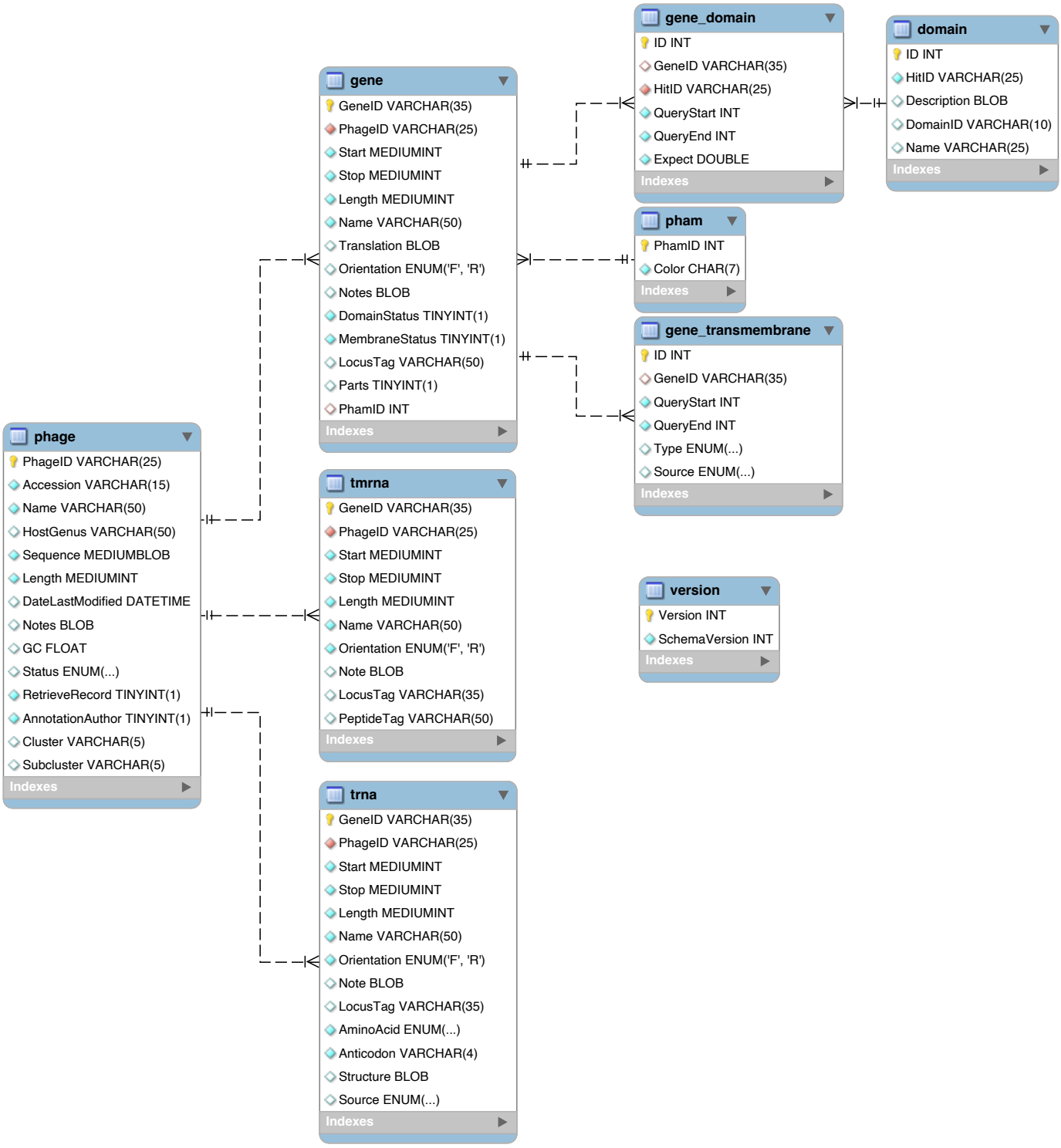
gene_domain
ID INT
GeneID VARCHAR(35)
HitID VARCHAR(25)
QueryStart INT
QueryEnd INT
Expect DOUBLE
Indexes
domain
ID INT
HitID VARCHAR(25)
Description BLOB
DomainID VARCHAR(10)
Name VARCHAR(25)
Indexes
gene
GeneID VARCHAR(35)
PhageID VARCHAR(25)
Start MEDIUMINT
Stop MEDIUMINT
Length MEDIUMINT
Name VARCHAR(50)
Translation BLOB
Orientation ENUM('F', 'R')
Notes BLOB
DomainStatus TINYINT(1)
MembraneStatus TINYINT(1)
LocusTag VARCHAR(50)
Parts TINYINT(1)
PhamID INT
Indexes
pham
PhamID INT
Color CHAR(7)
Indexes
gene_transmembrane
ID INT
GeneID VARCHAR(35)
QueryStart INT
QueryEnd INT
Type ENUM(...)
Source ENUM(...)
Indexes
phage
PhageID VARCHAR(25)
Accession VARCHAR(15)
Name VARCHAR(50)
HostGenus VARCHAR(50)
Sequence MEDIUMBLOB
Length MEDIUMINT
DateLastModified DATETIME
Notes BLOB
GC FLOAT
Status ENUM(...)
RetrieveRecord TINYINT(1)
AnnotationAuthor TINYINT(1)
Cluster VARCHAR(5)
Subcluster VARCHAR(5)
Indexes
tmrna
GeneID VARCHAR(35)
PhageID VARCHAR(25)
Start MEDIUMINT
Stop MEDIUMINT
Length MEDIUMINT
Name VARCHAR(50)
Orientation ENUM('F', 'R')
Note BLOB
LocusTag VARCHAR(35)
PeptideTag VARCHAR(50)
Indexes
version
Version INT
SchemaVersion INT
Indexes
trna
GeneID VARCHAR(35)
PhageID VARCHAR(25)
Start MEDIUMINT
Stop MEDIUMINT
Length MEDIUMINT
Name VARCHAR(50)
Orientation ENUM('F', 'R')
Note BLOB
LocusTag VARCHAR(35)
AminoAcid ENUM(...)
Anticodon VARCHAR(4)
Structure BLOB
Source ENUM(...)
Indexes