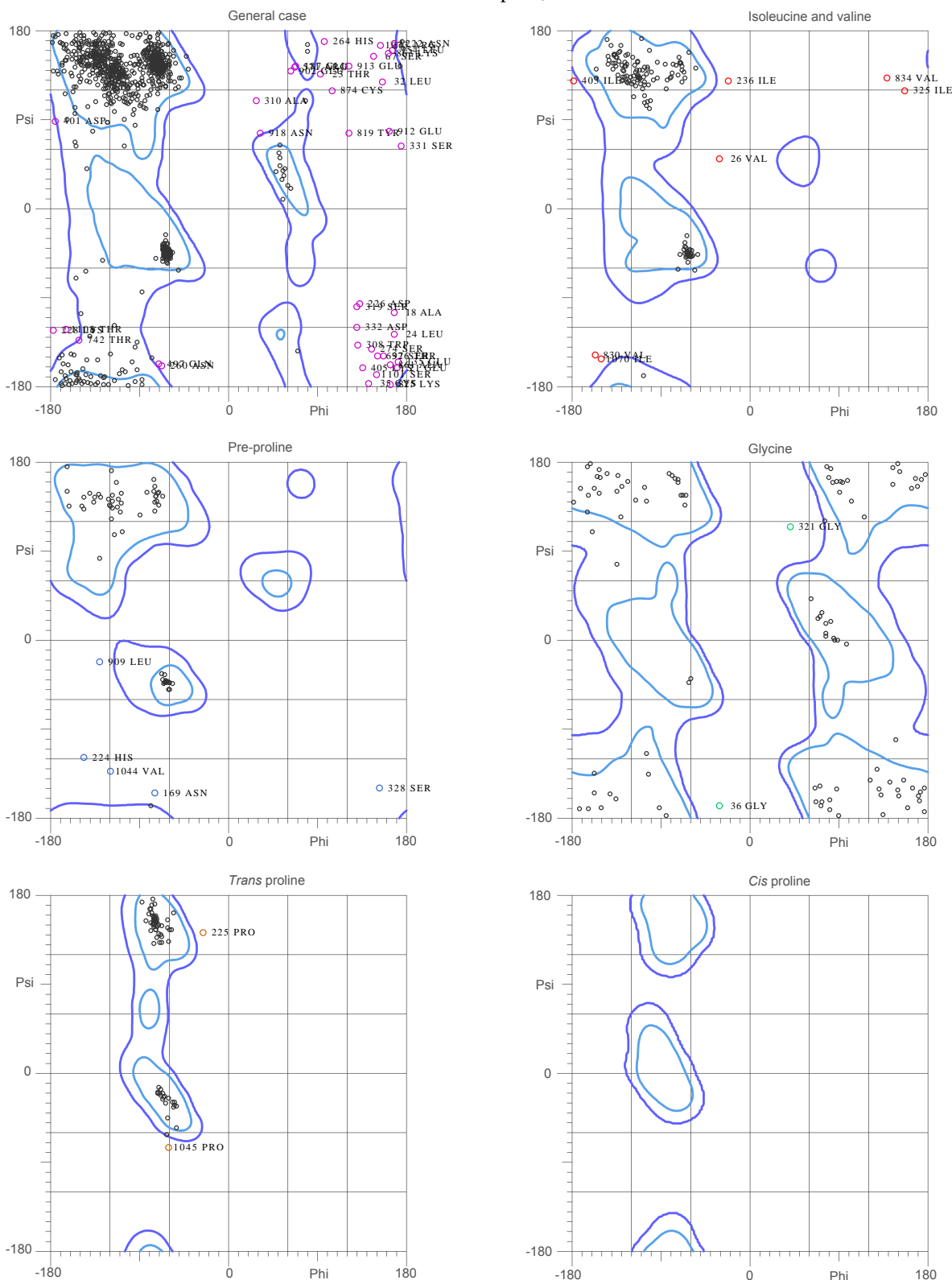


# MolProbity Ramachandran analysis

6abu.B99990025.pdb, model 1



90.7% (1176/1298) of all residues were in favored (90%) regions.

90.7% (1176/1298) of all residues were in allowed (9%) regions.

There were 30 outliers (2.3%):

11 ALA (108.0, -105.0)  
12 ARG (171.0, -104.0)  
13 THR (102.0, -117.0)  
14 ARG (108.0, -107.0)  
15 ARG (108.0, -107.0)  
16 ARG (108.0, -107.0)  
17 ARG (108.0, -107.0)  
18 ARG (108.0, -107.0)  
19 ARG (108.0, -107.0)  
20 ARG (108.0, -107.0)  
21 ARG (108.0, -107.0)  
22 ARG (108.0, -107.0)  
23 ARG (108.0, -107.0)  
24 ARG (108.0, -107.0)  
25 ARG (108.0, -107.0)  
26 ARG (108.0, -107.0)  
27 ARG (108.0, -107.0)  
28 ARG (108.0, -107.0)  
29 ARG (108.0, -107.0)  
30 ARG (108.0, -107.0)

31 THR (102.0, -117.0)  
32 ARG (108.0, -107.0)  
33 THR (102.0, -117.0)  
34 ARG (108.0, -107.0)  
35 THR (102.0, -117.0)  
36 ARG (108.0, -107.0)  
37 THR (102.0, -117.0)  
38 ARG (108.0, -107.0)  
39 THR (102.0, -117.0)  
40 ARG (108.0, -107.0)  
41 THR (102.0, -117.0)  
42 ARG (108.0, -107.0)  
43 THR (102.0, -117.0)  
44 ARG (108.0, -107.0)  
45 THR (102.0, -117.0)  
46 ARG (108.0, -107.0)  
47 THR (102.0, -117.0)  
48 ARG (108.0, -107.0)  
49 THR (102.0, -117.0)  
50 ARG (108.0, -107.0)