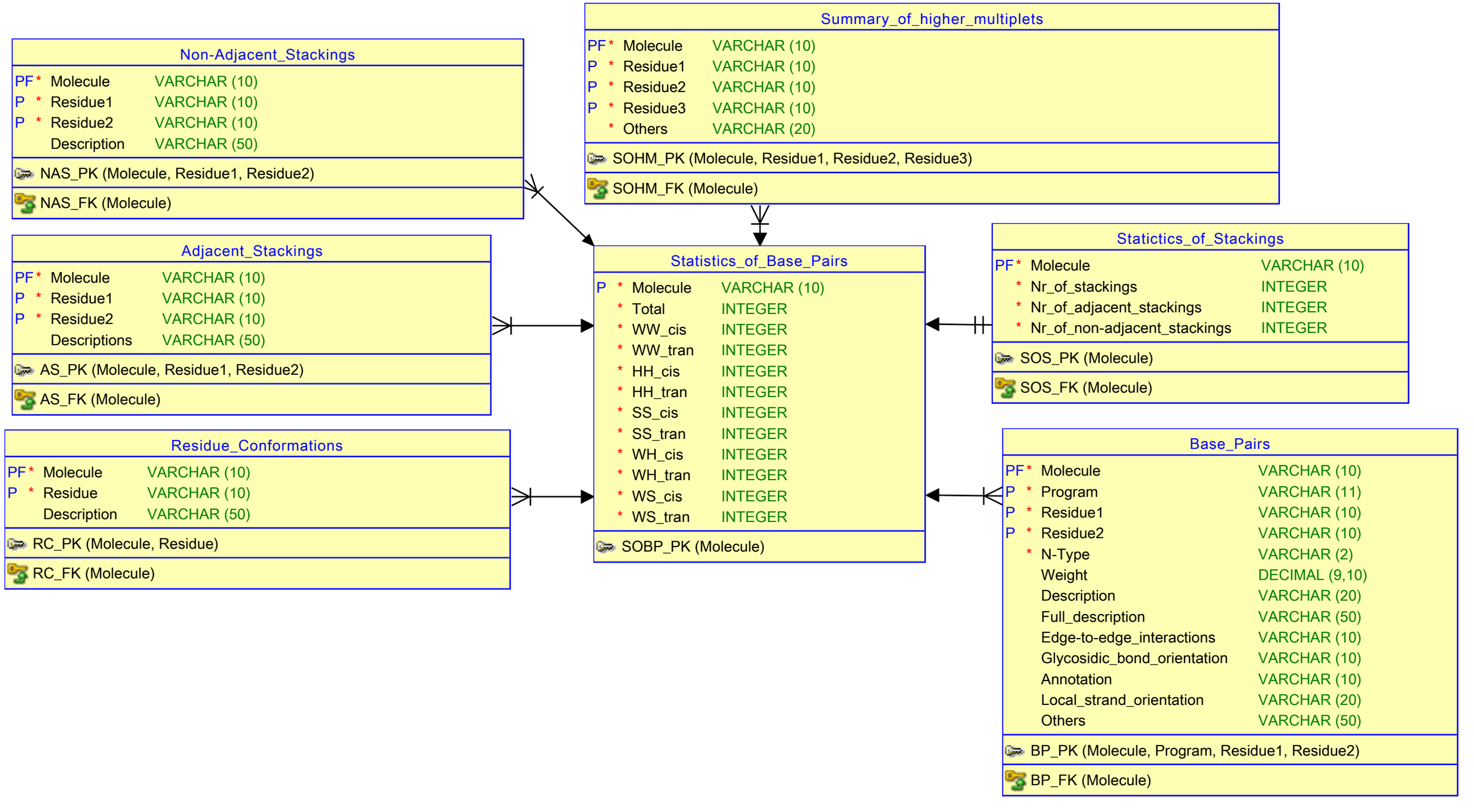
Summary_of_higher_multiplets
PF * Molecule VARCHAR (10)
P * Residue1 VARCHAR (10)
P * Residue2 VARCHAR (10)
P * Residue3 VARCHAR (10)
* Others VARCHAR (20)
SOHM_PK (Molecule, Residue1, Residue2, Residue3)
SOHM_FK (Molecule)
Non-Adjacent_Stackings
PF * Molecule VARCHAR (10)
P * Residue1 VARCHAR (10)
P * Residue2 VARCHAR (10)
Description VARCHAR (50)
NAS_PK (Molecule, Residue1, Residue2)
NAS_FK (Molecule)
Adjacent_Stackings
PF * Molecule VARCHAR (10)
P * Residue1 VARCHAR (10)
P * Residue2 VARCHAR (10)
Descriptions VARCHAR (50)
AS_PK (Molecule, Residue1, Residue2)
AS_FK (Molecule)
Statistics_of_Base_Pairs
P * Molecule VARCHAR (10)
* Total INTEGER
* WW_cis INTEGER
* WW_tran INTEGER
* HH_cis INTEGER
* HH_tran INTEGER
* SS_cis INTEGER
* SS_tran INTEGER
* WH_cis INTEGER
* WH_tran INTEGER
* WS_cis INTEGER
* WS_tran INTEGER
SOBP_PK (Molecule)
Statictics_of_Stackings
PF * Molecule VARCHAR (10)
* Nr_of_stackings INTEGER
* Nr_of_adjacent_stackings INTEGER
* Nr_of_non-adjacent_stackings INTEGER
SOS_PK (Molecule)
SOS_FK (Molecule)
Residue_Conformations
PF * Molecule VARCHAR (10)
P * Residue VARCHAR (10)
Description VARCHAR (50)
RC_PK (Molecule, Residue)
RC_FK (Molecule)
Base_Pairs
PF * Molecule VARCHAR (10)
P * Program VARCHAR (11)
P * Residue1 VARCHAR (10)
P * Residue2 VARCHAR (10)
* N-Type VARCHAR (2)
Weight DECIMAL (9,10)
Description VARCHAR (20)
Full_description VARCHAR (50)
Edge-to-edge_interactions VARCHAR (10)
Glycosidic_bond_orientation VARCHAR (10)
Annotation VARCHAR (10)
Local_strand_orientation VARCHAR (20)
Others VARCHAR (50)
BP_PK (Molecule, Program, Residue1, Residue2)
BP_FK (Molecule)