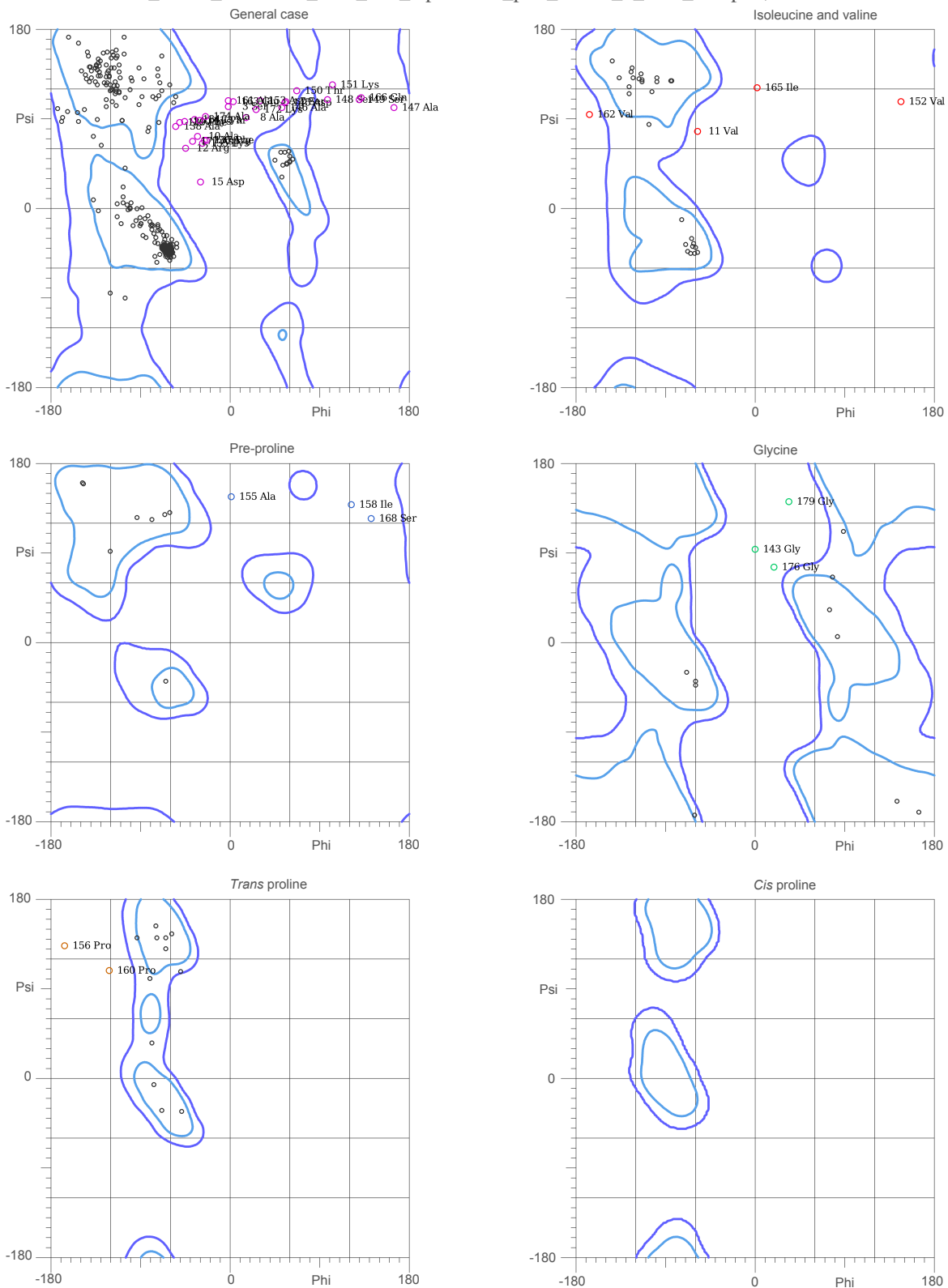


MolProbity Ramachandran analysis

test_76ac3_unrelaxed_rank_001_alphafold2_ptm_model_5_seed_000.pdb, model 1



82.0% (282/344) of all residues were in favored (98%) regions.
88.4% (304/344) of all residues were in allowed (>99.8%) regions.

There were 40 outliers (phi, psi):

3 Ser (-2.2, 103.3)	135 Lys (-28.8, 66.5)
7 Ser (56.5, 108.6)	136 Arg (-26.9, 69.2)
8 Ala (16.8, 92.4)	137 Phe (-24.4, 69.5)
10 Ala (-33.5, 73.4)	138 Ala (-55.4, 83.3)
11 Val (-58.8, 78.9)	139 Phe (-51.9, 87.2)
12 Arg (-45.6, 61.3)	140 Lys (-46.3, 88.0)
15 Asp (-30.5, 27.1)	141 Thr (-31.4, 89.3)
134 Lys (-37.0, 90.5)	143 Gly (-0.8, 94.3)
	145 Asp (55.7, 107.9)
	146 Ala (52.1, 102.3)
	147 Ala (165.3, 102.8)
	148 Ser (98.1, 110.5)
	149 Ser (130.7, 111.0)
	150 Thr (67.0, 119.5)
	151 Lys (103.3, 125.0)
	152 Val (147.3, 108.2)
	153 Asp (30.1, 109.6)
	155 Ala (1.2, 147.1)
	156 Pro (-167.1, 134.9)
	158 Ile (122.9, 139.4)

160 Pro (-122.8, 109.3)
161 Ala (-2.9, 110.0)
162 Val (-167.5, 95.8)
163 Glu (3.2, 108.6)
165 Ile (2.1, 122.6)
166 Gln (132.8, 112.2)
168 Ser (142.9, 125.1)
172 Lys (26.5, 101.0)
174 Ala (-25.6, 93.5)
176 Gly (19.3, 76.5)
177 Asp (-38.4, 68.7)
179 Gly (34.6, 142.4)