

srd.db.management	
+mgmt: VARCHAR(40) = PRIMARY KEY	
+country: VARCHAR(50)	
+managementauthority: VARCHAR(200)	

srd.db.assessor	
+assessorid: VARCHAR(40) = PRIMARY KEY	
+mgmt: VARCHAR(40) = REFERENCES mgmt	
+country: VARCHAR(50)	
+assessorfull: VARCHAR(200)	

srd.db.area	
+areaid: VARCHAR(70) = PRIMARY KEY	
+country: VARCHAR(50)	
+areatype: VARCHAR(40) = REFERENCES mgmt	
+areacode: VARCHAR(25)	
+areaname: VARCHAR(100)	
+alternatename: VARCHAR(40)	

srd.db.stock	
+stockid: VARCHAR(40) = PRIMARY KEY	
+tsn: INT = REFERENCES tsn	
+scientificname: VARCHAR(200)	
+commonname: VARCHAR(200)	
+areaid: VARCHAR(70) = REFERENCES areaid	
+stocklong: VARCHAR(200)	
+inmyersddb: INT	
+myersstockid: VARCHAR(40)	

srd.db.referencedoc	
+assessid: VARCHAR(200) = REFERENCES assessid	
+risfield: VARCHAR(50) = REFERENCES risfield	
+risentry: VARCHAR(200)	

srd.db.risfields	
+risfield: VARCHAR(50) = PRIMARY KEY	
+risdescription: VARCHAR(200)	

srd.db.taxonomy	
+tsn: INT = PRIMARY KEY	
+scientificname: VARCHAR(200)	
+kingdom: VARCHAR(200)	
+phylum: VARCHAR(200)	
+classname: VARCHAR(200)	
+ordername: VARCHAR(200)	
+family: VARCHAR(200)	
+genus: VARCHAR(200)	
+species: VARCHAR(200)	
+myersname: VARCHAR(20)	
+commonname1: VARCHAR(200)	
+commonname2: VARCHAR(200)	
+myersscientificname: VARCHAR(200)	
+myersfamily: VARCHAR(50)	

srd.db.assessmentmethod	
+category: VARCHAR(100)	
+methodshort: VARCHAR(20) = PRIMARY KEY	
+methodlong: VARCHAR(200)	

srd.db.assessment	
+assessid: VARCHAR(200)	
+assessorid: VARCHAR(40) REFERENCES assessorid	
+stockid: VARCHAR(40) REFERENCES stockid	
+recorder: VARCHAR(40)	
+daterecorded: DATE	
+assessyear: VARCHAR(9)	
+assesssource: VARCHAR(300)	
+contacts: VARCHAR(300)	
+notes: VARCHAR(1000)	
+pdffile: VARCHAR(300)	
+assess: INT	
+refpoints: INT	
+assessmentmethod: VARCHAR(200) = REFERENCES methodshort	
+assessmentcomments: VARCHAR(1000)	

srd.db.bioparams	
+bssessid: VARCHAR(200) REFERENCES assessid	
+bioid: VARCHAR(40) = REFERENCES biounique	
+biovalue: VARCHAR(100)	
+bioyear: VARCHAR(10)	

srd.db.biometrics	
+biounique: VARCHAR(70) = PRIMARY KEY	
+category: VARCHAR(100)	
+subcategory: VARCHAR(100)	
+bioshort: VARCHAR(40)	
+biolong: VARCHAR(300)	
+biounitsshort: VARCHAR(40)	
+biounitslong: VARCHAR(200)	

srd.db.timeseries	
+assessid: VARCHAR(200) REFERENCES assessid	
+tsid: VARCHAR(40) = REFERENCES tsunique	
+tsyear: INT	
+tsvalue: FLOAT	

srd.db.tsmetrics	
+tsunique: VARCHAR(70) = PRIMARY KEY	
+tscategory: VARCHAR(200)	
+tsshort: VARCHAR(40)	
+tslong: VARCHAR(200)	
+tsunitsshort: VARCHAR(40)	
+tsunitslong: VARCHAR(300)	