum_study_07r2 Show the field ONLY if: [ds_serum_stat_07r2] = '2'	PI/study name #07r2	dropdown 1 NIH DASH repository 2 Other  Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"
513	ds_serum_vol_07r2 Show the field ONLY if: [ds_serum_rem_07(1)] = '1'	Volume (ul) #07r2	text (integer)
514	ds_serum_rem_07r2 Show the field ONLY if: [ds_serum_rem_07(1)] = '1'	Volume remaining #07r2	checkbox 1 ds_serum_rem_07r2__1 Yes
515	ds_serum_label_07r3 Show the field ONLY if: [ds_serum_rem_07(1)] = '1'	Aliquot label #07r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'
516	ds_serum_usedt_07r3 Show the field ONLY if: [ds_serum_rem_07(1)] = '1'	Date aliquot used #07r3	text (date_ymd) Field Annotation: @HIDEBUTTON
517	ds_serum_stat_07r3 Show the field ONLY if: [ds_serum_rem_07(1)] = '1'	Aliquot status #07r3	dropdown 0 Available 1 Used for primary study 2 Used for ancillary study 3 Leftover 1st cycle 4 Leftover 2nd cycle  Custom alignment: RH
518	ds_serum_study_07r3 Show the field ONLY if: [ds_serum_stat_07r3] = '2'	PI/study name #07r3	dropdown 1 NIH DASH repository 2 Other  Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"
519	ds_serum_vol_07r3 Show the field ONLY if: [ds_serum_rem_07(1)] = '1'	Volume (ul) #07r3	text (integer)
520	ds_serum_rem_07r3 Show the field ONLY if: [ds_serum_rem_07(1)] = '1'	Volume remaining #07r3	checkbox 1 ds_serum_rem_07r3__1 Yes
521	ds_serum_label_08 Show the field ONLY if: [ds_aliqserum] >= 8	Aliquot label #08	text Field Annotation: @PLACEHOLDER = 'Aliquot 08'
522	ds_serum_usedt_08 Show the field ONLY if: [ds_aliqserum] >= 8	Date aliquot used #08	text (date_ymd) Field Annotation: @HIDEBUTTON
523	ds_serum_stat_08 Show the field ONLY if: [ds_aliqserum] >= 8	Aliquot status #08	dropdown 0 Available 1 Used for primary study 2 Used for ancillary study 3 Leftover 1st cycle 4 Leftover 2nd cycle  Custom alignment: RH
524	ds_serum_study_08 Show the field ONLY if: [ds_serum_stat_08] = '2'	PI/study name #08	dropdown 1 NIH DASH repository 2 Other  Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"



525	ds_serum_vol_08 Show the field ONLY if: [ds_aliqserum] >= 8	Volume (ul) #08	text (integer)
526	ds_serum_rem_08 Show the field ONLY if: [ds_aliqserum] >= 8	Volume remaining #08	checkbox 1 ds_serum_rem_08__1 Yes
527	ds_serum_label_08r1 Show the field ONLY if: [ds_serum_rem_08(1)] = '1'	Aliquot label #08R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'
528	ds_serum_usedt_08r1 Show the field ONLY if: [ds_serum_rem_08(1)] = '1'	Date aliquot used #08R1	text (date_ymd) Field Annotation: @HIDEBUTTON
529	ds_serum_stat_08r1 Show the field ONLY if: [ds_serum_rem_08(1)] = '1'	Aliquot status #08R1	dropdown 0 Available 1 Used for primary study 2 Used for ancillary study 3 Leftover 1st cycle 4 Leftover 2nd cycle Custom alignment: RH
530	ds_serum_study_08r1 Show the field ONLY if: [ds_serum_stat_08r1] = '2'	PI/study name #08R1	dropdown 1 NIH DASH repository 2 Other Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"
531	ds_serum_vol_08r1 Show the field ONLY if: [ds_serum_rem_08(1)] = '1'	Volume (ul) #08R1	text (integer)
532	ds_serum_rem_08r1 Show the field ONLY if: [ds_serum_rem_08(1)] = '1'	Volume remaining #08R1	checkbox 1 ds_serum_rem_08r1__1 Yes
533	ds_serum_label_08r2 Show the field ONLY if: [ds_serum_rem_08(1)] = '1'	Aliquot label #08r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'
534	ds_serum_usedt_08r2 Show the field ONLY if: [ds_serum_rem_08(1)] = '1'	Date aliquot used #08r2	text (date_ymd) Field Annotation: @HIDEBUTTON
535	ds_serum_stat_08r2 Show the field ONLY if: [ds_serum_rem_08(1)] = '1'	Aliquot status #08r2	dropdown 0 Available 1 Used for primary study 2 Used for ancillary study 3 Leftover 1st cycle 4 Leftover 2nd cycle Custom alignment: RH
536	ds_serum_study_08r2 Show the field ONLY if: [ds_serum_stat_08r2] = '2'	PI/study name #08r2	dropdown 1 NIH DASH repository 2 Other Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"
537	ds_serum_vol_08r2 Show the field ONLY if: [ds_serum_rem_08(1)] = '1'	Volume (ul) #08r2	text (integer)
538	ds_serum_rem_08r2 Show the field ONLY if: [ds_serum_rem_08(1)] = '1'	Volume remaining #08r2	checkbox 1 ds_serum_rem_08r2__1 Yes

539	ds_serum_label_08r3 Show the field ONLY if: [ds_serum_rem_08(1)] = '1'	Aliquot label #08r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'										
540	ds_serum_usedt_08r3 Show the field ONLY if: [ds_serum_rem_08(1)] = '1'	Date aliquot used #08r3	text (date_ymd) Field Annotation: @HIDEBUTTON										
541	ds_serum_stat_08r3 Show the field ONLY if: [ds_serum_rem_08(1)] = '1'	Aliquot status #08r3	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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2	Used for ancillary study												
3	Leftover 1st cycle												
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542	ds_serum_study_08r3 Show the field ONLY if: [ds_serum_stat_08r3] = '2'	PI/study name #08r3	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
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543	ds_serum_vol_08r3 Show the field ONLY if: [ds_serum_rem_08(1)] = '1'	Volume (ul) #08r3	text (integer)										
544	ds_serum_rem_08r3 Show the field ONLY if: [ds_serum_rem_08(1)] = '1'	Volume remaining #08r3	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_08r3__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_08r3__1	Yes							
1	ds_serum_rem_08r3__1	Yes											
545	ds_serum_label_09 Show the field ONLY if: [ds_aliqu serum] >= 9	Aliquot label #09	text Field Annotation: @PLACEHOLDER = 'Aliquot 09'										
546	ds_serum_usedt_09 Show the field ONLY if: [ds_aliqu serum] >= 9	Date aliquot used #09	text (date_ymd) Field Annotation: @HIDEBUTTON										
547	ds_serum_stat_09 Show the field ONLY if: [ds_aliqu serum] >= 9	Aliquot status #09	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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3	Leftover 1st cycle												
4	Leftover 2nd cycle												
548	ds_serum_study_09 Show the field ONLY if: [ds_serum_stat_09] = '2'	PI/study name #09	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
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549	ds_serum_vol_09 Show the field ONLY if: [ds_aliqu serum] >= 9	Volume (ul) #09	text (integer)										
550	ds_serum_rem_09 Show the field ONLY if: [ds_aliqu serum] >= 9	Volume remaining #09	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_09__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_09__1	Yes							
1	ds_serum_rem_09__1	Yes											
551	ds_serum_label_09r1 Show the field ONLY if: [ds_serum_rem_09(1)] = '1'	Aliquot label #09R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
552	ds_serum_usedt_09r1 Show the field ONLY if: [ds_serum_rem_09(1)] = '1'	Date aliquot used #09R1	text (date_ymd) Field Annotation: @HIDEBUTTON										

553	ds_serum_stat_09r1 Show the field ONLY if: [ds_serum_rem_09(1)] = '1'	Aliquot status #09R1	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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554	ds_serum_study_09r1 Show the field ONLY if: [ds_serum_stat_09r1] = '2'	PI/study name #09R1	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> <p>Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"</p>	1	NIH DASH repository	2	Other						
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555	ds_serum_vol_09r1 Show the field ONLY if: [ds_serum_rem_09(1)] = '1'	Volume (ul) #09R1	text (integer)										
556	ds_serum_rem_09r1 Show the field ONLY if: [ds_serum_rem_09(1)] = '1'	Volume remaining #09R1	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_09r1__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_09r1__1	Yes							
1	ds_serum_rem_09r1__1	Yes											
557	ds_serum_label_09r2 Show the field ONLY if: [ds_serum_rem_09(1)] = '1'	Aliquot label #09r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
558	ds_serum_usedt_09r2 Show the field ONLY if: [ds_serum_rem_09(1)] = '1'	Date aliquot used #09r2	text (date_ymd) Field Annotation: @HIDEBUTTON										
559	ds_serum_stat_09r2 Show the field ONLY if: [ds_serum_rem_09(1)] = '1'	Aliquot status #09r2	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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560	ds_serum_study_09r2 Show the field ONLY if: [ds_serum_stat_09r2] = '2'	PI/study name #09r2	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> <p>Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"</p>	1	NIH DASH repository	2	Other						
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561	ds_serum_vol_09r2 Show the field ONLY if: [ds_serum_rem_09(1)] = '1'	Volume (ul) #09r2	text (integer)										
562	ds_serum_rem_09r2 Show the field ONLY if: [ds_serum_rem_09(1)] = '1'	Volume remaining #09r2	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_09r2__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_09r2__1	Yes							
1	ds_serum_rem_09r2__1	Yes											
563	ds_serum_label_09r3 Show the field ONLY if: [ds_serum_rem_09(1)] = '1'	Aliquot label #09r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'										
564	ds_serum_usedt_09r3 Show the field ONLY if: [ds_serum_rem_09(1)] = '1'	Date aliquot used #09r3	text (date_ymd) Field Annotation: @HIDEBUTTON										
565	ds_serum_stat_09r3 Show the field ONLY if: [ds_serum_rem_09(1)] = '1'	Aliquot status #09r3	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> <p>Custom alignment: RH</p>	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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566	ds_serum_study_09r3 Show the field ONLY if: [ds_serum_stat_09r3] = '2'	PI/study name #09r3	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
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567	ds_serum_vol_09r3 Show the field ONLY if: [ds_serum_rem_09(1)] = '1'	Volume (ul) #09r3	text (integer)										
568	ds_serum_rem_09r3 Show the field ONLY if: [ds_serum_rem_09(1)] = '1'	Volume remaining #09r3	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_09r3__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_09r3__1	Yes							
1	ds_serum_rem_09r3__1	Yes											
569	ds_serum_label_10 Show the field ONLY if: [ds_aliqserum] >= 10	Aliquot label #10	text Field Annotation: @PLACEHOLDER = 'Aliquot 10'										
570	ds_serum_usedt_10 Show the field ONLY if: [ds_aliqserum] >= 10	Date aliquot used #10	text (date_ymd) Field Annotation: @HIDEBUTTON										
571	ds_serum_stat_10 Show the field ONLY if: [ds_aliqserum] >= 10	Aliquot status #10	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
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572	ds_serum_study_10 Show the field ONLY if: [ds_serum_stat_10] = '2'	PI/study name #10	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
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574	ds_serum_rem_10 Show the field ONLY if: [ds_aliqserum] >= 10	Volume remaining #10	checkbox <table border="1"> <tr><td>1</td><td>ds_serum_rem_10__1</td><td>Yes</td></tr> </table>	1	ds_serum_rem_10__1	Yes							
1	ds_serum_rem_10__1	Yes											
575	ds_serum_label_10r1 Show the field ONLY if: [ds_serum_rem_10(1)] = '1'	Aliquot label #10R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
576	ds_serum_usedt_10r1 Show the field ONLY if: [ds_serum_rem_10(1)] = '1'	Date aliquot used #10R1	text (date_ymd) Field Annotation: @HIDEBUTTON										
577	ds_serum_stat_10r1 Show the field ONLY if: [ds_serum_rem_10(1)] = '1'	Aliquot status #10R1	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
578	ds_serum_study_10r1 Show the field ONLY if: [ds_serum_stat_10r1] = '2'	PI/study name #10R1	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												

579	ds_serum_vol_10r1 Show the field ONLY if: [ds_serum_rem_10(1)] = '1'	Volume (ul) #10R1	text (integer)										
580	ds_serum_rem_10r1 Show the field ONLY if: [ds_serum_rem_10(1)] = '1'	Volume remaining #10R1	checkbox <table border="1"><tr><td>1</td><td>ds_serum_rem_10r1__1</td><td>Yes</td></tr></table>	1	ds_serum_rem_10r1__1	Yes							
1	ds_serum_rem_10r1__1	Yes											
581	ds_serum_label_10r2 Show the field ONLY if: [ds_serum_rem_10(1)] = '1'	Aliquot label #10r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
582	ds_serum_usedt_10r2 Show the field ONLY if: [ds_serum_rem_10(1)] = '1'	Date aliquot used #10r2	text (date_ymd) Field Annotation: @HIDEBUTTON										
583	ds_serum_stat_10r2 Show the field ONLY if: [ds_serum_rem_10(1)] = '1'	Aliquot status #10r2	dropdown <table border="1"><tr><td>0</td><td>Available</td></tr><tr><td>1</td><td>Used for primary study</td></tr><tr><td>2</td><td>Used for ancillary study</td></tr><tr><td>3</td><td>Leftover 1st cycle</td></tr><tr><td>4</td><td>Leftover 2nd cycle</td></tr></table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
584	ds_serum_study_10r2 Show the field ONLY if: [ds_serum_stat_10r2] = '2'	PI/study name #10r2	dropdown <table border="1"><tr><td>1</td><td>NIH DASH repository</td></tr><tr><td>2</td><td>Other</td></tr></table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
585	ds_serum_vol_10r2 Show the field ONLY if: [ds_serum_rem_10(1)] = '1'	Volume (ul) #10r2	text (integer)										
586	ds_serum_rem_10r2 Show the field ONLY if: [ds_serum_rem_10(1)] = '1'	Volume remaining #10r2	checkbox <table border="1"><tr><td>1</td><td>ds_serum_rem_10r2__1</td><td>Yes</td></tr></table>	1	ds_serum_rem_10r2__1	Yes							
1	ds_serum_rem_10r2__1	Yes											
587	ds_serum_label_10r3 Show the field ONLY if: [ds_serum_rem_10(1)] = '1'	Aliquot label #10r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'										
588	ds_serum_usedt_10r3 Show the field ONLY if: [ds_serum_rem_10(1)] = '1'	Date aliquot used #10r3	text (date_ymd) Field Annotation: @HIDEBUTTON										
589	ds_serum_stat_10r3 Show the field ONLY if: [ds_serum_rem_10(1)] = '1'	Aliquot status #10r3	dropdown <table border="1"><tr><td>0</td><td>Available</td></tr><tr><td>1</td><td>Used for primary study</td></tr><tr><td>2</td><td>Used for ancillary study</td></tr><tr><td>3</td><td>Leftover 1st cycle</td></tr><tr><td>4</td><td>Leftover 2nd cycle</td></tr></table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
590	ds_serum_study_10r3 Show the field ONLY if: [ds_serum_stat_10r3] = '2'	PI/study name #10r3	dropdown <table border="1"><tr><td>1</td><td>NIH DASH repository</td></tr><tr><td>2</td><td>Other</td></tr></table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
591	ds_serum_vol_10r3 Show the field ONLY if: [ds_serum_rem_10(1)] = '1'	Volume (ul) #10r3	text (integer)										
592	ds_serum_rem_10r3 Show the field ONLY if: [ds_serum_rem_10(1)] = '1'	Volume remaining #10r3	checkbox <table border="1"><tr><td>1</td><td>ds_serum_rem_10r3__1</td><td>Yes</td></tr></table>	1	ds_serum_rem_10r3__1	Yes							
1	ds_serum_rem_10r3__1	Yes											

593	emb_serum0200 Show the field ONLY if: [ds_aliqu serum] >= 1	Aliquot## Aliquot label Date aliquotused Aliquot status(study name if ancillary study) Volume(uL) Volume remaining	descriptive
594	emb_serum0201 Show the field ONLY if: [ds_aliqu serum] >= 1	01 {ds_serum_label_01} {ds_serum_usedt_01} {ds_serum_stat_01} {ds_serum_study_01} {ds_serum_vol_01} {ds_serum_rem_01}	descriptive
595	emb_serum0201r1 Show the field ONLY if: [ds_serum_rem_01(1)] = '1'	01 {ds_serum_label_01r1} {ds_serum_usedt_01r1} {ds_serum_stat_01r1}{ds_serum_study_01r1} {ds_serum_vol_01r1} {ds_serum_rem_01r1}	descriptive
596	emb_serum0201r2 Show the field ONLY if: [ds_serum_rem_01r1(1)] = '1'	01 {ds_serum_label_01r2} {ds_serum_usedt_01r2} {ds_serum_stat_01r2}{ds_serum_study_01r2} {ds_serum_vol_01r2} {ds_serum_rem_01r2}	descriptive
597	emb_serum0201r3 Show the field ONLY if: [ds_serum_rem_01r2(1)] = '1'	01 {ds_serum_label_01r3} {ds_serum_usedt_01r3} {ds_serum_stat_01r3}{ds_serum_study_01r3} {ds_serum_vol_01r3} {ds_serum_rem_01r3}	descriptive
598	emb_serum0202 Show the field ONLY if: [ds_aliqu serum] >= 2	02 {ds_serum_label_02} {ds_serum_usedt_02} {ds_serum_stat_02} {ds_serum_study_02} {ds_serum_vol_02} {ds_serum_rem_02}	descriptive
599	emb_serum0202r1 Show the field ONLY if: [ds_serum_rem_02(1)] = '1'	02 {ds_serum_label_02r1} {ds_serum_usedt_02r1} {ds_serum_stat_02r1}{ds_serum_study_02r1} {ds_serum_vol_02r1} {ds_serum_rem_02r1}	descriptive
600	emb_serum0202r2 Show the field ONLY if: [ds_serum_rem_02r1(1)] = '1'	02 {ds_serum_label_02r2} {ds_serum_usedt_02r2} {ds_serum_stat_02r2}{ds_serum_study_02r2} {ds_serum_vol_02r2} {ds_serum_rem_02r2}	descriptive
601	emb_serum0202r3 Show the field ONLY if: [ds_serum_rem_02r2(1)] = '1'	02 {ds_serum_label_02r3} {ds_serum_usedt_02r3} {ds_serum_stat_02r3}{ds_serum_study_02r3} {ds_serum_vol_02r3} {ds_serum_rem_02r3}	descriptive
602	emb_serum0203 Show the field ONLY if: [ds_aliqu serum] >= 3	03 {ds_serum_label_03} {ds_serum_usedt_03} {ds_serum_stat_03} {ds_serum_study_03} {ds_serum_vol_03} {ds_serum_rem_03}	descriptive
603	emb_serum0203r1 Show the field ONLY if: [ds_serum_rem_03(1)] = '1'	03 {ds_serum_label_03r1} {ds_serum_usedt_03r1} {ds_serum_stat_03r1}{ds_serum_study_03r1} {ds_serum_vol_03r1} {ds_serum_rem_03r1}	descriptive
604	emb_serum0203r2 Show the field ONLY if: [ds_serum_rem_03r1(1)] = '1'	03 {ds_serum_label_03r2} {ds_serum_usedt_03r2} {ds_serum_stat_03r2}{ds_serum_study_03r2} {ds_serum_vol_03r2} {ds_serum_rem_03r2}	descriptive
605	emb_serum0203r3 Show the field ONLY if: [ds_serum_rem_03r2(1)] = '1'	03 {ds_serum_label_03r3} {ds_serum_usedt_03r3} {ds_serum_stat_03r3}{ds_serum_study_03r3} {ds_serum_vol_03r3} {ds_serum_rem_03r3}	descriptive
606	emb_serum0204 Show the field ONLY if: [ds_aliqu serum] >= 4	04 {ds_serum_label_04} {ds_serum_usedt_04} {ds_serum_stat_04} {ds_serum_study_04} {ds_serum_vol_04} {ds_serum_rem_04}	descriptive
607	emb_serum0204r1 Show the field ONLY if: [ds_serum_rem_04(1)] = '1'	04 {ds_serum_label_04r1} {ds_serum_usedt_04r1} {ds_serum_stat_04r1}{ds_serum_study_04r1} {ds_serum_vol_04r1} {ds_serum_rem_04r1}	descriptive
608	emb_serum0204r2 Show the field ONLY if: [ds_serum_rem_04r1(1)] = '1'	04 {ds_serum_label_04r2} {ds_serum_usedt_04r2} {ds_serum_stat_04r2}{ds_serum_study_04r2} {ds_serum_vol_04r2} {ds_serum_rem_04r2}	descriptive
609	emb_serum0204r3 Show the field ONLY if: [ds_serum_rem_04r2(1)] = '1'	04 {ds_serum_label_04r3} {ds_serum_usedt_04r3} {ds_serum_stat_04r3}{ds_serum_study_04r3} {ds_serum_vol_04r3} {ds_serum_rem_04r3}	descriptive
610	emb_serum0205 Show the field ONLY if: [ds_aliqu serum] >= 5	05 {ds_serum_label_05} {ds_serum_usedt_05} {ds_serum_stat_05} {ds_serum_study_05} {ds_serum_vol_05} {ds_serum_rem_05}	descriptive
611	emb_serum0205r1 Show the field ONLY if: [ds_serum_rem_05(1)] = '1'	05 {ds_serum_label_05r1} {ds_serum_usedt_05r1} {ds_serum_stat_05r1}{ds_serum_study_05r1} {ds_serum_vol_05r1} {ds_serum_rem_05r1}	descriptive

612	emb_serum0205r2 Show the field ONLY if: [ds_serum_rem_05r1(1)] = '1'	05 {ds_serum_label_05r2} {ds_serum_usedt_05r2} {ds_serum_stat_05r2}{ds_serum_study_05r2} {ds_serum_vol_05r2} {ds_serum_rem_05r2}	descriptive
613	emb_serum0205r3 Show the field ONLY if: [ds_serum_rem_05r2(1)] = '1'	05 {ds_serum_label_05r3} {ds_serum_usedt_05r3} {ds_serum_stat_05r3}{ds_serum_study_05r3} {ds_serum_vol_05r3} {ds_serum_rem_05r3}	descriptive
614	emb_serum0206 Show the field ONLY if: [ds_aliqserum] >= 6	06 {ds_serum_label_06} {ds_serum_usedt_06} {ds_serum_stat_06} {ds_serum_study_06} {ds_serum_vol_06} {ds_serum_rem_06}	descriptive
615	emb_serum0206r1 Show the field ONLY if: [ds_serum_rem_06(1)] = '1'	06 {ds_serum_label_06r1} {ds_serum_usedt_06r1} {ds_serum_stat_06r1}{ds_serum_study_06r1} {ds_serum_vol_06r1} {ds_serum_rem_06r1}	descriptive
616	emb_serum0206r2 Show the field ONLY if: [ds_serum_rem_06r1(1)] = '1'	06 {ds_serum_label_06r2} {ds_serum_usedt_06r2} {ds_serum_stat_06r2}{ds_serum_study_06r2} {ds_serum_vol_06r2} {ds_serum_rem_06r2}	descriptive
617	emb_serum0206r3 Show the field ONLY if: [ds_serum_rem_06r2(1)] = '1'	06 {ds_serum_label_06r3} {ds_serum_usedt_06r3} {ds_serum_stat_06r3}{ds_serum_study_06r3} {ds_serum_vol_06r3} {ds_serum_rem_06r3}	descriptive
618	emb_serum0207 Show the field ONLY if: [ds_aliqserum] >= 7	07 {ds_serum_label_07} {ds_serum_usedt_07} {ds_serum_stat_07} {ds_serum_study_07} {ds_serum_vol_07} {ds_serum_rem_07}	descriptive
619	emb_serum0207r1 Show the field ONLY if: [ds_serum_rem_07(1)] = '1'	07 {ds_serum_label_07r1} {ds_serum_usedt_07r1} {ds_serum_stat_07r1}{ds_serum_study_07r1} {ds_serum_vol_07r1} {ds_serum_rem_07r1}	descriptive
620	emb_serum0207r2 Show the field ONLY if: [ds_serum_rem_07r1(1)] = '1'	07 {ds_serum_label_07r2} {ds_serum_usedt_07r2} {ds_serum_stat_07r2}{ds_serum_study_07r2} {ds_serum_vol_07r2} {ds_serum_rem_07r2}	descriptive
621	emb_serum0207r3 Show the field ONLY if: [ds_serum_rem_07r2(1)] = '1'	07 {ds_serum_label_07r3} {ds_serum_usedt_07r3} {ds_serum_stat_07r3}{ds_serum_study_07r3} {ds_serum_vol_07r3} {ds_serum_rem_07r3}	descriptive
622	emb_serum0208 Show the field ONLY if: [ds_aliqserum] >= 8	08 {ds_serum_label_08} {ds_serum_usedt_08} {ds_serum_stat_08} {ds_serum_study_08} {ds_serum_vol_08} {ds_serum_rem_08}	descriptive
623	emb_serum0208r1 Show the field ONLY if: [ds_serum_rem_08(1)] = '1'	08 {ds_serum_label_08r1} {ds_serum_usedt_08r1} {ds_serum_stat_08r1}{ds_serum_study_08r1} {ds_serum_vol_08r1} {ds_serum_rem_08r1}	descriptive
624	emb_serum0208r2 Show the field ONLY if: [ds_serum_rem_08r1(1)] = '1'	08 {ds_serum_label_08r2} {ds_serum_usedt_08r2} {ds_serum_stat_08r2}{ds_serum_study_08r2} {ds_serum_vol_08r2} {ds_serum_rem_08r2}	descriptive
625	emb_serum0208r3 Show the field ONLY if: [ds_serum_rem_08r2(1)] = '1'	08 {ds_serum_label_08r3} {ds_serum_usedt_08r3} {ds_serum_stat_08r3}{ds_serum_study_08r3} {ds_serum_vol_08r3} {ds_serum_rem_08r3}	descriptive
626	emb_serum0209 Show the field ONLY if: [ds_aliqserum] >= 9	09 {ds_serum_label_09} {ds_serum_usedt_09} {ds_serum_stat_09} {ds_serum_study_09} {ds_serum_vol_09} {ds_serum_rem_09}	descriptive
627	emb_serum0209r1 Show the field ONLY if: [ds_serum_rem_09(1)] = '1'	09 {ds_serum_label_09r1} {ds_serum_usedt_09r1} {ds_serum_stat_09r1}{ds_serum_study_09r1} {ds_serum_vol_09r1} {ds_serum_rem_09r1}	descriptive
628	emb_serum0209r2 Show the field ONLY if: [ds_serum_rem_09r1(1)] = '1'	09 {ds_serum_label_09r2} {ds_serum_usedt_09r2} {ds_serum_stat_09r2}{ds_serum_study_09r2} {ds_serum_vol_09r2} {ds_serum_rem_09r2}	descriptive
629	emb_serum0209r3 Show the field ONLY if: [ds_serum_rem_09r2(1)] = '1'	09 {ds_serum_label_09r3} {ds_serum_usedt_09r3} {ds_serum_stat_09r3}{ds_serum_study_09r3} {ds_serum_vol_09r3} {ds_serum_rem_09r3}	descriptive
630	emb_serum0210 Show the field ONLY if: [ds_aliqserum] >= 10	10 {ds_serum_label_10} {ds_serum_usedt_10} {ds_serum_stat_10} {ds_serum_study_10} {ds_serum_vol_10} {ds_serum_rem_10}	descriptive

631	emb_serum0210r1 Show the field ONLY if: [ds_serum_rem_10(1)] = '1'	10 {ds_serum_label_10r1} {ds_serum_usedt_10r1} {ds_serum_stat_10r1}{ds_serum_study_10r1} {ds_serum_vol_10r1} {ds_serum_rem_10r1}	descriptive										
632	emb_serum0210r2 Show the field ONLY if: [ds_serum_rem_10r1(1)] = '1'	10 {ds_serum_label_10r2} {ds_serum_usedt_10r2} {ds_serum_stat_10r2}{ds_serum_study_10r2} {ds_serum_vol_10r2} {ds_serum_rem_10r2}	descriptive										
633	emb_serum0210r3 Show the field ONLY if: [ds_serum_rem_10r2(1)] = '1'	10 {ds_serum_label_10r3} {ds_serum_usedt_10r3} {ds_serum_stat_10r3}{ds_serum_study_10r3} {ds_serum_vol_10r3} {ds_serum_rem_10r3}	descriptive										
634	ds_buffy_obtdt Show the field ONLY if: [ds_aliquibuffy] >= 1	Section Header: SECTION: SAMPLE LOG - BUFFY COAT <i>Only applicable for PreVAIKids study</i> Date aliquots obtained for sample	text (date_ymd) Field Annotation: @HIDEBUTTON										
635	ds_brkbuffy Show the field ONLY if: [ds_aliquibuffy] >= 1	INSTRUCTIONS: Date aliquot obtained should be at the sample level, unless it's a leftover sample. [ds_aliquibuffy] have been obtained for this patient. Enter each aliquot info in the log below. For each aliquot label, follow format "patientID_sampletype_aliquotnumber" e.g., "123-12_buffy_01" If aliquot volume remaining, please log a new aliquot with a revised sample number (suffix: _R1, _R2, etc.)	descriptive										
636	ds_buffy_label_01 Show the field ONLY if: [ds_aliquibuffy] >= 1	Aliquot label #01	text Field Annotation: @PLACEHOLDER = 'Aliquot 01'										
637	ds_buffy_usedt_01 Show the field ONLY if: [ds_aliquibuffy] >= 1	Date aliquot used #01	text (date_ymd) Field Annotation: @HIDEBUTTON										
638	ds_buffy_stat_01 Show the field ONLY if: [ds_aliquibuffy] >= 1	Aliquot status #01	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
639	ds_buffy_study_01 Show the field ONLY if: [ds_buffy_stat_01] = '2'	PI/study name #01	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
640	ds_buffy_vol_01 Show the field ONLY if: [ds_aliquibuffy] >= 1	Volume (ul) #01	text (integer)										
641	ds_buffy_rem_01 Show the field ONLY if: [ds_aliquibuffy] >= 1	Volume remaining #01	checkbox <table border="1"> <tr><td>1</td><td>ds_buffy_rem_01__1</td><td>Yes</td></tr> </table>	1	ds_buffy_rem_01__1	Yes							
1	ds_buffy_rem_01__1	Yes											
642	ds_buffy_label_01r1 Show the field ONLY if: [ds_buffy_rem_01(1)] = '1'	Aliquot label #01R1	text Field Annotation: @PLACEHOLDER = 'suffix "_R1"'										
643	ds_buffy_usedt_01r1 Show the field ONLY if: [ds_buffy_rem_01(1)] = '1'	Date aliquot used #01R1	text (date_ymd) Field Annotation: @HIDEBUTTON										
644	ds_buffy_stat_01r1 Show the field ONLY if: [ds_buffy_rem_01(1)] = '1'	Aliquot status #01R1	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												



645	ds_buffy_study_01r1 Show the field ONLY if: [ds_buffy_stat_01r1] = '2'	PI/study name #01R1	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
646	ds_buffy_vol_01r1 Show the field ONLY if: [ds_buffy_rem_01(1)] = '1'	Volume (ul) #01R1	text (integer)										
647	ds_buffy_rem_01r1 Show the field ONLY if: [ds_buffy_rem_01(1)] = '1'	Volume remaining #01R1	checkbox <table border="1"> <tr><td>1</td><td>ds_buffy_rem_01r1__1</td><td>Yes</td></tr> </table>	1	ds_buffy_rem_01r1__1	Yes							
1	ds_buffy_rem_01r1__1	Yes											
648	ds_buffy_label_01r2 Show the field ONLY if: [ds_buffy_rem_01(1)] = '1'	Aliquot label #01r2	text Field Annotation: @PLACEHOLDER = 'suffix "_R2"'										
649	ds_buffy_usedt_01r2 Show the field ONLY if: [ds_buffy_rem_01(1)] = '1'	Date aliquot used #01r2	text (date_ymd) Field Annotation: @HIDEBUTTON										
650	ds_buffy_stat_01r2 Show the field ONLY if: [ds_buffy_rem_01(1)] = '1'	Aliquot status #01r2	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
651	ds_buffy_study_01r2 Show the field ONLY if: [ds_buffy_stat_01r2] = '2'	PI/study name #01r2	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												
652	ds_buffy_vol_01r2 Show the field ONLY if: [ds_buffy_rem_01(1)] = '1'	Volume (ul) #01r2	text (integer)										
653	ds_buffy_rem_01r2 Show the field ONLY if: [ds_buffy_rem_01(1)] = '1'	Volume remaining #01r2	checkbox <table border="1"> <tr><td>1</td><td>ds_buffy_rem_01r2__1</td><td>Yes</td></tr> </table>	1	ds_buffy_rem_01r2__1	Yes							
1	ds_buffy_rem_01r2__1	Yes											
654	ds_buffy_label_01r3 Show the field ONLY if: [ds_buffy_rem_01(1)] = '1'	Aliquot label #01r3	text Field Annotation: @PLACEHOLDER = 'suffix "_R3"'										
655	ds_buffy_usedt_01r3 Show the field ONLY if: [ds_buffy_rem_01(1)] = '1'	Date aliquot used #01r3	text (date_ymd) Field Annotation: @HIDEBUTTON										
656	ds_buffy_stat_01r3 Show the field ONLY if: [ds_buffy_rem_01(1)] = '1'	Aliquot status #01r3	dropdown <table border="1"> <tr><td>0</td><td>Available</td></tr> <tr><td>1</td><td>Used for primary study</td></tr> <tr><td>2</td><td>Used for ancillary study</td></tr> <tr><td>3</td><td>Leftover 1st cycle</td></tr> <tr><td>4</td><td>Leftover 2nd cycle</td></tr> </table> Custom alignment: RH	0	Available	1	Used for primary study	2	Used for ancillary study	3	Leftover 1st cycle	4	Leftover 2nd cycle
0	Available												
1	Used for primary study												
2	Used for ancillary study												
3	Leftover 1st cycle												
4	Leftover 2nd cycle												
657	ds_buffy_study_01r3 Show the field ONLY if: [ds_buffy_stat_01r3] = '2'	PI/study name #01r3	dropdown <table border="1"> <tr><td>1</td><td>NIH DASH repository</td></tr> <tr><td>2</td><td>Other</td></tr> </table> Field Annotation: @PLACEHOLDER = "Ancillary study PI / study name"	1	NIH DASH repository	2	Other						
1	NIH DASH repository												
2	Other												

658	ds_buffy_vol_01r3 Show the field ONLY if: [ds_buffy_rem_01(1)] = '1'	Volume (ul) #01r3	text (integer)
659	ds_buffy_rem_01r3 Show the field ONLY if: [ds_buffy_rem_01(1)] = '1'	Volume remaining #01r3	checkbox 1 ds_buffy_rem_01r3__1 Yes
660	emb_buffy0200 Show the field ONLY if: [ds_aliquibuffy] >= 1	Aliquot## Aliquot label Date aliquotused Aliquot status(study name if ancillary study) Volume(uL) Volume remaining	descriptive
661	emb_buffy0201 Show the field ONLY if: [ds_aliquibuffy] >= 1	01 {ds_buffy_label_01} {ds_buffy_usedt_01} {ds_buffy_stat_01} {ds_buffy_study_01} {ds_buffy_vol_01} {ds_buffy_rem_01}	descriptive
662	emb_buffy0201r1 Show the field ONLY if: [ds_buffy_rem_01(1)] = '1'	01 {ds_buffy_label_01r1} {ds_buffy_usedt_01r1} {ds_buffy_stat_01r1} {ds_buffy_study_01r1} {ds_buffy_vol_01r1} {ds_buffy_rem_01r1}	descriptive
663	emb_buffy0201r2 Show the field ONLY if: [ds_buffy_rem_01r1(1)] = '1'	01 {ds_buffy_label_01r2} {ds_buffy_usedt_01r2} {ds_buffy_stat_01r2} {ds_buffy_study_01r2} {ds_buffy_vol_01r2} {ds_buffy_rem_01r2}	descriptive
664	emb_buffy0201r3 Show the field ONLY if: [ds_buffy_rem_01r2(1)] = '1'	01 {ds_buffy_label_01r3} {ds_buffy_usedt_01r3} {ds_buffy_stat_01r3} {ds_buffy_study_01r3} {ds_buffy_vol_01r3} {ds_buffy_rem_01r3}	descriptive
665	biological_sample_complete	Section Header: <i>Form Status</i> Complete?	dropdown 0 Incomplete 1 Unverified 2 Complete

Instrument: **Comorbidities** (comorbidities)

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666	brk_0202 Show the field ONLY if: [groupenrol] = '4'	Section Header: <i>COMORBIDITIES</i> THIS SECTION IS NOT APPLICABLE for KD patients (prior to Jan 01, 2020)	descriptive
667	brk0301 Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'		descriptive
668	heart Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Section Header: <i>SECTION: PRE-EXISTING CONDITIONS/COMORBIDITIES</i> only applicable for patients diagnosed on or after Jan 01, 2020 Heart failure	yesno 1 Yes 0 No
669	hearttyp Show the field ONLY if: [heart] = '1'	if Heart failure, NYHA	radio 1 NYHA I 2 NYHA II 3 NYHA III 4 NYHA IV
670	congenital Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Congenital heart disease	yesno 1 Yes 0 No
671	phtn Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Pulmonary hypertension	yesno 1 Yes 0 No
672	lung Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Chronic lung disease	yesno 1 Yes 0 No
673	asthma Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Asthma	yesno 1 Yes 0 No

674	diab Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Type I/II diabetes	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
675	htn Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Systemic hypertension	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
676	ckd Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Chronic kidney disease	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
677	malignancy Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Malignancy	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
678	malignancytx Show the field ONLY if: [malignancy] = '1'	Please specify the malignancy/malignancies	notes				
679	allergies Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Allergies	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
680	allergiestx Show the field ONLY if: [allergies] = '1'	Please specify allergies	notes				
681	othercomor Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Other comorbidities	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
682	comortx Show the field ONLY if: [othercomor] = '1'	Please specify other comorbidities	notes				
683	ds_deaf Show the field ONLY if: ([groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3') and [ds_substudy] = '1'	Deafness or serious difficulty hearing	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
684	ds_blind Show the field ONLY if: ([groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3') and [ds_substudy] = '1'	Blindness or serious eyesight impairment	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
685	ds_vape Show the field ONLY if: ([groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3') and [ds_substudy] = '1'	Vaping use documented in the EHR	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
686	ds_nicotine Show the field ONLY if: ([groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3') and [ds_substudy] = '1'	Nicotine use documented in the EHR	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
687	ds_alcohol Show the field ONLY if: ([groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3') and [ds_substudy] = '1'	Alcohol use document in the EHR	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
688	ds_cvs Show the field ONLY if: ([groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3') and [ds_substudy] = '1'	Cardiovascular disease	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						

689	ds_immuno Show the field ONLY if: ([groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3') and [ds_substudy] = '1'	Immunosuppressive condition	yesno 1 Yes 0 No
690	ds_mental Show the field ONLY if: ([groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3') and [ds_substudy] = '1'	Serious mental illness documented in the EHR	yesno 1 Yes 0 No
691	ds_sickle Show the field ONLY if: ([groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3') and [ds_substudy] = '1'	Sickle cell disease	yesno 1 Yes 0 No
692	ds_pregnant Show the field ONLY if: ([groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3') and [ds_substudy] = '1'	Active pregnancy documented in the EHR	yesno 1 Yes 0 No
693	ds_gestwk Show the field ONLY if: [ds_substudy] = '1'	Section Header: SECTION: PREVIOUS MEDICAL & FAMILY HISTORY <i>only applicable for PrevailKids study</i> Gestational age at birth (weeks) <i>If "term pregnancy", please enter "40" weeks</i>	text (integer)
694	ds_prevkd Show the field ONLY if: [ds_substudy] = '1'	Previous history of KD	yesno 1 Yes 0 No
695	ds_prevkddt Show the field ONLY if: [ds_prevkd] = '1'	Date of initial KD episode <i>Estimate date if no documentation available</i>	text (date_ymd) Field Annotation: @HIDEBUTTON
696	ds_familykd Show the field ONLY if: [ds_substudy] = '1'	Family history of KD	radio 0 None 1 1st degree relative 2 2nd degree relative 3 Unknown
697	ds_familyimm Show the field ONLY if: [ds_substudy] = '1'	Notable family history of allergic or autoimmune disease	radio 0 No 1 Yes 2 Unknown
698	ds_familyimmtx Show the field ONLY if: [ds_familyimm]='1'	Relevant comments on family history of allergic or autoimmune disease	notes
699	comorbidities_complete	Section Header: Form Status Complete?	dropdown 0 Incomplete 1 Unverified 2 Complete

Instrument: **Clinical findings** (clinical\_findings)

[^ Collapse](#)

700	sympdt	Section Header: CLINICAL FINDINGS Date of symptoms onset <i>Put "15" for the day field if you only have YYYY-MMe.g., 2010-05-15 and then select "estimated" below</i>	text (date_ymd) Field Annotation: @HIDEBUTTON
701	sympdt_typ	Date of symptoms onset	radio 1 Estimated date 2 Accurate date
702	deferdt	Date of defervescence	text (date_ymd) Field Annotation: @HIDEBUTTON
703	conjunc	Section Header: SECTION: SYMPTOMS & SIGNS Conjunctivitis	yesno 1 Yes 0 No  Custom alignment: RH

704	cervical	Cervical lymphadenopathy	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No		
1	Yes								
0	No								
705	rash	Rash	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No		
1	Yes								
0	No								
706	extrem	Erythema/ edema of extremities	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No		
1	Yes								
0	No								
707	oralmuco	Mucosal changes	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No		
1	Yes								
0	No								
708	criteria	Number of KD symptoms other than fever	calc Calculation: $\text{if}([\text{conjunc}] = 1, 1, \text{if}([\text{conjunc}] = 0, 0, \text{"NaN"})) + \text{if}([\text{cervical}] = 1, 1, \text{if}([\text{cervical}] = 0, 0, \text{"NaN"})) + \text{if}([\text{oralmuco}] = 1, 1, \text{if}([\text{oralmuco}] = 0, 0, \text{"NaN"})) + \text{if}([\text{rash}] = 1, 1, \text{if}([\text{rash}] = 0, 0, \text{"NaN"})) + \text{if}([\text{extrem}] = 1, 1, \text{if}([\text{extrem}] = 0, 0, \text{"NaN"}))$						
709	type	Type of KD <i>Complete KD: fever &gt;=5 days with &gt;=4 other KD symptoms; otherwise incomplete KD</i>	radio <table border="1"> <tr><td>1</td><td>Complete KD</td></tr> <tr><td>2</td><td>Incomplete KD</td></tr> <tr><td>3</td><td>Does not meet KD criteria</td></tr> </table>	1	Complete KD	2	Incomplete KD	3	Does not meet KD criteria
1	Complete KD								
2	Incomplete KD								
3	Does not meet KD criteria								
710	cough	Cough	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No		
1	Yes								
0	No								
711	sorethroat	Sore throat	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No		
1	Yes								
0	No								
712	rhinorrhea	Congestion/rhinorrhea	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No		
1	Yes								
0	No								
713	abdpain	Abdominal pain	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No		
1	Yes								
0	No								
714	diarr	Diarrhea	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No		
1	Yes								
0	No								
715	vomit	Vomiting	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No		
1	Yes								
0	No								

716	arthritis	Arthritis	yesno 1 Yes 0 No Custom alignment: RH
717	myalgias	Myalgia	yesno 1 Yes 0 No Custom alignment: RH
718	missedkd	Missed KD <i>Missed KD = patients given an alternative diagnosis by a medical professional while showing at least 2 signs of KD and fever</i>	yesno 1 Yes 0 No Custom alignment: RH
719	brk_0201 Show the field ONLY if: [groupenrol] = '4'	Section Header: SECTION: SYMPTOMS & SIGNS <i>only applicable for patients diagnosed on or after Jan 01, 2020</i> THIS SECTION IS NOT APPLICABLE for KD patients (prior to Jan 01, 2020)	descriptive
720	chills Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Chills	yesno 1 Yes 0 No Custom alignment: RH
721	dyspnea Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Dyspnea or shortness of breath	yesno 1 Yes 0 No Custom alignment: RH
722	anosmia Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Anosmia	yesno 1 Yes 0 No Custom alignment: RH
723	headache Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Headache	yesno 1 Yes 0 No Custom alignment: RH
724	confusion Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Altered mental status/confusion	yesno 1 Yes 0 No Custom alignment: RH
725	syncope Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Syncope	yesno 1 Yes 0 No Custom alignment: RH
726	shock Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Shock	yesno 1 Yes 0 No Custom alignment: RH
727	polyrash Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Polymorphous rash	yesno 1 Yes 0 No Custom alignment: RH

728	othercut Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Other cutaneous symptoms(areas of erythema-edema with vesicles or pustules lesions seen on toes, soles, fingers, extremities, and/or heel)	yesno 1 Yes 0 No Custom alignment: RH
729	othercuttx Show the field ONLY if: [othercut] = '1'	If other cutaneous symptoms, please specify	notes
730	irritability Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Irritability	yesno 1 Yes 0 No Custom alignment: RH
731	anorexia Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Anorexia	yesno 1 Yes 0 No Custom alignment: RH
732	lethargy Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Lethargy	yesno 1 Yes 0 No Custom alignment: RH
733	otitis Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Otitis media	yesno 1 Yes 0 No Custom alignment: RH
734	hepato Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Jaundice, splenomegaly or hepatomegaly	yesno 1 Yes 0 No Custom alignment: RH
735	ds_hospinf Show the field ONLY if: [ds_substudy]='1'	Infections at or during hospital admission (other than COVID)	yesno 1 Yes 0 No Custom alignment: RH
736	dsc_hospinf1 code Show the field ONLY if: [ds_hospinf]='1'	ICD-10 code of Infection at or during admission #1	text BIOPORTAL:ICD10CM BIOPORTAL:ICD10CM Custom alignment: RH
737	dsc_hospinf1 code_2 Show the field ONLY if: [ds_hospinf]='1'	ICD-10 code of Infection at or during admission #2	text BIOPORTAL:ICD10CM BIOPORTAL:ICD10CM Custom alignment: RH
738	dsc_hospinf1 code_3 Show the field ONLY if: [ds_hospinf]='1'	ICD-10 code of Infection at or during admission #3	text BIOPORTAL:ICD10CM BIOPORTAL:ICD10CM Custom alignment: RH
739	clinical_findings_complete	Section Header: <i>Form Status</i> Complete?	dropdown 0 Incomplete 1 Unverified 2 Complete
Instrument: <b>Clinical course</b> (clinical_course) <span style="float: right;"><a href="#">^ Collapse</a></span>			
740	brk4	Section Header: <i>INPATIENT CLINICAL COURSE</i>	descriptive
741	admdt	Section Header: <i>SECTION: ADMISSION INFORMATION</i> Date of hospital admission	text (date_ymd) Field Annotation: @HIDEBUTTON
742	dcdt	Date of primary hospital discharge	text (date_ymd) Field Annotation: @HIDEBUTTON

743	los	Length of initial hospital stay (days) <i>autocalculated</i>	calc Calculation: round(datediff([admdt],[dcdt],"d","ymd",true),3)+1 Field Annotation: @FIELD-NOTES-DISPLAY = "tooltip"
744	readm Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Was patient acutely readmitted?	yesno 1 Yes 0 No
745	admdt2 Show the field ONLY if: [readm] = '1'	Date of hospital readmission	text (date_ymd) Field Annotation: @HIDEBUTTON
746	dcdt2 Show the field ONLY if: [readm] = '1'	Date of hospital discharge after readmission	text (date_ymd) Field Annotation: @HIDEBUTTON
747	lostotal	Length of total hospital stay (days) <i>autocalculated</i>	calc Calculation: round(sum(datediff([admdt],[dcdt],"d","ymd",true)+1,datediff([admdt2],[dcdt2],"d","ymd",true)+1),3)
748	brk0501 Show the field ONLY if: [groupenrol] = '4'	Section Header: SECTION: ICU STAYonly applicable for patients diagnosed on or after Jan 01, 2020 THIS SECTION IS NOT APPLICABLEfor patients enrolled before Jan 01, 2020	descriptive
749	icu	ICU stay	yesno 1 Yes 0 No
750	icuin1 Show the field ONLY if: [icu] = '1'	ICU stay 1: date of admission	text (date_ymd) Field Annotation: @HIDEBUTTON
751	icout1 Show the field ONLY if: [icu] = '1'	ICU stay 1: date of discharge	text (date_ymd) Field Annotation: @HIDEBUTTON
752	icuin2 Show the field ONLY if: [icout1] <> "	ICU stay 2: date of admission	text (date_ymd) Field Annotation: @HIDEBUTTON
753	icout2 Show the field ONLY if: [icuin2] <> "	ICU stay 2: date of discharge	text (date_ymd) Field Annotation: @HIDEBUTTON
754	iculos	ICU length of stay (autocalculated)	calc Calculation: sum(datediff([icuin1],[icout1],"d","ymd",true),datediff([icuin2],[icout2],"d","ymd",true))
755	brk0502 Show the field ONLY if: [groupenrol] = '4'	Section Header: SECTION: RESPIRATORY COMPLICATIONSOnly applicable for Covid & PIMS groups THIS SECTION IS NOT APPLICABLEfor patients enrolled before Jan 01, 2020	descriptive
756	respdysfun Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Respiratory dysfunction <i>None: O2 saturation &gt;93% on room air, no dyspnea nor abnormal CXR)Moderate: tachypnea, +/- hypoxia O2 sat &lt; =93% on room air and abnormal chest imaging with less than 50% involvement of the lung parenchyma)Severe: tachypnea, with hypoxia O2 saturation &lt; =93 percent on room air or PaO2/FIO2 &lt; 300 mmHg and more than 50% involvement of the lung parenchyma on chest imaging</i>	radio 1 None 3 Moderate 4 Severe
757	respsupp Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Respiratory support	yesno 1 Yes 0 No
758	respsupp_typ Show the field ONLY if: [respsupp] = '1'	Type of respiratory support(check all that apply)	checkbox 1 respsupp_typ__1 NC 2 respsupp_typ__2 HFNC 3 respsupp_typ__3 NIV 4 respsupp_typ__4 Intubated/mechanical ventilation 5 respsupp_typ__5 Inhaled nitric oxide 6 respsupp_typ__6 ECMO (VV or VA)



759	cxr Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	CXR (performed on or shortly after the day of SARS CoV 2 infection diagnostic testing)	radio 1 Not done 2 Normal 3 Abnormal
760	cxr_typ Show the field ONLY if: [cxr] = '3'	CXR findings(check all that apply)	checkbox 1 cxr_typ__1 Lobar consolidation 2 cxr_typ__2 Multifocal or patchy opacity 3 cxr_typ__3 Interstitial abnormalities 4 cxr_typ__4 Other (please specify below)
761	cxr_tx Show the field ONLY if: [cxr_typ(4)] = '1'	if other CXR findings, please specify	notes
762	cct Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	CT chest (performed on or shortly after the day of SARS CoV 2 infection diagnostic testing)	radio 1 Not done 2 Normal 3 Abnormal
763	cct_typ Show the field ONLY if: [cct] = '3'	CT chest findings(check all that apply)	checkbox 1 cct_typ__1 Lobar consolidation 2 cct_typ__2 Multifocal or patchy opacity or ground glass 3 cct_typ__3 Interstitial abnormalities 4 cct_typ__4 Other (please specify below)
764	cct_tx Show the field ONLY if: [cct_typ(4)] = '1'	if other CT chest findings, please specify	notes
765	brk0503 Show the field ONLY if: [groupenrol] = '4'	Section Header: SECTION: OTHER COMPLICATIONS only applicable for patients diagnosed on or after Jan 01, 2020 THIS SECTION IS NOT APPLICABLE for patients enrolled before Jan 01, 2020	descriptive
766	card Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Cardiac arrest/need for cardiopulmonary resuscitation	yesno 1 Yes 0 No
767	arr Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Arrhythmia	yesno 1 Yes 0 No
768	arrtyp Show the field ONLY if: [arr] = '1'	Arrhythmia type(check all that apply)	checkbox 1 arrtyp__1 First degree AV block 2 arrtyp__2 Second degree AV block 3 arrtyp__3 Third degree AV block 4 arrtyp__4 Ventricular tachycardia 5 arrtyp__5 Premature ventricular contractions 6 arrtyp__6 Supraventricular tachycardia 7 arrtyp__7 Premature atrial contractions
769	renal Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Acute kidney injury/need for continuous renal replacement therapy	yesno 1 Yes 0 No
770	cvd Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Cerebrovascular stroke	yesno 1 Yes 0 No
771	pe Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Pulmonary embolism	yesno 1 Yes 0 No

772	dvt Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	DVT	yesno 1 Yes 0 No
773	othercomplics Show the field ONLY if: [groupenrol] = '1' or [groupenrol] = '2' or [groupenrol] = '3'	Other complications	yesno 1 Yes 0 No
774	othercomplicstx Show the field ONLY if: [othercomplics] = '1'	if other complications, please specify	notes
775	clinical_course_complete	Section Header: <i>Form Status</i> Complete?	dropdown 0 Incomplete 1 Unverified 2 Complete

Instrument: **Serial labwork** (serial\_labwork)[^ Collapse](#)

776	brk_0302	Section Header: <i>SERIAL LABWORK</i> REDCap DATA ENTRY NOT REQUIRED Site to upload "Serial labwork" datafile (.csv file) in the "file upload" section DCC to import serial labwork data (received from the sites) into REDCap if data not available for any labwork (or labwork not done), leave blank	descriptive
777	sr_labdt	Lab sample collection date/time	text (datetime_ymd) Field Annotation: @HIDEBUTTON
778	sr_labinterval Show the field ONLY if: [sr_labdt] <> "	Labwork interval (days) <i>autocalculated</i>	calc Calculation: round(datediff([admdt], [sr_labdt], "d", "ymd", true), 3)
779	sr_wbc Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Complete blood count (CBC)</i> White blood cells (RBC) (x109/L)	text (number) Field Annotation: @PLACEHOLDER = '10e9/L'
780	sr_rbc Show the field ONLY if: [sr_labdt] <> "	Red blood cells (RBC) (x1012/L)	text (number, Min: 2.93, Max: 5.74) Field Annotation: @PLACEHOLDER = '10e12/L'
781	sr_hgb Show the field ONLY if: [sr_labdt] <> "	Hemoglobin (g/L)	text (number, Min: 89, Max: 200) Field Annotation: @PLACEHOLDER = 'g/L'
782	sr_hgb_us Show the field ONLY if: [sr_labdt] <> "	Hemoglobin (g/dL)	text (number) Field Annotation: @PLACEHOLDER = 'g/dL'
783	sr_hct Show the field ONLY if: [sr_labdt] <> "	Hematocrit (%)	text (number) Field Annotation: @PLACEHOLDER = '%'
784	sr_hct_us Show the field ONLY if: [sr_labdt] <> "	Hematocrit (x.xxx)	text (number) Field Annotation: @PLACEHOLDER = 'x.xxx'
785	sr_plt Show the field ONLY if: [sr_labdt] <> "	Platelets (x109/L)	text (number) Field Annotation: @PLACEHOLDER = '10e9/L'
786	sr_neutr Show the field ONLY if: [sr_labdt] <> "	Neutrophils (x109/L)	text (number) Field Annotation: @PLACEHOLDER = '10e9/L'
787	sr_lymph Show the field ONLY if: [sr_labdt] <> "	Lymphocytes (x109/L)	text (number) Field Annotation: @PLACEHOLDER = '10e9/L'
788	sr_mono Show the field ONLY if: [sr_labdt] <> "	Macrophages (x109/L)	text (number) Field Annotation: @PLACEHOLDER = '10e9/L'
789	sr_eosino Show the field ONLY if: [sr_labdt] <> "	Eosinophils (x109/L)	text (number) Field Annotation: @PLACEHOLDER = '10e9/L'

790	sr_eosino_us Show the field ONLY if: [sr_labdt] <> "	Eosinophils (x103/uL)	text (number) Field Annotation: @PLACEHOLDER = '10e3/uL'
791	sr_baso Show the field ONLY if: [sr_labdt] <> "	Basophils (x109/L)	text (number) Field Annotation: @PLACEHOLDER = '10e9/L'
792	sr_esr Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Inflammatory markers</i> ESR (mm/hr)	text (number, Min: 2, Max: 37) Field Annotation: @PLACEHOLDER = 'mm/hr'
793	sr_alb Show the field ONLY if: [sr_labdt] <> "	Albumin (g/L)	text (number, Min: 23, Max: 53) Field Annotation: @PLACEHOLDER = 'g/L'
794	sr_alb_us Show the field ONLY if: [sr_labdt] <> "	Albumin (g/dL)	text (number) Field Annotation: @PLACEHOLDER = 'g/dL'
795	sr_regrcp Show the field ONLY if: [sr_labdt] <> "	C-reactive protein (CRP) (mg/L)	text (number, Min: 0.1, Max: 6.1) Field Annotation: @PLACEHOLDER = 'mg/L'
796	sr_hscrp Show the field ONLY if: [sr_labdt] <> "	High sensitivity C-reactive protein (hS CRP) (mg/L)	text (number, Min: 0.1, Max: 6.1) Field Annotation: @PLACEHOLDER = 'mg/L'
797	sr_regrcp_us Show the field ONLY if: [sr_labdt] <> "	C-reactive protein (CRP) (mg/dL)	text (number) Field Annotation: @PLACEHOLDER = 'mg/dL'
798	sr_hscrp_us Show the field ONLY if: [sr_labdt] <> "	High sensitivity C-reactive protein (hS CRP) (mg/dL)	text (number) Field Annotation: @PLACEHOLDER = 'mg/dL'
799	sr_ferr Show the field ONLY if: [sr_labdt] <> "	Ferritin (ug/L)	text (number) Field Annotation: @PLACEHOLDER = 'ug/L'
800	sr_ferr_us Show the field ONLY if: [sr_labdt] <> "	Ferritin (ng/mL)	text (number) Field Annotation: @PLACEHOLDER = 'ng/mL'
801	sr_trig Show the field ONLY if: [sr_labdt] <> "	Triglycerides (mmol/L)	text (number) Field Annotation: @PLACEHOLDER = 'mmol/L'
802	sr_trig_us Show the field ONLY if: [sr_labdt] <> "	Triglycerides (mg/dL)	text (number) Field Annotation: @PLACEHOLDER = 'mg/dL'
803	sr_amylase Show the field ONLY if: [sr_labdt] <> "	Amylase (U/L)	text (number) Field Annotation: @PLACEHOLDER = 'U/L'
804	sr_ldh Show the field ONLY if: [sr_labdt] <> "	LDH (U/L)	text (number) Field Annotation: @PLACEHOLDER = 'U/L'
805	sr_alt Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Liver function tests</i> ALT (U/L)	text (number) Field Annotation: @PLACEHOLDER = 'U/L'
806	sr_ast Show the field ONLY if: [sr_labdt] <> "	AST (U/L)	text (number) Field Annotation: @PLACEHOLDER = 'U/L'
807	sr_ggt Show the field ONLY if: [sr_labdt] <> "	GGT (U/L)	text (number) Field Annotation: @PLACEHOLDER = 'U/L'
808	sr_alp Show the field ONLY if: [sr_labdt] <> "	Alkaline phosphatase (U/L)	text (number) Field Annotation: @PLACEHOLDER = 'U/L'

809	sr_totbili Show the field ONLY if: [sr_labdt] <> "	Total bilirubin (umol/L)	text (number) Field Annotation: @PLACEHOLDER = 'umol/L'
810	sr_totbili_us Show the field ONLY if: [sr_labdt] <> "	Total bilirubin (mg/dL)	text (number) Field Annotation: @PLACEHOLDER = 'mg/dL'
811	sr_directbili Show the field ONLY if: [sr_labdt] <> "	Direct bilirubin (umol/L)	text (number) Field Annotation: @PLACEHOLDER = 'umol/L'
812	sr_directbili_us Show the field ONLY if: [sr_labdt] <> "	Direct bilirubin (mg/dL)	text (number) Field Annotation: @PLACEHOLDER = 'mg/dL'
813	sr_sod Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Renal function</i> Sodium (mmol/L)	text (number) Field Annotation: @PLACEHOLDER = 'mmol/L'
814	sr_pota Show the field ONLY if: [sr_labdt] <> "	Potassium (mmol/L)	text (number) Field Annotation: @PLACEHOLDER = 'mmol/L'
815	sr_chlor Show the field ONLY if: [sr_labdt] <> "	Chloride (mmol/L)	text (number) Field Annotation: @PLACEHOLDER = 'mmol/L'
816	sr_bicarb Show the field ONLY if: [sr_labdt] <> "	Bicarbonate (mmol/L)	text (number) Field Annotation: @PLACEHOLDER = 'mmol/L'
817	sr_bun Show the field ONLY if: [sr_labdt] <> "	BUN (mmol/L)	text (number) Field Annotation: @PLACEHOLDER = 'mmol/L'
818	sr_bun_us Show the field ONLY if: [sr_labdt] <> "	BUN (mg/dL)	text (number) Field Annotation: @PLACEHOLDER = 'mg/dL'
819	sr_creat Show the field ONLY if: [sr_labdt] <> "	S.Creatinine (umol/L)	text (number) Field Annotation: @PLACEHOLDER = 'umol/L'
820	sr_creat_us Show the field ONLY if: [sr_labdt] <> "	S.Creatinine (mg/dL)	text (number) Field Annotation: @PLACEHOLDER = 'mg/dL'
821	sr_proteinuria Show the field ONLY if: [sr_labdt] <> "	Urine analysis: proteinuria (mg/L) <i>if negative: put "0"; if trace: put "20" mg/dL for dipstick method, please refer to the picture below</i>	text (number) Field Annotation: @PLACEHOLDER = 'mg/L'
822	sr_proteinuria_us Show the field ONLY if: [sr_labdt] <> "	Urine analysis: proteinuria (mg/dL) <i>if negative: put "0"; if trace: put "20" mg/dL for dipstick method, please refer to the picture below</i>	text (number) Field Annotation: @PLACEHOLDER = 'mg/dL'
823	sr_inr Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Coagulation</i> INR	text (number)
824	sr_pt Show the field ONLY if: [sr_labdt] <> "	PT (sec)	text (number) Field Annotation: @PLACEHOLDER = 'sec'
825	sr_ptt Show the field ONLY if: [sr_labdt] <> "	PTT (sec)	text (number) Field Annotation: @PLACEHOLDER = 'sec'
826	sr_ddimer Show the field ONLY if: [sr_labdt] <> "	D-dimer (ug/mL DDU)	text (number) Field Annotation: @PLACEHOLDER = 'ug/mL DDU'
827	sr_ddimer_us Show the field ONLY if: [sr_labdt] <> "	D-dimer (ug FEU /mL)	text (number) Field Annotation: @PLACEHOLDER = 'ug FEU /mL'

828	sr_fibrin Show the field ONLY if: [sr_labdt] <> "	Fibrinogen (g/L)	text (number) Field Annotation: @PLACEHOLDER = 'g/L'
829	sr_fibrin_us Show the field ONLY if: [sr_labdt] <> "	Fibrinogen (mg/dL)	text (number) Field Annotation: @PLACEHOLDER = 'mg/dL'
830	sr_tropt Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Cardiac biomarkers</i> if Troponin values are reported as "< 10" ug/L, <i>please put "9.99"</i> Troponin-T (ng/L)	text (number) Field Annotation: @PLACEHOLDER = 'ng/L'
831	sr_hstropt Show the field ONLY if: [sr_labdt] <> "	High sensitivity troponin-T (ng/L)	text (number) Field Annotation: @PLACEHOLDER = 'ng/L'
832	sr_hstropt_us Show the field ONLY if: [sr_labdt] <> "	High sensitivity troponin-T (ng/mL)	text (number) Field Annotation: @PLACEHOLDER = 'ng/mL'
833	sr_tropi Show the field ONLY if: [sr_labdt] <> "	Troponin-I (ng/L)	text (number) Field Annotation: @PLACEHOLDER = 'ng/L'
834	sr_hstropi Show the field ONLY if: [sr_labdt] <> "	High sensitivity Troponin-I (ng/L)	text (number) Field Annotation: @PLACEHOLDER = 'ng/L'
835	sr_probnp Show the field ONLY if: [sr_labdt] <> "	Pro-BNP (ng/L)	text (number) Field Annotation: @PLACEHOLDER = 'ng/L'
836	sr_bnp Show the field ONLY if: [sr_labdt] <> "	BNP (ng/L)	text (number) Field Annotation: @PLACEHOLDER = 'ng/L'
837	sr_procalc Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Other</i> Procalcitonin (ug/L)	text (number) Field Annotation: @PLACEHOLDER = 'ug/L'
838	sr_il1 Show the field ONLY if: [sr_labdt] <> "	IL-1 (pg/mL)	text (number) Field Annotation: @PLACEHOLDER = 'pg/mL'
839	sr_il1ra Show the field ONLY if: [sr_labdt] <> "	IL-1Ra (pg/mL)	text (number) Field Annotation: @PLACEHOLDER = 'pg/mL'
840	sr_il6 Show the field ONLY if: [sr_labdt] <> "	IL-6 (pg/mL)	text (number) Field Annotation: @PLACEHOLDER = 'pg/mL'
841	sr_wbc_calc Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>CALCULATED UNITS Complete blood count (CBC)</i> White blood cells (RBC) (x109/L)	calc Calculation: if([sr_wbc] != "", [sr_wbc], "") Field Annotation: @PLACEHOLDER = '10e9/L'
842	sr_rbc_calc Show the field ONLY if: [sr_labdt] <> "	Red blood cells (RBC) (x1012/L)	calc Calculation: if([sr_rbc] != "", [sr_rbc], "") Field Annotation: @PLACEHOLDER = '10e12/L'
843	sr_hgb_calc Show the field ONLY if: [sr_labdt] <> "	Hemoglobin (g/L)	calc Calculation: round(if([sr_hgb] != "", [sr_hgb], [sr_hgb_us]*10),3) Field Annotation: @PLACEHOLDER = 'g/L'
844	sr_hct_calc Show the field ONLY if: [sr_labdt] <> "	Hematocrit (%)	calc Calculation: if([sr_hct] != "", [sr_hct], [sr_hct_us]*100) Field Annotation: @PLACEHOLDER = '%'
845	sr_plt_calc Show the field ONLY if: [sr_labdt] <> "	Platelets (x109/L)	calc Calculation: if([sr_plt] != "", [sr_plt], "") Field Annotation: @PLACEHOLDER = '10e9/L'
846	sr_neutr_calc Show the field ONLY if: [sr_labdt] <> "	Neutrophils (x109/L)	calc Calculation: if([sr_neutr] != "", [sr_neutr], "") Field Annotation: @PLACEHOLDER = '10e9/L'

847	sr_lymph_calc Show the field ONLY if: [sr_labdt] <> "	Lymphocytes (x109/L)	calc Calculation: if([sr_lymph] != "", [sr_lymph], "") Field Annotation: @PLACEHOLDER = '10e9/L'
848	sr_mono_calc Show the field ONLY if: [sr_labdt] <> "	Macrophages (x109/L)	calc Calculation: if([sr_mono] != "", [sr_mono], "") Field Annotation: @PLACEHOLDER = '10e9/L'
849	sr_eosino_calc Show the field ONLY if: [sr_labdt] <> "	Eosinophils (x109/L)	calc Calculation: round(if([sr_eosino] != "", [sr_eosino], [sr_eosino_us]),3) Field Annotation: @PLACEHOLDER = '10e9/L'
850	sr_baso_calc Show the field ONLY if: [sr_labdt] <> "	Basophils (x109/L)	calc Calculation: if([sr_baso] != "", [sr_baso], "") Field Annotation: @PLACEHOLDER = '10e9/L'
851	sr_esr_calc Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Inflammatory markers</i> ESR (mm/hr)	calc Calculation: if([sr_esr] != "", [sr_esr], "") Field Annotation: @PLACEHOLDER = 'mm/hr'
852	sr_alb_calc Show the field ONLY if: [sr_labdt] <> "	Albumin (g/L)	calc Calculation: round(if([sr_alb] != "", [sr_alb], [sr_alb_us]*10),3) Field Annotation: @PLACEHOLDER = 'g/L'
853	sr_regcrp_calc Show the field ONLY if: [sr_labdt] <> "	C-reactive protein (CRP) (mg/L)	calc Calculation: if([sr_regcrp] != "", [sr_regcrp], [sr_regcrp_us]*10) Field Annotation: @PLACEHOLDER = 'mg/L'
854	sr_hscrp_calc Show the field ONLY if: [sr_labdt] <> "	High sensitivity C-reactive protein (hs CRP) (mg/L)	calc Calculation: if([sr_hscrp] != "", [sr_hscrp], [sr_hscrp_us]*10) Field Annotation: @PLACEHOLDER = 'mg/L'
855	sr_ferr_calc Show the field ONLY if: [sr_labdt] <> "	Ferritin (ug/L)	calc Calculation: if([sr_ferr] != "", [sr_ferr], [sr_ferr_us]) Field Annotation: @PLACEHOLDER = 'ug/L'
856	sr_trig_calc Show the field ONLY if: [sr_labdt] <> "	Triglycerides (mmol/L)	calc Calculation: round(if([sr_trig] != "", [sr_trig], [sr_trig_us]*0.0113),3) Field Annotation: @PLACEHOLDER = 'mmol/L'
857	sr_amylase_calc Show the field ONLY if: [sr_labdt] <> "	Amylase (U/L)	calc Calculation: if([sr_amylase] != "", [sr_amylase], "") Field Annotation: @PLACEHOLDER = 'U/L'
858	sr_ldh_calc Show the field ONLY if: [sr_labdt] <> "	LDH (U/L)	calc Calculation: if([sr_ldh] != "", [sr_ldh], "") Field Annotation: @PLACEHOLDER = 'U/L'
859	sr_alt_calc Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Liver function tests</i> ALT (U/L)	calc Calculation: if([sr_alt] != "", [sr_alt], "") Field Annotation: @PLACEHOLDER = 'U/L'
860	sr_ast_calc Show the field ONLY if: [sr_labdt] <> "	AST (U/L)	calc Calculation: if([sr_ast] != "", [sr_ast], "") Field Annotation: @PLACEHOLDER = 'U/L'
861	sr_ggt_calc Show the field ONLY if: [sr_labdt] <> "	GGT (U/L)	calc Calculation: if([sr_ggt] != "", [sr_ggt], "") Field Annotation: @PLACEHOLDER = 'U/L'
862	sr_alp_calc Show the field ONLY if: [sr_labdt] <> "	Alkaline phosphatase (U/L)	calc Calculation: if([sr_alp] != "", [sr_alp], "") Field Annotation: @PLACEHOLDER = 'U/L'
863	sr_totbili_calc Show the field ONLY if: [sr_labdt] <> "	Total bilirubin (umol/L)	calc Calculation: round(if([sr_totbili] != "", [sr_totbili], [sr_totbili_us]*17.1),3) Field Annotation: @PLACEHOLDER = 'umol/L'
864	sr_directbili_calc Show the field ONLY if: [sr_labdt] <> "	Direct bilirubin (umol/L)	calc Calculation: round(if([sr_directbili] != "", [sr_directbili], [sr_directbili_us]*17.1),3) Field Annotation: @PLACEHOLDER = 'umol/L'
865	sr_sod_calc Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Renal function</i> Sodium (mmol/L)	calc Calculation: if([sr_sod] != "", [sr_sod], "") Field Annotation: @PLACEHOLDER = 'mmol/L'

866	sr_pota_calc Show the field ONLY if: [sr_labdt] <> "	Potassium (mmol/L)	calc Calculation: if([sr_pota] != "", [sr_pota], "") Field Annotation: @PLACEHOLDER = 'mmol/L'
867	sr_chlor_calc Show the field ONLY if: [sr_labdt] <> "	Chloride (mmol/L)	calc Calculation: if([sr_chlor] != "", [sr_chlor], "") Field Annotation: @PLACEHOLDER = 'mmol/L'
868	sr_bicarb_calc Show the field ONLY if: [sr_labdt] <> "	Bicarbonate (mmol/L)	calc Calculation: if([sr_bicarb] != "", [sr_bicarb], "") Field Annotation: @PLACEHOLDER = 'mmol/L'
869	sr_bun_calc Show the field ONLY if: [sr_labdt] <> "	BUN (mmol/L)	calc Calculation: round(if([sr_bun] != "", [sr_bun], [sr_bun_us]*0.357),3) Field Annotation: @PLACEHOLDER = 'mmol/L'
870	sr_creat_calc Show the field ONLY if: [sr_labdt] <> "	S.Creatinine (umol/L)	calc Calculation: round(if([sr_creat] != "", [sr_creat], [sr_creat_us]*88.4),3) Field Annotation: @PLACEHOLDER = 'umol/L'
871	sr_proteinuria_calc Show the field ONLY if: [sr_labdt] <> "	Urine analysis: proteinuria (mg/L) <i>if negative: put "0"; if trace: put "20" mg/dL for dipstick method, please refer to the picture below</i>	calc Calculation: if([sr_proteinuria] != "", [sr_proteinuria], [sr_proteinuria_us]*10) Field Annotation: @PLACEHOLDER = 'mg/L'
872	sr_inr_calc Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Coagulation</i> INR	calc Calculation: if([sr_inr] != "", [sr_inr], "")
873	sr_pt_calc Show the field ONLY if: [sr_labdt] <> "	PT (sec)	calc Calculation: if([sr_pt] != "", [sr_pt], "") Field Annotation: @PLACEHOLDER = 'sec'
874	sr_ptt_calc Show the field ONLY if: [sr_labdt] <> "	PTT (sec)	calc Calculation: if([sr_ptt] != "", [sr_ptt], "") Field Annotation: @PLACEHOLDER = 'sec'
875	sr_ddimer_calc Show the field ONLY if: [sr_labdt] <> "	D-dimer (ug/mL DDU)	calc Calculation: round(if([sr_ddimer] != "", [sr_ddimer], [sr_ddimer_us]*0.5),3) Field Annotation: @PLACEHOLDER = 'ug/mL DDU'
876	sr_fibrin_calc Show the field ONLY if: [sr_labdt] <> "	Fibrinogen (g/L)	calc Calculation: round(if([sr_fibrin] != "", [sr_fibrin], [sr_fibrin_us]/100),3) Field Annotation: @PLACEHOLDER = 'g/L'
877	sr_tropt_calc Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Cardiac biomarkers</i> Troponin-T (ng/L)	calc Calculation: if([sr_tropt] != "", [sr_tropt], "") Field Annotation: @PLACEHOLDER = 'ug/L'
878	sr_hstropt_calc Show the field ONLY if: [sr_labdt] <> "	High sensitivity troponin-T (ng/L)	calc Calculation: if([sr_hstropt] != "", [sr_hstropt], [sr_hstropt_us]*1000) Field Annotation: @PLACEHOLDER = 'ug/L'
879	sr_tropi_calc Show the field ONLY if: [sr_labdt] <> "	Troponin-I (ng/L)	calc Calculation: if([sr_tropi] != "", [sr_tropi], "") Field Annotation: @PLACEHOLDER = 'ug/L'
880	sr_hstropi_calc Show the field ONLY if: [sr_labdt] <> "	High sensitivity Troponin-I (ng/L)	calc Calculation: if([sr_hstropi] != "", [sr_hstropi], "") Field Annotation: @PLACEHOLDER = 'ug/L'
881	sr_probnp_calc Show the field ONLY if: [sr_labdt] <> "	Pro-BNP (ng/L)	calc Calculation: if([sr_probnp] != "", [sr_probnp], "") Field Annotation: @PLACEHOLDER = 'ng/L'
882	sr_bnp_calc Show the field ONLY if: [sr_labdt] <> "	BNP (ng/L)	calc Calculation: if([sr_bnp] != "", [sr_bnp], "") Field Annotation: @PLACEHOLDER = 'ng/L'
883	sr_procalc_calc Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Other</i> Procalcitonin (mcg/L)	calc Calculation: if([sr_procalc] != "", [sr_procalc], "") Field Annotation: @PLACEHOLDER = 'mcg/L'
884	sr_il1_calc Show the field ONLY if: [sr_labdt] <> "	IL-1 (pg/mL)	calc Calculation: if([sr_il1] != "", [sr_il1], "") Field Annotation: @PLACEHOLDER = 'pg/mL'

885	sr_il1ra_calc Show the field ONLY if: [sr_labdt] <> "	IL-1Ra (pg/mL)	calc Calculation: if([sr_il1ra] != "", [sr_il1ra], "") Field Annotation: @PLACEHOLDER = 'pg/mL'
886	sr_il6_calc Show the field ONLY if: [sr_labdt] <> "	IL-6 (pg/mL)	calc Calculation: if([sr_il6] != "", [sr_il6], "") Field Annotation: @PLACEHOLDER = 'pg/mL'
887	sr_desc_brk Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Complete blood count (CBC)</i> SI unit Alternate unit (SI unit)	descriptive
888	sr_wbc_embedbrk Show the field ONLY if: [sr_labdt] <> "	White blood cells {sr_wbc:icons} {sr_wbc_calc} (x109/L)	descriptive Field Annotation: embedded_field
889	sr_rbc_embedbrk Show the field ONLY if: [sr_labdt] <> "	Red blood cells {sr_rbc:icons} {sr_rbc_calc} (x1012/L)	descriptive Field Annotation: embedded_field
890	sr_hgb_embedbrk Show the field ONLY if: [sr_labdt] <> "	Hemoglobin {sr_hgb:icons} {sr_hgb_us:icons} {sr_hgb_calc} (g/L)	descriptive Field Annotation: embedded_field
891	sr_hct_embedbrk Show the field ONLY if: [sr_labdt] <> "	Hematocrit {sr_hct:icons} {sr_hct_us:icons} {sr_hct_calc} (%)	descriptive Field Annotation: embedded_field
892	sr_plt_embedbrk Show the field ONLY if: [sr_labdt] <> "	Platelets {sr_plt:icons} {sr_plt_calc} (x109/L)	descriptive Field Annotation: embedded_field
893	sr_neutr_embedbrk Show the field ONLY if: [sr_labdt] <> "	Neutrophils {sr_neutr:icons} {sr_neutr_calc} (x109/L)	descriptive Field Annotation: embedded_field
894	sr_lymph_embedbrk Show the field ONLY if: [sr_labdt] <> "	Lymphocytes {sr_lymph:icons} {sr_lymph_calc} (x109/L)	descriptive Field Annotation: embedded_field
895	sr_mono_embedbrk Show the field ONLY if: [sr_labdt] <> "	Macrophages {sr_mono:icons} {sr_mono_calc} (x109/L)	descriptive Field Annotation: embedded_field
896	sr_eosino_embedbrk Show the field ONLY if: [sr_labdt] <> "	Eosinophils {sr_eosino:icons} {sr_eosino_us:icons} {sr_eosino_calc} (x109/L)	descriptive Field Annotation: embedded_field
897	sr_baso_embedbrk Show the field ONLY if: [sr_labdt] <> "	Basophils {sr_baso:icons} {sr_baso_calc} (x109/L)	descriptive Field Annotation: embedded_field
898	sr_esr_embedbrk Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Inflammatory markers</i> ESR {sr_esr:icons} {sr_esr_calc} (mm/hr)	descriptive Field Annotation: embedded_field
899	sr_alb_embedbrk Show the field ONLY if: [sr_labdt] <> "	Albumin {sr_alb:icons} {sr_alb_us:icons} {sr_alb_calc} (g/L)	descriptive Field Annotation: embedded_field
900	sr_regcrp_embedbrk Show the field ONLY if: [sr_labdt] <> "	C-reactive protein {sr_regcrp:icons} {sr_regcrp_us:icons} {sr_regcrp_calc} (mg/L)	descriptive Field Annotation: embedded_field
901	sr_hscrp_embedbrk Show the field ONLY if: [sr_labdt] <> "	hS C-reactive protein {sr_hscrp:icons} {sr_hscrp_us:icons} {sr_hscrp_calc} (mg/L)	descriptive Field Annotation: embedded_field
902	sr_ferr_embedbrk Show the field ONLY if: [sr_labdt] <> "	Ferritin {sr_ferr:icons} {sr_ferr_us:icons} {sr_ferr_calc} (ug/L)	descriptive Field Annotation: embedded_field
903	sr_trig_embedbrk Show the field ONLY if: [sr_labdt] <> "	Triglycerides {sr_trig:icons} {sr_trig_us:icons} {sr_trig_calc} (umol/L)	descriptive Field Annotation: embedded_field



904	sr_amylase_embedbrk Show the field ONLY if: [sr_labdt] <> "	Amylase {sr_amylase:icons} {sr_amylase_calc} (U/L)	descriptive Field Annotation: embedded_field
905	sr_ldh_embedbrk Show the field ONLY if: [sr_labdt] <> "	LDH {sr_ldh:icons} {sr_ldh_calc} (U/L)	descriptive Field Annotation: embedded_field
906	sr_alt_embedbrk Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Liver function tests</i> ALT {sr_alt:icons} {sr_alt_calc} (U/L)	descriptive Field Annotation: embedded_field
907	sr_ast_embedbrk Show the field ONLY if: [sr_labdt] <> "	AST {sr_ast:icons} {sr_ast_calc} (U/L)	descriptive Field Annotation: embedded_field
908	sr_ggt_embedbrk Show the field ONLY if: [sr_labdt] <> "	GGT {sr_ggt:icons} {sr_ggt_calc} (U/L)	descriptive Field Annotation: embedded_field
909	sr_alp_embedbrk Show the field ONLY if: [sr_labdt] <> "	Alkaline phosphatase {sr_alp:icons} {sr_alp_calc} (U/L)	descriptive Field Annotation: embedded_field
910	sr_totbili_embedbrk Show the field ONLY if: [sr_labdt] <> "	Total bilirubin {sr_totbili:icons} {sr_totbili_us:icons} {sr_totbili_calc} (umol/L)	descriptive Field Annotation: embedded_field
911	sr_directbili_embedbrk Show the field ONLY if: [sr_labdt] <> "	Direct bilirubin {sr_directbili:icons} {sr_directbili_us:icons} {sr_directbili_calc} (umol/L)	descriptive Field Annotation: embedded_field
912	sr_sod_embedbrk Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Renal function</i> Sodium {sr_sod:icons} {sr_sod_calc} (mmol/L)	descriptive Field Annotation: embedded_field
913	sr_pota_embedbrk Show the field ONLY if: [sr_labdt] <> "	Potassium {sr_pota:icons} {sr_pota_calc} (mmol/L)	descriptive Field Annotation: embedded_field
914	sr_chlor_embedbrk Show the field ONLY if: [sr_labdt] <> "	Chloride {sr_chlor:icons} {sr_chlor_calc} (mmol/L)	descriptive Field Annotation: embedded_field
915	sr_bicarb_embedbrk Show the field ONLY if: [sr_labdt] <> "	Bicarbonate {sr_bicarb:icons} {sr_bicarb_calc} (mmol/L)	descriptive Field Annotation: embedded_field
916	sr_bun_embedbrk Show the field ONLY if: [sr_labdt] <> "	BUN (Urea Nitrogen) {sr_bun:icons} {sr_bun_us:icons} {sr_bun_calc} (mmol/L)	descriptive Field Annotation: embedded_field
917	sr_creat_embedbrk Show the field ONLY if: [sr_labdt] <> "	S.Creatinine {sr_creat:icons} {sr_creat_us:icons} {sr_creat_calc} (umol/L)	descriptive Field Annotation: embedded_field
918	sr_proteinuria_embedbrk Show the field ONLY if: [sr_labdt] <> "	Urine analysis: proteinuria {sr_proteinuria:icons} {sr_proteinuria_us:icons} {sr_proteinuria_calc} (mg/L) <i>if negative: put "0"; if trace: put "20" mg/dL for dipstick method, please refer to the picture below</i>	descriptive Field Annotation: embedded_field
919	brk0204 Show the field ONLY if: [sr_labdt] <> "		descriptive Field Annotation: embedded_field
920	sr_inr_embedbrk Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Coagulation</i> INR {sr_inr:icons} {sr_inr_calc} (g/L)	descriptive Field Annotation: embedded_field
921	sr_pt_embedbrk Show the field ONLY if: [sr_labdt] <> "	PT {sr_pt:icons} {sr_pt_calc} (sec)	descriptive Field Annotation: embedded_field
922	sr_ptt_embedbrk Show the field ONLY if: [sr_labdt] <> "	PTT {sr_ptt:icons} {sr_ptt_calc} (sec)	descriptive Field Annotation: embedded_field

923	sr_ddimer_embedbrk Show the field ONLY if: [sr_labdt] <> "	D-dimer {sr_ddimer:icons} {sr_ddimer_us:icons} {sr_ddimer_calc} (ug/mL DDU)	descriptive Field Annotation: embedded_field
924	sr_fibrin_embedbrk Show the field ONLY if: [sr_labdt] <> "	Fibrinogen {sr_fibrin:icons} {sr_fibrin_us:icons} {sr_fibrin_calc} (g/L)	descriptive Field Annotation: embedded_field
925	sr_tropt_embedbrk Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Cardiac biomarkers if Troponin values are reported as "&lt; 10" ug/L, please put "9.99"</i> Troponin-T {sr_tropt:icons} {sr_tropt_calc} (ng/L)	descriptive Field Annotation: embedded_field
926	sr_hstropt_embedbrk Show the field ONLY if: [sr_labdt] <> "	High sensit. Troponin-T {sr_hstropt:icons} {sr_hstropt_us:icons} {sr_hstropt_calc} (ng/L)	descriptive Field Annotation: embedded_field
927	sr_tropi_embedbrk Show the field ONLY if: [sr_labdt] <> "	Troponin-I {sr_tropi:icons} {sr_tropi_calc} (ng/L)	descriptive Field Annotation: embedded_field
928	sr_hstropi_embedbrk Show the field ONLY if: [sr_labdt] <> "	High sensit. Troponin-I {sr_hstropi:icons} {sr_hstropi_calc} (ng/L)	descriptive Field Annotation: embedded_field
929	sr_probnp_embedbrk Show the field ONLY if: [sr_labdt] <> "	Pro-BNP {sr_probnp:icons} {sr_probnp_calc} (ng/L)	descriptive Field Annotation: embedded_field
930	sr_bnp_embedbrk Show the field ONLY if: [sr_labdt] <> "	BNP {sr_bnp:icons} {sr_bnp_calc} (ng/L)	descriptive Field Annotation: embedded_field
931	sr_procalc_embedbrk Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>Other</i> Procalcitonin {sr_procalc:icons} {sr_procalc_calc} (ug/L)	descriptive Field Annotation: embedded_field
932	sr_il1_embedbrk Show the field ONLY if: [sr_labdt] <> "	IL-1 {sr_il1:icons} {sr_il1_calc} (pg/L)	descriptive Field Annotation: embedded_field
933	sr_il1ra_embedbrk Show the field ONLY if: [sr_labdt] <> "	IL-1Ra {sr_il1ra:icons} {sr_il1ra_calc} (pg/L)	descriptive Field Annotation: embedded_field
934	sr_il6_embedbrk Show the field ONLY if: [sr_labdt] <> "	IL-6 {sr_il6:icons} {sr_il6_calc} (pg/L)	descriptive Field Annotation: embedded_field
935	sr_lab_name1 Show the field ONLY if: [sr_labdt] <> "	Section Header: <i>ADDITIONAL LABWORK</i> Other lab test 1: name	text
936	sr_lab_result1 Show the field ONLY if: [sr_lab_name1] <> "	[sr_lab_name1]: result	text Field Annotation: @PLACEHOLDER = "result"
937	sr_lab_unit1 Show the field ONLY if: [sr_lab_name1] <> "	[sr_lab_name1]: unit	text Field Annotation: @PLACEHOLDER = "unit"
938	sr_lab_embed1 Show the field ONLY if: [sr_lab_name1] <> "	[sr_lab_name1]: result & unit {sr_lab_result1} {sr_lab_unit1}	descriptive
939	sr_lab_name2 Show the field ONLY if: [sr_labdt] <> "	Other lab test 2: name	text
940	sr_lab_result2 Show the field ONLY if: [sr_lab_name2] <> "	[sr_lab_name2]: result	text Field Annotation: @PLACEHOLDER = "result"
941	sr_lab_unit2 Show the field ONLY if: [sr_lab_name2] <> "	[sr_lab_name2]: unit	text Field Annotation: @PLACEHOLDER = "unit"

942	sr_lab_embed2 Show the field ONLY if: [sr_lab_name2] <> "	[sr_lab_name2]: result & unit {sr_lab_result2} {sr_lab_unit2}	descriptive						
943	sr_lab_name3 Show the field ONLY if: [sr_labdt] <> "	Other lab test 3: name	text						
944	sr_lab_result3 Show the field ONLY if: [sr_lab_name3] <> "	[sr_lab_name3]: result	text Field Annotation: @PLACEHOLDER = "result"						
945	sr_lab_unit3 Show the field ONLY if: [sr_lab_name3] <> "	[sr_lab_name3]: unit	text Field Annotation: @PLACEHOLDER = "unit"						
946	sr_lab_embed3 Show the field ONLY if: [sr_lab_name3] <> "	[sr_lab_name3]: result & unit {sr_lab_result3} {sr_lab_unit3}	descriptive						
947	sr_lab_name4 Show the field ONLY if: [sr_labdt] <> "	Other lab test 4: name	text						
948	sr_lab_result4 Show the field ONLY if: [sr_lab_name4] <> "	[sr_lab_name4]: result	text Field Annotation: @PLACEHOLDER = "result"						
949	sr_lab_unit4 Show the field ONLY if: [sr_lab_name4] <> "	[sr_lab_name4]: unit	text Field Annotation: @PLACEHOLDER = "unit"						
950	sr_lab_embed4 Show the field ONLY if: [sr_lab_name4] <> "	[sr_lab_name4]: result & unit {sr_lab_result4} {sr_lab_unit4}	descriptive						
951	sr_lab_name5 Show the field ONLY if: [sr_labdt] <> "	Other lab test 5: name	text						
952	sr_lab_result5 Show the field ONLY if: [sr_lab_name5] <> "	[sr_lab_name5]: result	text Field Annotation: @PLACEHOLDER = "result"						
953	sr_lab_unit5 Show the field ONLY if: [sr_lab_name5] <> "	[sr_lab_name5]: unit	text Field Annotation: @PLACEHOLDER = "unit"						
954	sr_lab_embed5 Show the field ONLY if: [sr_lab_name5] <> "	[sr_lab_name5]: result & unit {sr_lab_result5} {sr_lab_unit5}	descriptive						
955	serial_labwork_complete	Section Header: <i>Form Status</i> Complete?	dropdown <table border="1"> <tr><td>0</td><td>Incomplete</td></tr> <tr><td>1</td><td>Unverified</td></tr> <tr><td>2</td><td>Complete</td></tr> </table>	0	Incomplete	1	Unverified	2	Complete
0	Incomplete								
1	Unverified								
2	Complete								

Instrument: **Serial culture** (serial\_culture)

[^ Collapse](#)

956	brk_0402	Section Header: <i>SERIAL CULTURE</i> Only applicable for patients diagnosed on or after Jan 01, 2020  REDCap DATA ENTRY NOT REQUIRED Site to upload "Serial labwork" datafile (.csv file) in the "file upload" section DCC to import serial labwork data (received from the sites) into REDCap	descriptive										
957	sr_culturedt	Specimen collection date/time for culture	text (date_ymd) Field Annotation: @HIDEBUTTON										
958	sr_culturetyp Show the field ONLY if: [sr_culturedt] <> "	Culture type	radio <table border="1"> <tr><td>1</td><td>Blood culture</td></tr> <tr><td>2</td><td>Urine culture</td></tr> <tr><td>3</td><td>Stool culture</td></tr> <tr><td>4</td><td>Throat culture</td></tr> <tr><td>5</td><td>Other</td></tr> </table>	1	Blood culture	2	Urine culture	3	Stool culture	4	Throat culture	5	Other
1	Blood culture												
2	Urine culture												
3	Stool culture												
4	Throat culture												
5	Other												

959	sr_culturetypx Show the field ONLY if: [sr_culturetyp] = '5'	Specify culture type	text
960	sr_cultureres Show the field ONLY if: [sr_culturedt] <> "	Culture result	radio 1 No growth 2 Positive culture
961	sr_culturetx Show the field ONLY if: [sr_cultureres] = '2'	Organism(s) identified in the positive culture	notes
962	serial_culture_complete	Section Header: <i>Form Status</i> Complete?	dropdown 0 Incomplete 1 Unverified 2 Complete

Instrument: **Serial panel** (serial\_panel)[^ Collapse](#)

963	brk_0502	Section Header: <i>SERIAL PANEL</i> Only applicable for patients diagnosed on or after Jan 01, 2020 REDCap DATA ENTRY NOT REQUIRED Site to upload "Serial panel" datafile (.csv file) in the "file upload" section DCC to import serial panel data (received from the sites) into REDCap	descriptive
964	sr_paneldt	Specimen collection date/time for panel	text (datetime_ymd) Field Annotation: @HIDEBUTTON
965	sr_panelyp Show the field ONLY if: [sr_paneldt] <> "	Specimen type <i>e.g., nasal swab</i>	text
966	sr_panetyt Show the field ONLY if: [sr_paneldt] <> "	Specimen type	radio 1 Nasal swab 2 Throat swab 3 Blood 4 Stool 5 Other
967	sr_panetyptx Show the field ONLY if: [sr_panetyt] = '5'	Specify the specimen	text
968	sr_paneltx Show the field ONLY if: [sr_paneldt] <> "	Panel result	notes
969	serial_panel_complete	Section Header: <i>Form Status</i> Complete?	dropdown 0 Incomplete 1 Unverified 2 Complete

Instrument: **Serial PCR** (serial\_pcr)[^ Collapse](#)

970	brk_0602	Section Header: <i>SERIAL PCR</i> Only applicable for patients diagnosed on or after Jan 01, 2020 REDCap DATA ENTRY NOT REQUIRED Site to upload "Serial PCR" datafile (.csv file) in the "file upload" section DCC to import serial PCR data (received from the sites) into REDCap	descriptive
971	sr_pcrdt	PCR date/time	text (datetime_ymd) Field Annotation: @HIDEBUTTON
972	sr_pcrtyt Show the field ONLY if: [sr_pcrdt] <> "	PCR specimen type	radio 1 Swab 2 Serum 3 Other
973	sr_pcrtyptx Show the field ONLY if: [sr_pcrtyt] = '3'	Specify the specimen	text

974	sr_pcrresult Show the field ONLY if: [sr_pcrdt] <> "	Result of PCR	radio 1 Covid +ve 2 Covid -ve 3 Other
975	sr_pcrcomments Show the field ONLY if: [sr_pcrdt] <> "	Comments	notes
976	serial_pcr_complete	Section Header: <i>Form Status</i> Complete?	dropdown 0 Incomplete 1 Unverified 2 Complete
Instrument: <b>Acute phase meds</b> (acute_phase_meds) <span style="float: right;">^ Collapse</span>			
977	ap_wt	Section Header: <i>ACUTE PHASE TREATMENT Medications used during acute episode</i> Weight during acute phase (kg)	text (number, Min: 1, Max: 200) Field Annotation: @PLACEHOLDER = 'kg'
978	ap_ht	Height during acute phase (cm)	text (number, Min: 1, Max: 500) Field Annotation: @PLACEHOLDER = 'cm'
979	ds_highesttemp Show the field ONLY if: [ds_substudy] = '1'	Highest fever recorded before start of IVIG (degree C)	text (number, Max: 45) Field Annotation: @PLACEHOLDER = 'centigrade)
980	ap_ivgg	Section Header: <i>SECTION: IVIG/IVGG</i> Intravenous gamma Immunoglobulin (IVIG/IVGG)	yesno 1 Yes 0 No
981	ap_ivggdoses Show the field ONLY if: [ap_ivgg] = '1'	Number of IVIG doses <i>1 dose is the equivalent of 2g/kg (2000mg/kg)(i.e. a split infusion of 2 x 1000mg/kg counts as a single infusion)</i>	radio 1 1 2 2 3 3 4 4  Custom alignment: RH
982	ap_ivggdt1 Show the field ONLY if: [ap_ivggdoses] >= 1	1st IVIG treatment - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON
983	ap_ivggdose1 Show the field ONLY if: [ap_ivggdoses] >= 1	1st IVIG treatment - dose (mg/kg)	text (number, Min: 1000, Max: 5000) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'mg/kg'
984	ds_ap_ivggstarttm1 Show the field ONLY if: [ap_ivggdoses] >= 1	1st IVIG treatment - start time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = "start"
985	ds_ap_ivggenddt1 Show the field ONLY if: [ap_ivggdoses] >= 1	1st IVIG treatment - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON
986	ds_ap_ivggendtm1 Show the field ONLY if: [ap_ivggdoses] >= 1	1st IVIG treatment - end time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = "end"
987	ds_ap_ivggbrand1 Show the field ONLY if: [ap_ivggdoses] >= 1	1st IVIG treatment - brand	text Custom alignment: RH
988	ds_ap_ivggconc1 Show the field ONLY if: [ap_ivggdoses] >= 1	1st IVIG treatment - concentration	dropdown 1 5% concentration 2 10% concentration
989	ds_ap_ivggrate1 Show the field ONLY if: [ap_ivggdoses] >= 1	1st IVIG treatment - infusion rate (ml/hr)	text (number) Custom alignment: RH
990	emb_1001 Show the field ONLY if: [ap_ivggdoses] >= 1	1st IVIG	descriptive

991	emb_1002 Show the field ONLY if: [ap_ivggdoses] >= 1	Start date: {ap_ivggt1} Dose: {ap_ivggdose1}	descriptive				
992	emb_1003 Show the field ONLY if: [ap_ivggdoses] >= 1 and [ds_substudy] = '1'	Start time: {ds_ap_ivggstarttm1} Brand: {ds_ap_ivggbrand1} End date: {ds_ap_ivggenddt1} Concentration: {ds_ap_ivggconc1} End time: {ds_ap_ivggendtm1} Infusion rate (ml/hr): {ds_ap_ivggtrate1}	descriptive				
993	brk_1001 Show the field ONLY if: [ap_ivggdoses] >= 1		descriptive				
994	ap_ivggt2 Show the field ONLY if: [ap_ivggdoses] >= 2	2nd IVIG treatment - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
995	ap_ivggdose2 Show the field ONLY if: [ap_ivggdoses] >= 2	2nd IVIG treatment - dose (mg/kg)	text (number, Min: 1000, Max: 5000) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'mg/kg'				
996	ds_ap_ivggstarttm2 Show the field ONLY if: [ap_ivggdoses] >= 2	2nd IVIG treatment - start time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = "start"				
997	ds_ap_ivggenddt2 Show the field ONLY if: [ap_ivggdoses] >= 2	2nd IVIG treatment - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
998	ds_ap_ivggendtm2 Show the field ONLY if: [ap_ivggdoses] >= 2	2nd IVIG treatment - end time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = "end"				
999	ds_ap_ivggbrand2 Show the field ONLY if: [ap_ivggdoses] >= 2	2nd IVIG treatment - brand	text Custom alignment: RH				
1000	ds_ap_ivggconc2 Show the field ONLY if: [ap_ivggdoses] >= 2	2nd IVIG treatment - concentration	dropdown <table border="1"> <tr> <td>1</td> <td>5% concentration</td> </tr> <tr> <td>2</td> <td>10% concentration</td> </tr> </table>	1	5% concentration	2	10% concentration
1	5% concentration						
2	10% concentration						
1001	ds_ap_ivggtrate2 Show the field ONLY if: [ap_ivggdoses] >= 2	2nd IVIG treatment - infusion rate (ml/hr)	text (number) Custom alignment: RH				
1002	emb_1004 Show the field ONLY if: [ap_ivggdoses] >= 2	2nd IVIG	descriptive				
1003	emb_1005 Show the field ONLY if: [ap_ivggdoses] >= 2	Start date: {ap_ivggt2} Dose: {ap_ivggdose2}	descriptive				
1004	emb_1006 Show the field ONLY if: [ap_ivggdoses] >= 2 and [ds_substudy] = '1'	Start time: {ds_ap_ivggstarttm2} Brand: {ds_ap_ivggbrand2} End date: {ds_ap_ivggenddt2} Concentration: {ds_ap_ivggconc2} End time: {ds_ap_ivggendtm2} Infusion rate (ml/hr): {ds_ap_ivggtrate2}	descriptive				
1005	brk_1002 Show the field ONLY if: [ap_ivggdoses] >= 2		descriptive				
1006	ap_ivggt3 Show the field ONLY if: [ap_ivggdoses] >= 3	3rd IVIG treatment - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1007	ap_ivggdose3 Show the field ONLY if: [ap_ivggdoses] >= 3	3rd IVIG treatment - dose (mg/kg)	text (number, Min: 1000, Max: 5000) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'mg/kg'				
1008	ds_ap_ivggstarttm3 Show the field ONLY if: [ap_ivggdoses] >= 3	3rd IVIG treatment - start time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = "start"				
1009	ds_ap_ivggenddt3 Show the field ONLY if: [ap_ivggdoses] >= 3	3rd IVIG treatment - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				

1010	ds_ap_ivggendtm3 Show the field ONLY if: [ap_ivggdoses] >= 3	3rd IVIG treatment - end time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = "end"				
1011	ds_ap_ivggbrand3 Show the field ONLY if: [ap_ivggdoses] >= 3	3rd IVIG treatment - brand	text Custom alignment: RH				
1012	ds_ap_ivggconc3 Show the field ONLY if: [ap_ivggdoses] >= 3	3rd IVIG treatment - concentration	dropdown <table border="1"> <tr> <td>1</td> <td>5% concentration</td> </tr> <tr> <td>2</td> <td>10% concentration</td> </tr> </table>	1	5% concentration	2	10% concentration
1	5% concentration						
2	10% concentration						
1013	ds_ap_ivggrate3 Show the field ONLY if: [ap_ivggdoses] >= 3	3rd IVIG treatment - infusion rate (ml/hr)	text (number) Custom alignment: RH				
1014	emb_1007 Show the field ONLY if: [ap_ivggdoses] >= 3	3rd IVIG	descriptive				
1015	emb_1008 Show the field ONLY if: [ap_ivggdoses] >= 3	Start date: {ap_ivggt3} Dose: {ap_ivggdose3}	descriptive				
1016	emb_1009 Show the field ONLY if: [ap_ivggdoses] >= 3 and [ds_substudy] = '1'	Start time: {ds_ap_ivggstarttm3} Brand: {ds_ap_ivggbrand3} End date: {ds_ap_ivggenddt3} Concentration: {ds_ap_ivggconc3} End time: {ds_ap_ivggendtm3} Infusion rate (ml/hr): {ds_ap_ivggrate3}	descriptive				
1017	brk_1003 Show the field ONLY if: [ap_ivggdoses] >= 3		descriptive				
1018	ap_ivggt4 Show the field ONLY if: [ap_ivggdoses] >= 4	4th IVIG treatment - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1019	ap_ivggdose4 Show the field ONLY if: [ap_ivggdoses] >= 4	4th IVIG treatment - dose (mg/kg)	text (number, Min: 1000, Max: 5000) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'mg/kg'				
1020	ds_ap_ivggstarttm4 Show the field ONLY if: [ap_ivggdoses] >= 4	4th IVIG treatment - start time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = "start"				
1021	ds_ap_ivggenddt4 Show the field ONLY if: [ap_ivggdoses] >= 4	4th IVIG treatment - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1022	ds_ap_ivggendtm4 Show the field ONLY if: [ap_ivggdoses] >= 4	4th IVIG treatment - end time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = "end"				
1023	ds_ap_ivggbrand4 Show the field ONLY if: [ap_ivggdoses] >= 4	4th IVIG treatment - brand	text Custom alignment: RH				
1024	ds_ap_ivggconc4 Show the field ONLY if: [ap_ivggdoses] >= 4	4th IVIG treatment - concentration	dropdown <table border="1"> <tr> <td>1</td> <td>5% concentration</td> </tr> <tr> <td>2</td> <td>10% concentration</td> </tr> </table>	1	5% concentration	2	10% concentration
1	5% concentration						
2	10% concentration						
1025	ds_ap_ivggrate4 Show the field ONLY if: [ap_ivggdoses] >= 4	4th IVIG treatment - infusion rate (ml/hr)	text (number) Custom alignment: RH				
1026	emb_1010 Show the field ONLY if: [ap_ivggdoses] >= 4	4th IVIG	descriptive				
1027	emb_1011 Show the field ONLY if: [ap_ivggdoses] >= 4	Start date: {ap_ivggt4} Dose: {ap_ivggdose4}	descriptive				
1028	emb_1012 Show the field ONLY if: [ap_ivggdoses] >= 4 and [ds_substudy] = '1'	Start time: {ds_ap_ivggstarttm4} Brand: {ds_ap_ivggbrand4} End date: {ds_ap_ivggenddt4} Concentration: {ds_ap_ivggconc4} End time: {ds_ap_ivggendtm4} Infusion rate (ml/hr): {ds_ap_ivggrate4}	descriptive				

1029	brk_1004 Show the field ONLY if: [ap_ivggdoses] >= 4		descriptive
1030	ap_ivggresist Show the field ONLY if: [ap_ivgg] = '1'	Section Header: SECTION: IVIG/IVGG EFFECTS IVIG resistance <i>Resistant KD: failure to respond within 36 h following the first dose of IVIG</i>	yesno 1 Yes 0 No
1031	ds_ap_ivggresisttyp Show the field ONLY if: [ap_ivggresist] = '1' and [ds_substudy] = '1'	Type of IVIG resistance	radio 1 Partial (fever reduced during IVIG) 2 Complete (fever unaffected by IVIG)
1032	ds_ap_ivggadvers Show the field ONLY if: [ap_ivgg] = '1' and [ds_substudy] = '1'	Adverse events associated with IVIG	checkbox 0 ds_ap_ivggadvers__0 None 1 ds_ap_ivggadvers__1 Abdominal pain 2 ds_ap_ivggadvers__2 Anaphylaxis 3 ds_ap_ivggadvers__3 Arrhythmia 4 ds_ap_ivggadvers__4 Chest tightness 5 ds_ap_ivggadvers__5 Chills 6 ds_ap_ivggadvers__6 Headaches 7 ds_ap_ivggadvers__7 Hives 8 ds_ap_ivggadvers__8 Hypertension 9 ds_ap_ivggadvers__9 Hypotension 10 ds_ap_ivggadvers__10 Hypotonicity 11 ds_ap_ivggadvers__11 Itching 12 ds_ap_ivggadvers__12 Lethargy 13 ds_ap_ivggadvers__13 New onset myalgia/arthralgia 14 ds_ap_ivggadvers__14 Rigors 15 ds_ap_ivggadvers__15 Tachypnea 16 ds_ap_ivggadvers__16 Volume overload/generalized edema 17 ds_ap_ivggadvers__17 Vomiting/nausea 18 ds_ap_ivggadvers__18 Hemolysis 19 ds_ap_ivggadvers__19 Critical anemia 20 ds_ap_ivggadvers__20 Other  Field Annotation: @NONEOFHEABOVE='0'
1033	ds_ap_ivggadverstx Show the field ONLY if: [ds_ap_ivggadvers(20)] = '1'	if other adverse events, please specify	text
1034	ap_antiplat Show the field ONLY if: [ap_antiplat] = '1'	Section Header: SECTION: ANTIPLATELETS Antiplatelets	yesno 1 Yes 0 No
1035	ap_antiplattyp Show the field ONLY if: [ap_antiplat] = '1'	Antiplatelets: agents(check all that apply)	checkbox 1 ap_antiplattyp__1 Aspirin/ASA 2 ap_antiplattyp__2 Clopidogrel 3 ap_antiplattyp__3 Other agents
1036	ap_antiplattx Show the field ONLY if: [ap_antiplattyp(3)] = '1'	if other antiplatelet agents, please specify	text Field Annotation: @PLACEHOLDER = "Other agent name"
1037	ap_antiplattxdt Show the field ONLY if: [ap_antiplattx] <> ""	[ap_antiplattx] start date	text (date_ymd) Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start date'
1038	ap_antiplatdis Show the field ONLY if: [ap_antiplattx] <> ""	[ap_antiplattx] at discharge	yesno 1 Yes 0 No  Custom alignment: RH



1039	ap_clopiddt Show the field ONLY if: [ap_antiplattyp(2)] = '1'	Clopidogrel start date	text (date_ymd) Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON				
1040	ap_clopiddis Show the field ONLY if: [ap_antiplattyp(2)] = '1'	Clopidogrel at discharge	yesno <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>No</td> </tr> </table> Custom alignment: RH Field Annotation: @HIDEBUTTON	1	Yes	0	No
1	Yes						
0	No						
1041	ap_aspirinsta1 Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin 1 start	text (date_ymd) Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON				
1042	ap_aspirinsta2 Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin 2 start	text (date_ymd) Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON				
1043	ap_aspirinsta3 Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin 3 start	text (date_ymd) Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON				
1044	ap_aspirinsta4 Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin 4 start	text (date_ymd) Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON				
1045	ap_aspirinsta5 Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin 5 start	text (date_ymd) Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON				
1046	ap_aspirinend1 Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin 1 end date	text (date_ymd) Field Annotation: @PLACEHOLDER = 'end date' @HIDEBUTTON				
1047	ap_aspirinend2 Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin 2 end date	text (date_ymd) Field Annotation: @PLACEHOLDER = 'end date' @HIDEBUTTON				
1048	ap_aspirinend3 Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin 3 end date	text (date_ymd) Field Annotation: @PLACEHOLDER = 'end date' @HIDEBUTTON				
1049	ap_aspirinend4 Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin 4 end date	text (date_ymd) Field Annotation: @PLACEHOLDER = 'end date' @HIDEBUTTON				
1050	ap_aspirinend5 Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin 5 end date	text (date_ymd) Field Annotation: @PLACEHOLDER = 'end date' @HIDEBUTTON				
1051	ap_aspirindose1 Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin 1 dose (mg/kg)	text (number) Field Annotation: @PLACEHOLDER = 'dose (mg/kg/day)'				
1052	ap_aspirindose2 Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin 2 dose (mg/kg)	text (number) Field Annotation: @PLACEHOLDER = 'dose (mg/kg/day)'				
1053	ap_aspirindose3 Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin 3 dose (mg/kg)	text (number) Field Annotation: @PLACEHOLDER = 'dose (mg/kg/day)'				
1054	ap_aspirindose4 Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin 4 dose (mg/kg)	text (number) Field Annotation: @PLACEHOLDER = 'dose (mg/kg/day)'				
1055	ap_aspirindose5 Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin 5 dose (mg/kg)	text (number) Field Annotation: @PLACEHOLDER = 'dose (mg/kg/day)'				
1056	ap_aspirindis Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin at discharge	radio <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>No</td> </tr> </table> Custom alignment: RH Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON	1	Yes	0	No
1	Yes						
0	No						

1057	ap_aspirinembed Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin 1 {ap_aspirinsta1} {ap_aspirinend1} {ap_aspirindose1} Aspirin 2 {ap_aspirinsta2} {ap_aspirinend2} {ap_aspirindose2} Aspirin 3 {ap_aspirinsta3} {ap_aspirinend3} {ap_aspirindose3} Aspirin 4 {ap_aspirinsta4} {ap_aspirinend4} {ap_aspirindose4} Aspirin 5 {ap_aspirinsta5} {ap_aspirinend5} {ap_aspirindose5}	descriptive
1058	emb_1013 Show the field ONLY if: [ap_antiplattyp(1)] = '1' or [ap_antiplattyp(2)] = '1' or [ap_antiplattyp(3)] = '1'	Given at discharge?	descriptive
1059	emb_1014 Show the field ONLY if: [ap_antiplattyp(1)] = '1'	Aspirin {ap_aspirindis}	descriptive
1060	emb_1015 Show the field ONLY if: [ap_antiplattyp(2)] = '1'	Clopidogrel {ap_clopiddt} {ap_clopiddis}	descriptive
1061	emb_1016 Show the field ONLY if: [ap_antiplattyp(3)] = '1'	{ap_antiplattx} {ap_antiplattxd} {ap_antiplatdis}	descriptive
1062	ap_uhf	Section Header: <i>SECTION: ANTICOAGULANTS</i> Unfractionated heparin (UFH)	yesno 1 Yes 0 No
1063	ap_ufhstdt1 Show the field ONLY if: [ap_uhf] = '1'	UFH 1 - start date	text (date_ymd) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON
1064	ap_ufhstdt2 Show the field ONLY if: [ap_ufhstdt1] <> "	UFH 2 - start date	text (date_ymd) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON
1065	ap_ufhstdt3 Show the field ONLY if: [ap_ufhstdt2] <> "	UFH 3 - start date	text (date_ymd) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON
1066	ap_anticoag	Other anticoagulants (LMWH/Warfarin/Bivalirudin)	yesno 1 Yes 0 No
1067	ap_anticoagtyp Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulants: agents(check all that apply) <i>LMWH: Enoxaparin, Tinzaparin, Dalteparin</i> For DOACs (select DOAC field below)	checkbox 1 ap_anticoagtyp__1 UFH 2 ap_anticoagtyp__2 LMWH 3 ap_anticoagtyp__3 Warfarin 4 ap_anticoagtyp__4 Bivalirudin Field Annotation: @HIDECHOICE='1'
1068	ap_doac	Direct acting anticoagulants (DOACs) <i>DOACs: Apixaban, Edoxaban, Rivaroxaban, Dabigatran</i>	yesno 1 Yes 0 No
1069	ap_doactyp Show the field ONLY if: [ap_doac] = '1'	DOAC: agents(check all that apply)	checkbox 1 ap_doactyp__1 Apixaban 2 ap_doactyp__2 Edoxaban 3 ap_doactyp__3 Rivaroxaban 4 ap_doactyp__4 Dabigatran
1070	ds_ap_actyp1 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulants: agent 1	dropdown 2 LMWH 3 Warfarin 4 Bivalirudin
1071	ds_ap_acstdt1 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulant start date 1	text (date_ymd) Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON

1072	ds_ap_acenddt1 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulant end date 1	text (date_ymd) Field Annotation: @PLACEHOLDER = 'end date' @HIDEBUTTON						
1073	ds_ap_acdis1 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulant at discharge 1	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No		
1	Yes								
0	No								
1074	ds_ap_actyp2 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulants: agent 2	dropdown <table border="1"> <tr><td>2</td><td>LMWH</td></tr> <tr><td>3</td><td>Warfarin</td></tr> <tr><td>4</td><td>Bivalirudin</td></tr> </table>	2	LMWH	3	Warfarin	4	Bivalirudin
2	LMWH								
3	Warfarin								
4	Bivalirudin								
1075	ds_ap_acstdt2 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulant start date 2	text (date_ymd) Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON						
1076	ds_ap_acenddt2 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulant end date 2	text (date_ymd) Field Annotation: @PLACEHOLDER = 'end date' @HIDEBUTTON						
1077	ds_ap_acdis2 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulant at discharge 2	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No		
1	Yes								
0	No								
1078	ds_ap_actyp3 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulants: agent 3	dropdown <table border="1"> <tr><td>2</td><td>LMWH</td></tr> <tr><td>3</td><td>Warfarin</td></tr> <tr><td>4</td><td>Bivalirudin</td></tr> </table>	2	LMWH	3	Warfarin	4	Bivalirudin
2	LMWH								
3	Warfarin								
4	Bivalirudin								
1079	ds_ap_acstdt3 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulant start date 3	text (date_ymd) Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON						
1080	ds_ap_acenddt3 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulant end date 3	text (date_ymd) Field Annotation: @PLACEHOLDER = 'end date' @HIDEBUTTON						
1081	ds_ap_acdis3 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulant at discharge 3	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No		
1	Yes								
0	No								
1082	ds_ap_actyp4 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulants: agent 4	dropdown <table border="1"> <tr><td>2</td><td>LMWH</td></tr> <tr><td>3</td><td>Warfarin</td></tr> <tr><td>4</td><td>Bivalirudin</td></tr> </table>	2	LMWH	3	Warfarin	4	Bivalirudin
2	LMWH								
3	Warfarin								
4	Bivalirudin								
1083	ds_ap_acstdt4 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulant start date 4	text (date_ymd) Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON						
1084	ds_ap_acenddt4 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulant end date 4	text (date_ymd) Field Annotation: @PLACEHOLDER = 'end date' @HIDEBUTTON						
1085	ds_ap_acdis4 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulant at discharge 4	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No		
1	Yes								
0	No								
1086	ds_ap_actyp5 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulants: agent 5	dropdown <table border="1"> <tr><td>2</td><td>LMWH</td></tr> <tr><td>3</td><td>Warfarin</td></tr> <tr><td>4</td><td>Bivalirudin</td></tr> </table>	2	LMWH	3	Warfarin	4	Bivalirudin
2	LMWH								
3	Warfarin								
4	Bivalirudin								

1087	ds_ap_acstdt5 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulant start date 5	text (date_ymd) Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON								
1088	ds_ap_acenddt5 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulant end date 5	text (date_ymd) Field Annotation: @PLACEHOLDER = 'end date' @HIDEBUTTON								
1089	ds_ap_acdis5 Show the field ONLY if: [ap_anticoag] = '1'	Anticoagulant at discharge 5	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1090	emb_1018 Show the field ONLY if: [ap_anticoag] = '1'	Agent StartDate EndDate Given at discharge? Anticoagulants {ds_ap_actyp1} {ds_ap_acstdt1} {ds_ap_acenddt1} {ds_ap_acdis1} {ds_ap_actyp2} {ds_ap_acstdt2} {ds_ap_acenddt2} {ds_ap_acdis2} {ds_ap_actyp3} {ds_ap_acstdt3} {ds_ap_acenddt3} {ds_ap_acdis3} {ds_ap_actyp4} {ds_ap_acstdt4} {ds_ap_acenddt4} {ds_ap_acdis4} {ds_ap_actyp5} {ds_ap_acstdt5} {ds_ap_acenddt5} {ds_ap_acdis5}	descriptive								
1091	ds_ap_doactyp1 Show the field ONLY if: [ap_doac] = '1'	DOACs: agent 1	dropdown <table border="1"> <tr><td>1</td><td>Apixaban</td></tr> <tr><td>2</td><td>Edoxaban</td></tr> <tr><td>3</td><td>Rivaroxaban</td></tr> <tr><td>4</td><td>Dabigatran</td></tr> </table>	1	Apixaban	2	Edoxaban	3	Rivaroxaban	4	Dabigatran
1	Apixaban										
2	Edoxaban										
3	Rivaroxaban										
4	Dabigatran										
1092	ds_ap_doacstdt1 Show the field ONLY if: [ap_doac] = '1'	DOAC start date 1	text (date_ymd) Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON								
1093	ds_ap_doacenddt1 Show the field ONLY if: [ap_doac] = '1'	DOAC end date 1	text (date_ymd) Field Annotation: @PLACEHOLDER = 'end date' @HIDEBUTTON								
1094	ds_ap_doacdis1 Show the field ONLY if: [ap_doac] = '1'	DOAC given at discharge 1	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1095	ds_ap_doactyp2 Show the field ONLY if: [ap_doac] = '1'	DOACs: agent 2	dropdown <table border="1"> <tr><td>1</td><td>Apixaban</td></tr> <tr><td>2</td><td>Edoxaban</td></tr> <tr><td>3</td><td>Rivaroxaban</td></tr> <tr><td>4</td><td>Dabigatran</td></tr> </table> Field Annotation: @HIDDEN	1	Apixaban	2	Edoxaban	3	Rivaroxaban	4	Dabigatran
1	Apixaban										
2	Edoxaban										
3	Rivaroxaban										
4	Dabigatran										
1096	ds_ap_doacstdt2 Show the field ONLY if: [ap_doac] = '1'	DOAC start date 2	text (date_ymd) Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON								
1097	ds_ap_doacenddt2 Show the field ONLY if: [ap_doac] = '1'	DOAC end date 2	text (date_ymd) Field Annotation: @PLACEHOLDER = 'end date' @HIDEBUTTON								
1098	ds_ap_doacdis2 Show the field ONLY if: [ap_doac] = '1'	DOAC given at discharge 2	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1099	ds_ap_doactyp3 Show the field ONLY if: [ap_doac] = '1'	DOACs: agent 3	dropdown <table border="1"> <tr><td>1</td><td>Apixaban</td></tr> <tr><td>2</td><td>Edoxaban</td></tr> <tr><td>3</td><td>Rivaroxaban</td></tr> <tr><td>4</td><td>Dabigatran</td></tr> </table>	1	Apixaban	2	Edoxaban	3	Rivaroxaban	4	Dabigatran
1	Apixaban										
2	Edoxaban										
3	Rivaroxaban										
4	Dabigatran										
1100	ds_ap_doacstdt3 Show the field ONLY if: [ap_doac] = '1'	DOAC start date 3	text (date_ymd) Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON								

1101	ds_ap_doacenddt3 Show the field ONLY if: [ap_doac] = '1'	DOAC end date 3	text (date_ymd) Field Annotation: @PLACEHOLDER = 'end date' @HIDEBUTTON								
1102	ds_ap_doacdis3 Show the field ONLY if: [ap_doac] = '1'	DOAC given at discharge 3	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1103	ds_ap_doactyp4 Show the field ONLY if: [ap_doac] = '1'	DOACs: agent 4	dropdown <table border="1"> <tr><td>1</td><td>Apixaban</td></tr> <tr><td>2</td><td>Edoxaban</td></tr> <tr><td>3</td><td>Rivaroxaban</td></tr> <tr><td>4</td><td>Dabigatran</td></tr> </table>	1	Apixaban	2	Edoxaban	3	Rivaroxaban	4	Dabigatran
1	Apixaban										
2	Edoxaban										
3	Rivaroxaban										
4	Dabigatran										
1104	ds_ap_doacstdt4 Show the field ONLY if: [ap_doac] = '1'	DOAC start date 4	text (date_ymd) Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON								
1105	ds_ap_doacenddt4 Show the field ONLY if: [ap_doac] = '1'	DOAC end date 4	text (date_ymd) Field Annotation: @PLACEHOLDER = 'end date' @HIDEBUTTON								
1106	ds_ap_doacdis4 Show the field ONLY if: [ap_doac] = '1'	DOAC given at discharge 4	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1107	ds_ap_doactyp5 Show the field ONLY if: [ap_doac] = '1'	DOACs: agent 5	dropdown <table border="1"> <tr><td>1</td><td>Apixaban</td></tr> <tr><td>2</td><td>Edoxaban</td></tr> <tr><td>3</td><td>Rivaroxaban</td></tr> <tr><td>4</td><td>Dabigatran</td></tr> </table>	1	Apixaban	2	Edoxaban	3	Rivaroxaban	4	Dabigatran
1	Apixaban										
2	Edoxaban										
3	Rivaroxaban										
4	Dabigatran										
1108	ds_ap_doacstdt5 Show the field ONLY if: [ap_doac] = '1'	DOAC start date 5	text (date_ymd) Field Annotation: @PLACEHOLDER = 'start date' @HIDEBUTTON								
1109	ds_ap_doacenddt5 Show the field ONLY if: [ap_doac] = '1'	DOAC end date 5	text (date_ymd) Field Annotation: @PLACEHOLDER = 'end date' @HIDEBUTTON								
1110	ds_ap_doacdis5 Show the field ONLY if: [ap_doac] = '1'	DOAC given at discharge 5	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1111	emb_1017 Show the field ONLY if: [ap_doac] = '1'	Agent StartDate EndDate Given at discharge? DOACs {ds_ap_doactyp1} {ds_ap_doacstdt1} {ds_ap_doacenddt1} {ds_ap_doacdis1} {ds_ap_doactyp2} {ds_ap_doacstdt2} {ds_ap_doacenddt2} {ds_ap_doacdis2} {ds_ap_doactyp3} {ds_ap_doacstdt3} {ds_ap_doacenddt3} {ds_ap_doacdis3} {ds_ap_doactyp4} {ds_ap_doacstdt4} {ds_ap_doacenddt4} {ds_ap_doacdis4} {ds_ap_doactyp5} {ds_ap_doacstdt5} {ds_ap_doacenddt5} {ds_ap_doacdis5}	descriptive								
1112	ap_thrombol	Section Header: SECTION: THROMBOLYTICS Thrombolytics <i>Thrombolytics: tissue-plasminogen activator (tPA), Streptokinase, Urokinase, Alteplase</i>	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table>	1	Yes	0	No				
1	Yes										
0	No										
1113	ap_thromboldt1 Show the field ONLY if: [ap_thrombol] = '1'	Thrombolytics 1 - date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON								
1114	ap_thromboldt2 Show the field ONLY if: [ap_thromboldt1] <> "	Thrombolytics 2 - date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON								
1115	ap_thromboldt3 Show the field ONLY if: [ap_thromboldt2] <> "	Thrombolytics 3 - date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON								

1116	ap_steriv	Section Header: SECTION: IV STEROIDS Steroids IV	yesno <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>No</td> </tr> </table>	1	Yes	0	No						
1	Yes												
0	No												
1117	ap_sterivtyp1 Show the field ONLY if: [ap_steriv] = '1'	IV steroid 1: type	dropdown <table border="1"> <tr> <td>1</td> <td>Prednisone</td> </tr> <tr> <td>2</td> <td>Methylprednisolone</td> </tr> <tr> <td>3</td> <td>Dexamethasone</td> </tr> <tr> <td>4</td> <td>Hydrocortisone</td> </tr> <tr> <td>5</td> <td>Other</td> </tr> </table>	1	Prednisone	2	Methylprednisolone	3	Dexamethasone	4	Hydrocortisone	5	Other
1	Prednisone												
2	Methylprednisolone												
3	Dexamethasone												
4	Hydrocortisone												
5	Other												
1118	ap_sterivdt1 Show the field ONLY if: [ap_sterivtyp1] <> "	IV steroid 1: start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = "start date"										
1119	ds_ap_sterivstarttm1 Show the field ONLY if: [ap_sterivtyp1] <> " and [ds_substudy] = 1	IV steroid 1: start time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = "start time"										
1120	ds_ap_sterivenddt1 Show the field ONLY if: [ap_sterivtyp1] <> " and [ds_substudy] = 1	IV steroid 1: end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = "end date"										
1121	ds_ap_sterivendtm1 Show the field ONLY if: [ap_sterivtyp1] <> " and [ds_substudy] = 1	IV steroid 1: end time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = "end time"										
1122	ds_ap_sterivdose1 Show the field ONLY if: [ap_sterivtyp1] <> " and [ds_substudy] = 1	IV steroid 1 total dose: (mg/kg)	text (number) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'dose (mg/kg)'										
1123	ap_sterivtyp2 Show the field ONLY if: [ap_steriv] = '1'	IV steroid 2: type	dropdown <table border="1"> <tr> <td>1</td> <td>Prednisone</td> </tr> <tr> <td>2</td> <td>Methylprednisolone</td> </tr> <tr> <td>3</td> <td>Dexamethasone</td> </tr> <tr> <td>4</td> <td>Hydrocortisone</td> </tr> <tr> <td>5</td> <td>Other</td> </tr> </table>	1	Prednisone	2	Methylprednisolone	3	Dexamethasone	4	Hydrocortisone	5	Other
1	Prednisone												
2	Methylprednisolone												
3	Dexamethasone												
4	Hydrocortisone												
5	Other												
1124	ds_ap_sterivdose2 Show the field ONLY if: [ap_sterivtyp2] <> "" and [ds_substudy] = 1	IV steroid 2 total dose: (mg/kg)	text (number) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'dose (mg/kg)'										
1125	ap_sterivdt2 Show the field ONLY if: [ap_sterivtyp2] <> ""	IV steroid 2: start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start date'										
1126	ds_ap_sterivstarttm2 Show the field ONLY if: [ap_sterivtyp2] <> "" and [ds_substudy] = 1	IV steroid 2: start time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start time'										
1127	ds_ap_sterivenddt2 Show the field ONLY if: [ap_sterivtyp2] <> "" and [ds_substudy] = 1	IV steroid 2: end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'end date'										
1128	ds_ap_sterivendtm2 Show the field ONLY if: [ap_sterivtyp2] <> "" and [ds_substudy] = 1	IV steroid 2: end time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'end time'										

1129	ap_sterivtyp3 Show the field ONLY if: [ap_steriv] = '1'	IV steroid 3: type	dropdown <table border="1"> <tr><td>1</td><td>Prednisone</td></tr> <tr><td>2</td><td>Methylprednisolone</td></tr> <tr><td>3</td><td>Dexamethasone</td></tr> <tr><td>4</td><td>Hydrocortisone</td></tr> <tr><td>5</td><td>Other</td></tr> </table>	1	Prednisone	2	Methylprednisolone	3	Dexamethasone	4	Hydrocortisone	5	Other
1	Prednisone												
2	Methylprednisolone												
3	Dexamethasone												
4	Hydrocortisone												
5	Other												
1130	ds_ap_sterivdose3 Show the field ONLY if: [ap_sterivtyp3] <> "" and [ds_substudy] = 1	IV steroid 3 total dose: (mg/kg)	text (number) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'dose (mg/kg)'										
1131	ap_sterivdt3 Show the field ONLY if: [ap_sterivtyp3] <> ""	IV steroid 3: start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start date'										
1132	ds_ap_sterivstarttm3 Show the field ONLY if: [ap_sterivtyp3] <> "" and [ds_substudy] = 1	IV steroid 3: start time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start time'										
1133	ds_ap_sterivenddt3 Show the field ONLY if: [ap_sterivtyp3] <> "" and [ds_substudy] = 1	IV steroid 3: end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'end date'										
1134	ds_ap_sterivendtm3 Show the field ONLY if: [ap_sterivtyp3] <> "" and [ds_substudy] = 1	IV steroid 3: end time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'end time'										
1135	ap_sterivtyp4 Show the field ONLY if: [ap_steriv] = '1'	IV steroid 4: type	dropdown <table border="1"> <tr><td>1</td><td>Prednisone</td></tr> <tr><td>2</td><td>Methylprednisolone</td></tr> <tr><td>3</td><td>Dexamethasone</td></tr> <tr><td>4</td><td>Hydrocortisone</td></tr> <tr><td>5</td><td>Other</td></tr> </table>	1	Prednisone	2	Methylprednisolone	3	Dexamethasone	4	Hydrocortisone	5	Other
1	Prednisone												
2	Methylprednisolone												
3	Dexamethasone												
4	Hydrocortisone												
5	Other												
1136	ds_ap_sterivdose4 Show the field ONLY if: [ap_sterivtyp4] <> "" and [ds_substudy] = 1	IV steroid 4 total dose: (mg/kg)	text (number) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'dose (mg/kg)'										
1137	ap_sterivdt4 Show the field ONLY if: [ap_sterivtyp4] <> ""	IV steroid 4: start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start date'										
1138	ds_ap_sterivstarttm4 Show the field ONLY if: [ap_sterivtyp4] <> "" and [ds_substudy] = 1	IV steroid 4: start time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start time'										
1139	ds_ap_sterivenddt4 Show the field ONLY if: [ap_sterivtyp4] <> "" and [ds_substudy] = 1	IV steroid 4: end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'end date'										
1140	ds_ap_sterivendtm4 Show the field ONLY if: [ap_sterivtyp4] <> "" and [ds_substudy] = 1	IV steroid 4: end time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'end time'										
1141	emb_1019 Show the field ONLY if: [ap_steriv] = '1' and [ds_substudy] = '1'	Agent Dose Start/End date Start/End time	descriptive										
1142	emb_1020 Show the field ONLY if: [ap_steriv] = '1' and [ds_substudy] = '0'	Agent Start date	descriptive										

1143	emb_1022	{ap_sterivtyp1} {ds_ap_sterivdose1} {ap_sterivdt1} {ds_ap_sterivstarttm1} {ds_ap_sterivenddt1} {ds_ap_sterivendtm1} {ap_sterivtyp2} {ds_ap_sterivdose2} {ap_sterivdt2} {ds_ap_sterivstarttm2} {ds_ap_sterivenddt2} {ds_ap_sterivendtm2} {ap_sterivtyp3} {ds_ap_sterivdose3} {ap_sterivdt3} {ds_ap_sterivstarttm3} {ds_ap_sterivenddt3} {ds_ap_sterivendtm3} {ap_sterivtyp4} {ds_ap_sterivdose4} {ap_sterivdt4} {ds_ap_sterivstarttm4} {ds_ap_sterivenddt4} {ds_ap_sterivendtm4} <i>Repeated for steroid dose 1-5</i>	descriptive
1144	ap_steroral	Section Header: SECTION: ORAL STEROIDS Steroids oral	yesno 1 Yes 0 No
1145	ap_steroraltyp Show the field ONLY if: [ap_steroral] = '1'	Oral steroid: type(check all that were used during acute phase)	checkbox 1 ap_steroraltyp__1 Prednisone 2 ap_steroraltyp__2 Methylprednisolone 3 ap_steroraltyp__3 Dexamethasone 4 ap_steroraltyp__4 Hydrocortisone 5 ap_steroraltyp__5 Other
1146	ds_ap_steroraldose Show the field ONLY if: [ap_steroral] = '1'	Steroids oral: daily dose	text (number) Field Annotation: @PLACEHOLDER = 'daily dose (mg/kg)'
1147	ds_ap_steroralstartdt Show the field ONLY if: [ap_steroral] = '1'	Steroids oral: start date	text (date_ymd) Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start date'
1148	ds_ap_steroralstarttm Show the field ONLY if: [ap_steroral] = '1'	Steroids oral: start time	text (time) Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start time'
1149	ds_ap_steroralenddt Show the field ONLY if: [ap_steroral] = '1'	Steroids oral: end date	text (date_ymd) Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'end date'
1150	ds_ap_steroralendtm Show the field ONLY if: [ap_steroral] = '1'	Steroids oral: end time	text (time) Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'end time'
1151	emb_1025 Show the field ONLY if: [ap_steroral] = '1' and [ds_substudy] = '1'	Daily dose Start/End date Start/End time	descriptive
1152	emb_1023 Show the field ONLY if: [ap_steroral] = '1' and [ds_substudy] = '1'	{ds_ap_steroraldose} {ds_ap_steroralstartdt} {ds_ap_steroralstarttm} {ds_ap_steroralenddt} {ds_ap_steroralendtm} <i>Repeated for steroid dose 1-5</i>	descriptive
1153	ap_tnfa	Section Header: SECTION: TNF- $\alpha$ INHIBITORS TNF- $\alpha$ inhibitors	yesno 1 Yes 0 No
1154	ap_tnfa_typ Show the field ONLY if: [ap_tnfa] = '1'	TNF- $\alpha$ inhibitors(variable to be decommissioned following data migration)	checkbox 1 ap_tnfa_typ__1 Infliximab/Remicaid 2 ap_tnfa_typ__2 Etanercept  Field Annotation: @READONLY
1155	ap_inflixdt1 Show the field ONLY if: [ap_tnfa_typ(1)] = '1'	Infliximab 1 - start date (variable to be decommissioned following data migration)	text (date_ymd) Custom alignment: RH Field Annotation: @READONLY @HIDEBUTTON @PLACEHOLDER = 'start date'
1156	ap_inflixdose1 Show the field ONLY if: [ap_inflixdt1] <> "	Infliximab 1 - dose (mg/kg) (variable to be decommissioned following data migration)	text (number) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'mg/kg' @READONLY
1157	ap_inflixdt2 Show the field ONLY if: [ap_inflixdt1] <> "	Infliximab 2 - start date (variable to be decommissioned following data migration)	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start date' @READONLY



1158	ap_inflixdose2 Show the field ONLY if: [ap_inflixdt2] <> "	Infliximab 2 - dose (mg/kg) (variable to be decommissioned following data migration)	text (number) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'mg/kg' @READONLY				
1159	ap_brk Show the field ONLY if: [ap_tnfa_typ(1)] = '1' and [ap_tnfa_typ(2)] = '1'		descriptive				
1160	ap_etanerdtd1 Show the field ONLY if: [ap_tnfa_typ(2)] = '1'	Etanercept 1 - start date (variable to be decommissioned following data migration)	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start date' @READONLY				
1161	ap_etanerdose1 Show the field ONLY if: [ap_etanerdtd1] <> "	Etanercept 1 - dose (mg/kg) (variable to be decommissioned following data migration)	text (number) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'mg/kg' @READONLY				
1162	ap_etanerdtd2 Show the field ONLY if: [ap_etanerdtd1] <> "	Etanercept 2 - start date (variable to be decommissioned following data migration)	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start date' @READONLY				
1163	ap_etanerdose2 Show the field ONLY if: [ap_etanerdtd2] <> "	Etanercept 2 - dose (mg/kg) (variable to be decommissioned following data migration)	text (number) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'mg/kg' @READONLY				
1164	ap_tnftyp1 Show the field ONLY if: [ap_tnfa] = '1'	TNF-a inhibitors 1	dropdown <table border="1"> <tr> <td>1</td> <td>Infliximab/Remicaid</td> </tr> <tr> <td>2</td> <td>Etanercept</td> </tr> </table>	1	Infliximab/Remicaid	2	Etanercept
1	Infliximab/Remicaid						
2	Etanercept						
1165	ap_tnfdtd1 Show the field ONLY if: [ap_tnftyp1] <> "	TNF-a blocker 1 - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start date'				
1166	ds_ap_tnfstarttm1 Show the field ONLY if: [ap_tnftyp1] <> " and [ds_substudy] = '1'	TNF-a blocker 1 - start time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start time'				
1167	ds_ap_tnfendtd1 Show the field ONLY if: [ap_tnftyp1] <> " and [ds_substudy] = '1'	TNF-a blocker 1 - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'end date'				
1168	ds_ap_tnfendtm1 Show the field ONLY if: [ap_tnftyp1] <> " and [ds_substudy] = '1'	TNF-a blocker 1 - end time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'end time'				
1169	ap_tnfdose1 Show the field ONLY if: [ap_tnftyp1] <> "	TNF-a blocker 1 - dose (mg/kg)	text (number) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'dose (mg/kg)'				
1170	ap_tnftyp2 Show the field ONLY if: [ap_tnfa] = '1'	TNF-a inhibitors 2	dropdown <table border="1"> <tr> <td>1</td> <td>Infliximab/Remicaid</td> </tr> <tr> <td>2</td> <td>Etanercept</td> </tr> </table>	1	Infliximab/Remicaid	2	Etanercept
1	Infliximab/Remicaid						
2	Etanercept						
1171	ap_tnfdtd2 Show the field ONLY if: [ap_tnftyp2] <> "	TNF-a blocker 2 - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start date'				
1172	ds_ap_tnfstarttm2 Show the field ONLY if: [ap_tnftyp2] <> " and [ds_substudy] = '1'	TNF-a blocker 2 - start time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start time'				
1173	ds_ap_tnfendtd2 Show the field ONLY if: [ap_tnftyp2] <> " and [ds_substudy] = '1'	TNF-a blocker 2 - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'end date'				
1174	ds_ap_tnfendtm2 Show the field ONLY if: [ap_tnftyp2] <> " and [ds_substudy] = '1'	TNF-a blocker 2 - end time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'end time'				

1175	ap_tnfdose2 Show the field ONLY if: [ap_tnftyp2] <> "	TNF-a blocker 2 - dose (mg/kg)	text (number) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'dose (mg/kg)'				
1176	ap_tnftyp3 Show the field ONLY if: [ap_tnfa] = '1'	TNF-a inhibitors 3	dropdown <table border="1"> <tr> <td>1</td> <td>Infliximab/Remicaid</td> </tr> <tr> <td>2</td> <td>Etanercept</td> </tr> </table>	1	Infliximab/Remicaid	2	Etanercept
1	Infliximab/Remicaid						
2	Etanercept						
1177	ap_tnfdt3 Show the field ONLY if: [ap_tnftyp3] <> "	TNF-a blocker 3 - start date	text (date_yumd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start date'				
1178	ds_ap_tnfstarttm3 Show the field ONLY if: [ap_tnftyp3] <> " and [ds_substudy] = '1'	TNF-a blocker 3 - start time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start time'				
1179	ds_ap_tnfenddt3 Show the field ONLY if: [ap_tnftyp3] <> " and [ds_substudy] = '1'	TNF-a blocker 3 - end date	text (date_yumd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'end date'				
1180	ds_ap_tnfendtm3 Show the field ONLY if: [ap_tnftyp3] <> " and [ds_substudy] = '1'	TNF-a blocker 3 - end time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'end time'				
1181	ap_tnfdose3 Show the field ONLY if: [ap_tnftyp3] <> "	TNF-a blocker 3 - dose (mg/kg)	text (number) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'dose (mg/kg)'				
1182	ap_tnfdose4 Show the field ONLY if: [ap_tnftyp4] <> "	TNF-a blocker 4 - dose (mg/kg)	text (number) Custom alignment: RH Field Annotation: @PLACEHOLDER = 'dose (mg/kg)'				
1183	ap_tnfdt4 Show the field ONLY if: [ap_tnftyp4] <> "	TNF-a blocker 4 - date	text (date_yumd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start date'				
1184	ds_ap_tnfstarttm4 Show the field ONLY if: [ap_tnftyp4] <> " and [ds_substudy] = '1'	TNF-a blocker 4 - start time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start time'				
1185	ds_ap_tnfenddt4 Show the field ONLY if: [ap_tnftyp4] <> " and [ds_substudy] = '1'	TNF-a blocker 4 - end date	text (date_yumd) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'end date'				
1186	ap_tnftyp4 Show the field ONLY if: [ap_tnfa] = '1'	TNF-a inhibitors 4	dropdown <table border="1"> <tr> <td>1</td> <td>Infliximab/Remicaid</td> </tr> <tr> <td>2</td> <td>Etanercept</td> </tr> </table>	1	Infliximab/Remicaid	2	Etanercept
1	Infliximab/Remicaid						
2	Etanercept						
1187	ds_ap_tnfendtm4 Show the field ONLY if: [ap_tnftyp4] <> " and [ds_substudy] = '1'	TNF-a blocker 4 - end time	text (time) Custom alignment: RH Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'end time'				
1188	emb_1021 Show the field ONLY if: [ap_tnfa] = '1' and [ds_substudy] = '1'	Agent Dose Start/End date Start/End time	descriptive				
1189	emb_1024 Show the field ONLY if: [ap_tnfa] = '1' and [ds_substudy] = '0'	Agent Dose Start date	descriptive				
1190	emb_1031	{ap_tnftyp1} {ap_tnfdose1} {ap_tnfdt1} {ds_ap_tnfstarttm1} {ds_ap_tnfenddt1} {ds_ap_tnfendtm1} {ap_tnftyp2} {ap_tnfdose2} {ap_tnfdt2} {ds_ap_tnfstarttm2} {ds_ap_tnfenddt2} {ds_ap_tnfendtm2} {ap_tnftyp3} {ap_tnfdose3} {ap_tnfdt3} {ds_ap_tnfstarttm3} {ds_ap_tnfenddt3} {ds_ap_tnfendtm3} {ap_tnftyp4} {ap_tnfdose4} {ap_tnfdt4} {ds_ap_tnfstarttm4} {ds_ap_tnfenddt4} {ds_ap_tnfendtm4}	descriptive				

Repeated for steroid dose 1-5

1191	ap_il	Section Header: SECTION: OTHER IMMUNOSUPPRESSANTS Interleukin pathway blockers <i>Sarilumab, Tocilizumab, Siltuximab</i>	yesno 1 Yes 0 No
1192	ap_il_typ Show the field ONLY if: [ap_il] = '1'	IL blockers: agents(check all that apply)	checkbox 1 ap_il_typ__1 Sarilumab 2 ap_il_typ__2 Tocilizumab 3 ap_il_typ__3 Siltuximab 4 ap_il_typ__4 Anakinra
1193	ap_jak	Janus Kinase (JAK) inhibitors <i>Baricitinib, Ruxolitinib, Fedratinib, Tofacitinib</i>	yesno 1 Yes 0 No
1194	ap_jak_typ Show the field ONLY if: [ap_jak] = '1'	JAK inhibitors: type(check all that apply)	checkbox 1 ap_jak_typ__1 Baricitinib 2 ap_jak_typ__2 Ruxolitinib 3 ap_jak_typ__3 Fedratinib 4 ap_jak_typ__4 Tofacitinib
1195	ap_ifn	Interferon Alpha	yesno 1 Yes 0 No
1196	ds_immunother	Other immunosuppressive	yesno 1 Yes 0 No
1197	ds_immunotherx Show the field ONLY if: [ds_immunother] = '1'	if other immunosuppressive, provide details	notes
1198	ap_antivir	Section Header: SECTION: ANTIVIRALS Anitvirals <i>Remdesivir (GS-5735) or another protease inhibitor, Lopinavir/Ritonavir, Sofosbuvir, Osetamivir, Baloxavir, Ribavirin</i>	yesno 1 Yes 0 No
1199	ap_antivir_typ Show the field ONLY if: [ap_antivir] = '1'	Anitvirals: type(check all that apply)	checkbox 1 ap_antivir_typ__1 Remdesivir (GS-5735) or another protease inhibitor 2 ap_antivir_typ__2 Lopinavir/Ritonavir 3 ap_antivir_typ__3 Sofosbuvir 4 ap_antivir_typ__4 Osetamivir 5 ap_antivir_typ__5 Baloxavir 6 ap_antivir_typ__6 Ribavirin 7 ap_antivir_typ__7 Monoclonal antibodies 8 ap_antivir_typ__8 COVID plasma
1200	ap_nsaid	Section Header: SECTION: NSAIDs NSAIDs	yesno 1 Yes 0 No
1201	ap_nsaid_typ Show the field ONLY if: [ap_nsaid] = '1'	NSAIDs: type(check all that apply) <i>Ibuprofen, Naproxen, Celecoxib, Diclofenac, Indomethacin, Piroxicam Do not enter "Aspirin" here. For Aspirin, please complete the dose table above.</i>	checkbox 1 ap_nsaid_typ__1 Ibuprofen 3 ap_nsaid_typ__3 Naproxen 4 ap_nsaid_typ__4 Celecoxib 5 ap_nsaid_typ__5 Diclofenac 6 ap_nsaid_typ__6 Indomethacin 7 ap_nsaid_typ__7 Piroxicam 8 ap_nsaid_typ__8 Ketorolac
1202	ds_ap_nsaidstartdt Show the field ONLY if: [ap_nsaid] = '1' and [ds_substud y] = '1'	NSAIDs start date	text (date_ymd) Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start date'

1203	ds_ap_antibiopre Show the field ONLY if: [ds_substudy] = '1'	Section Header: SECTION: CHEMOTHERAPEUTICS/ANTIBIOTICS Antibiotics at admission but prescribed before admission	yesno <table border="1" style="width: 100%;"> <tr> <td style="width: 20px; text-align: center;">1</td> <td style="width: 50px;">Yes</td> </tr> <tr> <td style="text-align: center;">0</td> <td>No</td> </tr> </table>	1	Yes	0	No																																																																							
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0	No																																																																													
1204	ds_ap_antibiopretyp Show the field ONLY if: [ds_ap_antibiopre] = '1'	Type of antibiotics before admission(select all that apply)	checkbox <table border="1" style="width: 100%;"> <tr><td style="width: 20px; text-align: center;">1</td><td style="width: 50px;">ds_ap_antibiopretyp__1</td><td style="width: 50%;">Beta-lactams (Penicillin G/Cepl</td></tr> <tr><td style="text-align: center;">2</td><td>ds_ap_antibiopretyp__2</td><td>Semisyntheticbeta-lactams (Ampicillin/Amoxicillin)</td></tr> <tr><td style="text-align: center;">3</td><td>ds_ap_antibiopretyp__3</td><td>Clavulanic Acid (Clavulin)</td></tr> <tr><td style="text-align: center;">4</td><td>ds_ap_antibiopretyp__4</td><td>Monobactams (Aztreonam)</td></tr> <tr><td style="text-align: center;">5</td><td>ds_ap_antibiopretyp__5</td><td>Carboxypenems (Imipenem)</td></tr> <tr><td style="text-align: center;">6</td><td>ds_ap_antibiopretyp__6</td><td>Aminoglycosides (Streptomyci</td></tr> <tr><td style="text-align: center;">7</td><td>ds_ap_antibiopretyp__7</td><td>Aminoglycosides (Gentamicin)</td></tr> <tr><td style="text-align: center;">8</td><td>ds_ap_antibiopretyp__8</td><td>Glycopeptides (Vancomycin)</td></tr> <tr><td style="text-align: center;">9</td><td>ds_ap_antibiopretyp__9</td><td>Lincomycins (Clindamycin)</td></tr> <tr><td style="text-align: center;">10</td><td>ds_ap_antibiopretyp__10</td><td>Macrolides (Erythromycin/Azithromycin)</td></tr> <tr><td style="text-align: center;">11</td><td>ds_ap_antibiopretyp__11</td><td>Polypeptides (Polymyxin)</td></tr> <tr><td style="text-align: center;">12</td><td>ds_ap_antibiopretyp__12</td><td>Polypeptides (Bacitracin)</td></tr> <tr><td style="text-align: center;">13</td><td>ds_ap_antibiopretyp__13</td><td>Polyenes (Amphotericin)</td></tr> <tr><td style="text-align: center;">14</td><td>ds_ap_antibiopretyp__14</td><td>Polyenes (Nystatin)</td></tr> <tr><td style="text-align: center;">15</td><td>ds_ap_antibiopretyp__15</td><td>Rifamycins (Rifampicin)</td></tr> <tr><td style="text-align: center;">16</td><td>ds_ap_antibiopretyp__16</td><td>Tetracyclines (Tetracycline)</td></tr> <tr><td style="text-align: center;">17</td><td>ds_ap_antibiopretyp__17</td><td>Semisynthetic tetracycline (Dox</td></tr> <tr><td style="text-align: center;">18</td><td>ds_ap_antibiopretyp__18</td><td>Chloramphenicol (Chloramph</td></tr> <tr><td style="text-align: center;">19</td><td>ds_ap_antibiopretyp__19</td><td>Quinolones (Nalidixic acid)</td></tr> <tr><td style="text-align: center;">20</td><td>ds_ap_antibiopretyp__20</td><td>Fluoroquinolones (Ciprofloxac</td></tr> <tr><td style="text-align: center;">21</td><td>ds_ap_antibiopretyp__21</td><td>Growth factor analogs (Sulfanilamide/Gantrisin/Trime</td></tr> <tr><td style="text-align: center;">22</td><td>ds_ap_antibiopretyp__22</td><td>Growth factor analogs (Isoniaz</td></tr> <tr><td style="text-align: center;">23</td><td>ds_ap_antibiopretyp__23</td><td>Growth factor analogs (PAS)</td></tr> <tr><td style="text-align: center;">24</td><td>ds_ap_antibiopretyp__24</td><td>Other</td></tr> <tr><td style="text-align: center;">25</td><td>ds_ap_antibiopretyp__25</td><td>Unknown</td></tr> </table>	1	ds_ap_antibiopretyp__1	Beta-lactams (Penicillin G/Cepl	2	ds_ap_antibiopretyp__2	Semisyntheticbeta-lactams (Ampicillin/Amoxicillin)	3	ds_ap_antibiopretyp__3	Clavulanic Acid (Clavulin)	4	ds_ap_antibiopretyp__4	Monobactams (Aztreonam)	5	ds_ap_antibiopretyp__5	Carboxypenems (Imipenem)	6	ds_ap_antibiopretyp__6	Aminoglycosides (Streptomyci	7	ds_ap_antibiopretyp__7	Aminoglycosides (Gentamicin)	8	ds_ap_antibiopretyp__8	Glycopeptides (Vancomycin)	9	ds_ap_antibiopretyp__9	Lincomycins (Clindamycin)	10	ds_ap_antibiopretyp__10	Macrolides (Erythromycin/Azithromycin)	11	ds_ap_antibiopretyp__11	Polypeptides (Polymyxin)	12	ds_ap_antibiopretyp__12	Polypeptides (Bacitracin)	13	ds_ap_antibiopretyp__13	Polyenes (Amphotericin)	14	ds_ap_antibiopretyp__14	Polyenes (Nystatin)	15	ds_ap_antibiopretyp__15	Rifamycins (Rifampicin)	16	ds_ap_antibiopretyp__16	Tetracyclines (Tetracycline)	17	ds_ap_antibiopretyp__17	Semisynthetic tetracycline (Dox	18	ds_ap_antibiopretyp__18	Chloramphenicol (Chloramph	19	ds_ap_antibiopretyp__19	Quinolones (Nalidixic acid)	20	ds_ap_antibiopretyp__20	Fluoroquinolones (Ciprofloxac	21	ds_ap_antibiopretyp__21	Growth factor analogs (Sulfanilamide/Gantrisin/Trime	22	ds_ap_antibiopretyp__22	Growth factor analogs (Isoniaz	23	ds_ap_antibiopretyp__23	Growth factor analogs (PAS)	24	ds_ap_antibiopretyp__24	Other	25	ds_ap_antibiopretyp__25	Unknown
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1206	dsc_ap_antibiopretypx Show the field ONLY if: [ds_ap_antibiopretyp(24)] = '1'	if other antibiotics prescribed before admission, please specify	text																																																																											
1207	ds_ap_antibiopost Show the field ONLY if: [ds_substudy] = '1'	Antibiotics prescribed after admission	yesno <table border="1" style="width: 100%;"> <tr> <td style="width: 20px; text-align: center;">1</td> <td style="width: 50px;">Yes</td> </tr> <tr> <td style="text-align: center;">0</td> <td>No</td> </tr> </table>	1	Yes	0	No																																																																							
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1208	ds_ap_antibiopostdtm Show the field ONLY if: [ds_ap_antibiopost] = '1'	Date antibiotics started after admission	text (date_ymd) Field Annotation: @HIDEBUTTON																																																																											

1209	ds_ap_antibioposttyp Show the field ONLY if: [ds_ap_antibiopost] = '1'	Type of antibiotics prescribed after admission(select all that apply)	checkbox <table border="1"> <tr><td>1</td><td>ds_ap_antibioposttyp__1</td><td>Beta-lactams (Penicillin G/Cef</td></tr> <tr><td>2</td><td>ds_ap_antibioposttyp__2</td><td>Semisyntheticbeta-lactams (Ampicillin/Amoxicillin)</td></tr> <tr><td>3</td><td>ds_ap_antibioposttyp__3</td><td>Clavulanic Acid (Clavulin)</td></tr> <tr><td>4</td><td>ds_ap_antibioposttyp__4</td><td>Monobactams (Aztreonam)</td></tr> <tr><td>5</td><td>ds_ap_antibioposttyp__5</td><td>Carboxypenems (Imipenem)</td></tr> <tr><td>6</td><td>ds_ap_antibioposttyp__6</td><td>Aminoglycosides (Streptomyc</td></tr> <tr><td>7</td><td>ds_ap_antibioposttyp__7</td><td>Aminoglycosides (Gentamicin</td></tr> <tr><td>8</td><td>ds_ap_antibioposttyp__8</td><td>Glycopeptides (Vancomycin)</td></tr> <tr><td>9</td><td>ds_ap_antibioposttyp__9</td><td>Lincomycins (Clindamycin)</td></tr> <tr><td>10</td><td>ds_ap_antibioposttyp__10</td><td>Macrolides (Erythromycin/Azithromycin)</td></tr> <tr><td>11</td><td>ds_ap_antibioposttyp__11</td><td>Polypeptides (Polymyxin)</td></tr> <tr><td>12</td><td>ds_ap_antibioposttyp__12</td><td>Polypeptides (Bacitracin)</td></tr> <tr><td>13</td><td>ds_ap_antibioposttyp__13</td><td>Polyenes (Amphotericin)</td></tr> <tr><td>14</td><td>ds_ap_antibioposttyp__14</td><td>Polyenes (Nystatin)</td></tr> <tr><td>15</td><td>ds_ap_antibioposttyp__15</td><td>Rifamycins (Rifampicin)</td></tr> <tr><td>16</td><td>ds_ap_antibioposttyp__16</td><td>Tetracyclines (Tetracycline)</td></tr> <tr><td>17</td><td>ds_ap_antibioposttyp__17</td><td>Semisynthetic tetracycline (Do</td></tr> <tr><td>18</td><td>ds_ap_antibioposttyp__18</td><td>Chloramphenicol (Chloramph</td></tr> <tr><td>19</td><td>ds_ap_antibioposttyp__19</td><td>Quinolones (Nalidixic acid)</td></tr> <tr><td>20</td><td>ds_ap_antibioposttyp__20</td><td>Fluoroquinolones (Ciproflox</td></tr> <tr><td>21</td><td>ds_ap_antibioposttyp__21</td><td>Growth factor analogs (Sulfanilamide/Gantrisin/Tripr</td></tr> <tr><td>22</td><td>ds_ap_antibioposttyp__22</td><td>Growth factor analogs (Isonia</td></tr> <tr><td>23</td><td>ds_ap_antibioposttyp__23</td><td>Growth factor analogs (PAS)</td></tr> <tr><td>24</td><td>ds_ap_antibioposttyp__24</td><td>Other</td></tr> <tr><td>25</td><td>ds_ap_antibioposttyp__25</td><td>Unknown</td></tr> </table>	1	ds_ap_antibioposttyp__1	Beta-lactams (Penicillin G/Cef	2	ds_ap_antibioposttyp__2	Semisyntheticbeta-lactams (Ampicillin/Amoxicillin)	3	ds_ap_antibioposttyp__3	Clavulanic Acid (Clavulin)	4	ds_ap_antibioposttyp__4	Monobactams (Aztreonam)	5	ds_ap_antibioposttyp__5	Carboxypenems (Imipenem)	6	ds_ap_antibioposttyp__6	Aminoglycosides (Streptomyc	7	ds_ap_antibioposttyp__7	Aminoglycosides (Gentamicin	8	ds_ap_antibioposttyp__8	Glycopeptides (Vancomycin)	9	ds_ap_antibioposttyp__9	Lincomycins (Clindamycin)	10	ds_ap_antibioposttyp__10	Macrolides (Erythromycin/Azithromycin)	11	ds_ap_antibioposttyp__11	Polypeptides (Polymyxin)	12	ds_ap_antibioposttyp__12	Polypeptides (Bacitracin)	13	ds_ap_antibioposttyp__13	Polyenes (Amphotericin)	14	ds_ap_antibioposttyp__14	Polyenes (Nystatin)	15	ds_ap_antibioposttyp__15	Rifamycins (Rifampicin)	16	ds_ap_antibioposttyp__16	Tetracyclines (Tetracycline)	17	ds_ap_antibioposttyp__17	Semisynthetic tetracycline (Do	18	ds_ap_antibioposttyp__18	Chloramphenicol (Chloramph	19	ds_ap_antibioposttyp__19	Quinolones (Nalidixic acid)	20	ds_ap_antibioposttyp__20	Fluoroquinolones (Ciproflox	21	ds_ap_antibioposttyp__21	Growth factor analogs (Sulfanilamide/Gantrisin/Tripr	22	ds_ap_antibioposttyp__22	Growth factor analogs (Isonia	23	ds_ap_antibioposttyp__23	Growth factor analogs (PAS)	24	ds_ap_antibioposttyp__24	Other	25	ds_ap_antibioposttyp__25	Unknown
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1210	ds_ap_antibioposttyp24 Show the field ONLY if: [ds_ap_antibioposttyp(24)] = '1'	if other antibiotics prescribed after admission, please specify	text																																																																											
1211	ap_abcixi	Section Header: SECTION: OTHER MEDS USED DURING ACUTE EPISODE Abciximab	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table>	1	Yes	0	No																																																																							
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1212	ds_ap_abcixidt Show the field ONLY if: [ap_abcixi] = '1' and [ds_substudy] = '1'	Abciximab date	text (date_ymd) Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'date'																																																																											
1213	ap_pentox	Pentoxifylline/Trental	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table>	1	Yes	0	No																																																																							
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1214	ds_ap_pentoxdt Show the field ONLY if: [ap_pentox] = '1' and [ds_substudy] = '1'	Pentoxifylline/Trental start date	text (date_ymd) Field Annotation: @HIDEBUTTON @PLACEHOLDER = 'start date'																																																																											
1215	ap_ino	Inotropes Epinephrine, Dopamine, Dobutamine, Vasopressin	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table>	1	Yes	0	No																																																																							
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1217	brk0401		descriptive						
1218	ap_medother	Other cardiac support meds during acute treatment	yesno <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>No</td> </tr> </table>	1	Yes	0	No		
1	Yes								
0	No								
1219	ap_medother1 Show the field ONLY if: [ap_medother] = '1'	Other cardiac med 1 - Name	text Custom alignment: RH						
1220	ap_medother2 Show the field ONLY if: [ap_medother] = '1'	Other cardiac med 2 - Name	text Custom alignment: RH						
1221	ap_mednotes	Acute care medication comments (if any)	notes						
1222	acute_phase_meds_complete	Section Header: <i>Form Status</i> Complete?	dropdown <table border="1"> <tr> <td>0</td> <td>Incomplete</td> </tr> <tr> <td>1</td> <td>Unverified</td> </tr> <tr> <td>2</td> <td>Complete</td> </tr> </table>	0	Incomplete	1	Unverified	2	Complete
0	Incomplete								
1	Unverified								
2	Complete								

Instrument: **Follow-up meds** (followup\_meds)

[^ Collapse](#)

1223	brk32	Section Header: <i>FOLLOW-UP MEDICATIONS Medications used during follow-up/long-term period</i>	descriptive				
1224	fp_asa	Section Header: <i>SECTION: ANTITHROMBOTICS</i> Aspirin	yesno <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>No</td> </tr> </table>	1	Yes	0	No
1	Yes						
0	No						
1225	fp_asastdt Show the field ONLY if: [fp_asa] = '1'	Aspirin - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1226	fp_asalfu Show the field ONLY if: [fp_asa] = '1'	Aspirin at last follow-up <i>if discontinued at follow-up, select "No" and put the date below</i>	yesno <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>No</td> </tr> </table>	1	Yes	0	No
1	Yes						
0	No						
1227	fp_asaendt Show the field ONLY if: [fp_asalfu] = '0'	Aspirin - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1228	fp_brk35 Show the field ONLY if: [fp_asa] = '1'		descriptive				
1229	fp_clopid	Clopidogrel/Plavix	yesno <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>No</td> </tr> </table>	1	Yes	0	No
1	Yes						
0	No						
1230	fp_clopidstdt Show the field ONLY if: [fp_clopid] = '1'	Clopidogrel - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1231	fp_clopidlfu Show the field ONLY if: [fp_clopid] = '1'	Clopidogrel at last follow-up <i>if discontinued at follow-up, select "No" and put the date below</i>	yesno <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>No</td> </tr> </table>	1	Yes	0	No
1	Yes						
0	No						
1232	fp_clopidendt Show the field ONLY if: [fp_clopidlfu] = '0'	Clopidogrel - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1233	fp_brk36 Show the field ONLY if: [fp_clopid] = '1'		descriptive				
1234	fp_dipyrid	Dipyridamole/Persantine	yesno <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>No</td> </tr> </table>	1	Yes	0	No
1	Yes						
0	No						
1235	fp_dipyridstdt Show the field ONLY if: [fp_dipyrid] = '1'	Dipyridamole - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				

1236	fp_dipyridfu Show the field ONLY if: [fp_dipyrid] = '1'	Dipyridamole at last follow-up <i>if discontinued at follow-up, select "No" and put the date below</i>	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
1237	fp_dipyridendt Show the field ONLY if: [fp_dipyridfu] = '0'	Dipyridamole - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1238	fp_brk37 Show the field ONLY if: [fp_dipyrid] = '1'		descriptive				
1239	fp_enox	LMWH - Enxaparin	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
1240	fp_enoxstdt1 Show the field ONLY if: [fp_enox] = '1'	Enoxaparin 1 - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1241	fp_enoxenddt1 Show the field ONLY if: [fp_enoxstdt1] <> "	Enoxaparin 1 - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1242	fp_enoxstdt2 Show the field ONLY if: [fp_enoxstdt1] <> "	Enoxaparin 2 - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1243	fp_enoxenddt2 Show the field ONLY if: [fp_enoxstdt2] <> "	Enoxaparin 2 - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1244	fp_enoxstdt3 Show the field ONLY if: [fp_enoxstdt2] <> "	Enoxaparin 3 - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1245	fp_enoxdtend3 Show the field ONLY if: [fp_enoxstdt3] <> "	Enoxaparin 3 - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1246	fp_enoxlfu Show the field ONLY if: [fp_enox] = '1'	Enoxaparin at last follow-up <i>if discontinued at follow-up, select "No" and put the date below</i>	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
1247	fp_enoxenddt Show the field ONLY if: [fp_enoxlfu] = '0'	Enoxaparin - end date	text (date_ymd) Field Annotation: @HIDEBUTTON				
1248	fp_brk39 Show the field ONLY if: [fp_enox] = '1'		descriptive				
1249	fp_tinza	LMWH - Tinzaparin	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
1250	fp_tinzastdt1 Show the field ONLY if: [fp_tinza] = '1'	Tinzaparin 1 - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1251	fp_tinzaenddt1 Show the field ONLY if: [fp_tinzastdt1] <> "	Tinzaparin 1 - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1252	fp_tinzastdt2 Show the field ONLY if: [fp_tinzastdt1] <> "	Tinzaparin 2 - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1253	fp_tinzaenddt2 Show the field ONLY if: [fp_tinzastdt2] <> "	Tinzaparin 2 - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1254	fp_tinzastdt3 Show the field ONLY if: [fp_tinzastdt2] <> "	Tinzaparin 3 - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				

1255	fp_tinzaendt3 Show the field ONLY if: [fp_tinzastdt3] <> "	Tinzaparin 3 - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1256	fp_tinzalfu Show the field ONLY if: [fp_tinza] = '1'	Tinzaparin at last follow-up <i>if discontinued at follow-up, select "No" and put the date below</i>	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
1257	fp_tinzaenddt Show the field ONLY if: [fp_tinzalfu] = '0'	Tinzaparin - end date	text (date_ymd) Field Annotation: @HIDEBUTTON				
1258	fp_brk40 Show the field ONLY if: [fp_tinza] = '1'		descriptive				
1259	fp_warf	Warfarin/Coumadin	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
1260	fp_warfst1 Show the field ONLY if: [fp_warf] = '1'	Warfarin 1 - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1261	fp_warfenddt1 Show the field ONLY if: [fp_warfst1] <> "	Warfarin 1 - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1262	fp_warflead1 Show the field ONLY if: [fp_warfst1] <> "	Warfarin 1 - heparin (UFH/LMWH) lead-in	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table> Custom alignment: RH	1	Yes	0	No
1	Yes						
0	No						
1263	fp_warfst2 Show the field ONLY if: [fp_warfst1] <> "	Warfarin 2 - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1264	fp_warfenddt2 Show the field ONLY if: [fp_warfst2] <> "	Warfarin 2 - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1265	fp_warflead2 Show the field ONLY if: [fp_warfst2] <> "	Warfarin 2 - heparin (UFH/LMWH) lead-in	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table> Custom alignment: RH	1	Yes	0	No
1	Yes						
0	No						
1266	fp_warfst3 Show the field ONLY if: [fp_warfst2] <> "	Warfarin 3 - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1267	fp_warfenddt3 Show the field ONLY if: [fp_warfst3] <> "	Warfarin 3 - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON				
1268	fp_warflead3 Show the field ONLY if: [fp_warfst3] <> "	Warfarin 3 - heparin (UFH/LMWH) lead-in	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table> Custom alignment: RH	1	Yes	0	No
1	Yes						
0	No						
1269	fp_warffu Show the field ONLY if: [fp_warf] = '1'	Warfarin at last follow-up <i>if discontinued at follow-up, select "No" and put the date below</i>	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No
1	Yes						
0	No						
1270	fp_warfenddt Show the field ONLY if: [fp_warffu] = '0'	Warfarin - end date	text (date_ymd) Field Annotation: @HIDEBUTTON				
1271	fp_brk41 Show the field ONLY if: [fp_warf] = '1'		descriptive				



1272	fp_doac	DOACs <i>DOACs: dabigatran, rivaroxaban, apixaban, edoxaban</i>	yesno 1 Yes 0 No
1273	fp_doacstdt Show the field ONLY if: [fp_doac] = '1'	DOACs - start date	text (date_ymd) Field Annotation: @HIDEBUTTON
1274	fp_doacifu Show the field ONLY if: [fp_doac] = '1'	DOACs at last follow-up <i>if discontinued at follow-up, select "No" and put the date below</i>	yesno 1 Yes 0 No
1275	fp_doacenddt Show the field ONLY if: [fp_doacifu] = '0'	DOACs - end date	text (date_ymd) Field Annotation: @HIDEBUTTON
1276	fp_statin	Section Header: SECTION: LONG-TERM TREATMENT Statins	yesno 1 Yes 0 No
1277	fp_statinstdt Show the field ONLY if: [fp_statin] = '1'	Statins - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON
1278	fp_statinifu Show the field ONLY if: [fp_statin] = '1'	Statin at last follow-up <i>if discontinued at follow-up, select "No" and put the date below</i>	yesno 1 Yes 0 No
1279	fp_statinenddt Show the field ONLY if: [fp_statinifu] = '0'	Statins - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON
1280	fp_bblock	Beta blockers	yesno 1 Yes 0 No
1281	fp_bblockstdt Show the field ONLY if: [fp_bblock] = '1'	Beta blockers - start date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON
1282	fp_bblockifu Show the field ONLY if: [fp_bblock] = '1'	Beta blocker at last follow-up <i>if discontinued at follow-up, select "No" and put the date below</i>	yesno 1 Yes 0 No
1283	fp_bblockenddt Show the field ONLY if: [fp_bblockifu] = '0'	Beta blockers - end date	text (date_ymd) Custom alignment: RH Field Annotation: @HIDEBUTTON
1284	fp_mednotes	Follow-up/Long term medication comments (if any)	notes
1285	followup_meds_complete	Section Header: Form Status Complete?	dropdown 0 Incomplete 1 Unverified 2 Complete

Instrument: **Follow-up status** (followup\_status)

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1286	crfdt	Section Header: FOLLOW-UP STATUS Date of follow-up data entry <i>if you are completing this form today, please enter today's date</i>	text (date_ymd)
1287	clinic	Patient still followed at KD clinic <i>for acute patients, select "no"</i>	yesno 1 Yes 0 No
1288	lfudt	Date of last follow-up	text (date_ymd)
1289	dischargersn Show the field ONLY if: [clinic] = '0'	Discharge from KD clinic reason	radio 1 No follow-up needed 2 Transferred to other hospital 3 Lost to follow-up 5 Transition to adult care 4 Death

1290	dischargehosp Show the field ONLY if: [dischargersn] = '2'	Name of the other hospital	text
1291	status	Current status	radio 0 Deceased 1 Alive 2 Unknown
1292	deathrsn Show the field ONLY if: [status] = '0'	Primary cause of death	radio 1 Cardiac MI 2 Cardiac other 3 Non-cardiac
1293	deathtx Show the field ONLY if: [status] = '0'	Please provide details on cause of death	notes
1294	fupnotes	Other relevant information	notes Custom alignment: RH
1295	followup_status_complete	Section Header: <i>Form Status</i> Complete?	dropdown 0 Incomplete 1 Unverified 2 Complete

Instrument: **Complications Thrombi** (complications\_thrombi)[^ Collapse](#)

1296	brk0701	Section Header: <i>COMPLICATIONS: THROMBOSIS</i>	descriptive
1297	thromb	Section Header: <i>SECTION: CORONARY ARTERY (CA) THROMBOSIS</i> Coronary artery thrombosis	radio 1 Yes 0 No
1298	thrombdt1 Show the field ONLY if: [thromb] = '1'	CA thrombus 1 - date of diagnosis	text (date_ymd) Custom alignment: RH
1299	thrombloc1 Show the field ONLY if: [thrombdt1] <> "	Thrombus 1: location	text Custom alignment: RH
1300	thrombdeg1 Show the field ONLY if: [thrombdt1] <> "	Thrombus 1: worst degree of occlusion	radio 1 Non-occlusive 2 Partially occlusive 3 Occlusive
1301	thrombsym1 Show the field ONLY if: [thrombdt1] <> "	Thrombus 1: associated symptom	radio 1 Symptomatic 2 Asymptomatic
1302	thrombsymtx1 Show the field ONLY if: [thrombsym1] = '1'	Thrombus 1: if symptomatic, please specify	notes Custom alignment: RH
1303	thrombrx1 Show the field ONLY if: [thrombdt1] <> "	Thrombus 1: treatment	notes Field Annotation: @PLACEHOLDER = 'what agents the thrombus treated with'
1304	thrombdt2 Show the field ONLY if: [thrombdt1] <> "	CA thrombus 2: date of diagnosis	text (date_ymd) Custom alignment: RH
1305	thrombloc2 Show the field ONLY if: [thrombdt2] <> "	Thrombus 2: location	text Custom alignment: RH
1306	thrombdeg2 Show the field ONLY if: [thrombdt2] <> "	Thrombus 2: worst degree of occlusion	radio 1 Non-occlusive 2 Partially occlusive 3 Occlusive

1307	thrombsym2 Show the field ONLY if: [thrombdt2] <> "	Thrombus 2: associated symptom	radio 1 Symptomatic 2 Asymptomatic
1308	thrombsymtx2 Show the field ONLY if: [thrombsym2] = '1'	Thrombus 2: if symptomatic, please specify	notes Custom alignment: RH
1309	thrombrx2 Show the field ONLY if: [thrombdt2] <> "	Thrombus 2: treatment	notes Field Annotation: @PLACEHOLDER = 'what agents the thrombus treated with'
1310	thrombdt3 Show the field ONLY if: [thrombdt2] <> "	CA thrombus 3: date of diagnosis	text (date_ymd) Custom alignment: RH
1311	thrombloc3 Show the field ONLY if: [thrombdt3] <> "	Thrombus 3: location	text Custom alignment: RH
1312	thrombdeg3 Show the field ONLY if: [thrombdt3] <> "	Thrombus 3: worst degree of occlusion	radio 1 Non-occlusive 2 Partially occlusive 3 Occlusive
1313	thrombsym3 Show the field ONLY if: [thrombdt3] <> "	Thrombus 3: associated symptom	radio 1 Symptomatic 2 Asymptomatic
1314	thrombsymtx3 Show the field ONLY if: [thrombsym3] = '1'	Thrombus 3: if symptomatic, please specify	notes Custom alignment: RH
1315	thrombrx3 Show the field ONLY if: [thrombdt3] <> "	Thrombus 3: treatment	notes Field Annotation: @PLACEHOLDER = 'what agents the thrombus treated with'
1316	thrombdt4 Show the field ONLY if: [thrombdt3] <> "	CA thrombus 4 - date of diagnosis	text (date_ymd) Custom alignment: RH
1317	thrombloc4 Show the field ONLY if: [thrombdt4] <> "	Thrombus 4: location	text Custom alignment: RH
1318	thrombdeg4 Show the field ONLY if: [thrombdt4] <> "	Thrombus 4: worst degree of occlusion	radio 1 Non-occlusive 2 Partially occlusive 3 Occlusive
1319	thrombsym4 Show the field ONLY if: [thrombdt4] <> "	Thrombus 4: associated symptom	radio 1 Symptomatic 2 Asymptomatic
1320	thrombsymtx4 Show the field ONLY if: [thrombsym4] = '4'	Thrombus 4: if symptomatic, please specify	notes Custom alignment: RH
1321	thrombrx4 Show the field ONLY if: [thrombdt4] <> "	Thrombus 4: treatment	notes Field Annotation: @PLACEHOLDER = 'what agents the thrombus treated with'
1322	thrombdt5 Show the field ONLY if: [thrombdt4] <> "	CA thrombus 5: date of diagnosis	text (date_ymd) Custom alignment: RH
1323	thrombloc5 Show the field ONLY if: [thrombdt5] <> "	Thrombus 5: location	text Custom alignment: RH
1324	thrombdeg5 Show the field ONLY if: [thrombdt5] <> "	Thrombus 5: worst degree of occlusion	radio 1 Non-occlusive 2 Partially occlusive 3 Occlusive

1325	thrombsym5 Show the field ONLY if: [thrombdt5] <> "	Thrombus 5: associated symptom	radio 1 Symptomatic 2 Asymptomatic
1326	thrombsymtx5 Show the field ONLY if: [thrombsym5] = '4'	Thrombus 5: if symptomatic, please specify	notes Custom alignment: RH
1327	thrombrx5 Show the field ONLY if: [thrombdt5] <> "	Thrombus 5: treatment	notes Field Annotation: @PLACEHOLDER = 'what agents the thrombus treated with'
1328	thrombdt6 Show the field ONLY if: [thrombdt5] <> "	CA thrombus 6: date of diagnosis	text (date_ymd) Custom alignment: RH
1329	thrombloc6 Show the field ONLY if: [thrombdt6] <> "	Thrombus 6: location	text Custom alignment: RH
1330	thrombdeg6 Show the field ONLY if: [thrombdt6] <> "	Thrombus 6: worst degree of occlusion	radio 1 Non-occlusive 2 Partially occlusive 3 Occlusive
1331	thrombsym6 Show the field ONLY if: [thrombdt6] <> "	Thrombus 6: associated symptom	radio 1 Symptomatic 2 Asymptomatic
1332	thrombsymtx6 Show the field ONLY if: [thrombsym6] = '4'	Thrombus 6: if symptomatic, please specify	notes Custom alignment: RH
1333	thrombrx6 Show the field ONLY if: [thrombdt6] <> "	Thrombus 6: treatment	notes Field Annotation: @PLACEHOLDER = 'what agents the thrombus treated with'
1334	thrombdt7 Show the field ONLY if: [thrombdt6] <> "	CA thrombus 7 - date of diagnosis	text (date_ymd) Custom alignment: RH
1335	thrombloc7 Show the field ONLY if: [thrombdt7] <> "	Thrombus 7: location	text Custom alignment: RH
1336	thrombdeg7 Show the field ONLY if: [thrombdt7] <> "	Thrombus 7: worst degree of occlusion	radio 1 Non-occlusive 2 Partially occlusive 3 Occlusive
1337	thrombsym7 Show the field ONLY if: [thrombdt7] <> "	Thrombus 7: associated symptom	radio 1 Symptomatic 2 Asymptomatic
1338	thrombsymtx7 Show the field ONLY if: [thrombsym7] = '7'	Thrombus 7: if symptomatic, please specify	notes Custom alignment: RH
1339	thrombrx7 Show the field ONLY if: [thrombdt7] <> "	Thrombus 7: treatment	notes Field Annotation: @PLACEHOLDER = 'what agents the thrombus treated with'
1340	thrombdt8 Show the field ONLY if: [thrombdt7] <> "	CA thrombus 8: date of diagnosis	text (date_ymd) Custom alignment: RH
1341	thrombloc8 Show the field ONLY if: [thrombdt8] <> "	Thrombus 8: location	text Custom alignment: RH
1342	thrombdeg8 Show the field ONLY if: [thrombdt8] <> "	Thrombus 8: worst degree of occlusion	radio 1 Non-occlusive 2 Partially occlusive 3 Occlusive

1343	thrombsym8 Show the field ONLY if: [thrombdt8] <> "	Thrombus 8: associated symptom	radio 1 Symptomatic 2 Asymptomatic
1344	thrombsymtx8 Show the field ONLY if: [thrombsym8] = '7'	Thrombus 8: if symptomatic, please specify	notes Custom alignment: RH
1345	thrombrx8 Show the field ONLY if: [thrombdt8] <> "	Thrombus 8: treatment	notes Field Annotation: @PLACEHOLDER = 'what agents the thrombus treated with'
1346	thrombdt9 Show the field ONLY if: [thrombdt8] <> "	CA thrombus 9: date of diagnosis	text (date_ymd) Custom alignment: RH
1347	thrombloc9 Show the field ONLY if: [thrombdt9] <> "	Thrombus 9: location	text Custom alignment: RH
1348	thrombdeg9 Show the field ONLY if: [thrombdt9] <> "	Thrombus 9: worst degree of occlusion	radio 1 Non-occlusive 2 Partially occlusive 3 Occlusive
1349	thrombsym9 Show the field ONLY if: [thrombdt9] <> "	Thrombus 9: associated symptom	radio 1 Symptomatic 2 Asymptomatic
1350	thrombsymtx9 Show the field ONLY if: [thrombsym9] = '7'	Thrombus 9: if symptomatic, please specify	notes Custom alignment: RH
1351	thrombrx9 Show the field ONLY if: [thrombdt9] <> "	Thrombus 9: treatment	notes Field Annotation: @PLACEHOLDER = 'what agents the thrombus treated with'
1352	complications_thrombi_complete	Section Header: <i>Form Status</i> Complete?	dropdown 0 Incomplete 1 Unverified 2 Complete

Instrument: **Complications Other** (complications\_other)[^ Collapse](#)

1353	brk0801	Section Header: <i>COMPLICATIONS: OTHER</i>	descriptive
1354	rcasteno	Section Header: <i>SECTION: CORONARY ARTERY STENOSIS</i> RCA stenosis <i>RCA: Right coronary artery</i>	radio 1 Yes 0 No
1355	rcastenodt1 Show the field ONLY if: [rcasteno] = '1'	RCA stenosis 1: date	text (date_ymd)
1356	rcastenodt2 Show the field ONLY if: [rcastenodt1] <> "	RCA stenosis 2: date	text (date_ymd)
1357	rcastenodt3 Show the field ONLY if: [rcastenodt2] <> "	RCA stenosis 3: date	text (date_ymd)
1358	lcasteno	LCA stenosis <i>LCA: Left main coronary artery</i>	radio 1 Yes 0 No
1359	lcastenodt1 Show the field ONLY if: [lcasteno] = '1'	LCA stenosis 1: date	text (date_ymd)
1360	lcastenodt2 Show the field ONLY if: [lcastenodt1] <> "	LCA stenosis 2: date	text (date_ymd)

1361	lcastenodt3 Show the field ONLY if: [lcastenodt2] <> "	LCA stenosis 3: date	text (date_ymd)
1362	ladsteno	LAD stenosis <i>LAD: Left anterior descending artery</i>	radio 1 Yes 0 No
1363	ladstenodt1 Show the field ONLY if: [ladsteno] = '1'	LAD stenosis 1: date	text (date_ymd)
1364	ladstenodt2 Show the field ONLY if: [ladstenodt1] <> "	LAD stenosis 2: date	text (date_ymd)
1365	ladstenodt3 Show the field ONLY if: [ladstenodt2] <> "	LAD stenosis 3: date	text (date_ymd)
1366	ccxsteno	CCX/LCX stenosis <i>CCX/LCX: Coronary circumflex artery</i>	radio 1 Yes 0 No
1367	ccxstenodt1 Show the field ONLY if: [ccxsteno] = '1'	CCX stenosis 1: date	text (date_ymd)
1368	ccxstenodt2 Show the field ONLY if: [ccxstenodt1] <> "	CCX stenosis 2: date	text (date_ymd)
1369	ccxstenodt3 Show the field ONLY if: [ccxstenodt2] <> "	CCX stenosis 3: date	text (date_ymd)
1370	cath	Section Header: <i>SECTION: CATH + CABG + MI</i> Catheter revascularization	radio 1 Yes 0 No
1371	cathdt1 Show the field ONLY if: [cath] = '1'	Cath 1: date	text (date_ymd) Custom alignment: RH
1372	cath_desc1 Show the field ONLY if: [cathdt1] <> "	Cath 1: description	notes
1373	cathdt2 Show the field ONLY if: [cathdt1] <> "	Cath 2: date	text (date_ymd) Custom alignment: RH
1374	cath_desc2 Show the field ONLY if: [cathdt2] <> "	Cath 2: description	notes
1375	cathdt3 Show the field ONLY if: [cathdt2] <> "	Cath 3: date	text (date_ymd) Custom alignment: RH
1376	cath_desc3 Show the field ONLY if: [cathdt3] <> "	Cath 3: description	notes
1377	cabg	CABG	radio 1 Yes 0 No
1378	cabgdt1 Show the field ONLY if: [cabg] = '1'	CABG 1: date	text (date_ymd) Custom alignment: RH
1379	cabg_desc1 Show the field ONLY if: [cabgdt1] <> "	CABG 1: description	notes

1380	cabgdt2 Show the field ONLY if: [cabgdt1] <> "	CABG 2: date	text (date_ymd) Custom alignment: RH						
1381	cabg_desc2 Show the field ONLY if: [cabgdt2] <> "	CABG 2: description	notes						
1382	cabgdt3 Show the field ONLY if: [cabgdt2] <> "	CABG 3: date	text (date_ymd) Custom alignment: RH						
1383	cabg_desc3 Show the field ONLY if: [cabgdt3] <> "	CABG 3: description	notes						
1384	mi	Myocardial infarction	radio <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>No</td> </tr> </table>	1	Yes	0	No		
1	Yes								
0	No								
1385	midt1 Show the field ONLY if: [mi] = '1'	MI 1: date	text (date_ymd) Custom alignment: RH						
1386	midt2 Show the field ONLY if: [midt1] <> "	MI 2: date	text (date_ymd) Custom alignment: RH						
1387	midt3 Show the field ONLY if: [midt2] <> "	MI 3: date	text (date_ymd) Custom alignment: RH						
1388	bleeding	Section Header: <i>SECTION: BLEEDING ASSOCIATED WITH ANTICOAGULATION</i> Any bleeding event associated with anticoagulation <i>Do not include bleeding complications associated with a surgical/catheter procedures. Only include major complications. Major complications: Intracranial hemorrhage, pulmonary hemorrhage, hematuria, intraocular, retroperitoneal and/or gastrointestinal bleed or any bleeding episode requiring blood transfusion are classified as major bleeding complications.</i>	radio <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>None at last follow-up</td> </tr> </table>	1	Yes	0	None at last follow-up		
1	Yes								
0	None at last follow-up								
1389	bleedingdt1 Show the field ONLY if: [bleeding] = '1'	Bleeding event 1: date of diagnosis	text (date_ymd) Custom alignment: RH						
1390	bleedingnotes1 Show the field ONLY if: [bleedingdt1] <> "	Bleeding event 1: notes	notes Custom alignment: RH						
1391	bleedingdt2 Show the field ONLY if: [bleedingdt1] <> "	Bleeding event 2: date of diagnosis	text (date_ymd) Custom alignment: RH						
1392	bleedingnotes2 Show the field ONLY if: [bleedingdt2] <> "	Bleeding event 2: notes	notes Custom alignment: RH						
1393	bleedingdt3 Show the field ONLY if: [bleedingdt2] <> "	Bleeding event 3: date of diagnosis	text (date_ymd) Custom alignment: RH						
1394	bleedingnotes3 Show the field ONLY if: [bleedingdt3] <> "	Bleeding event 3: notes	notes Custom alignment: RH						
1395	skok	Section Header: <i>NAKDR VERIFICATION</i> SickKids verified <i>For NAKDR staff only</i>	yesno <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>No</td> </tr> </table> Custom alignment: RH	1	Yes	0	No		
1	Yes								
0	No								
1396	complications_other_complete	Section Header: <i>Form Status</i> Complete?	dropdown <table border="1"> <tr> <td>0</td> <td>Incomplete</td> </tr> <tr> <td>1</td> <td>Unverified</td> </tr> <tr> <td>2</td> <td>Complete</td> </tr> </table>	0	Incomplete	1	Unverified	2	Complete
0	Incomplete								
1	Unverified								
2	Complete								
Instrument: <b>File upload</b> (file_upload)			<a href="#">^ Collapse</a>						
1397	brk5	Section Header: <i>FILE UPLOAD</i>	descriptive						

1398	brk6	<p>FILE UPLOAD INSTRUCTIONS:</p> <p>GENERAL INSTRUCTIONS: All reports must be deidentified before the upload. Include research ID on each page. File can be in any document or image format and cannot exceed 32MB in size. The file is viewable following upload. In case of incorrect file upload or report with identifiers, you can always remove the file. Please combine reports with multiple pages (e.g., page 1 &amp; 2 of the same echo report should be merged into 1 file). Use option 1 or 2 (see below) below to upload the reports.</p> <p>Option 1: ~ Use any file merge tool (e.g., Adobe Professional) to combine all available reports (echo, angio, MRI) ~ Upload the single file using the link below to attach your file.</p> <p>Option 2: ~ Use the Echo 01-20 slots to upload echocardiography reports. File can be in any document or image format. ~ Use the Angio 01-10 slots to upload angiogram reports. File can be in any document or image format. ~ Use the MRI 01-05 slots to upload cardiac MRI reports. File can be in any document or image format.</p>	descriptive
1399	brk7		descriptive
1400	allfileupload	<p>Section Header: <i>OPTION 1 : COMBINED FILE UPLOAD</i></p> <p>Attach de-identified copies of ALL of the following reports(Please combine all reports and prepare one file to upload) Serial echocardiography Serial cardiac MRI Serial angiography Serial labwork Serial blood culture Serial ECGs (including rhythm strips) Holter on follow-up Note: DCC to abstract the serial values (from the files uploaded by the sites) for future analysis purposes</p>	file
1401	brk8		descriptive
1402	fileuploadecho01	<p>Section Header: <i>OPTION 2 : ECHOCARDIOGRAM REPORT UPLOAD</i></p> <p>ECHO 01 REPORT: Attach de-identified copies of report here</p>	file
1403	fileuploadecho02 Show the field ONLY if: [fileuploadecho01] != "	ECHO 02 REPORT: Attach de-identified copies of report here	file
1404	fileuploadecho03 Show the field ONLY if: [fileuploadecho02] != "	ECHO 03 REPORT: Attach de-identified copies of report here	file
1405	fileuploadecho04 Show the field ONLY if: [fileuploadecho03] != "	ECHO 04 REPORT: Attach de-identified copies of report here	file
1406	fileuploadecho05 Show the field ONLY if: [fileuploadecho04] != "	ECHO 05 REPORT: Attach de-identified copies of report here	file
1407	fileuploadecho06 Show the field ONLY if: [fileuploadecho05] != "	ECHO 06 REPORT: Attach de-identified copies of report here	file
1408	fileuploadecho07 Show the field ONLY if: [fileuploadecho06] != "	ECHO 07 REPORT: Attach de-identified copies of report here	file
1409	fileuploadecho08 Show the field ONLY if: [fileuploadecho07] != "	ECHO 08 REPORT: Attach de-identified copies of report here	file
1410	fileuploadecho09 Show the field ONLY if: [fileuploadecho08] != "	ECHO 09 REPORT: Attach de-identified copies of report here	file
1411	fileuploadecho10 Show the field ONLY if: [fileuploadecho09] != "	ECHO 10 REPORT: Attach de-identified copies of report here	file
1412	fileuploadecho11 Show the field ONLY if: [fileuploadecho10] != "	ECHO 11 REPORT: Attach de-identified copies of report here	file



1413	fileuploadecho12 Show the field ONLY if: [fileuploadecho11] != "	ECHO 12 REPORT: Attach de-identified copies of report here	file
1414	fileuploadecho13 Show the field ONLY if: [fileuploadecho12] != "	ECHO 13 REPORT: Attach de-identified copies of report here	file
1415	fileuploadecho14 Show the field ONLY if: [fileuploadecho13] != "	ECHO 14 REPORT: Attach de-identified copies of report here	file
1416	fileuploadecho15 Show the field ONLY if: [fileuploadecho14] != "	ECHO 15 REPORT: Attach de-identified copies of report here	file
1417	fileuploadecho16 Show the field ONLY if: [fileuploadecho15] != "	ECHO 16 REPORT: Attach de-identified copies of report here	file
1418	fileuploadecho17 Show the field ONLY if: [fileuploadecho16] != "	ECHO 17 REPORT: Attach de-identified copies of report here	file
1419	fileuploadecho18 Show the field ONLY if: [fileuploadecho17] != "	ECHO 18 REPORT: Attach de-identified copies of report here	file
1420	fileuploadecho19 Show the field ONLY if: [fileuploadecho18] != "	ECHO 19 REPORT: Attach de-identified copies of report here	file
1421	fileuploadecho20 Show the field ONLY if: [fileuploadecho19] != "	ECHO 20 REPORT: Attach de-identified copies of report here	file
1422	fileuploadecho21 Show the field ONLY if: [fileuploadecho20] != "	ECHO 21 REPORT: Attach de-identified copies of report here	file
1423	fileuploadecho22 Show the field ONLY if: [fileuploadecho21] != "	ECHO 22 REPORT: Attach de-identified copies of report here	file
1424	fileuploadecho23 Show the field ONLY if: [fileuploadecho22] != "	ECHO 23 REPORT: Attach de-identified copies of report here	file
1425	fileuploadecho24 Show the field ONLY if: [fileuploadecho23] != "	ECHO 24 REPORT: Attach de-identified copies of report here	file
1426	fileuploadecho25 Show the field ONLY if: [fileuploadecho24] != "	ECHO 25 REPORT: Attach de-identified copies of report here	file
1427	fileuploadecho26 Show the field ONLY if: [fileuploadecho25] != "	ECHO 26 REPORT: Attach de-identified copies of report here	file
1428	fileuploadecho27 Show the field ONLY if: [fileuploadecho26] != "	ECHO 27 REPORT: Attach de-identified copies of report here	file
1429	fileuploadecho28 Show the field ONLY if: [fileuploadecho27] != "	ECHO 28 REPORT: Attach de-identified copies of report here	file
1430	fileuploadecho29 Show the field ONLY if: [fileuploadecho28] != "	ECHO 29 REPORT: Attach de-identified copies of report here	file
1431	fileuploadecho30 Show the field ONLY if: [fileuploadecho29] != "	ECHO 30 REPORT: Attach de-identified copies of report here	file

1432	fileuploadecho31 Show the field ONLY if: [fileuploadecho30] != ""	ECHO 31 REPORT: Attach de-identified copies of report here	file
1433	fileuploadecho32 Show the field ONLY if: [fileuploadecho31] != ""	ECHO 32 REPORT: Attach de-identified copies of report here	file
1434	fileuploadecho33 Show the field ONLY if: [fileuploadecho32] != ""	ECHO 33 REPORT: Attach de-identified copies of report here	file
1435	fileuploadecho34 Show the field ONLY if: [fileuploadecho33] != ""	ECHO 34 REPORT: Attach de-identified copies of report here	file
1436	fileuploadecho35 Show the field ONLY if: [fileuploadecho34] != ""	ECHO 35 REPORT: Attach de-identified copies of report here	file
1437	fileuploadecho36 Show the field ONLY if: [fileuploadecho35] != ""	ECHO 36 REPORT: Attach de-identified copies of report here	file
1438	fileuploadecho37 Show the field ONLY if: [fileuploadecho36] != ""	ECHO 37 REPORT: Attach de-identified copies of report here	file
1439	fileuploadecho38 Show the field ONLY if: [fileuploadecho37] != ""	ECHO 38 REPORT: Attach de-identified copies of report here	file
1440	fileuploadecho39 Show the field ONLY if: [fileuploadecho38] != ""	ECHO 39 REPORT: Attach de-identified copies of report here	file
1441	fileuploadecho40 Show the field ONLY if: [fileuploadecho39] != ""	ECHO 40 REPORT: Attach de-identified copies of report here	file
1442	brk9		descriptive
1443	fileuploadangio01	Section Header: <i>OPTION 2 : CORONARY ANGIOGRAM REPORT UPLOAD</i> ANGIO 01 REPORT: Attach de-identified copies of report here	file
1444	fileuploadangio02 Show the field ONLY if: [fileuploadangio01] != ""	ANGIO 02 REPORT: Attach de-identified copies of report here	file
1445	fileuploadangio03 Show the field ONLY if: [fileuploadangio02] != ""	ANGIO 03 REPORT: Attach de-identified copies of report here	file
1446	fileuploadangio04 Show the field ONLY if: [fileuploadangio03] != ""	ANGIO 04 REPORT: Attach de-identified copies of report here	file
1447	fileuploadangio05 Show the field ONLY if: [fileuploadangio04] != ""	ANGIO 05 REPORT: Attach de-identified copies of report here	file
1448	fileuploadangio06 Show the field ONLY if: [fileuploadangio05] != ""	ANGIO 06 REPORT: Attach de-identified copies of report here	file
1449	fileuploadangio07 Show the field ONLY if: [fileuploadangio06] != ""	ANGIO 07 REPORT: Attach de-identified copies of report here	file
1450	fileuploadangio08 Show the field ONLY if: [fileuploadangio07] != ""	ANGIO 08 REPORT: Attach de-identified copies of report here	file
1451	fileuploadangio09 Show the field ONLY if: [fileuploadangio08] != ""	ANGIO 09 REPORT: Attach de-identified copies of report here	file

1452	fileuploadangio10 Show the field ONLY if: [fileuploadangio09] != ""	ANGIO 10 REPORT: Attach de-identified copies of report here	file
1453	fileuploadangio11 Show the field ONLY if: [fileuploadangio10] != ""	ANGIO 11 REPORT: Attach de-identified copies of report here	file
1454	fileuploadangio12 Show the field ONLY if: [fileuploadangio11] != ""	ANGIO 12 REPORT: Attach de-identified copies of report here	file
1455	fileuploadangio13 Show the field ONLY if: [fileuploadangio12] != ""	ANGIO 13 REPORT: Attach de-identified copies of report here	file
1456	fileuploadangio14 Show the field ONLY if: [fileuploadangio13] != ""	ANGIO 14 REPORT: Attach de-identified copies of report here	file
1457	fileuploadangio15 Show the field ONLY if: [fileuploadangio14] != ""	ANGIO 15 REPORT: Attach de-identified copies of report here	file
1458	fileuploadangio16 Show the field ONLY if: [fileuploadangio15] != ""	ANGIO 16 REPORT: Attach de-identified copies of report here	file
1459	fileuploadangio17 Show the field ONLY if: [fileuploadangio16] != ""	ANGIO 17 REPORT: Attach de-identified copies of report here	file
1460	fileuploadangio18 Show the field ONLY if: [fileuploadangio17] != ""	ANGIO 18 REPORT: Attach de-identified copies of report here	file
1461	fileuploadangio19 Show the field ONLY if: [fileuploadangio18] != ""	ANGIO 19 REPORT: Attach de-identified copies of report here	file
1462	fileuploadangio20 Show the field ONLY if: [fileuploadangio19] != ""	ANGIO 20 REPORT: Attach de-identified copies of report here	file
1463	brk10		descriptive
1464	fileuploadmri01	Section Header: <i>OPTION 2 : MRI REPORT UPLOAD</i> MRI 01 REPORT: Attach de-identified copies of report here	file
1465	fileuploadmri02 Show the field ONLY if: [fileuploadmri01] != ""	MRI 02 REPORT: Attach de-identified copies of report here	file
1466	fileuploadmri03 Show the field ONLY if: [fileuploadmri02] != ""	MRI 03 REPORT: Attach de-identified copies of report here	file
1467	fileuploadmri04 Show the field ONLY if: [fileuploadmri03] != ""	MRI 04 REPORT: Attach de-identified copies of report here	file
1468	fileuploadmri05 Show the field ONLY if: [fileuploadmri04] != ""	MRI 05 REPORT: Attach de-identified copies of report here	file
1469	fileuploadmri06 Show the field ONLY if: [fileuploadmri05] != ""	MRI 06 REPORT: Attach de-identified copies of report here	file
1470	fileuploadmri07 Show the field ONLY if: [fileuploadmri06] != ""	MRI 07 REPORT: Attach de-identified copies of report here	file
1471	fileuploadmri08 Show the field ONLY if: [fileuploadmri07] != ""	MRI 08 REPORT: Attach de-identified copies of report here	file

1472	fileuploadmri09 Show the field ONLY if: [fileuploadmri08] != ""	MRI 09 REPORT: Attach de-identified copies of report here	file
1473	fileuploadmri10 Show the field ONLY if: [fileuploadmri09] != ""	MRI 10 REPORT: Attach de-identified copies of report here	file
1474	fileuploadmri11 Show the field ONLY if: [fileuploadmri10] != ""	MRI 11 REPORT: Attach de-identified copies of report here	file
1475	fileuploadmri12 Show the field ONLY if: [fileuploadmri11] != ""	MRI 12 REPORT: Attach de-identified copies of report here	file
1476	fileuploadmri13 Show the field ONLY if: [fileuploadmri12] != ""	MRI 13 REPORT: Attach de-identified copies of report here	file
1477	fileuploadmri14 Show the field ONLY if: [fileuploadmri13] != ""	MRI 14 REPORT: Attach de-identified copies of report here	file
1478	fileuploadmri15 Show the field ONLY if: [fileuploadmri14] != ""	MRI 15 REPORT: Attach de-identified copies of report here	file
1479	fileuploadmri16 Show the field ONLY if: [fileuploadmri15] != ""	MRI 16 REPORT: Attach de-identified copies of report here	file
1480	fileuploadmri17 Show the field ONLY if: [fileuploadmri16] != ""	MRI 17 REPORT: Attach de-identified copies of report here	file
1481	fileuploadmri18 Show the field ONLY if: [fileuploadmri17] != ""	MRI 18 REPORT: Attach de-identified copies of report here	file
1482	fileuploadmri19 Show the field ONLY if: [fileuploadmri18] != ""	MRI 19 REPORT: Attach de-identified copies of report here	file
1483	fileuploadmri20 Show the field ONLY if: [fileuploadmri19] != ""	MRI 20 REPORT: Attach de-identified copies of report here	file
1484	brk11		descriptive
1485	fileuploadecg01	Section Header: <i>OPTION 2 : SERIAL EKG UPLOAD</i> ECG 01 STRIP & REPORT: Attach de-identified copies	file
1486	fileuploadecg02 Show the field ONLY if: [fileuploadecg01] != ""	ECG 02 STRIP & REPORT: Attach de-identified copies	file
1487	fileuploadecg03 Show the field ONLY if: [fileuploadecg02] != ""	ECG 03 STRIP & REPORT: Attach de-identified copies	file
1488	fileuploadecg04 Show the field ONLY if: [fileuploadecg03] != ""	ECG 04 STRIP & REPORT: Attach de-identified copies	file
1489	fileuploadecg05 Show the field ONLY if: [fileuploadecg04] != ""	ECG 05 STRIP & REPORT: Attach de-identified copies	file
1490	fileuploadecg06 Show the field ONLY if: [fileuploadecg05] != ""	ECG 06 STRIP & REPORT: Attach de-identified copies	file
1491	fileuploadecg07 Show the field ONLY if: [fileuploadecg06] != ""	ECG 07 STRIP & REPORT: Attach de-identified copies	file

1492	fileuploaddecg08 Show the field ONLY if: [fileuploaddecg07] != ""	ECG 08 STRIP & REPORT: Attach de-identified copies	file						
1493	fileuploaddecg09 Show the field ONLY if: [fileuploaddecg08] != ""	ECG 09 STRIP & REPORT: Attach de-identified copies	file						
1494	fileuploadlab01	Section Header: <i>OPTION 2 : SERIAL LABWORK</i> LABWORK 01: Attach de-identified spreadsheet/PDF	file						
1495	fileuploadlab02	LABWORK 02: Attach de-identified spreadsheet/PDF	file						
1496	fileuploadlab03	LABWORK 03: Attach de-identified spreadsheet/PDF	file						
1497	fileuploadlab04	LABWORK 04: Attach de-identified spreadsheet/PDF	file						
1498	fileuploadlab05	LABWORK 05: Attach de-identified spreadsheet/PDF	file						
1499	fileuploadlab06	LABWORK 06: Attach de-identified spreadsheet/PDF	file						
1500	fileuploadlab07	LABWORK 07: Attach de-identified spreadsheet/PDF	file						
1501	fileuploadlab08	LABWORK 08: Attach de-identified spreadsheet/PDF	file						
1502	fileuploadlab09	LABWORK 09: Attach de-identified spreadsheet/PDF	file						
1503	fileuploadlab10	LABWORK 10: Attach de-identified spreadsheet/PDF	file						
1504	fileuploadlab11	LABWORK 11: Attach de-identified spreadsheet/PDF	file						
1505	fileuploadculture01	Section Header: <i>OPTION 2 : SERIAL CULTURE</i> Serial CULTURE 01: Attach de-identified spreadsheet/PDF	file						
1506	fileuploadculture02	Serial CULTURE 02: Attach de-identified spreadsheet/PDF	file						
1507	fileuploadculture03	Serial CULTURE 03: Attach de-identified spreadsheet/PDF	file						
1508	fileuploadculture04	Serial CULTURE 04: Attach de-identified spreadsheet/PDF	file						
1509	fileuploadculture05	Serial CULTURE 05: Attach de-identified spreadsheet/PDF	file						
1510	fileuploadculture06	Serial CULTURE 06: Attach de-identified spreadsheet/PDF	file						
1511	fileuploadculture07	Serial CULTURE 07: Attach de-identified spreadsheet/PDF	file						
1512	fileuploadculture08	Serial CULTURE 08: Attach de-identified spreadsheet/PDF	file						
1513	fileuploadculture09	Serial CULTURE 09: Attach de-identified spreadsheet/PDF	file						
1514	fileuploadculture10	Serial CULTURE 10: Attach de-identified spreadsheet/PDF	file						
1515	fileuploadculture11	Serial CULTURE 11: Attach de-identified spreadsheet/PDF	file						
1516	fileuploadpanel01	Section Header: <i>OPTION 2 : SERIAL VIRAL PANEL</i> Serial Panel 01: Attach de-identified spreadsheet/PDF	file						
1517	fileuploadpanel02	Serial Panel 02: Attach de-identified spreadsheet/PDF	file						
1518	fileuploadpanel03	Serial Panel 03: Attach de-identified spreadsheet/PDF	file						
1519	fileuploadpanel04	Serial Panel 04: Attach de-identified spreadsheet/PDF	file						
1520	fileuploadpanel05	Serial Panel 05: Attach de-identified spreadsheet/PDF	file						
1521	fileuploadpanel06	Serial Panel 06: Attach de-identified spreadsheet/PDF	file						
1522	fileuploadpanel07	Serial Panel 07: Attach de-identified spreadsheet/PDF	file						
1523	fileuploadpanel08	Serial Panel 08: Attach de-identified spreadsheet/PDF	file						
1524	fileuploadpanel09	Serial Panel 09: Attach de-identified spreadsheet/PDF	file						
1525	fileuploadpanel10	Serial Panel 10: Attach de-identified spreadsheet/PDF	file						
1526	fileuploadpanel11	Serial Panel 11: Attach de-identified spreadsheet/PDF	file						
1527	fileuploadholter01	Section Header: <i>OPTION 2 : FOLLOW-UP HOLTER</i> Follow-up Holter: Attach de-identified report	file						
1528	file_upload_complete	Section Header: <i>Form Status</i> Complete?	dropdown <table border="1"> <tr> <td>0</td> <td>Incomplete</td> </tr> <tr> <td>1</td> <td>Unverified</td> </tr> <tr> <td>2</td> <td>Complete</td> </tr> </table>	0	Incomplete	1	Unverified	2	Complete
0	Incomplete								
1	Unverified								
2	Complete								
Instrument: <b>Adjudication</b> (adjudication) <span style="float: right;"><a href="#">^ Collapse</a></span>									
1529	brk15	Section Header: <i>NAKDR ADJUDICATION: CORONARY ARTERY INVOLVEMENT (SUBMITTING SITE: PLEASE LEAVE SECTION BLANK - TO BE COMPLETED BY NAKDR STAFF FOR ALL PATIENTS THAT ARE SUBMITTED IN THE REGISTRY)</i>	descriptive						
1530	adj_caadt	Section Header: <i>SECTION: CORONARY ARTERY INVOLVEMENT SUMMARY</i> Adjudication date	text (date_ymd)						
1531	time0dt	Date of entry	text (date_ymd)						

1532	cca_lv	CAA involvement summary	dropdown 1 DLTN 2 NGCAA 3 GCAA
1533	caamaxz	Maximum CAA z-score	text (number, Min: -5, Max: 80)
1534	rca_lv	RCA involvement	dropdown 1 NOINVL 2 DLTN 3 NGCAA 4 GCAA
1535	rcamaxz	Maximum RCA z-score	text (number, Min: -5, Max: 80)
1536	lca_lv	LCA involvement	dropdown 1 NOINVL 2 DLTN 3 NGCAA 4 GCAA
1537	lcamaxz	Maximum LCA z-score	text (number, Min: -5, Max: 80)
1538	lad_lv	LAD involvement	dropdown 1 NOINVL 2 DLTN 3 NGCAA 4 GCAA
1539	ladmaxz	Maximum LAD z-score	text (number, Min: -5, Max: 80)
1540	ccx_lv	CCX involvement	dropdown 1 NOINVL 2 DLTN 3 NGCAA 4 GCAA
1541	ccxmaxz	Maximum CCX z-score	text (number, Min: -5, Max: 80)
1542	numechos	Number of echos submitted	text (integer, Min: 0, Max: 50) Custom alignment: RH
1543	numangio	Number of angio submitted	text (integer, Min: 0, Max: 50) Custom alignment: RH
1544	nummri	Number of MRI submitted	text (integer, Min: 0, Max: 50) Custom alignment: RH
1545	numct	Number of CT submitted	text (integer, Min: 0, Max: 50) Custom alignment: RH
1546	regstatus	Patient's enrollment in the registry (to be completed by SK staff only)	radio, Required 1 Included 2 Excluded
1547	exclusionrsn Show the field ONLY if: [regstatus] = '2'	Reason for exclusion	radio, Required 1 Diagnosed before Jan 1, 1999 2 No CAA 3 Limited information 4 No imaging
1548	brk66		descriptive
1549	adj_thro	Section Header: SECTION: THROMBUS Thrombosis	yesno 1 Yes 0 No
1550	adj_throbrk		descriptive
1551	adj_throm1loc Show the field ONLY if: [adj_thro] = '1'	Thrombus 1: Location	dropdown 1 RCA 2 LMCA 3 LAD 4 LCX/CCX

1552	adj_throm1dt Show the field ONLY if: [adj_thro] = '1'	Thrombus 1: Date first identified	text (date_ymd)								
1553	adj_throm1occ Show the field ONLY if: [adj_thro] = '1'	Thrombus 1: Worst level of occlusion	dropdown <table border="1"> <tr><td>1</td><td>Non-occlusive</td></tr> <tr><td>2</td><td>Partial occlusion</td></tr> <tr><td>3</td><td>Complete occlusion</td></tr> </table>	1	Non-occlusive	2	Partial occlusion	3	Complete occlusion		
1	Non-occlusive										
2	Partial occlusion										
3	Complete occlusion										
1554	adj_throm1occdt Show the field ONLY if: [adj_thro] = '1'	Thrombus 1: Date of worst level of occlusion	text (date_ymd)								
1555	adj_throm1proc Show the field ONLY if: [adj_thro] = '1'	Thrombus 1: Procedure related event	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1556	adj_throm1sym Show the field ONLY if: [adj_thro] = '1'	Thrombus 1: Symptomatic	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1557	adj_throm1res Show the field ONLY if: [adj_thro] = '1'	Thrombus 1: Resolved	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1558	adj_throm1resdt Show the field ONLY if: [adj_throm1res] = '1'	Thrombus 1: Date of resolution	text (date_ymd)								
1559	adj_throm1notes Show the field ONLY if: [adj_thro] = '1'	Thrombus 1: Notes	notes Custom alignment: RH								
1560	adj_throm1brk Show the field ONLY if: [adj_thro] = '1'		descriptive								
1561	adj_throm2 Show the field ONLY if: [adj_thro] = '1'	Another thrombus <i>Please treat thrombotic recurrence as new clot</i>	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1562	adj_throm2loc Show the field ONLY if: [adj_throm2] = '1'	Thrombus 2: Location	dropdown <table border="1"> <tr><td>1</td><td>RCA</td></tr> <tr><td>2</td><td>LMCA</td></tr> <tr><td>3</td><td>LAD</td></tr> <tr><td>4</td><td>LCX/CCX</td></tr> </table>	1	RCA	2	LMCA	3	LAD	4	LCX/CCX
1	RCA										
2	LMCA										
3	LAD										
4	LCX/CCX										
1563	adj_throm2dt Show the field ONLY if: [adj_throm2] = '1'	Thrombus 2: Date first identified	text (date_ymd)								
1564	adj_throm2occ Show the field ONLY if: [adj_throm2] = '1'	Thrombus 2: Worst level of occlusion	dropdown <table border="1"> <tr><td>1</td><td>Non-occlusive</td></tr> <tr><td>2</td><td>Partial occlusion</td></tr> <tr><td>3</td><td>Complete occlusion</td></tr> </table>	1	Non-occlusive	2	Partial occlusion	3	Complete occlusion		
1	Non-occlusive										
2	Partial occlusion										
3	Complete occlusion										
1565	adj_throm2occdt Show the field ONLY if: [adj_throm2] = '1'	Thrombus 2: Date of worst level of occlusion	text (date_ymd)								
1566	adj_throm2proc Show the field ONLY if: [adj_throm2] = '1'	Thrombus 2: Procedure related event	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										

1567	adj_throm2sym Show the field ONLY if: [adj_throm2] = '1'	Thrombus 2: Symptomatic	yesno 1 Yes 0 No Custom alignment: RH
1568	adj_throm2res Show the field ONLY if: [adj_throm2] = '1'	Thrombus 2: Resolved	yesno 1 Yes 0 No Custom alignment: RH
1569	adj_throm2resdt Show the field ONLY if: [adj_throm2res] = '1'	Thrombus 2: Date of resolution	text (date_ymd)
1570	adj_throm2notes Show the field ONLY if: [adj_throm2] = '1'	Thrombus 2: Notes	notes Custom alignment: RH
1571	adj_throm2brk Show the field ONLY if: [adj_throm2] = '1'		descriptive
1572	adj_throm3 Show the field ONLY if: [adj_throm2] = '1'	Another thrombus <i>Please treat thrombotic recurrence as new clot</i>	yesno 1 Yes 0 No Custom alignment: RH
1573	adj_throm3loc Show the field ONLY if: [adj_throm3] = '1'	Thrombus 3: Location	dropdown 1 RCA 2 LMCA 3 LAD 4 LCX/CCX
1574	adj_throm3dt Show the field ONLY if: [adj_throm3] = '1'	Thrombus 3: Date first identified	text (date_ymd)
1575	adj_throm3occ Show the field ONLY if: [adj_throm3] = '1'	Thrombus 3: Worst level of occlusion	dropdown 1 Non-occlusive 2 Partial occlusion 3 Complete occlusion
1576	adj_throm3occdt Show the field ONLY if: [adj_throm3] = '1'	Thrombus 3: Date of worst level of occlusion	text (date_ymd)
1577	adj_throm3proc Show the field ONLY if: [adj_throm3] = '1'	Thrombus 3: Procedure related event	yesno 1 Yes 0 No Custom alignment: RH
1578	adj_throm3sym Show the field ONLY if: [adj_throm3] = '1'	Thrombus 3: Symptomatic	yesno 1 Yes 0 No Custom alignment: RH
1579	adj_throm3res Show the field ONLY if: [adj_throm3] = '1'	Thrombus 3: Resolved	yesno 1 Yes 0 No Custom alignment: RH
1580	adj_throm3resdt Show the field ONLY if: [adj_throm3res] = '1'	Thrombus 3: Date of resolution	text (date_ymd)



1581	adj_throm3notes Show the field ONLY if: [adj_throm3] = '1'	Thrombus 3: Notes	notes Custom alignment: RH								
1582	adj_throm3brk Show the field ONLY if: [adj_throm3] = '1'		descriptive								
1583	adj_throm4 Show the field ONLY if: [adj_throm3] = '1'	Another thrombus <i>Please treat thrombotic recurrence as new clot</i>	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1584	adj_throm4loc Show the field ONLY if: [adj_throm4] = '1'	Thrombus 4: Location	dropdown <table border="1"> <tr><td>1</td><td>RCA</td></tr> <tr><td>2</td><td>LMCA</td></tr> <tr><td>3</td><td>LAD</td></tr> <tr><td>4</td><td>LCX/CCX</td></tr> </table>	1	RCA	2	LMCA	3	LAD	4	LCX/CCX
1	RCA										
2	LMCA										
3	LAD										
4	LCX/CCX										
1585	adj_throm4dt Show the field ONLY if: [adj_throm4] = '1'	Thrombus 4: Date first identified	text (date_ymd)								
1586	adj_throm4occ Show the field ONLY if: [adj_throm4] = '1'	Thrombus 4: Worst level of occlusion	dropdown <table border="1"> <tr><td>1</td><td>Non-occlusive</td></tr> <tr><td>2</td><td>Partial occlusion</td></tr> <tr><td>3</td><td>Complete occlusion</td></tr> </table>	1	Non-occlusive	2	Partial occlusion	3	Complete occlusion		
1	Non-occlusive										
2	Partial occlusion										
3	Complete occlusion										
1587	adj_throm4occdt Show the field ONLY if: [adj_throm4] = '1'	Thrombus 4: Date of worst level of occlusion	text (date_ymd)								
1588	adj_throm4proc Show the field ONLY if: [adj_throm4] = '1'	Thrombus 4: Procedure related event	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1589	adj_throm4sym Show the field ONLY if: [adj_throm4] = '1'	Thrombus 4: Symptomatic	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1590	adj_throm4res Show the field ONLY if: [adj_throm4] = '1'	Thrombus 4: Resolved	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1591	adj_throm4resdt Show the field ONLY if: [adj_throm4res] = '1'	Thrombus 4: Date of resolution	text (date_ymd)								
1592	adj_throm4notes Show the field ONLY if: [adj_throm4] = '1'	Thrombus 4: Notes	notes Custom alignment: RH								
1593	adj_throm4brk Show the field ONLY if: [adj_throm4] = '1'		descriptive								
1594	adj_throm5 Show the field ONLY if: [adj_throm4] = '1'	Another thrombus <i>Please treat thrombotic recurrence as new clot</i>	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										

1595	adj_throm5loc Show the field ONLY if: [adj_throm5] = '1'	Thrombus 5: Location	dropdown <table border="1"> <tr><td>1</td><td>RCA</td></tr> <tr><td>2</td><td>LMCA</td></tr> <tr><td>3</td><td>LAD</td></tr> <tr><td>4</td><td>LCX/CCX</td></tr> </table>	1	RCA	2	LMCA	3	LAD	4	LCX/CCX
1	RCA										
2	LMCA										
3	LAD										
4	LCX/CCX										
1596	adj_throm5dt Show the field ONLY if: [adj_throm5] = '1'	Thrombus 5: Date first identified	text (date_ymd)								
1597	adj_throm5occ Show the field ONLY if: [adj_throm5] = '1'	Thrombus 5: Worst level of occlusion	dropdown <table border="1"> <tr><td>1</td><td>Non-occlusive</td></tr> <tr><td>2</td><td>Partial occlusion</td></tr> <tr><td>3</td><td>Complete occlusion</td></tr> </table>	1	Non-occlusive	2	Partial occlusion	3	Complete occlusion		
1	Non-occlusive										
2	Partial occlusion										
3	Complete occlusion										
1598	adj_throm5occdt Show the field ONLY if: [adj_throm5] = '1'	Thrombus 5: Date of worst level of occlusion	text (date_ymd)								
1599	adj_throm5proc Show the field ONLY if: [adj_throm5] = '1'	Thrombus 5: Procedure related event	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1600	adj_throm5sym Show the field ONLY if: [adj_throm5] = '1'	Thrombus 5: Symptomatic	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1601	adj_throm5res Show the field ONLY if: [adj_throm5] = '1'	Thrombus 5: Resolved	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1602	adj_throm5resdt Show the field ONLY if: [adj_throm5res] = '1'	Thrombus 5: Date of resolution	text (date_ymd)								
1603	adj_throm5notes Show the field ONLY if: [adj_throm5] = '1'	Thrombus 5: Notes	notes Custom alignment: RH								
1604	adj_throm5brk Show the field ONLY if: [adj_throm5] = '1'		descriptive								
1605	adj_throm6 Show the field ONLY if: [adj_throm5] = '1'	Another thrombus <i>Please treat thrombotic recurrence as new clot</i>	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table> Custom alignment: RH	1	Yes	0	No				
1	Yes										
0	No										
1606	adj_throm6loc Show the field ONLY if: [adj_throm6] = '1'	Thrombus 6: Location	dropdown <table border="1"> <tr><td>1</td><td>RCA</td></tr> <tr><td>2</td><td>LMCA</td></tr> <tr><td>3</td><td>LAD</td></tr> <tr><td>4</td><td>LCX/CCX</td></tr> </table>	1	RCA	2	LMCA	3	LAD	4	LCX/CCX
1	RCA										
2	LMCA										
3	LAD										
4	LCX/CCX										
1607	adj_throm6dt Show the field ONLY if: [adj_throm6] = '1'	Thrombus 6: Date first identified	text (date_ymd)								
1608	adj_throm6occ Show the field ONLY if: [adj_throm6] = '1'	Thrombus 6: Worst level of occlusion	dropdown <table border="1"> <tr><td>1</td><td>Non-occlusive</td></tr> <tr><td>2</td><td>Partial occlusion</td></tr> <tr><td>3</td><td>Complete occlusion</td></tr> </table>	1	Non-occlusive	2	Partial occlusion	3	Complete occlusion		
1	Non-occlusive										
2	Partial occlusion										
3	Complete occlusion										

1609	adj_throm6occdt Show the field ONLY if: [adj_throm6] = '1'	Thrombus 6: Date of worst level of occlusion	text (date_ymd)
1610	adj_throm6proc Show the field ONLY if: [adj_throm6] = '1'	Thrombus 6: Procedure related event	yesno 1 Yes 0 No Custom alignment: RH
1611	adj_throm6sym Show the field ONLY if: [adj_throm6] = '1'	Thrombus 6: Symptomatic	yesno 1 Yes 0 No Custom alignment: RH
1612	adj_throm6res Show the field ONLY if: [adj_throm6] = '1'	Thrombus 6: Resolved	yesno 1 Yes 0 No Custom alignment: RH
1613	adj_throm6resdt Show the field ONLY if: [adj_throm6res] = '1'	Thrombus 6: Date of resolution	text (date_ymd)
1614	adj_throm6notes Show the field ONLY if: [adj_throm6] = '1'	Thrombus 6: Notes	notes Custom alignment: RH
1615	adj_throm6brk Show the field ONLY if: [adj_throm6] = '1'		descriptive
1616	adj_throm7 Show the field ONLY if: [adj_throm6] = '1'	Another thrombus <i>Please treat thrombotic recurrence as new clot</i>	yesno 1 Yes 0 No Custom alignment: RH
1617	adj_throm7loc Show the field ONLY if: [adj_throm7] = '1'	Thrombus 7: Location	dropdown 1 RCA 2 LMCA 3 LAD 4 LCX/CCX
1618	adj_throm7dt Show the field ONLY if: [adj_throm7] = '1'	Thrombus 7: Date first identified	text (date_ymd)
1619	adj_throm7occ Show the field ONLY if: [adj_throm7] = '1'	Thrombus 7: Worst level of occlusion	dropdown 1 Non-occlusive 2 Partial occlusion 3 Complete occlusion
1620	adj_throm7occdt Show the field ONLY if: [adj_throm7] = '1'	Thrombus 7: Date of worst level of occlusion	text (date_ymd)
1621	adj_throm7proc Show the field ONLY if: [adj_throm7] = '1'	Thrombus 7: Procedure related event	yesno 1 Yes 0 No Custom alignment: RH
1622	adj_throm7sym Show the field ONLY if: [adj_throm7] = '1'	Thrombus 7: Symptomatic	yesno 1 Yes 0 No Custom alignment: RH

1623	adj_throm7res Show the field ONLY if: [adj_throm7] = '1'	Thrombus 7: Resolved	yesno 1 Yes 0 No Custom alignment: RH
1624	adj_throm7resdt Show the field ONLY if: [adj_throm7res] = '1'	Thrombus 7: Date of resolution	text (date_ymd)
1625	adj_throm7notes Show the field ONLY if: [adj_throm7] = '1'	Thrombus 7: Notes	notes Custom alignment: RH
1626	adj_throm7brk Show the field ONLY if: [adj_throm7] = '1'		descriptive
1627	adj_throm8 Show the field ONLY if: [adj_throm7] = '1'	Another thrombus <i>Please treat thrombotic recurrence as new clot</i>	yesno 1 Yes 0 No Custom alignment: RH
1628	adj_throm8loc Show the field ONLY if: [adj_throm8] = '1'	Thrombus 8: Location	dropdown 1 RCA 2 LMCA 3 LAD 4 LCX/CCX
1629	adj_throm8dt Show the field ONLY if: [adj_throm8] = '1'	Thrombus 8: Date first identified	text (date_ymd)
1630	adj_throm8occ Show the field ONLY if: [adj_throm8] = '1'	Thrombus 8: Worst level of occlusion	dropdown 1 Non-occlusive 2 Partial occlusion 3 Complete occlusion
1631	adj_throm8occdt Show the field ONLY if: [adj_throm8] = '1'	Thrombus 8: Date of worst level of occlusion	text (date_ymd)
1632	adj_throm8proc Show the field ONLY if: [adj_throm8] = '1'	Thrombus 8: Procedure related event	yesno 1 Yes 0 No Custom alignment: RH
1633	adj_throm8sym Show the field ONLY if: [adj_throm8] = '1'	Thrombus 8: Symptomatic	yesno 1 Yes 0 No Custom alignment: RH
1634	adj_throm8res Show the field ONLY if: [adj_throm8] = '1'	Thrombus 8: Resolved	yesno 1 Yes 0 No Custom alignment: RH
1635	adj_throm8resdt Show the field ONLY if: [adj_throm8res] = '1'	Thrombus 8: Date of resolution	text (date_ymd)
1636	adj_throm8notes Show the field ONLY if: [adj_throm8] = '1'	Thrombus 8: Notes	notes Custom alignment: RH
1637	adj_throm8brk Show the field ONLY if: [adj_throm8] = '1'		descriptive

1638	adj_throm9 Show the field ONLY if: [adj_throm8] = '1'	Another thrombus <i>Please treat thrombotic recurrence as new clot</i>	yesno 1 Yes 0 No  Custom alignment: RH
1639	adj_throm9loc Show the field ONLY if: [adj_throm9] = '1'	Thrombus 9: Location	dropdown 1 RCA 2 LMCA 3 LAD 4 LCX/CCX
1640	adj_throm9dt Show the field ONLY if: [adj_throm9] = '1'	Thrombus 9: Date first identified	text (date_ymd)
1641	adj_throm9occ Show the field ONLY if: [adj_throm9] = '1'	Thrombus 9: Worst level of occlusion	dropdown 1 Non-occlusive 2 Partial occlusion 3 Complete occlusion
1642	adj_throm9occdt Show the field ONLY if: [adj_throm9] = '1'	Thrombus 9: Date of worst level of occlusion	text (date_ymd)
1643	adj_throm9proc Show the field ONLY if: [adj_throm9] = '1'	Thrombus 9: Procedure related event	yesno 1 Yes 0 No  Custom alignment: RH
1644	adj_throm9sym Show the field ONLY if: [adj_throm9] = '1'	Thrombus 9: Symptomatic	yesno 1 Yes 0 No  Custom alignment: RH
1645	adj_throm9res Show the field ONLY if: [adj_throm9] = '1'	Thrombus 9: Resolved	yesno 1 Yes 0 No  Custom alignment: RH
1646	adj_throm9resdt Show the field ONLY if: [adj_throm9res] = '1'	Thrombus 9: Date of resolution	text (date_ymd)
1647	adj_throm9notes Show the field ONLY if: [adj_throm9] = '1'	Thrombus 9: Notes	notes Custom alignment: RH
1648	adj_throm9brk Show the field ONLY if: [adj_throm9] = '1'		descriptive
1649	adj_cath	Section Header: <i>SECTION: CATHETER INTERVENTIONS</i> Cath intervention	yesno 1 Yes 0 No
1650	adj_cathbrk		descriptive
1651	adj_cathrca1 Show the field ONLY if: [adj_cath] = '1'	RCA cath interventions	yesno 1 Yes 0 No
1652	adj_cathrca1dt Show the field ONLY if: [adj_cathrca1] = '1'	RCA cath intervention 1: date	text (date_ymd)

1653	adj_cathrca1typ Show the field ONLY if: [adj_cathrca1] = '1'	RCA cath intervention 1: type <i>Angioplasty includes balloon or laser angioplastyAtherectomy includes rotational (rotablation)/extraction/directional atherectomy</i>	dropdown 1 Angioplasty without stent 2 Angioplasty with stent 3 Angioplasty with atherectomy 4 Angioplasty with stent and atherectomy
1654	adj_cathrca2 Show the field ONLY if: [adj_cathrca1] = '1'	Another RCA cath intervention	yesno 1 Yes 0 No
1655	adj_cathrca2dt Show the field ONLY if: [adj_cathrca2] = '1'	RCA cath intervention 2: date	text (date_ymd)
1656	adj_cathrca2typ Show the field ONLY if: [adj_cathrca2] = '1'	RCA cath intervention 2: type	dropdown 1 Angioplasty without stent 2 Angioplasty with stent 3 Angioplasty with atherectomy 4 Angioplasty with stent and atherectomy
1657	adj_cathrca3 Show the field ONLY if: [adj_cathrca2] = '1'	Another RCA cath intervention	yesno 1 Yes 0 No
1658	adj_cathrca3dt Show the field ONLY if: [adj_cathrca3] = '1'	RCA cath intervention 3: date	text (date_ymd)
1659	adj_cathrca3typ Show the field ONLY if: [adj_cathrca3] = '1'	RCA cath intervention 3: type	dropdown 1 Angioplasty without stent 2 Angioplasty with stent 3 Angioplasty with atherectomy 4 Angioplasty with stent and atherectomy
1660	adj_cathrcatx Show the field ONLY if: [adj_cathrca1] = '1'	RCA cath intervention: notes	notes
1661	adj_cathlca1 Show the field ONLY if: [adj_cath] = '1'	LCA cath interventions	yesno 1 Yes 0 No
1662	adj_cathlca1dt Show the field ONLY if: [adj_cathlca1] = '1'	LCA cath intervention 1: date	text (date_ymd)
1663	adj_cathlca1typ Show the field ONLY if: [adj_cathlca1] = '1'	LCA cath intervention 1: type <i>Angioplasty includes balloon or laser angioplastyAtherectomy includes rotational (rotablation)/extraction/directional atherectomy</i>	dropdown 1 Angioplasty without stent 2 Angioplasty with stent 3 Angioplasty with atherectomy 4 Angioplasty with stent and atherectomy
1664	adj_cathlca2 Show the field ONLY if: [adj_cathlca1] = '1'	Another LCA cath intervention	yesno 1 Yes 0 No
1665	adj_cathlca2dt Show the field ONLY if: [adj_cathlca2] = '1'	LCA cath intervention 2: date	text (date_ymd)
1666	adj_cathlca2typ Show the field ONLY if: [adj_cathlca2] = '1'	LCA cath intervention 2: type	dropdown 1 Angioplasty without stent 2 Angioplasty with stent 3 Angioplasty with atherectomy 4 Angioplasty with stent and atherectomy
1667	adj_cathlca3 Show the field ONLY if: [adj_cathlca2] = '1'	Another LCA cath intervention	yesno 1 Yes 0 No

1668	adj_cathlca3dt Show the field ONLY if: [adj_cathlca3] = '1'	LCA cath intervention 3: date	text (date_ymd)								
1669	adj_cathlca3typ Show the field ONLY if: [adj_cathlca3] = '1'	LCA cath intervention 3: type	dropdown <table border="1"> <tr><td>1</td><td>Angioplasty without stent</td></tr> <tr><td>2</td><td>Angioplasty with stent</td></tr> <tr><td>3</td><td>Angioplasty with atherectomy</td></tr> <tr><td>4</td><td>Angioplasty with stent and atherectomy</td></tr> </table>	1	Angioplasty without stent	2	Angioplasty with stent	3	Angioplasty with atherectomy	4	Angioplasty with stent and atherectomy
1	Angioplasty without stent										
2	Angioplasty with stent										
3	Angioplasty with atherectomy										
4	Angioplasty with stent and atherectomy										
1670	adj_cathlca1tx Show the field ONLY if: [adj_cathlca1] = '1'	LCA cath intervention: notes	notes								
1671	adj_cathlad1 Show the field ONLY if: [adj_cath] = '1'	LAD cath interventions	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table>	1	Yes	0	No				
1	Yes										
0	No										
1672	adj_cathlad1dt Show the field ONLY if: [adj_cathlad1] = '1'	LAD cath intervention 1: date	text (date_ymd)								
1673	adj_cathlad1typ Show the field ONLY if: [adj_cathlad1] = '1'	LAD cath intervention 1: type <i>Angioplasty includes balloon or laser angioplasty/Atherectomy includes rotational (rotablation)/extraction/directional atherectomy</i>	dropdown <table border="1"> <tr><td>1</td><td>Angioplasty without stent</td></tr> <tr><td>2</td><td>Angioplasty with stent</td></tr> <tr><td>3</td><td>Angioplasty with atherectomy</td></tr> <tr><td>4</td><td>Angioplasty with stent and atherectomy</td></tr> </table>	1	Angioplasty without stent	2	Angioplasty with stent	3	Angioplasty with atherectomy	4	Angioplasty with stent and atherectomy
1	Angioplasty without stent										
2	Angioplasty with stent										
3	Angioplasty with atherectomy										
4	Angioplasty with stent and atherectomy										
1674	adj_cathlad2 Show the field ONLY if: [adj_cathlad1] = '1'	Another LAD cath intervention	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table>	1	Yes	0	No				
1	Yes										
0	No										
1675	adj_cathlad2dt Show the field ONLY if: [adj_cathlad2] = '1'	LAD cath intervention 2: date	text (date_ymd)								
1676	adj_cathlad2typ Show the field ONLY if: [adj_cathlad2] = '1'	LAD cath intervention 2: type	dropdown <table border="1"> <tr><td>1</td><td>Angioplasty without stent</td></tr> <tr><td>2</td><td>Angioplasty with stent</td></tr> <tr><td>3</td><td>Angioplasty with atherectomy</td></tr> <tr><td>4</td><td>Angioplasty with stent and atherectomy</td></tr> </table>	1	Angioplasty without stent	2	Angioplasty with stent	3	Angioplasty with atherectomy	4	Angioplasty with stent and atherectomy
1	Angioplasty without stent										
2	Angioplasty with stent										
3	Angioplasty with atherectomy										
4	Angioplasty with stent and atherectomy										
1677	adj_cathlad3 Show the field ONLY if: [adj_cathlad2] = '1'	Another LAD cath intervention	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table>	1	Yes	0	No				
1	Yes										
0	No										
1678	adj_cathlad3dt Show the field ONLY if: [adj_cathlad3] = '1'	LAD cath intervention 3: date	text (date_ymd)								
1679	adj_cathlad3typ Show the field ONLY if: [adj_cathlad3] = '1'	LAD cath intervention 3: type	dropdown <table border="1"> <tr><td>1</td><td>Angioplasty without stent</td></tr> <tr><td>2</td><td>Angioplasty with stent</td></tr> <tr><td>3</td><td>Angioplasty with atherectomy</td></tr> <tr><td>4</td><td>Angioplasty with stent and atherectomy</td></tr> </table>	1	Angioplasty without stent	2	Angioplasty with stent	3	Angioplasty with atherectomy	4	Angioplasty with stent and atherectomy
1	Angioplasty without stent										
2	Angioplasty with stent										
3	Angioplasty with atherectomy										
4	Angioplasty with stent and atherectomy										
1680	adj_cathladtx Show the field ONLY if: [adj_cathlad1] = '1'	LAD cath intervention: notes	notes								
1681	adj_cathccx1 Show the field ONLY if: [adj_cath] = '1'	CCX cath interventions	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table>	1	Yes	0	No				
1	Yes										
0	No										
1682	adj_cathccx1dt Show the field ONLY if: [adj_cathccx1] = '1'	CCX cath intervention 1: date	text (date_ymd)								

1683	adj_cathccx1typ Show the field ONLY if: [adj_cathccx1] = '1'	CCX cath intervention 1: type <i>Angioplasty includes balloon or laser angioplasty/Atherectomy includes rotational (rotablation)/extraction/directional atherectomy</i>	dropdown <table border="1"> <tr><td>1</td><td>Angioplasty without stent</td></tr> <tr><td>2</td><td>Angioplasty with stent</td></tr> <tr><td>3</td><td>Angioplasty with atherectomy</td></tr> <tr><td>4</td><td>Angioplasty with stent and atherectomy</td></tr> </table>	1	Angioplasty without stent	2	Angioplasty with stent	3	Angioplasty with atherectomy	4	Angioplasty with stent and atherectomy
1	Angioplasty without stent										
2	Angioplasty with stent										
3	Angioplasty with atherectomy										
4	Angioplasty with stent and atherectomy										
1684	adj_cathccx2 Show the field ONLY if: [adj_cathccx1] = '1'	Another CCX cath intervention	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table>	1	Yes	0	No				
1	Yes										
0	No										
1685	adj_cathccx2dt Show the field ONLY if: [adj_cathccx2] = '1'	CCX cath intervention 2: date	text (date_ymd)								
1686	adj_cathccx2typ Show the field ONLY if: [adj_cathccx2] = '1'	CCX cath intervention 2: type	dropdown <table border="1"> <tr><td>1</td><td>Angioplasty without stent</td></tr> <tr><td>2</td><td>Angioplasty with stent</td></tr> <tr><td>3</td><td>Angioplasty with atherectomy</td></tr> <tr><td>4</td><td>Angioplasty with stent and atherectomy</td></tr> </table>	1	Angioplasty without stent	2	Angioplasty with stent	3	Angioplasty with atherectomy	4	Angioplasty with stent and atherectomy
1	Angioplasty without stent										
2	Angioplasty with stent										
3	Angioplasty with atherectomy										
4	Angioplasty with stent and atherectomy										
1687	adj_cathccx3 Show the field ONLY if: [adj_cathccx2] = '1'	Another CCX cath intervention	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table>	1	Yes	0	No				
1	Yes										
0	No										
1688	adj_cathccx3dt Show the field ONLY if: [adj_cathccx3] = '1'	CCX cath intervention 3: date	text (date_ymd)								
1689	adj_cathccx3typ Show the field ONLY if: [adj_cathccx3] = '1'	CCX cath intervention 3: type	dropdown <table border="1"> <tr><td>1</td><td>Angioplasty without stent</td></tr> <tr><td>2</td><td>Angioplasty with stent</td></tr> <tr><td>3</td><td>Angioplasty with atherectomy</td></tr> <tr><td>4</td><td>Angioplasty with stent and atherectomy</td></tr> </table>	1	Angioplasty without stent	2	Angioplasty with stent	3	Angioplasty with atherectomy	4	Angioplasty with stent and atherectomy
1	Angioplasty without stent										
2	Angioplasty with stent										
3	Angioplasty with atherectomy										
4	Angioplasty with stent and atherectomy										
1690	adj_cathccctx Show the field ONLY if: [adj_cathccx1] = '1'	CCX cath intervention: notes	notes								
1691	adj_cathccxbrk Show the field ONLY if: [adj_cath] = '1'		descriptive								
1692	adj_steno	Section Header: SECTION: CORONARY STENOSIS Stenosis	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table>	1	Yes	0	No				
1	Yes										
0	No										
1693	adj_stenorca Show the field ONLY if: [adj_steno] = '1'	RCA stenosis	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table>	1	Yes	0	No				
1	Yes										
0	No										
1694	adj_stenorcadt Show the field ONLY if: [adj_stenorca] = '1'	RCA stenosis: date	text (date_ymd)								
1695	adj_stenorcctx Show the field ONLY if: [adj_stenorca] = '1'	RCA stenosis: notes	notes								
1696	adj_stenolca Show the field ONLY if: [adj_steno] = '1'	LCA stenosis	yesno <table border="1"> <tr><td>1</td><td>Yes</td></tr> <tr><td>0</td><td>No</td></tr> </table>	1	Yes	0	No				
1	Yes										
0	No										
1697	adj_stenolcadt Show the field ONLY if: [adj_stenolca] = '1'	LCA stenosis: date	text (date_ymd)								
1698	adj_stenolcctx Show the field ONLY if: [adj_stenolca] = '1'	LCA stenosis: notes	notes								



1699	adj_stenolad Show the field ONLY if: [adj_steno] = '1'	LAD stenosis	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No		
1	Yes								
0	No								
1700	adj_stenoladdt Show the field ONLY if: [adj_stenolad] = '1'	LAD stenosis: date	text (date_ymd)						
1701	adj_stenoladtx Show the field ONLY if: [adj_stenolad] = '1'	LAD stenosis: notes	notes						
1702	adj_stenoccx Show the field ONLY if: [adj_steno] = '1'	CCX/LCX stenosis	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No		
1	Yes								
0	No								
1703	adj_stenoccxdt Show the field ONLY if: [adj_stenoccx] = '1'	CCX/LCX stenosis: date	text (date_ymd)						
1704	adj_stenoccxtx Show the field ONLY if: [adj_stenoccx] = '1'	CCX/LCX stenosis: notes	notes						
1705	brk61		descriptive						
1706	adj_mi	Section Header: <i>SECTION: MYOCARDIAL INFARCTION</i> MI	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No		
1	Yes								
0	No								
1707	adj_minum Show the field ONLY if: [adj_mi] = '1'	Number of MI events	dropdown <table border="1"><tr><td>1</td><td>1</td></tr><tr><td>2</td><td>2</td></tr><tr><td>3</td><td>3</td></tr></table>	1	1	2	2	3	3
1	1								
2	2								
3	3								
1708	adj_mi1dt Show the field ONLY if: [adj_minum] = '1' or [adj_minum] = '2' or [adj_minum] = '3'	MI event 1: date	text (date_ymd) Custom alignment: RH						
1709	adj_mi1tx Show the field ONLY if: [adj_minum] = '1' or [adj_minum] = '2' or [adj_minum] = '3'	MI event 1: notes	notes Custom alignment: RH						
1710	adj_mi2dt Show the field ONLY if: [adj_minum] = '2' or [adj_minum] = '3'	MI event 2: date	text (date_ymd) Custom alignment: RH						
1711	adj_mi2tx Show the field ONLY if: [adj_minum] = '2' or [adj_minum] = '3'	MI event 2: notes	notes Custom alignment: RH						
1712	adj_mi3dt Show the field ONLY if: [adj_minum] = '3'	MI event 3: date	text (date_ymd) Custom alignment: RH						
1713	adj_mi3tx Show the field ONLY if: [adj_minum] = '3'	MI event 3: notes	notes Custom alignment: RH						
1714	brk62		descriptive						
1715	adj_cabg	Section Header: <i>SECTION: CABG</i> CABG	yesno <table border="1"><tr><td>1</td><td>Yes</td></tr><tr><td>0</td><td>No</td></tr></table>	1	Yes	0	No		
1	Yes								
0	No								
1716	adj_cabgdt Show the field ONLY if: [adj_cabg] = '1'	CABG: date	text (date_ymd) Custom alignment: RH						
1717	adj_cabgtx Show the field ONLY if: [adj_cabg] = '1'	CABG: notes	notes Custom alignment: RH						

1718	brk63		descriptive						
1719	adj_bleed	Section Header: <i>SECTION: BLEEDING</i> Bleeding associated with anticoagulation	yesno <table border="1"> <tr> <td>1</td> <td>Yes</td> </tr> <tr> <td>0</td> <td>No</td> </tr> </table>	1	Yes	0	No		
1	Yes								
0	No								
1720	adj_bleednum Show the field ONLY if: [adj_bleed] = '1'	Bleeding: number of events	dropdown <table border="1"> <tr> <td>1</td> <td>1</td> </tr> <tr> <td>2</td> <td>2</td> </tr> <tr> <td>3</td> <td>3</td> </tr> </table> Custom alignment: RH	1	1	2	2	3	3
1	1								
2	2								
3	3								
1721	adj_bleed1dt Show the field ONLY if: [adj_bleednum] = '1' or [adj_bleednum] = '2' or [adj_bleednum] = '3'	Bleeding event 1: date	text (date_ymd) Custom alignment: RH						
1722	adj_bleed1tx Show the field ONLY if: [adj_bleednum] = '1' or [adj_bleednum] = '2' or [adj_bleednum] = '3'	Bleeding event 1: notes	notes Custom alignment: RH						
1723	adj_bleed2dt Show the field ONLY if: [adj_bleednum] = '2' or [adj_bleednum] = '3'	Bleeding event 2: date	text (date_ymd) Custom alignment: RH						
1724	adj_bleed2tx Show the field ONLY if: [adj_bleednum] = '2' or [adj_bleednum] = '3'	Bleeding event 2: notes	notes Custom alignment: RH						
1725	adj_bleed3dt Show the field ONLY if: [adj_bleednum] = '3'	Bleeding event 3: date	text (date_ymd) Custom alignment: RH						
1726	adj_bleed3tx Show the field ONLY if: [adj_bleednum] = '3'	Bleeding event 3: notes	notes Custom alignment: RH						
1727	brk64		descriptive						
1728	adjudication_complete	Section Header: <i>Form Status</i> Complete?	dropdown <table border="1"> <tr> <td>0</td> <td>Incomplete</td> </tr> <tr> <td>1</td> <td>Unverified</td> </tr> <tr> <td>2</td> <td>Complete</td> </tr> </table>	0	Incomplete	1	Unverified	2	Complete
0	Incomplete								
1	Unverified								
2	Complete								