

dbSNP
bin: smallint(5)
chrom:varchar(31)
chromStart:int(10)
chromEnd:int(10)
name:varchar(15)
score:smalint(5)
strand:enum('?', '+', '-')
refNCBI:blob
refUCSC:blob
observed:varchar(255)
moltype:genomic:enum('unknown','genomic','cDNA','mito')
class:enum(('unknown', 'single', 'in-del', 'het', 'microsatellite', 'named', 'no var', 'mixed', 'mnp', 'insertion', 'deletion'))
valid:set('unknown', 'by-cluster', 'by-frequency', 'by-submitter', 'by-2hit-2allele', 'by-hapmap')
avHet:float
avHetSE:float
func:set('unknown', 'locus', 'coding', 'coding-synon', 'coding-nonsynon', 'untranslated', 'intron', 'splice-site', 'cds-reference', 'coding-synonymy-unknown', 'gene-segment', 'near-gene-3', 'near-gene-5', 'nonsense', 'missense', 'frameshift', 'untranslated-3', 'untranslated-5', 'splice-3', 'splice-5')
locType:enum('unknown', 'range', 'exact', 'between', 'rangeInsertion', 'rangeSubstitution', 'rangeDeletion')
weight:int(10)