

cellmeta	
subjectname	CHARACTER VARYING(10)
training	CHARACTER VARYING(40)
region	CHARACTER VARYING(80)
engmode	CHARACTER VARYING
minquality	DOUBLE PRECISION
maxquality	DOUBLE PRECISION
hc_rostral	INTEGER
hc_lateral	INTEGER
hc_ventral	INTEGER
cellid	INTEGER
subjectid	INTEGER
trainingtypeid	INTEGER
penetrationid	INTEGER
siteid	INTEGER
regionid	INTEGER
engmodeid	INTEGER
sortmarkercode	CHARACTER VARYING(5)
byeaud	INTEGER

penetrationmeta	
penetrationid	INTEGER
subjectid	INTEGER
subjectname	CHARACTER VARYING(10)
hemispherename	CHARACTER VARYING(10)
rostral	INTEGER
lateral	INTEGER
serialnumber	CHARACTER VARYING(50)
alphaangle	DOUBLE PRECISION
betaangle	DOUBLE PRECISION
rotationangle	DOUBLE PRECISION
histologycorrectionrostral	INTEGER
histologycorrectionlateral	INTEGER
histologycorrectionventral	INTEGER
cmtop	INTEGER
cmbottom	INTEGER

cellmetab	
subjectname	CHARACTER VARYING(10)
training	CHARACTER VARYING(40)
region	CHARACTER VARYING(80)
engmode	CHARACTER VARYING
minquality	DOUBLE PRECISION
maxquality	DOUBLE PRECISION
cellid	INTEGER
subjectid	INTEGER
trainingtypeid	INTEGER
penetrationid	INTEGER
siteid	INTEGER
regionid	INTEGER
engmodeid	INTEGER
sortmarkercode	CHARACTER VARYING(5)
byeaud	INTEGER

trialeventmeta	
trialtime	TIMESTAMP(6) WITHOUT TIME ZONE
trialid	INTEGER
spiketrainid	INTEGER
eventtime	DOUBLE PRECISION
trialeventid	INTEGER
trialeventtypeid	INTEGER
eventcode1	INTEGER
eventcode2	INTEGER
eventcode3	INTEGER
eventcode4	INTEGER
cellid	INTEGER
epochid	INTEGER
stimulusid	INTEGER

spiketrainmeta	
trialtime	TIMESTAMP(6) WITHOUT TIME ZONE
spiketrainid	INTEGER
spikecount	INTEGER
spiketimes	DOUBLE PRECISION[]
relstarttime	DOUBLE PRECISION
relendtime	DOUBLE PRECISION
trialid	INTEGER
cellid	INTEGER
sortid	INTEGER
epochid	INTEGER
stimulusid	INTEGER

sst	
sstrialid	INTEGER
trialid	INTEGER
sstrialtime	TIMESTAMP(6) WITHOUT TIME ZONE
stimulusid	INTEGER
sseffectiveclassid	INTEGER
sortquality	DOUBLE PRECISION
spiketrainid	INTEGER
cellid	INTEGER

sortmeta_temp	
cellid	INTEGER
sortid	INTEGER
sortquality	DOUBLE PRECISION
isolationtypename	CHARACTER VARYING(100)
isolationtypeid	INTEGER
subjectid	INTEGER
subjectname	CHARACTER VARYING(10)
concatfilename	CHARACTER VARYING(500)

sortmeta	
cellid	INTEGER
sortid	INTEGER
sortquality	DOUBLE PRECISION
isolationtypename	CHARACTER VARYING(100)
isolationtypeid	INTEGER
concatfilename	CHARACTER VARYING(500)

spiketrainmetab	
spiketrainid	INTEGER
trialid	INTEGER
cellid	INTEGER
sortid	INTEGER
sortquality	DOUBLE PRECISION