

CRF Specification for Test Trial abcd

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Protocol Name: Test Trial

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AE/CM/CP/Other events involving IMP Events <i>Design note: This form is only used in adolescent trials, where a subject might reach a country specific legal age that require them to sign an informed consent during the conduct of the trial.</i>			SDTM Annotation
1.1	Did the subject have any adverse events since screening	<input type="radio"/> Yes <input type="radio"/> No	NOT SUBMITTED
1.2	Did the subject take any concomitant medications?	<input type="radio"/> Yes <input type="radio"/> No	NOT SUBMITTED
1.3	Did the subject have any concurrent procedures/surgeries since screening?	<input type="radio"/> Yes <input type="radio"/> No	NOT SUBMITTED
1.4	Did the subject have any other events involving IMP since screening? <i>Internal note: If the code Lack of efficacy is added to the item 'other event involving IMP' on the form 'other event involving IMP' then adjust the CRF guidance text accordingly</i>	<input type="radio"/> Yes <input type="radio"/> No	

Demographics			SDTM Annotation
1.1	Date of birth	<input type="text" value="mm/dd/yyyy"/>	BRTHDTC
2.1	Age	<input type="text"/>	QVAL, SUPPDM.QNAM=AGEn. Set SUPPDM.QNAM=AGEnVIS where QVAL=the visit where age is collected
2.2	Age unit	<input type="radio"/> Year(s)	QVAL, SUPPDM.QNAM=AGEnU n is a consecutive counter starting from 1
3.1	Sex	<input type="radio"/> Male <input type="radio"/> Female	SEX
3.2	Ethnicity	<input type="radio"/> Hispanic or Latino <input type="radio"/> Not Hispanic or Latino	ETHNIC
3.3	Race	<input type="radio"/> White <input type="radio"/> Black or African American <input type="radio"/> Asian <input type="radio"/> American Indian or Alaska Native <input type="radio"/> Native Hawaiian or Other Pacific Islander <input type="radio"/> Other	RACE, if RACE=OTHER then value is set to NULL
3.4	Race	<input type="radio"/> White <input type="radio"/> Black or African American <input type="radio"/> Asian Japanese <input type="radio"/> Asian Other <input type="radio"/> American Indian or Alaska Native <input type="radio"/> Native Hawaiian or Other Pacific Islander <input type="radio"/> Other	RACE, if RACE=OTHER then value is set to NULL. If one of the two Asian races are indicated, set RACE=ASIAN and store original value in SUPPDM.QNAM=RACEC
4.1	[If Other,]Specify	<input type="text"/>	QVAL, SUPPDM.QNAM=RACEOTH

Height and weight			SDTM Annotation
1.1	Height	<input type="text"/>	VSORRES, VSTESTCD=HEIGHT
1.2	Unit height	<input type="radio"/> cm <input type="radio"/> in	VSORRESU, VSTESTCD=HEIGHT
1.3	Weight	<input type="text"/>	VSORRES, VSTESTCD=WEIGHT
1.4	Unit weight	<input type="radio"/> kg <input type="radio"/> lb	VSORRESU, VSTESTCD=WEIGHT
1.5	Body Mass Index (BMI)	<input type="text"/>	VSORRES, VSTESTCD=BMI
1.6	BMI unit	<input type="radio"/> kg/m2	VSORRESU, VSTESTCD=BMI

Urine pregnancy test			SDTM Annotation
1.1	Is the female of child-bearing potential?	<input type="radio"/> Yes <input type="radio"/> No	RPORRES, RPTESTCD=CHILDPOT
1.2	[If Yes,]Was a urine pregnancy test performed?	<input type="radio"/> Yes <input type="radio"/> No	RPORRES, RPTESTCD=HCGDONE. If No then PRSTAT=NOT DONE
1.3	[If No,]Specify reason		RPREASND
2.1	[If Yes,]Urine pregnancy test date	mm/dd/yyyy	LBDTC, LBCAT=URINALYSIS, LBTESTCD=HCG, LBSPEC=URINE, LBSPID=COLLECTED
2.2	[If Yes,]Result	<input type="radio"/> Positive <input type="radio"/> Negative	LBORRES, LBCAT=URINALYSIS, LBTESTCD=HCG, LBSPEC=URINE, LBSPID=COLLECTED

Adverse event			SDTM Annotation
1.1	AE identifier	<input type="text"/>	AESPID
2.1	Adverse event	<input type="text"/>	AETERM
3.1	Location of AE relative to treatment area	<input type="radio"/> Non-cutaneous <input type="radio"/> Lesional/perilesional <input type="radio"/> Distant <input type="radio"/> Inside treatment area <input type="radio"/> Outside treatment area <input type="radio"/> Application area	QVAL, SUPPAE.QNAM=AERELTRT
4.1	Location of AE	<input type="radio"/> NA (Non-cutaneous) <input type="radio"/> Back <input type="radio"/> Scalp <input type="radio"/> Face <input type="radio"/> Chest <input type="radio"/> Arm <input type="radio"/> Leg <input type="radio"/> Trunk <input type="radio"/> Limb <input type="radio"/> Site 1 <input type="radio"/> Site 2 <input type="radio"/> Site 3 <input type="radio"/> Site 4	AELOC, If value does not exist on LOC codelist then place in SUPPAE.QVAL where QNAM=ALTLOC.
4.2	Start date	<input type="text" value="mm/dd/yyyy"/>	AESTDTC
5.1	If the adverse event started on the SAME DAY a dose was administered, did the adverse event start before or after dosing of IMP?	<input type="radio"/> Before dosing <input type="radio"/> After dosing <input type="radio"/> Not applicable	QVAL, SUPPAE.QNAM=AEOCCUR

Adverse event			SDTM Annotation
5.2	<p>Did the adverse event start before the first administration of IMP?</p> <p><i>Internal note: Used in e.g. AK trials and Test drug trials. Only one of the questions 'If the adverse event started on the SAME DAY a dose was administered, did the adverse event start before or after dosing of IMP?' or 'Did the adverse event start before the first administration of IMP?' is used.</i></p> <p><i>(tralokinumab/placebo) only to be added to question in trials where additional IMP's than the randomized treatment are included. Align the text in the brackets to the protocol. CRF Guidance text: Note the first administration = the administration at week 0/Visit 3. Adjust CRF Guidance to protocol.</i></p>	<input type="radio"/> Yes <input type="radio"/> No	QVAL, SUPPAE.QNAM=AETRTEM
5.3	Did the adverse event start before the administration of the vaccines?	<input type="radio"/> Yes <input type="radio"/> No	QVAL, SUPPAE.QNAM=AEOCCUR
6.1	<p>Stop date</p> <p><i>Design note: Day can be unknown but month and year must be filled out</i></p>	mm/dd/yyyy	AEENDTC
6.2	Ongoing	<input type="radio"/> Yes <input type="radio"/> No	AEENRTPT, If ONGOING then AEENTPT=END OF TRIAL
6.3	Severity	<input type="radio"/> Mild <input type="radio"/> Moderate <input type="radio"/> Severe	AESEV
6.4	Was the AE related to IMP?	<input type="radio"/> Probably related <input type="radio"/> Possibly related <input type="radio"/> Not related	AEREL
6.5	Action taken with IMP	<input type="radio"/> Dose not changed <input type="radio"/> Dose reduced <input type="radio"/> Dose increased <input type="radio"/> Drug interrupted <input type="radio"/> Drug withdrawn <input type="radio"/> Not applicable <input type="radio"/> Unknown	AEACN

Adverse event			SDTM Annotation
6.6	Was the AE related to AxMP (TCS)?	<input type="radio"/> Probably related <input type="radio"/> Possibly related <input type="radio"/> Not related	AERELNST
7.1	Was the AE related to IMP (Tdap vaccine)?	<input type="radio"/> Probably related <input type="radio"/> Possibly related <input type="radio"/> Not related	QVAL, SUPPAE.QNAM=RELTDAP
7.2	Was the AE related to IMP (meningococcal vaccine)? <i>Internal note: Item only to be added in trials where additional IMP's than the randomized treatment are included. Align the text in the brackets to the protocol.</i>	<input type="radio"/> Probably related <input type="radio"/> Possibly related <input type="radio"/> Not related	QVAL, SUPPAE.QNAM=RELMENI
8.1	Did the AE start at the same day as CYP cocktail was given?	<input type="radio"/> Yes <input type="radio"/> No	QVAL, SUPPAE.QNAM=CYPTRTEM
8.2	[if Yes,]Start time of AE	--:-- --	QVAL, SUPPAE.QNAM=AESTDTC
8.3	[If Yes,]Stop time of AE	--:-- --	QVAL, SUPPAE.QNAM=AEENDTC
9.1	Action taken with AxMP (TCS)	<input type="radio"/> Dose not changed <input type="radio"/> Dose reduced <input type="radio"/> Dose increased <input type="radio"/> Drug interrupted <input type="radio"/> Drug withdrawn <input type="radio"/> Not applicable <input type="radio"/> Unknown	QVAL, SUPPAE.QNAM=AEACNNST
10.1	Other action taken (tick all that apply)	<input type="checkbox"/> None <input type="checkbox"/> Concomitant medication <input type="checkbox"/> Concurrent procedure	AEACNOTH, IF MULTIPLE ANSWERS ARE GIVEN SET AEACNOTH TO MULTIPLE AND STORE INDIVIDUAL ANSWERS IN SUPPAE
11.1	Withdrawn from trial due to this AE?	<input type="radio"/> Yes <input type="radio"/> No	QVAL, SUPPAE.QNAM=AEAWDR
12.1	Was the AE serious?	<input type="radio"/> Yes <input type="radio"/> No	AESER
13.1	SAE criteria (Tick all that apply)	<input type="checkbox"/> Death <input type="checkbox"/> Life-threatening <input type="checkbox"/> In-patient hospitalisation/prolongation of existing hospitalisation <input type="checkbox"/> Persistent or significant disability/incapacity <input type="checkbox"/> Congenital anomaly/birth defect <input type="checkbox"/> Other medically important condition	If ticked, set the relevant variable in AE to Y. If Fatal, set AESDTH. If Life-threatening set AESLIFE. If Hospitalisation set AESHOSP.

Adverse event			SDTM Annotation
14.1	Outcome	<input type="radio"/> Fatal <input type="radio"/> Not recovered/not resolved <input type="radio"/> Recovering/resolving <input type="radio"/> Recovered/resolved <input type="radio"/> Recovered/resolved with sequelae <input type="radio"/> Unknown	AEOUT
15.1	Is the event an adverse event of special interest?	<input type="radio"/> Yes <input type="radio"/> No	QVAL, SUPPAE.QNAM=AESI

Urine dipstick			SDTM Annotation
1.1	Was a urine sample tested with a dipstick?	<input type="radio"/> Yes <input type="radio"/> No	LBORRES, LBCAT=URINALYSIS, LBTESTCD=SAMPLE, LBSPEC=URINE, LBSPID=COLLECTED, LBMETHOD=DIPSTICK. If No, then LBSTAT=NOT DONE and populate LBREASND. If Yes set LBORRES to Y.
1.2	[If No,]Specify reason	<input type="text"/>	LBREASND, LBCAT=URINALYSIS, LBTESTCD=SAMPLE, LBSPEC=URINE, LBSPID=COLLECTED, LBMETHOD=DIPSTICK.
1.3	[If Yes,]Result	<input type="radio"/> Normal <input type="radio"/> Abnormal	LBORRES, LBCAT=URINALYSIS, LBTESTCD=INTP, LBSPEC=URINE, LBSPID=COLLECTED, LBMETHOD=DIPSTICK.

Concomitant medication			SDTM Annotation
1.1	CM Number	<input type="text"/>	CMSPID
1.2	Medication or therapy (generic or brand name)	<input type="text"/>	CMTRT
2.1	Is this medication or therapy a rescue medication for chronic hand eczema?	<input type="radio"/> Yes <input type="radio"/> No	QVAL, SUPPCM.QNAM=RESCMED
3.1	Indication	<input type="text"/>	CMINDC
3.2	Dose per administration	<input type="text"/>	CMDOSTXT, If CMDOSTXT is a numeric value, then transfer the number to the numeric field CMDOSE, and set CMDOSTXT to null
3.3	Unit	<input type="radio"/> mcg <input type="radio"/> mg <input type="radio"/> g <input type="radio"/> MI <input type="radio"/> Application <input type="radio"/> International Unit (IU) <input type="radio"/> Tablet <input type="radio"/> Capsule <input type="radio"/> Other	CMDOSU
4.1	[If Other,]Specify	<input type="text"/>	QVAL, SUPPCM.QNAM = DOSUOTH
5.1	Frequency	<input type="radio"/> As needed (PRN) <input type="radio"/> Daily (QD) <input type="radio"/> Twice daily (BID) <input type="radio"/> Three times per day (TID) <input type="radio"/> Four times per day <input type="radio"/> Weekly (QS) <input type="radio"/> Monthly (QM) <input type="radio"/> Once <input type="radio"/> Other	CMDOSFRQ
6.1	[If Other,]Specify	<input type="text"/>	QVAL, SUPPCM.QNAM = FRQOTH
7.1	Route	<input type="radio"/> Oral <input type="radio"/> Topical <input type="radio"/> Subcutaneous <input type="radio"/> Transdermal <input type="radio"/> Intraocular <input type="radio"/> Intramuscular <input type="radio"/> Respiratory (inhalation) <input type="radio"/> Intralesional <input type="radio"/> Intraperitoneal <input type="radio"/> Nasal <input type="radio"/> Vaginal <input type="radio"/> Rectal <input type="radio"/> Intravenous <input type="radio"/> Other	CMROUTE

Concomitant medication			SDTM Annotation
8.1	[If Topical,]Dose form	<input type="radio"/> Cream <input type="radio"/> Lotion <input type="radio"/> Ointment <input type="radio"/> Other	CMDOSFRM
9.1	[If Other,]Specify	<input type="text"/>	QVAL, SUPPCM.QNAM=FRMOTH
10.1	Start date	<input type="text" value="mm/dd/yyyy"/>	CMSTDTC
10.2	Stop date	<input type="text" value="mm/dd/yyyy"/>	CMENDTC
10.3	Ongoing	<input type="radio"/> Yes <input type="radio"/> No	CMENRTPT, CMENPTPT=END OF TRIAL

Blood biomarkers			SDTM Annotation
1.1	Did the subject sign the additional informed consent for biomarkers?	<input type="radio"/> Yes <input type="radio"/> No	DSDECOD, DSCAT=PROTOCOL MILESTONE. If No then create no record. If Yes then DSTERM=INFORMED CONSENT FOR ADDITIONAL BIOMARKERS, DSDECOD=INFORMED CONSENT OBTAINED
1.2	[If Yes,]Provide date the consent was signed	mm/dd/yyyy	DSSTDTC
2.1	Was sampling performed for blood biomarkers?	<input type="radio"/> Yes <input type="radio"/> No <input type="radio"/> Not applicable	LBORRES, LBCAT=BIOMARKER, LBTESTCD=SAMPLE, LBSPEC=SERUM, LBSPID=COLLECTED. If No, then LBSTAT=NOT DONE, LBORRES and populate LBREASND. If Yes set LBORRES to Y. If NA set LBORRES to NA.
2.2	[If No,]Specify reason		LBREASND, LBCAT=BIOMARKER, LBTESTCD=SAMPLE, LBSPEC=SERUM, LBSPID=COLLECTED.

Concurrent procedures			SDTM Annotation
1.1	Procedure number	<input type="text"/>	PRSPID
1.2	Procedure name (include anatomical area if relevant)	<input type="text"/>	PRTRT
1.3	Procedure name	<input type="text"/>	PRTRT
1.4	Indication	<input type="text"/>	PRINDC
1.5	Start date	<input type="text" value="mm/dd/yyyy"/>	PRSTDTC
1.6	Stop date	<input type="text" value="mm/dd/yyyy"/>	PRENDTC
1.7	Ongoing	<input type="radio"/> Yes <input type="radio"/> No	PRENRTPT, PRENTPT=END OF TRIAL
1.8	Body location	<input type="radio"/> Head <input type="radio"/> Trunk <input type="radio"/> Upper limb <input type="radio"/> Lower limb	PRLOC
2.1	Inside treatment area?	<input type="radio"/> Yes <input type="radio"/> No <input type="radio"/> Unknown <input type="radio"/> Not applicable	QVAL, SUPPPR.QNAM=INTRTAR

Central laboratory			SDTM Annotation
1.1	Was blood sampling performed?	<input type="radio"/> Yes <input type="radio"/> No	LBORRES, LBCAT=BLOOD ANALYSIS, LBTESTCD=SAMPLE, LBSPEC=BLOOD, LBSPID=COLLECTED. If No, then LBSTAT=NOT DONE and populate LBREASND. If Yes set LBORRES to Y.
1.2	[If No,]Specify reason	<input type="text"/>	LBREASND, LBCAT=BLOOD ANALYSIS, LBTESTCD=SAMPLE, LBSPEC=BLOOD, LBSPID=COLLECTED.
1.3	[If Yes,]Date of sampling	<input type="text" value="mm/dd/yyyy"/>	LBDTC, LBCAT=BLOOD ANALYSIS, LBTESTCD=SAMPLE, LBSPEC=BLOOD, LBSPID=COLLECTED.
1.4	[If Yes,]Time of sampling	<input type="text" value="--:-- --"/>	LBDTC, LBCAT=BLOOD ANALYSIS, LBTESTCD=SAMPLE, LBSPEC=BLOOD, LBSPID=COLLECTED.
1.5	[If Yes,]Result	<input type="radio"/> Normal <input type="radio"/> Abnormal, not clinically significant <input type="radio"/> Abnormal, clinically significant	LBORRES, LBCAT=BLOOD ANALYSIS, LBTESTCD=INTP, LBSPEC=BLOOD, LBSPID=COLLECTED. If Normal LBORRES=NORMAL, if Abnormal LBORRES=ABNORMAL.
1.6	Was a urine sample sent for urinalysis?	<input type="radio"/> Yes <input type="radio"/> No <input type="radio"/> Not applicable	LBORRES, LBCAT=URINALYSIS, LBTESTCD=SAMPLE, LBSPEC=URINE, LBSPID=COLLECTED. If No, then LBSTAT=NOT DONE, LBORRES set as blank and populate LBREASND. If Yes set LBORRES to Y.
1.7	[If No,]Specify reason	<input type="text"/>	LBREASND, LBCAT=URINALYSIS, LBTESTCD=SAMPLE, LBSPEC=URINE, LBSPID=COLLECTED.
1.8	[If Yes,]Date of sampling	<input type="text" value="mm/dd/yyyy"/>	LBDTC, LBCAT=URINALYSIS, LBTESTCD=SAMPLE, LBSPEC=URINE, LBSPID=COLLECTED.
1.9	[If Yes,]Time of sampling	<input type="text" value="--:-- --"/>	LBDTC, LBCAT=URINALYSIS, LBTESTCD=SAMPLE, LBSPEC=URINE, LBSPID=COLLECTED.

Central laboratory			SDTM Annotation
1.10	[If Yes,]Result	<input type="radio"/> Normal <input type="radio"/> Abnormal, not clinically significant <input type="radio"/> Abnormal, clinically significant	LBORRES, If Clinically Significant SUPPLB.QVAL=Y(QNAM=LBCLSIG),if Not Clinically significant SUPPLB.QVAL=N(QNAM=LBCLSIG)

ECG			SDTM Annotation
1.1	Was the ECG performed?	<input type="radio"/> Yes <input type="radio"/> No	EGORRES, EGTESTCD=INTP. If Yes then use EGORRES for the result. If No then EGSTAT=NOT DONE and populate EGREASND.
1.2	[If No,]Specify reason	<input type="text"/>	EGREASND, EGTESTCD=INTP
1.3	Date of ECG	<input type="text" value="mm/dd/yyyy"/>	EGDTC, EGTESTCD=INTP
1.4	[If Yes,]Results	<input type="radio"/> Normal <input type="radio"/> Abnormal, not clinically significant <input type="radio"/> Abnormal, clinically significant	EGORRES, EGTESTCD=INTP. If Normal EGORRES= NORMAL, if Abnormal EGORRES= ABNORMAL. If Clinically Significant SUPPEG.QVAL=Y(QNAM=EGCLSIG), if Not Clinically significant SUPPEG.QVAL=N(QNAM=EGCLSIG)

