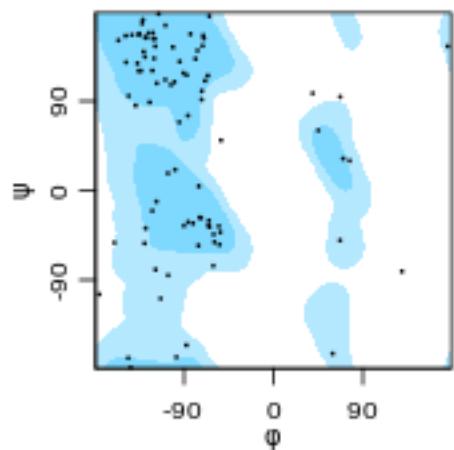
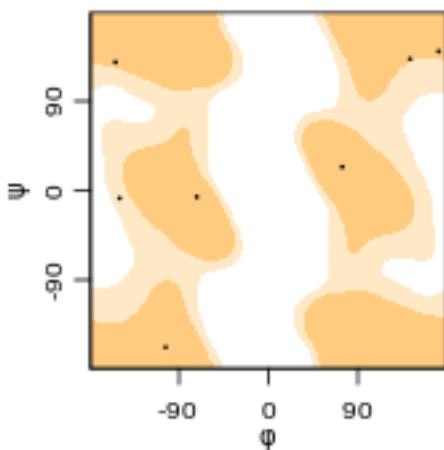


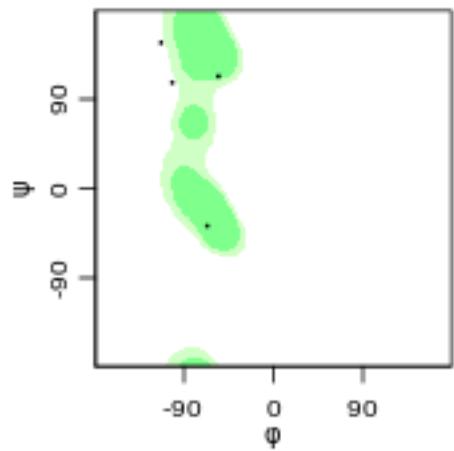
**General**



**Glycine (Symmetric)**



**Proline**

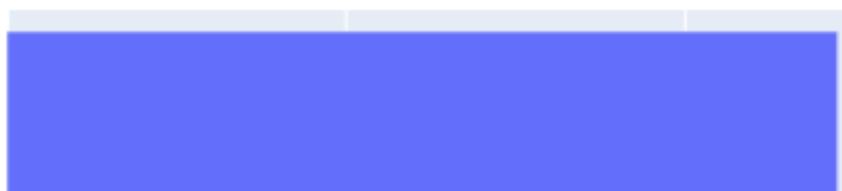


**Pre-Proline**

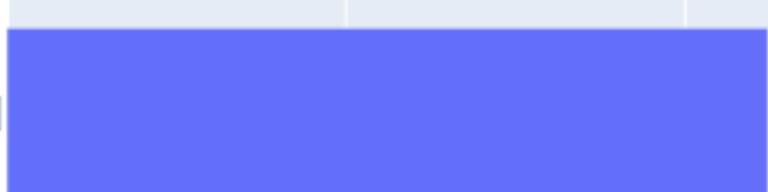


cases

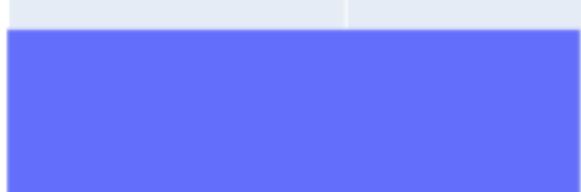
21 GLY HA2-95 PHE HE1



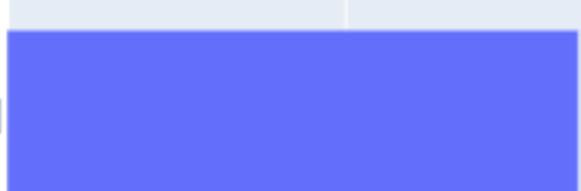
81 ASN HA-85 GLY HN



84 THR HB-86 LYS HG1



21 GLY HA2-93 ALA HN



20 TRP HE3-92 ARG HG1



22 TRP HA-93 ALA HN



84 THR HB-86 LYS HD1



94 PRO HA-95 PHE HD1



weak - violation fraction:0.01

medium - violation fraction:0

strong - violation fraction:0

0 0.5 1  
violations(Å deviation)