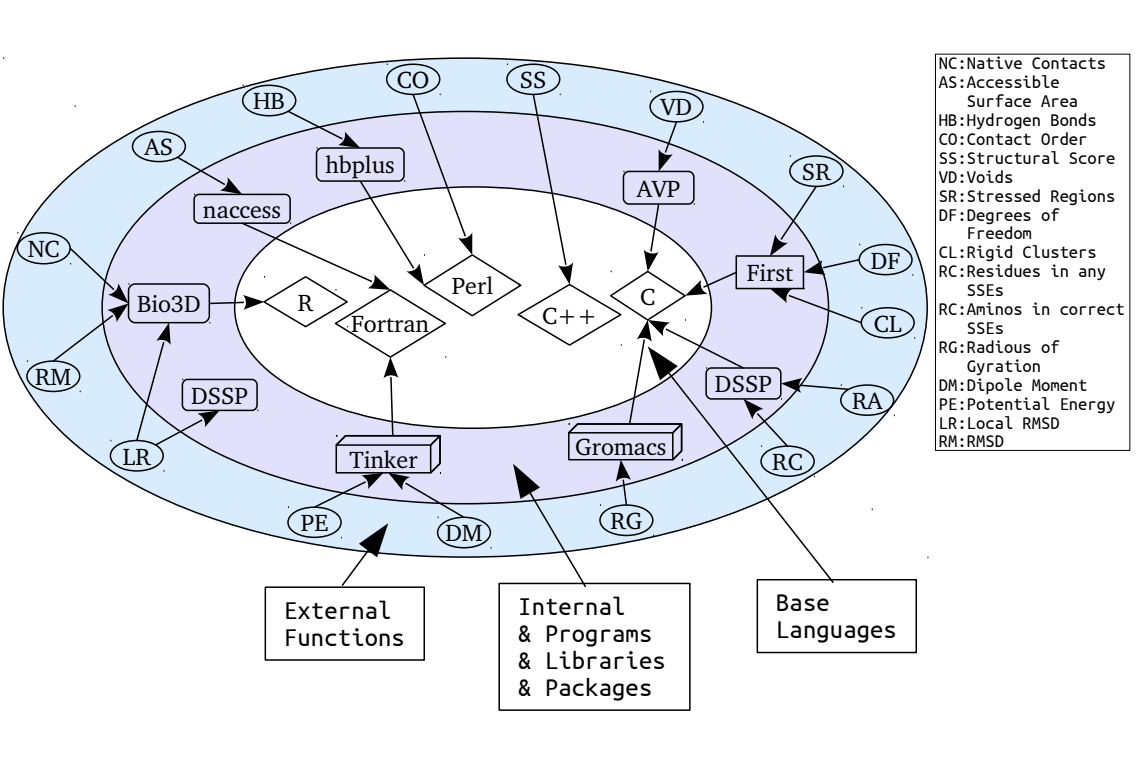
NC:Native Contacts
AS:Accessible Surface Area
HB:Hydrogen Bonds
CO:Contact Order
SS:Structural Score
VD:Voids
SR:Stressed Regions
DF:Degrees of Freedom
CL:Rigid Clusters
RC:Residues in any SSEs
RC:Aminos in correct SSEs
RG:Radious of Gyration
DM:Dipole Moment
PE:Potential Energy
LR:Local RMSD
RM:RMSD
HB
CO
SS
VD
AS
hbplus
AVP
SR
naccess
NC
Bio3D
R
Perl
C++
C
First
DF
Fortran
CL
RM
DSSP
DSSP
RA
LR
Tinker
Gromacs
RC
PE
DM
RG
External Functions
Internal & Programs & Libraries & Packages
Base Languages