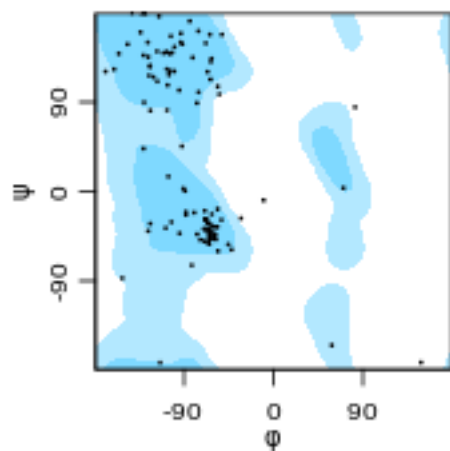
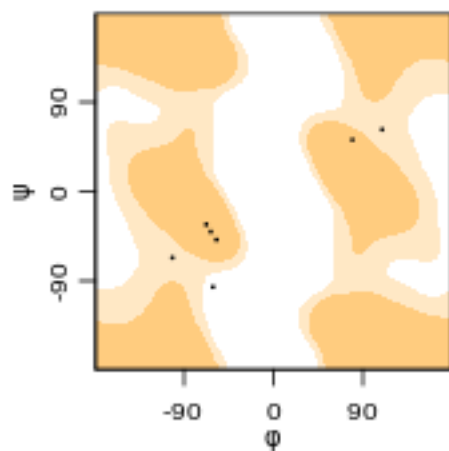
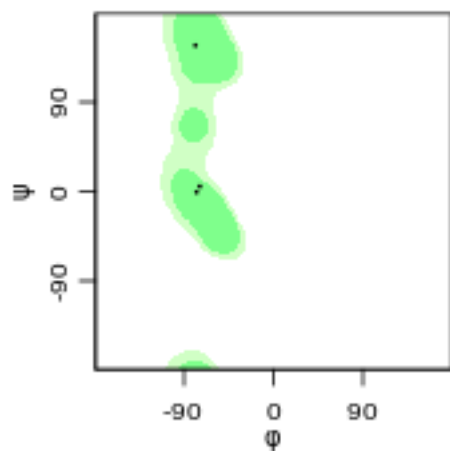


General**Glycine (Symmetric)****Proline****Pre-Proline**

cases

378 ARG HN-426 LEU HN

weak - violation fraction:0

medium - violation fraction:0

strong - violation fraction:0

0 0.2 0.4 0.6

violations(\AA deviation)

