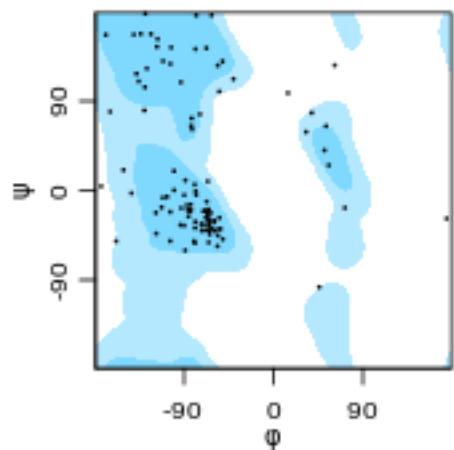
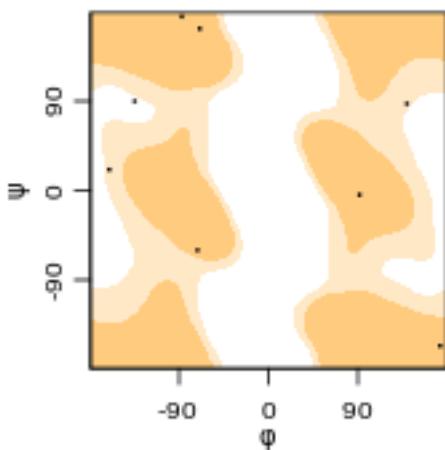


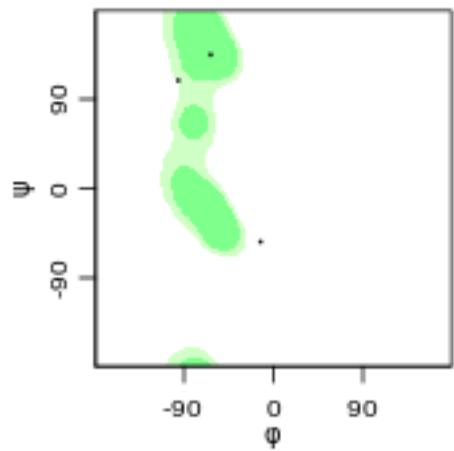
**General**



**Glycine (Symmetric)**



**Proline**



**Pre-Proline**



cases

8 ILE HA-12 ASP HN

26 LYS HA-27 VAL HB

26 LYS HB2-29 ASP HN

3 LYS HN-4 GLU HN

weak - violation fraction:0.01

medium - violation fraction:0

110 ARG HA-111 PRO HD2

strong - violation fraction:0.2

0

0.5

1

violations(Å deviation)