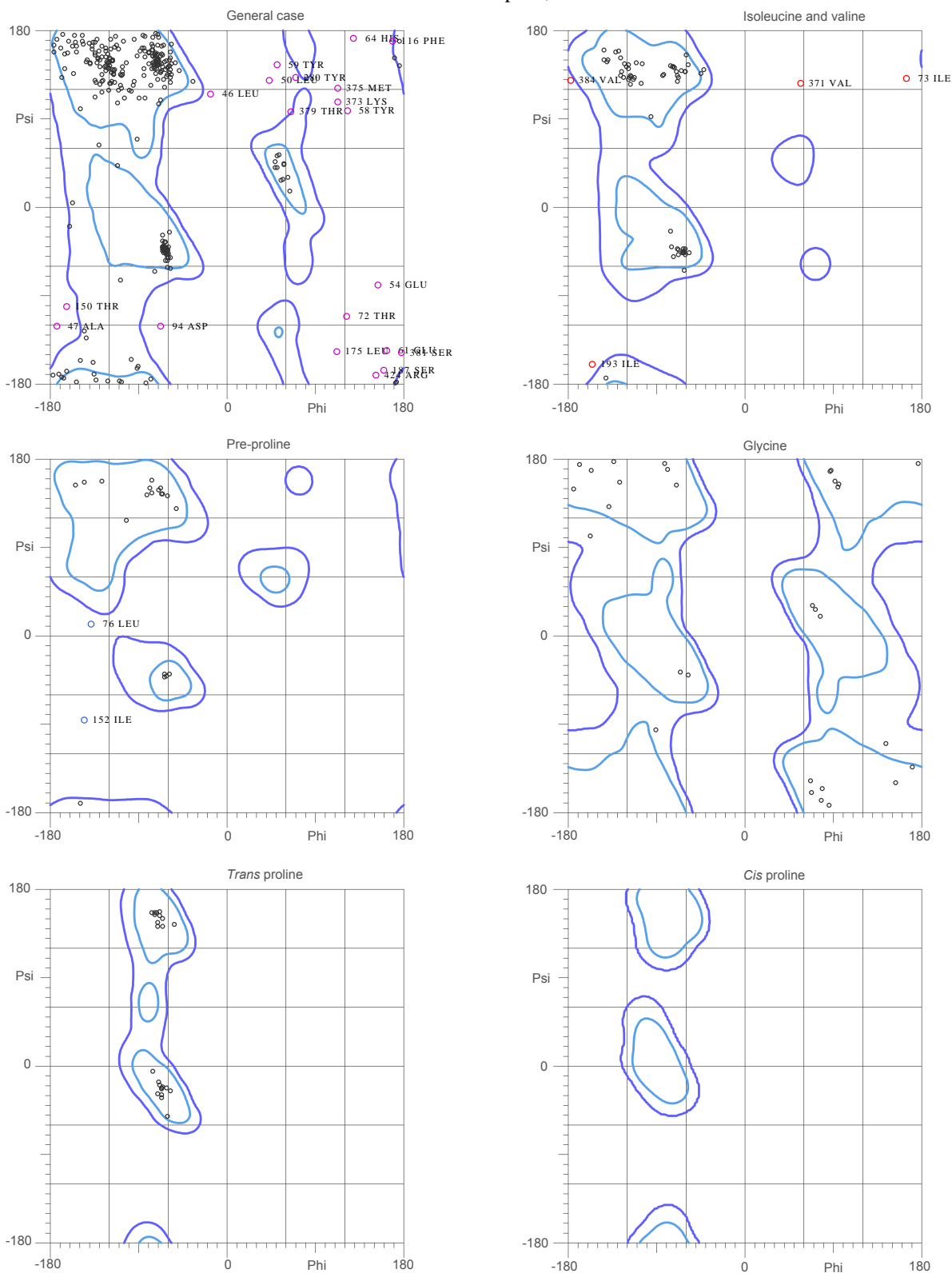


MolProbity Ramachandran analysis

1ad3A.B99990020.pdb, model 1



86.7% (101/116) of all residues were in General (90%) regions.
10.7% (12/116) of all residues were in allowed (100%) regions.

There were 20 outliers (Phi, Psi):

40 ALA (-17.0, 130.7)
41 ALA (-17.0, 130.7)
42 ALA (-17.0, 130.7)
43 ALA (-17.0, 130.7)
44 ALA (-17.0, 130.7)
45 ALA (-17.0, 130.7)
46 ALA (-17.0, 130.7)
47 ALA (-17.0, 130.7)
48 ALA (-17.0, 130.7)
49 ALA (-17.0, 130.7)
50 ALA (-17.0, 130.7)
51 ALA (-17.0, 130.7)
52 ALA (-17.0, 130.7)
53 ALA (-17.0, 130.7)
54 ALA (-17.0, 130.7)
55 ALA (-17.0, 130.7)
56 ALA (-17.0, 130.7)
57 ALA (-17.0, 130.7)
58 ALA (-17.0, 130.7)
59 ALA (-17.0, 130.7)
60 ALA (-17.0, 130.7)
61 ALA (-17.0, 130.7)
62 ALA (-17.0, 130.7)
63 ALA (-17.0, 130.7)
64 ALA (-17.0, 130.7)
65 ALA (-17.0, 130.7)
66 ALA (-17.0, 130.7)
67 ALA (-17.0, 130.7)
68 ALA (-17.0, 130.7)
69 ALA (-17.0, 130.7)
70 ALA (-17.0, 130.7)
71 ALA (-17.0, 130.7)
72 ALA (-17.0, 130.7)
73 ALA (-17.0, 130.7)
74 ALA (-17.0, 130.7)
75 ALA (-17.0, 130.7)
76 ALA (-17.0, 130.7)
77 ALA (-17.0, 130.7)
78 ALA (-17.0, 130.7)
79 ALA (-17.0, 130.7)
80 ALA (-17.0, 130.7)
81 ALA (-17.0, 130.7)
82 ALA (-17.0, 130.7)
83 ALA (-17.0, 130.7)
84 ALA (-17.0, 130.7)
85 ALA (-17.0, 130.7)
86 ALA (-17.0, 130.7)
87 ALA (-17.0, 130.7)
88 ALA (-17.0, 130.7)
89 ALA (-17.0, 130.7)
90 ALA (-17.0, 130.7)
91 ALA (-17.0, 130.7)
92 ALA (-17.0, 130.7)
93 ALA (-17.0, 130.7)
94 ALA (-17.0, 130.7)
95 ALA (-17.0, 130.7)
96 ALA (-17.0, 130.7)
97 ALA (-17.0, 130.7)
98 ALA (-17.0, 130.7)
99 ALA (-17.0, 130.7)
100 ALA (-17.0, 130.7)