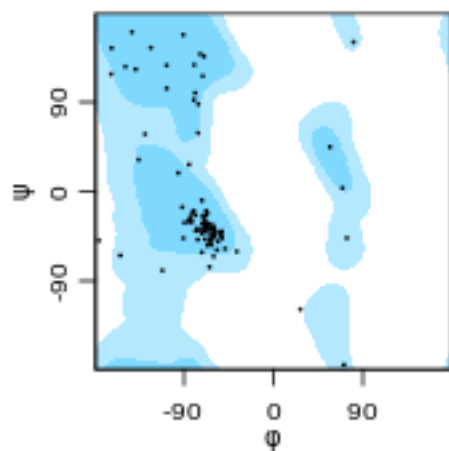
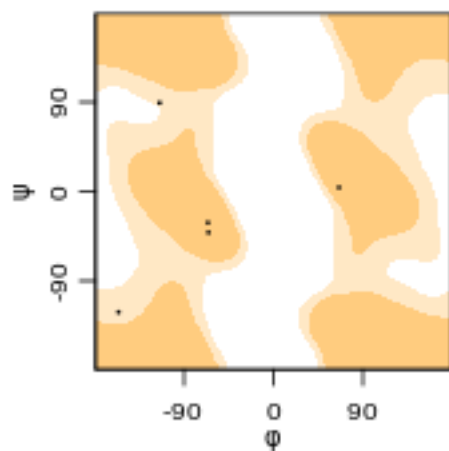
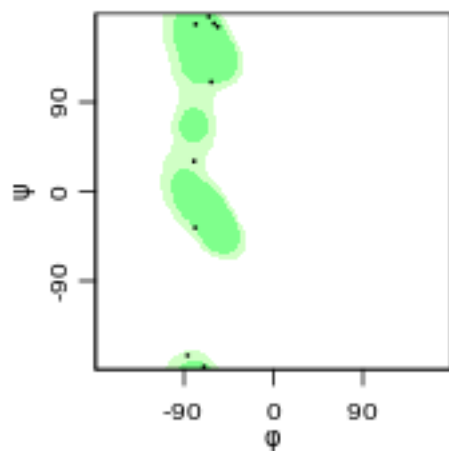


General**Glycine (Symmetric)****Proline****Pre-Proline**

cases

378 TRP HH2-391 LEU HB2

weak - violation fraction:0

medium - violation fraction:0

strong - violation fraction:0

0

0.2

0.4

violations(\AA deviation)

