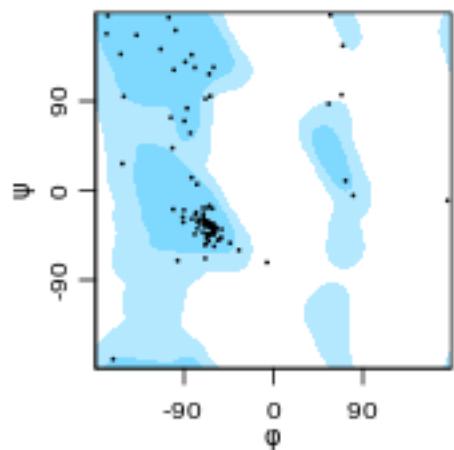
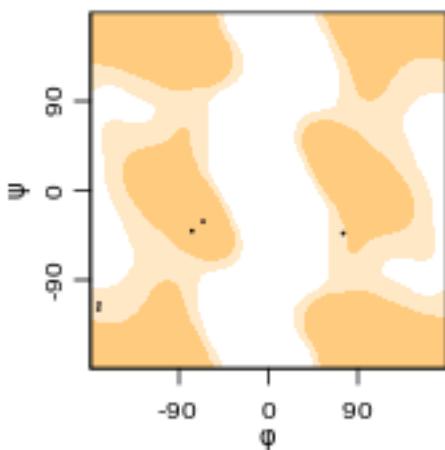


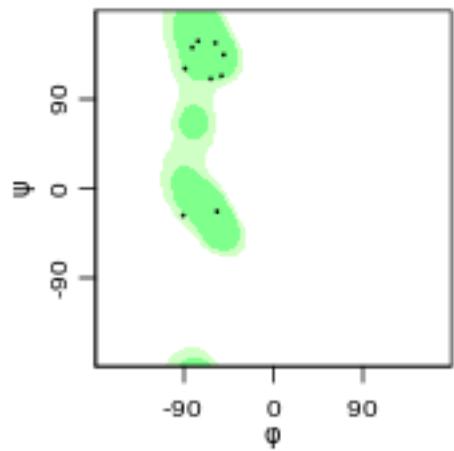
General



Glycine (Symmetric)



Proline

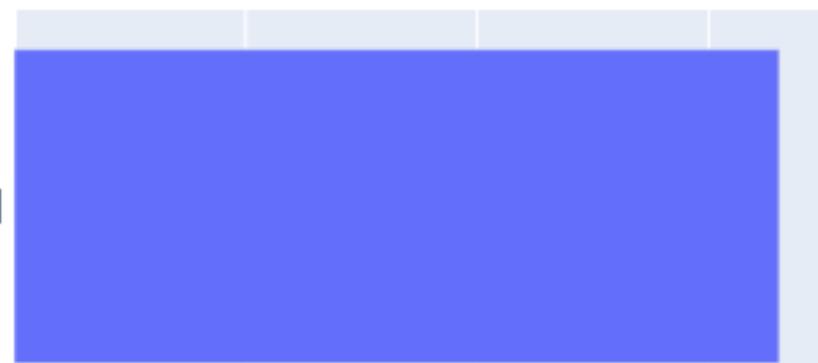


Pre-Proline

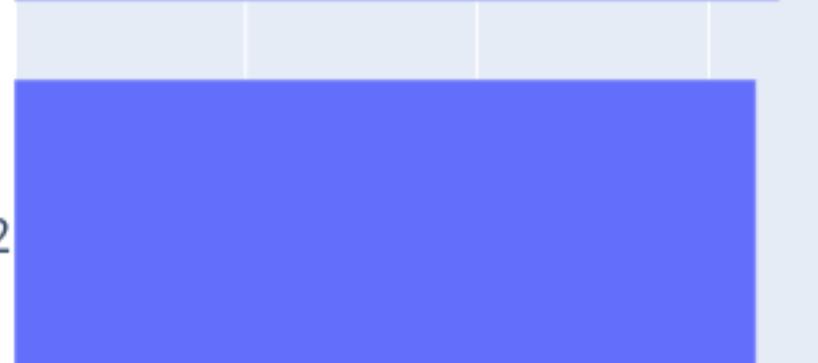


cases

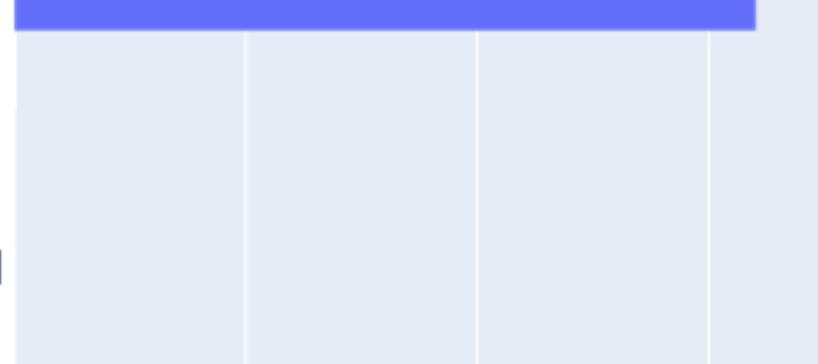
330 LEU HA-375 LEU HB1



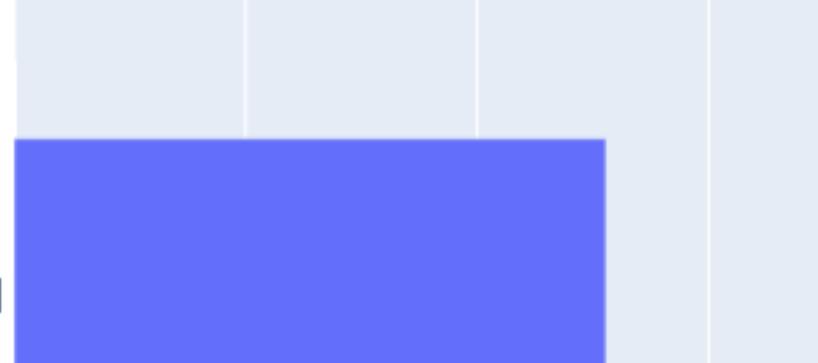
378 TRP HH2-391 LEU HB2



weak - violation fraction:0.01



389 LEU HA-392 LEU HB1



medium - violation fraction:0.02

strong - violation fraction:0

violations(Å deviation)

0 0.2 0.4 0.6