Data Dictionary Codebook

CRECOOLT 3.0 (PID: 40)

06/05/2025 3:10pm

Instruments		
Instrument	Form Name	Events
Demographic Data	demographic_data	olt_date_of_transp_arm_1
Comorbidities	comorbidities	olt_date_of_transp_arm_1
Underlying Liver Disease	underlying_liver_disease	olt_date_of_transp_arm_1
Information About Transplant	information_about_transplant	olt_date_of_transp_arm_1
Immunosuppressive Medication	immunosuppressive_medication	olt_date_of_transp_arm_
CRE Issues	cre_issues	olt_date_of_transp_arm_
Susceptibility Of CRE Colonizing Strain	susceptibility_of_cre_colonizing_strain	olt_date_of_transp_arm_^
Post-OLT Complications	postolt_complications	olt_date_of_transp_arm_
Management Of CRE Colonization and CRE infection	management_of_cre_colonization_and_cre_infection	olt_date_of_transp_arm_^
CRECOOLT score evaluation 🕏	crecoolt_score_evaluation	olt_date_of_transp_arm_r week_1_arm_1 week_2_arm_1 week_3_arm_1 week_4_arm_1 week_5_arm_1 week_6_arm_1 week_7_arm_1 week_8_arm_1 week_8_arm_1
Microbiological Screening	microbiological_screening	olt_date_of_transp_arm_1
Exposure To CRE Antibiotics 🕏	exposure_to_cre_antibiotics	olt_date_of_transp_arm_′
Outcome at 180 days	outcome_at_180_days	olt_date_of_transp_arm_

Events	
Event Name	Unique event name
OLT: [date_of_transplant]	olt_date_of_transp_arm_1
Week 1	week_1_arm_1
Week 2	week_2_arm_1
Week 3	week_3_arm_1
Week 4	week_4_arm_1
Week 5	week_5_arm_1
Week 6	week_6_arm_1
Week 7	week_7_arm_1
Week 8	week_8_arm_1

	#	Variable / Field Name	Field Label Field Note	Field Attributes (Field Type, Validation, Choices, Calculations, etc.)		
Ins	Instrument: Demographic Data (demographic_data)					
	1	[record_id]	Record ID	text		

	2	<pre>[note]</pre>	Dear Investigator, we kindly remind you that: • Retrospective study: OLT patients transplanted in the study period 2018-2022 will be included in the retrospective study. • Prospective study: recent OLT patients followed prospectively, according to the study protocol, will be included in the prospective study.	des	scriptive	
	3	[retro_or_pros]	According to the definitions above, is this patient to be included in:	rad 1 2	io, Required Retrospective study Prospective study	
	4	[date_of_birth]	Date of birth:	tex	t (date_dmy), Require	d
	5	[gender]	Gender:	rad 1 2	Male Female	
	6	[date_of_admission]	Date of hospital admission prior transplantation:	tex	t (date_dmy), Require	d
	7	[age]	Age at time of transplantation:	([da	c culation: round (date ate_of_admission], [da ny"))	
	8	<pre>[demographic_data_com plete]</pre>	Section Header: Form Status Complete?	0 1 2	Incomplete Unverified Complete	
Inst	rum	ent: Comorbidities	(comorbidities)			
	9	[comorbidities1]	Section Header: Charlson Comorbidity Index (CCI):	che	eckbox	
			Comorbidities (+1 point):	1	comorbidities11	Myocardial infarction
				2	comorbidities12	Congestive heart failure
				3	comorbidities13	Peripheral vascular disease
				4	comorbidities14	CVA or TIA
				5	comorbidities15	Dementia
				6	comorbidities16	COPD
				7	comorbidities17	Connective tissue disease

			8 comorbidities18 Peptic ulcer disease
10	[comorbidities2]	Comorbidities (+2 points):	checkbox
			1 comorbidities21 Hemiplegia
			2 comorbidities22 Moderate to severe CKD
			3 comorbidities23 Leukemia
			4 comorbidities24 Lymphoma
11	[liver_dis]	Liver disease:	checkbox
			1 liver_dis1 mild (+1 point)
			2 liver_dis2 moderate to severe (+3 points)
12	[diabetes_mellitus]	Diabetes mellitus:	checkbox
			1 diabetes_mellitus1 none or diet- controlled
			2 diabetes_mellitus2 uncomplicated (+1 point)
			3 diabetes_mellitus3 end-organ damage (+2 points)
13	[solid_tumor]	Solid tumor:	checkbox
			1 solid_tumor1 none
			2 solid_tumor2 localized (+2 points)
			3 solid_tumor3 metastatic (+6 points)
14	[aids]	AIDS (+6 points)	yesno
			1 Yes
			0 No
15	[cci]	Charlson Comordibities Index (CCI):	calc
		Adjusted for [age] years old patient.	Calculation: round (if(([age]>=50 and [age] < 60),1,0) + if(([age]>=60 and [age] < 70),2,0) + if(([age]>=70 and [age] < 80),3,0) + if(([age]>=80),4,0) + if(([age]>=80),4,0) + if(([comorbidities1(1)]=1,1,0) + if(([comorbidities1(2)]=1,1,0) + if(([comorbidities1(3)]=1,1,0) + if(([comorbidities1(4)]=1,1,0) + if(([comorbidities1(5)]=1,1,0) + if(([comorbidities1(6)]=1,1,0) + if(([comorbidities1(7)]=1,1,0) + if(([comorbidities2(1)]=1,2,0) + if(([comorbidities2(2)]=1,2,0) + if(([comorbidities2(4)]=1,2,0) + if(([comorbidities2(4)]

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				if([li if([d if([d if([s	ver_dis(1)]=1 ver_dis(2)]=1 liabetes_mell liabetes_mell olid_tumor(2 olid_tumor(3	,3,0) + litus(2)]= litus(3)]= 2)]=1,2,0)	=1,2,0) +
	16	[comorbidities_comple	Section Header: Form Status	dro	pdown		
		te]	Complete?	0	Incomplete		
			·	l 	Unverified		
				2	Complete		
Inst	rum	ent: Underlying Live	er Disease (underlying_liver_disease)				
	17	[liver_disease]	UNDERLYING LIVER DISEASE	che	ckbox		
			- Please check all that apply:	1	liver_diseas	se1	HCV infection
				2	liver_diseas	se2	HBV infection
				3	liver_diseas	se3	HIV infection
				4	liver_diseas	se4	Alcohol abuse
				5	liver_diseas	se5	Metabolic disease
				6	liver_diseas	se6	Autoimmune disease
				7	liver_diseas	se7	Other (please, specify below)
				8	liver_diseas	se8	Fulminant hepatitis
				9	liver_diseas	se9	Prior transplant
				10	liver_diseas	se10	НСС
	18	[metabolic_disease]	Metabolic disease - Please, specify:	text	:		
		Show the field ONLY i					
		f: [liver_disease(5)] = '1'					
	19	[autoimmune_disease]	Autoimmune disease - Please, specify:	text	-		
		Show the field ONLY i f:					
		[liver_disease(6)] = '1'					
	20	[other_specify]	Other - Please, specify:	text			
		Show the field ONLY i f:					
		[liver_disease(7)] = '1'					
	21	<pre>[type_prior_transplan t]</pre>	Prior transplant - Please, specify type:	text			
		Show the field ONLY i f:					
		[liver_disease(9)] = '1'					

	22	<pre>[year_prior_transplan t]</pre>	Prior transplant - Please, specify the year:	text (integer)
		Show the field ONLY i		
		f: [liver_disease(9)] = '1'		
	23	[date_waiting_list]	Date of inclusion in waiting list:	text (date_dmy)
	24	[meld_waiting_list]	MELD at inclusion in waiting list:	text
	25	<pre>[underlying_liver_dis ease_complete]</pre>	Section Header: Form Status Complete?	dropdown 0 Incomplete 1 Unverified 2 Complete
Ins	trum	ent: Information Ab	out Transplant (information_about_tran	nsplant)
	26	[date_of_transplant]	Date of transplant:	text (date_dmy)
	27	[date_admission_icu]	Date of admission in ICU:	text (date_dmy)
	28	[liver_split]	Liver split?	yesno 1 Yes 0 No
	29	<pre>[combined_transplan t]</pre>	Combined transplant:	yesno 1 Yes 0 No
	30	[meld_score]	MELD score at OLT:	text
	31	<pre>[biliary_anastomosi s]</pre>	Type of biliary anastomosis:	checkbox 1 biliary_anastomosis1 Duct-duct with Kher 2 biliary_anastomosis2 Duct-duct
				without Kher 3 biliary_anastomosis3 Coledocojejunal
	32	[prolonged_surgery]	Prolonged surgery (more than 8 hours):	yesno 1 Yes 0 No
	33	[donor_age]	Donor age:	text (integer)
	34	[deceased_donor]	Deceased donor?	yesno 1 Yes 0 No
	35	[time_cold_ischemia]	Duration of cold ischemia of the graft (hh:mm):	text (time)
	36	<pre>[intraoperative_bleed ing]</pre>	Intraoperative bleeding with need of maxive transfusion:	yesno 1 Yes

				0 No
	37	<pre>[specify_number_of_ b] Show the field ONLY i f: [intraoperative_bleedi ng] = '1'</pre>	If yes, specify number of blood trasfusions:	text
	38	[date_discharge_icu]	Date of discharge from ICU:	text (date_dmy)
	39	<pre>[information_about_tr ansplant_complete]</pre>	Section Header: Form Status Complete?	dropdown 0 Incomplete 1 Unverified 2 Complete
Inst	rum	ent: Immunosuppre	essive Medication (immunosuppressive_	medication)
	40	[induction_regimen]	Induction regimen	checkbox
			- Please check all that apply:	1 induction_regimen1 Bolus of steroids
				2 induction_regimen2 Use of antilymphocyt globulin
				3 induction_regimen3 Use of basiliximab
				4 induction_regimen4 Use of rituximab
				5 induction_regimen5 Other (please, specify below)
	41	<pre>[other_induction_regi men] Show the field ONLY i f: [induction_regimen(5)] = '1'</pre>	Other, please specify:	text
	42	[maintenance_regime	Maintenance regimen:	checkbox
		n]		1 maintenance_regimen1 Prednison
				2 maintenance_regimen2 Cyclospori
				3 maintenance_regimen3 Tacrolimus
				4 maintenance_regimen4 Mycophen mofetil
				5 maintenance_regimen5 Azathiopri
				6 maintenance_regimen6 Sirolimus
				7 maintenance_regimen7 Everolimus
				8 maintenance_regimen8 Other (pleasure) specify bel
	43	[drug_dose]	Other, specify drug dose (mg):	text (number)

	44	Show the field ONLY i f: [maintenance_regimen (8)] = '1' [immunosuppressive_me	indicate in mg Section Header: Form Status	dropdown
		dication_complete]	Complete?	0 Incomplete 1 Unverified 2 Complete
Inst	trum	ent: CRE Issues (cre	_issues)	
	45	[cre_colon]	CRE colonization:	radio 1 Pre-OLT (before [date_of_transplant]) 2 Post-OLT (after [date_of_transplant])
	46	<pre>[date_positive_pre_ol t] Show the field ONLY i f: [cre_colon] = '1'</pre>	Pre-OLT, please specify date of first positive sample:	text (date_dmy)
	47	<pre>[date_positive_post_o lt] Show the field ONLY i f: [cre_colon] = '2'</pre>	Post-OLT, please specify date of first positive sample:	text (date_dmy)
	48	[type_of_sample]	CRE colonization - Please, specify the type of	checkbox
			sample: Check all that apply	1 type_of_sample1 Rectal swab 2 type_of_sample2 Respiratory sample
				3 type_of_sample3 Urine
				4 type_of_sample4 Other (please, specify below)
	49	<pre>[other_sample] Show the field ONLY i f: [type_of_sample(4)] = '1'</pre>	Other, please specify:	text
	50	<pre>[multisite_colonizati on]</pre>	Multisite colonization 0=No 1=Yes	calc Calculation: if (([type_of_sample(1)]=1 and [type_of_sample(2)]=1),1,0) or if (([type_of_sample(1)]=1 and [type_of_sample(3)]=1),1,0) or if (([type_of_sample(1)]=1 and [type_of_sample(4)]=1),1,0) or if (([type_of_sample(2)]=1 and [type_of_sample(3)]=1),1,0) or if (([type_of_sample(2)]=1 and [type_of_sample(2)]=1 and [type_of_sample(4)]=1),1,0) or if

				(([type_of_sample(3)]=1 and [type_of_sample(4)]=1),1,0)
	51	[mec_of_carbapenem_re		checkbox
		si]	-Please specify mechanisms of carbapenem resistance:	1 mec_of_carbapenem_resi1 KPC
				2 mec_of_carbapenem_resi2 OXA-48
				3 mec_of_carbapenem_resi3 NDM
				4 mec_of_carbapenem_resi4 VIM
				5 mec_of_carbapenem_resi5 Other (please specify below)
	52	[other_mec_carb_res]	Other, please specify:	text
		Show the field ONLY i		
		f:		
		[mec_of_carbapenem_ resi(5)] = '1'		
	53	[cre_issues_complet	Section Header: Form Status	dropdown
		e]	Complete?	0 Incomplete
				1 Unverified
				2 Complete
Ins	trum	ent: Susceptibility C	Of CRE Colonizing Strain (susceptibility	_of_cre_colonizing_strain)
	54	[erta]	ERTAPENEM	radio (Matrix)
				1 R
				2 S
				3 1
				4 NA
	55	[mero]	MEROPENEM	radio (Matrix)
				1 R
				2 S
				3 1
				4 NA
	56	[amik]	AMIKACIN	radio (Matrix)
				1 R
				2 S
				3 1
				4 NA
	57	[gent]	GENTAMICIN	radio (Matrix)
				1 R
				2 S
				3 1
1				

			4 NA
58	[tmp_smx]	TMP/SMX	radio (Matrix) 1 R 2 S 3 I 4 NA
59	[tig]	TIGECYCLINE	radio (Matrix) 1 R 2 S 3 I 4 NA
60	[col]	COLISTIN	radio (Matrix) 1 R 2 S 3 I 4 NA
61	[cef_avi]	CEFTAZIDIME-AVIBACTAM	radio (Matrix) 1 R 2 S 3 I 4 NA
62	[mero_vabo]	MEROPENEM/VABORBACTAM	radio (Matrix) 1 R 2 S 3 I 4 NA
63	[imi_rel]	IMIPENEM/RELEBACTAM	radio (Matrix) 1 R 2 S 3 I 4 NA
64	[cef]	CEFIDEROCOL	radio (Matrix) 1 R 2 S 3 I 4 NA
65	[other_ant]	Other	radio (Matrix)

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			2 S 3 I 4 NA
66	Show the field ONLY i f: [erta] = '1' or [erta] =	ERTAPENEM-MIC:	text
67	'2' or [erta] = '3' [mero_mic] Show the field ONLY i f: [mero] = '1' or [mero] = '2' or [mero] = '3'	MEROPENEM-MIC:	text
68		AMIKACIN-MIC:	text
69	[genta_mic] Show the field ONLY i f: [gent] = '1' or [gent] = '2' or [gent] = '3'	GENTAMICIN-MIC:	text
70	[tmp_smx_mic] Show the field ONLY i f: [tmp_smx] = '1' or [tm p_smx] = '2' or [tmp_s mx] = '3'	TMP/SMX-MIC:	text
71	<pre>[tige_mic] Show the field ONLY i f: [tig] = '1' or [tig] = '2' or [tig] = '3'</pre>	TIGECYCLINE-MIC:	text
72	[col_mic] Show the field ONLY i f: [col] = '1' or [col] = '2' o r [col] = '3'	COLISTIN-MIC:	text
73	[cef_avi_mic] Show the field ONLY i f: [cef_avi] = '1' or [cef_av i] = '2' or [cef_avi] = '3'	CEFTAZIDIME-AVIBACTAM-MIC:	text
74	[mero_vab_mic]	MEROPENEM/VABORBACTAM-MIC:	text

	Show the field ONLY i f: [mero_vabo] = '1' or [m ero_vabo] = '2' or [mer o_vabo] = '3'				
75	<pre>[imi_rel_mic] Show the field ONLY i f: [imi_rel] = '1' or [imi_re l] = '2' or [imi_rel] = '3'</pre>	IMIPENEM/RELEBACTAM-MIC:	text		
76	[cef_mic] Show the field ONLY i f: [cef] = '1' or [cef] = '2' o r [cef] = '3'	CEFIDEROCOL-MIC:	text		
77	[other_drug_specify] Show the field ONLY i f: [other_ant] = '1' or [oth er_ant] = '2' or [other_a nt] = '3' or [other_ant] = '4'	Other - Please, specify:	text		
78	[other_mic] Show the field ONLY i f: [other_ant] = '1' or [oth er_ant] = '2' or [other_a nt] = '3'	Other-MIC:	text		
79	<pre>[susceptibility_of_cr e_colonizing_strain_c omplete]</pre>	Section Header: Form Status Complete?	0	odown Incomplete Unverified Complete	
Instrum	ent: Post-OLT Comp	lications (postolt_complications)			
80	[post_olt_compli]	POST-OLT COMPLICATIONS (after [date_of_transplant]) Please check all that apply up to the first CRE infection, death or 2 months after OLT	1	post_olt_compli1	Acute renal failure (RIFLE criteria)
		whichever occurred first	2	post_olt_compli2	Need of renal replacement therapy
			3	post_olt_compli3	Mechanical ventilation > 4 h
			4	post_olt_compli4	Primary or Secondary Graft Non- Function

post_olt_compli___5

					intervention
			6	post_olt_compli6	Re- transplantation
			7	post_olt_compli7	Rejection
			8	post_olt_compli8	Biopsy proven rejection
			9	post_olt_compli9	Histological recurrence of HCV infection
			10	post_olt_compli10	Treatment for CMV infection and/or disease
			11	post_olt_compli1	Other (Please, specify below)
81	[other_compli] Show the field ONLY i f: [post_olt_compli(11)]= 1	Other - Please, specify:	text	t	
82	<pre>[postolt_complication s_complete]</pre>	Section Header: Form Status Complete?	0	pdown Incomplete Unverified	
			2	Complete	
fecti	on (management_of_c	of CRE Colonization and CRE re_colonization_and_cre_infection)	2	Complete	
	_		2		No strategy adopted
fecti	on (management_of_c	re_colonization_and_cre_infection) Management of CRE colonization:	che	Complete	
fecti	on (management_of_c	re_colonization_and_cre_infection) Management of CRE colonization:	che 1 2	ckbox cre_colonization1	adopted Targeted perioperative
fecti	on (management_of_c	re_colonization_and_cre_infection) Management of CRE colonization:	che 1 2	ckbox cre_colonization1 cre_colonization2 cre_colonization3	adopted Targeted perioperative prophylaxis Pre-emptive
83	<pre>[cre_colonization] [prophylaxis_drugs] Show the field ONLY i f:</pre>	Management of CRE colonization: Check all that apply. Targeted peri-operative prophylaxis - Please,	chee 1 3	ckbox cre_colonization1 cre_colonization2 cre_colonization3	adopted Targeted perioperative prophylaxis Pre-emptive

87	Show the field ONLY i f: [cre_colonization(3)]=1 [other_pre_emptive] Show the field ONLY i f:	Other pre-emptive strategy - Please, specify:	1 Early start of anti-CRE coverage upon infection symptoms onset 2 Early start of anti-CRE coverage according to CRECOOLT score 3 Other (Please, specify below) text
88	<pre>[pre_emptive]=3 [cre_infection]</pre>	CRE infection	yesno 1 Yes 0 No
89	[cre_infection_date] Show the field ONLY i f: [cre_infection]=1	CRE infection - Please, specify the date:	text (date_dmy)
90	[pao2] Show the field ONLY i f: [cre_infection] = '1'	Section Header: Clinical severity according to SOFA score at CRE infection PaO2/FiO2 (mmHg)	checkbox 0 pao20 ≥ 400 1 pao21 300-399 2 pao22 200-299 3 pao23 100-199 and mechanically ventilated 4 pao24 < 100 and mechanically ventilated
91	[gcs] Show the field ONLY i f: [cre_infection] = '1'	Glasgow Coma Scale (GCS):	checkbox 0 gcs0 15 1 gcs1 13-14 2 gcs2 10-12 3 gcs3 6-9 4 gcs4 < 6
92	[map] Show the field ONLY i f: [cre_infection] = '1'	MAP or administration of vasopressors:	checkbox 0 map0 >70mmHg 1 map1 < 70mmHg 2 map2 dop ≤5 or dob any dose 3 map3 dop >5 or epi/nor≤0.1 4 map4 dop >15 or epi /nor>0.1
93	[bilirubin] Show the field ONLY i f: [cre_infection] = '1'	Bilirubin (mg/dL):	checkbox 0 bilirubin0 < 1.2 1 bilirubin1 1.2-1.9 2 bilirubin2 2.0-5.9

Solirubin_3 6.0-11.9 4	00/20, 1			ONEGGET 3.0 NEBGE	
94 [platelets] Platelets: Checkbox O platelets					
Show the field ONLY i f.					4 01111 4 2 1 2
1 platelets_1 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150 150		94		Platelets:	
3 platelets_3			[cre_infection] = '1'		
Show the field ONLY in the field on					
Show the field ONLY f:					
1		95	[cr_sofa]	Creatinine (mg/dL) or urine output:	checkbox
[cre_infection] = "1"			Show the field ONLY i		0 cr_sofa0 0.5-1.2
2 cr_sofa					1 cr_sofa1 1.2-1.9
Sofa			[cre_iniection] = 1		2 cr_sofa2 2.0-3.4
Sofa					- -
Sofa					
Calculation: if([cre_infection]=1,(iff [pao2(0)]=1, 0, (iff [pao2(1)]=1, 1, (iff [pao2(0)]=1, 0, (iff [pao2(0)]=					- -
Show the field ONLY f. [pao2(0)]=1, 0, (if([pao2(1)]=1, 1, (if([pao2(2)]=1, 2, (ift [pao2(3)]=1, 3, (if([pao2(3)]=1, 2, (ift [pao2(3)]=1, 3, (if([pao2(3)]=1, 2, (ift [pao2(3)]=1, 0, (ift [gcs(1)]=1, 1, (ift [gcs(1)]=1, 1, (ift [gcs(1)]=1, 1, (ift [gcs(1)]=1, 1, (ift [gcs(2)]=1, 2, (ift [gcs(3)]=1, 3, (ift [gcs(3)]=1, 3, (ift [gcs(3)]=1, 2, (ift [map(0)]=1, 0, (ift [map(0)]=1, 2, (ift [map(0)]=1, 3, (ift [map(0)]=1, 0, (ift [patelets(0)]=1, 0, (ift [patelets(0)]=1, 2, (ift [pat		96	[sofa]	SOFA score:	
if([pao2(3)]=1, 2, (if([pao2(3)]=1, 3, (if([pao2(3)]=1, 0, (if([pao2(4)]=1, 4, 0))))))) + if([gcs(0)]=1, 0, (if([gcs(0)]=1, 1, (if([gcs(0)]=1, 2, (if([gcs(0)]=1, 2, (if([gcs(0)]=1, 3, (if([gcs(0)]=1, 2, (if([gcs(0)]=1, 2, (if([gcs(0)]=1, 2, (if([gcs(0)]=1, 2, (if([gcs(0)]=1, 2, (if([map(0)]=1, 2, (if([map(0)]=1, 2, (if([map(0)]=1, 2, (if([bilirubin(0)]=1, 2, (if([bilirubin(0)]=1, 3, (if([bilirubin(0)]=1, 0, (if([bilirubin(0)]=1, 3, (if([bilirubin(0)]=1, 2, (if([platelets(0)]=1, 2, (if([platelets(0)]=1, 2, (if([platelets(0)]=1, 2, (if([platelets(0)]=1, 2, (if([cr_csofa(0)]=1, 3, (if([cr_csofa(0)]=1, 3, (if([cr					
					if([pao2(2)]=1, 2, (if([pao2(3)]=1, 3, (
[gcs(3)]=1, 3, (if([gcs(4)]=1, 4, 0)))))))) + if([map(0)]=1, 0, (iff [map(1)]=1, 1, (if([map(2)]=1, 2, (iff [map(3)]=1, 3, (if([map(4)]=1, 4, 0)))))))) + if([bilirubin(0)]=1, 0, (iff [bilirubin(1)]=1, 1, (iff([bilirubin(4)]=1, 4, 0)))))))) + if([platelets(0)]=1, 0, (iff [bilirubin(3)]=1, 3, (iff([bilirubin(4)]=1, 4, 0))))))) + if([platelets(0)]=1, 0, (iff [bilirubin(3)]=1, 3, (iff([platelets(2)]=1, 2, (iff [platelets(3)]=1, 3, (iff([platelets(2)]=1, 2, (iff [cr_sofa(1)]=1, 1, (iff([cr_sofa(0)]=1, 0, (iff [cr_sofa(3)]=1, 3, (iff([cr_sofa(0)]=1, 0, (iff [cr_sofa(3)]=1, 3, (iff([platelets(2)]=1, 2, (iff [cr_sofa(3)]=1, 3, (iff([cr_sofa(0)]=1, 0, (iff [bilirubin(1)]=1, 3, (iff([platelets(2)]=1, 2, (iff [platelets(3)]=1, 3, (iff([platelets(3)]=1, (iff([platelets(3)]=1, 3, (iff([platelets(3)]=			[e.e_m.eedon]		
if([map(2)]=1, 2, (if([map(3)]=1, 3, ([gcs(3)]=1, 3, (if([gcs(4)]=1, 4, 0)))))))) + if(
iff([map(4)]=1, 4, 0)))))))) + if([bilirubin(0)]=1, 0, (iff ([bilirubin(1)]=1, 1, (iff([bilirubin(2)]=1, 2, (iff ([bilirubin(3)]=1, 3, (iff([bilirubin(2)]=1, 2, (iff ([bilirubin(3)]=1, 3, (iff([bilirubin(4)]=1, 4, 0)))))))) + if([platelets(0)]=1, 0, (iff ([platelets(3)]=1, 1, (iff([platelets(4)]=1, 4, 0))))))))) + if([cr_sofa(0)]=1, 2, (iff ([cr_sofa(1)]=1, 1, (iff([cr_sofa(2)]=1, 2, (iff ([cr_sofa(3)]=1, 3, (iff([cr_sofa(4)]=1, 4, 0))))))))) + if([cr_sofa(0)]=1, 2, (iff ([cr_sofa(3)]=1, 3, (iff([cr_sofa(4)]=1, 4, 0))))))))) + if([cr_sofa(0)]=1, 2, (iff ([cr_sofa(3)]=1, 3, (iff([cr_sofa(4)]=1, 4, 0))))))))) + if([platelets(2)]=1, 2, (iff ([cr_sofa(3)]=1, 3, (iff([cr_sofa(4)]=1, 4, 0)))))))))) + if([platelets(2)]=1, 2, (iff ([cr_sofa(3)]=1, 3, (iff([cr_sofa(4)]=1, 4, 0))))))))))) + if([platelets(2)]=1, 2, (iff ([cr_sofa(3)]=1, 3, (iff([cr_sofa(4)]=1, 4, 0))))))))))))))))))))))))))))))))))					
if([bilirubin(2)]=1, 2, (if([bilirubin(3)]=1, 3, (if([bilirubin(2)]=1, 2, (if([bilirubin(3)]=1, 3, (if([bilirubin(4)]=1, 4, 0)))))))) + if([platelets(0)]=1, 0, (if([platelets(1)]=1, 1, (if([platelets(2)]=1, 2, (if([platelets(3)]=1, 3, (if([platelets(2)]=1, 2, (if([platelets(3)]=1, 3, (if([platelets(4)]=1, 4, 0)))))))) + if([cr_sofa(0)]=1, 0, (if([cr_sofa(2)]=1, 2, (if([cr_sofa(3)]=1, 1, (if([cr_sofa(4)]=1, 4, 0))))))))) + if([cr_sofa(2)]=1, 2, (if([platelets(3)]=1, 2, (i					if([map(4)]=1, 4, 0))))))))) + if(
(if([bilirubin(4)]=1, 4, 0)))))))) + if([platelets(0)]=1, 0, (if([platelets(1)]=1, 1, (
if([platelets(2)]=1, 2, (if([platelets(3)]=1, 3, (if([platelets(4)]=1, 4, 0)))))))) + if((if([bilirubin(4)]=1, 4, 0)))))))) + if(
(if([platelets(4)]=1, 4, 0)))))))) + if([cr_sofa(0)]=1, 0, (if([cr_sofa(1)]=1, 1, (1 '
if([cr_sofa(2)]=1, 2, (if([cr_sofa(3)]=1, 3, (if([cr_sofa(4)]=1, 4, 0)))))))),"") 97 [sofa_import]					(if([platelets(4)]=1, 4, 0)))))))) + if(
if([cr_sofa(4)]=1, 4, 0)))))))),"") 97 [sofa_import]					
Show the field ONLY i f: [cre_infection]='1' AND [cre_infection_date]<'3 1-12-2016' 98 [infection_source] Show the field ONLY i f: [cre_infection]=1 Infection source/type: checkbox 1 infection_source1 BSI 2 infection_source2 Lower respiratory infection					
f: [cre_infection]='1' AND [cre_infection_date]<'3 1-12-2016' 98 [infection_source] Show the field ONLY i f: [cre_infection]=1 f:		97	[sofa_import]	SOFA score:	text (integer)
[cre_infection]='1' AND [cre_infection_date]<'3					
[cre_infection_date]<'3 1-12-2016' 98 [infection_source] Show the field ONLY i f: [cre_infection]=1 Checkbox infection_source1 BSI 2 infection_source2 Lower respiratory infection					
98 [infection_source] Show the field ONLY i f: [cre_infection]=1			[cre_infection_date]<'3		
f: [cre_infection]=1 2 infection_source2 Lower respiratory infection		98		Infection source/type:	checkbox
[cre_infection]=1					1 infection_source1 BSI
infection					
3 infection_source3 IAI			[c. c_nnection]		
					3 infection_source3 IAI

			4 infection_source4 UTI
			5 infection_source5 SSI
			6 infection_source6 Other (please, specify below)
100	[bsi_specify] Show the field ONLY i f: [infection_source(1)]=1 [other_sources] Show the field ONLY i f:	BSI - Please, specify: Other sources - Please, specify:	radio 1 Primary 2 Secondary 3 CVC related text
101	<pre>[infection_source(6)]=1 [cre_isolate] Show the field ONLY i f: [cre_infection] = '1'</pre>	Molecular characteristics of CRE infection isolate Check all that apply.	checkbox 1 cre_isolate1 KPC 2 cre_isolate2 OXA-48 3 cre_isolate3 VIM 4 cre_isolate4 NDM 5 cre_isolate5 Other (please, specify below)
102	[other_isolate] Show the field ONLY i f: [cre_isolate(5)]=1	Other - Please, specify:	text
103	[erta_2] Show the field ONLY i f: [cre_infection] = '1'	Section Header: Susceptibility of CRE infection isolate: ERTAPENEM	radio (Matrix) 1 R 2 S 3 I 4 NA
104	[mero_2] Show the field ONLY i f: [cre_infection] = '1'	MEROPENEM	radio (Matrix) 1 R 2 S 3 I 4 NA
105	[amik_2] Show the field ONLY i f: [cre_infection] = '1'	AMIKACIN	radio (Matrix) 1 R 2 S 3 I 4 NA
106	[gent_2]	GENTAMICIN	radio (Matrix)

	Show the field ONLY i f: [cre_infection] = '1'		2 S 3 I 4 NA
107	[tmp_smx_2] Show the field ONLY i f: [cre_infection] = '1'	TMP/SMX	radio (Matrix) 1 R 2 S 3 I 4 NA
108	<pre>[tig_2] Show the field ONLY i f: [cre_infection] = '1'</pre>	TIGECYCLINE	radio (Matrix) 1 R 2 S 3 I 4 NA
109	[col_2] Show the field ONLY i f: [cre_infection] = '1'	COLISTIN	radio (Matrix) 1 R 2 S 3 I 4 NA
110	[cef_avi_2] Show the field ONLY i f: [cre_infection] = '1'	CEFTAZIDIME-AVIBACTAM	radio (Matrix) 1 R 2 S 3 I 4 NA
111	[mero_vabo_2] Show the field ONLY if: [cre_infection] = '1'	MEROPENEM/VABORBACTAM	radio (Matrix) 1 R 2 S 3 I 4 NA
112	<pre>[imi_rel_2] Show the field ONLY i f: [cre_infection] = '1'</pre>	IMIPENEM/RELEBACTAM	radio (Matrix) 1 R 2 S 3 I 4 NA
113	[cef_2] Show the field ONLY i f: [cre_infection] = '1'	CEFIDEROCOL	radio (Matrix) 1 R 2 S 3 I 4 NA

		•	
114	[other_ant_2] Show the field ONLY i f: [cre_infection] = '1'	Other	radio (Matrix) 1 R 2 S 3 I 4 NA
115	[erta_mic_2] Show the field ONLY i f: [erta_2] = '1' or [erta_2] = '2' or [erta_2] = '3'	ERTAPENEM-MIC:	text
116	[mero_mic_2] Show the field ONLY i f: [mero_2] = '1' or [mero _2] = '2' or [mero_2] = '3'	MEROPENEM-MIC:	text
117	[ami_mic_2] Show the field ONLY i f: [amik_2] = '1' or [amik_ 2] = '2' or [amik_2] = '3'	AMIKACIN-MIC:	text
118	[genta_mic_2] Show the field ONLY i f: [gent_2] = '1' or [gent_ 2] = '2' or [gent_2] = '3'	GENTAMICIN-MIC:	text
119	[tmp_smx_mic_2] Show the field ONLY i f: [tmp_smx_2] = '1' or [t mp_smx_2] = '2' or [tm p_smx_2] = '3'	TMP/SMX-MIC:	text
120	[tige_mic_2] Show the field ONLY i f: [tig_2] = '1' or [tig_2] = '2' or [tig_2] = '3'	TIGECYCLINE-MIC:	text
121	[col_mic_2] Show the field ONLY i f: [col_2] = '1' or [col_2] = '2' or [col_2] = '3'	COLISTIN-MIC:	text
122	[cef_avi_mic_2] Show the field ONLY i f:	CEFTAZIDIME-AVIBACTAM-MIC:	text

		[cef_avi_2] = '1' or [cef_ avi_2] = '2' or [cef_avi_ 2] = '3'		
	123	[mero_vab_mic_2] Show the field ONLY i f: [mero_vabo_2] = '1' or [mero_vabo_2] = '2' or [mero_vabo_2] = '3'	MEROPENEM/VABORBACTAM-MIC:	text
	124	[imi_rel_mic_2] Show the field ONLY i f: [imi_rel_2] = '1' or [imi_ rel_2] = '2' or [imi_rel_ 2] = '3'	IMIPENEM/RELEBACTAM-MIC:	text
	125	[cef_mic_2] Show the field ONLY i f: [cef_2] = '1' or [cef_2] = '2' or [cef_2] = '3'	CEFIDEROCOL-MIC:	text
	126	[other_drug_specify_2] Show the field ONLY if: [other_ant_2] = '1' or [other_ant_2] = '2' or [other_ant_2] = '3' or [other_ant_2] = '4'	Other - Please, specify:	text
	127	[other_mic_2] Show the field ONLY i f: [other_ant_2] = '1' or [o ther_ant_2] = '2' or [ot her_ant_2] = '3'	Other-MIC:	text
	128	[source_control] Show the field ONLY if: [cre_infection] = '1'	Source control?	radio 0 No 1 Yes 2 NA
	129	<pre>[management_of_cre_co lonization_and_cre_in fection_complete]</pre>	Section Header: Form Status Complete?	dropdown 0 Incomplete 1 Unverified 2 Complete
Inst	rum	ent: CRECOOLT scor	e evaluation (crecoolt_score_evaluation)	
	130	[new_events] Show the field ONLY if:	Section Header: CRECOOLT score should be collected during the prospective period weekly and/or upon to new events (see list below), until day 60 after OLT	descriptive

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	[olt_date_of_transp_ar m_1][retro_or_pros]=2	([olt_date_of_transp_arm_1][date_of_transplant]) or hospital discharge				
		NEW EVENTS: - CRE colonization detected post-transplant - Multisite colonization - Need for prolonged (≥48 hours) mechanical ventilation - Acute renal failure - Reintervention				
		>>> Go to the CRECOOLT score calculator				
1:	Show the field ONLY i f: [olt_date_of_transp_ar m_1][retro_or_pros]=1	Since the OLT ([olt_date_of_transp_arm_1] [date_of_transplant]) has been performed before this study protocol, you don't need to fill in this form. Please, skip it and go to the next form.	des	scriptive		
1:	32 [score_date]	CRECOOLT score evaluation date:	tex	t (date_dmy)		
	Show the field ONLY i f: [olt_date_of_transp_ar m_1][retro_or_pros]=2					
1:	Show the field ONLY i f: [olt_date_of_transp_ar m_1][retro_or_pros]=2	Have any events occurred?	yes 1 0	no Yes No		
1:	34 [event_type]		checkbox			
	Show the field ONLY i f:	Check all that apply	1	event_type1	Surgical reinterventation	
	[olt_date_of_transp_ar m_1][retro_or_pros]=2		2	event_type2	CRE colonization post-transplant	
	and [events]=1		3	event_type3	Acute renal failure	
			4	event_type4	Mechanical ventilation	
			5	event_type5	CRE colonization 60-days prior to transplant	
			6	event_type6	Multi-site colonization post- transplant	
			7	event_type7	Other (please, specify below)	

	135	[other_event]	Other event - Please, specify:	text
		Show the field ONLY i		
		f:		
		[olt_date_of_transp_ar m_1][retro_or_pros]=2		
		and [event_type(7)]=1		
	136	[score]	Estimated probability of CRE infections (%)	text (number, Min: 0, Max: 100)
		Show the field ONLY i	according to the CRECOOLT score: Range: 0-100%	
		f: [olt_date_of_transp_ar		
		m_1][retro_or_pros]=2		
	137	[score_impact]	Did the CRECOOLT score impact on patient	checkbox
		Show the field ONLY i	management?	1 score_impact1 No impact
		f: [olt_date_of_transp_ar		2 score_impact2 Yes, an anti-CRE
		m_1][retro_or_pros]=2		therapy was started
				3 score_impact3 Yes, diagnostic
				tests were performed
	138	[cre_therapy]	Which type of anti-CRE therapy? New drugs regimen: ceftazidime/avibactam,	checkbox
		Show the field ONLY i f:	meropenem/vaborbactam, cefiderocol, ceftolozane/tazobactam	1 cre_therapy1 New drugs regimen
		[score_impact(2)]=1	ecjeolozumortazobactam	2 cre_therapy2 Old drugs regimen
	139	[diagno_test]	Which type of diagnostic testing?	checkbox
		Show the field ONLY i		1 diagno_test1 Imaging
		f:		2 diagno_test2 Microbiological
		[score_impact(3)]=1		testing
	140	[crecoolt_score_evalu	Section Header: Form Status	dropdown
		ation_complete]	Complete?	0 Incomplete
				1 Unverified
				2 Complete
Inst	rum	ent: Microbiological	Screening (microbiological_screening)	
	141	[micro_screen_blood]	Section Header: In the 60 days from OLT ([date_of_transplant]), please indicate:	text (integer)
			Overall number of blood cultures performed:	
	142	[micro_screen_resp]	Overall number of respiratory samples performed:	text (integer)
			(e.g. BAL)	
	143	[micro_screen_urino]	Overall number of urinocultures performed:	text (integer)
	144	<pre>[micro_screen_intra_a bd]</pre>	Overall number of intra-abdominal samples performed:	text (integer)
	145	[microbiological_scre	Section Header: Form Status	dropdown
		<pre>ening_complete]</pre>	Complete?	0 Incomplete

				1	Unverified		
				2	Complete		
Inst	rum	ent: Exposure To CR	E Antibiotics (exposure_to_cre_antibiotic	s)			
	146	[drug]	Section Header: Exposure to CRE antibiotics within 180	dro	opdown (autocor	mplete)	
			days after OLT ([date_of_transplant]) Drug:	1 ERTAPENEM			
			Drug.	2	MEROPENEM		
				3	AMIKACIN		
				4	GENTAMICIN		
				5	TMP/SMX		
				6	TIGECYCLINE		
				7	COLISTIN		
				8	CEFTAZIDIME-	AVIBACTAM	
				9	MEROPENEM	/VABORBACT	AM
				10	IMIPENEM/RE	LEBACTAM	
				11	1 CEFIDEROCOL	-	
				12	2 FOSFOMYCIN		
				13	3 Other		
	147	[other_drug]	Other drug - Please, specify:	tex	άt		
		Show the field ONLY i					
		f:					
	1/18	[drug]=13 [combination_treat]	Drug administered in combination with	VO	sno		
	140	[compination_treat]	other anti-Gram-negative antibiotics?	1	Yes		
				0	No		
	149	[daily_dosage]	Total daily dosage (mg):	tov	tt (number)		
	143	[uaily_uosage]	In milligrams (mg)	ic.	at (Halliber)		
	150	[daily_dosage_adjuste	Adjustment daily dosage	che	eckbox		Г
		d]		1	daily_dosage_a	djusted1	Renal function
				2	daily_dosage_a	diustad 2	TDM
				3	daily_dosage_a		Other
					duny_dosage_a	ajastea5	(please,
							specify below)
	454		OIL DI V	<u> </u>			below)
	151		Other - Please, specify:	tex	τ		
		Show the field ONLY i f:					
		[daily_dosage_adjuste					
		d(3)]=1					
	152	<pre>[administration_sched ule]</pre>	Administration schedule	rac			
		~=~]		1	Intermittent		

				2 Continuous 3 Prolonged
	153	[start_date]	Start date	text (date_dmy)
	154	[end_date]	End date	text (date_dmy)
	155	[indication]	Indication	radio 1 Symptoms onset 2 CRECOOLT score 3 Documented infection
	156	<pre>[exposure_to_cre_anti biotics_complete]</pre>	Section Header: Form Status Complete?	dropdown 0 Incomplete 1 Unverified 2 Complete
Inst	rum	ent: Outcome at 180	days (outcome_at_180_days)	
	157	[death]	All-cause death whitin 180-days from [date_of_transplant]?	yesno 1 Yes 0 No
	158	[death_date] Show the field ONLY i f: [death]=1	Please, specify the date:	text (date_dmy)
	159	[discharge_date]	Date of hospital discharge after transplantation:	text (date_dmy)
	160	[lenght_of_icu_stay]	Lenght of ICU stay after transplantation: days	calc Calculation: round(datediff([date_discharge_icu], [date_admission_icu],'d','dmy'))
	161	<pre>[hospital_readmission s]</pre>	Number of hospital readmissions within 6 months after OLT:	text (number)
	162	[cre_clearence]	Documentation of CRE colonization clearance? CRE colonization clearance: three consecutive surveillance culture negative results.	yesno 1 Yes 0 No
	163	[clearence_date] Show the field ONLY i f: [cre_clearence]=1	Please, specify the date:	text (date_dmy)
	164	[cre_recurrence]	CRE infection recurrence?	yesno 1 Yes 0 No
	165	[relapse_date] Show the field ONLY i f:	Please, specify the date:	text (date_dmy)

	[cre_recurrence]=1		
166	[further_resistance]	Emergence of further resistance? This could apply to either persistent colonization or relapse isolates.	yesno 1 Yes 0 No
167	<pre>[new_resistance_dat e] Show the field ONLY i f: [further_resistance]=1</pre>	Please, specify the date of isolation resistant strain with the new resistance profile:	text (date_dmy)
168	<pre>[new_resistance_mole c] Show the field ONLY i f: [further_resistance]=1</pre>	Molecular characteristics: Check all that apply.	checkbox 1 new_resistance_molec1 KPC 2 new_resistance_molec2 Oxa-48 3 new_resistance_molec3 VIM 4 new_resistance_molec4 NDM 5 new_resistance_molec5 Other (please, specify below)
169	[other_mol] Show the field ONLY i f: [new_resistance_molec (5)]=1	Other - Please, specify:	text
170	[erta_3]	Section Header: Susceptibility of CRE infection strain: ERTAPENEM	radio (Matrix) 1 R 2 S 3 I 4 NA
171	[mero_3]	MEROPENEM	radio (Matrix) 1 R 2 S 3 I 4 NA
172	[amik_3]	AMIKACIN	radio (Matrix) 1 R 2 S 3 I 4 NA
173	[gent_3]	GENTAMICIN	radio (Matrix) 1 R 2 S

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				3 I 4 NA
	174	[tmp_smx_3]	TMP/SMX	radio (Matrix) 1 R 2 S 3 I 4 NA
	175	[tig_3]	TIGECYCLINE	radio (Matrix) 1 R 2 S 3 I 4 NA
	176	[col_3]	COLISTIN	radio (Matrix) 1 R 2 S 3 I 4 NA
	177	[cef_avi_3]	CEFTAZIDIME-AVIBACTAM	radio (Matrix) 1 R 2 S 3 I 4 NA
	178	[mero_vabo_3]	MEROPENEM/VABORBACTAM	radio (Matrix) 1 R 2 S 3 I 4 NA
	179	[imi_rel_3]	IMIPENEM/RELEBACTAM	radio (Matrix) 1 R 2 S 3 I 4 NA
	180	[cef_3]	CEFIDEROCOL	radio (Matrix) 1 R 2 S 3 I 4 NA
	181	[other_ant_3]	Other	radio (Matrix)

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			1 R 2 S 3 I 4 NA
18	S2 [erta_mic_3] Show the field ONLY i f: [erta_3] = '1' or [erta_3] = '2' or [erta_3] = '3'	ERTAPENEM-MIC:	text
18	Show the field ONLY i f: [mero_3] = '1' or [mero _3] = '2' or [mero_3] = '3'	MEROPENEM-MIC:	text
18	Show the field ONLY i f: [amik_3] = '1' or [amik_ 3] = '2' or [amik_3] = '3'	AMIKACIN-MIC:	text
18	Show the field ONLY i f: [gent_3] = '1' or [gent_ 3] = '2' or [gent_3] = '3'	GENTAMICIN-MIC:	text
18	Show the field ONLY i f: [tmp_smx_3] = '1' or [t mp_smx_3] = '2' or [tm p_smx_3] = '3'	TMP/SMX-MIC:	text
18	Show the field ONLY i f: [tig_3] = '1' or [tig_3] = '2' or [tig_3] = '3'	TIGECYCLINE-MIC:	text
18	Show the field ONLY i f: [col_3] = '1' or [col_3] = '2' or [col_3] = '3'	COLISTIN-MIC:	text

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	189	[cef_avi_mic_3] Show the field ONLY i	CEFTAZIDIME-AVIBACTAM-MIC:	text
		f: [cef_avi_3] = '1' or [cef_ avi_3] = '2' or [cef_avi_ 3] = '3'		
	190	[mero_vab_mic_3] Show the field ONLY i f: [mero_vabo_3] = '1' or [mero_vabo_3] = '2' or [mero_vabo_3] = '3'	MEROPENEM/VABORBACTAM-MIC:	text
	191	<pre>[imi_rel_mic_3] Show the field ONLY i f: [imi_rel_3] = '1' or [imi_ rel_3] = '2' or [imi_rel_ 3] = '3'</pre>	IMIPENEM/RELEBACTAM-MIC:	text
	192	[cef_mic_3] Show the field ONLY i f: [cef_3] = '1' or [cef_3] = '2' or [cef_3] = '3'	CEFIDEROCOL-MIC:	text
	193	[other_drug_specify_3] Show the field ONLY i f: [other_ant_3] = '1' or [o ther_ant_3] = '2' or [ot her_ant_3] = '3' or [oth er_ant_3] = '4'	Other - Please, specify:	text
	194	[other_mic_3] Show the field ONLY i f: [other_ant_3] = '1' or [o ther_ant_3] = '2' or [ot her_ant_3] = '3'	Other-MIC:	text
	195	[outcome_at_180_days_ complete]	Section Header: Form Status Complete?	dropdown 0 Incomplete 1 Unverified 2 Complete