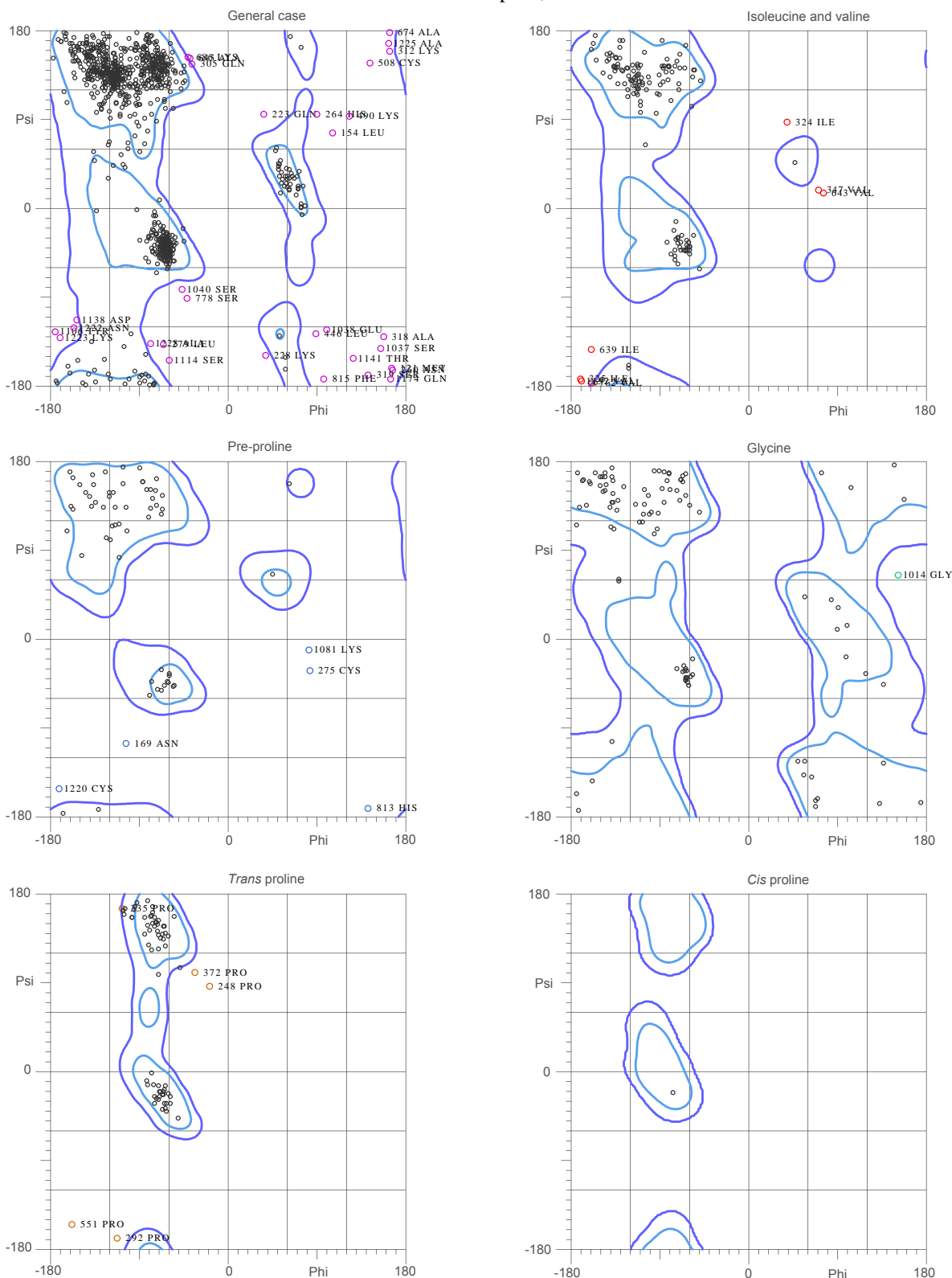


MolProbity Ramachandran analysis

6abu.B99990044.pdb, model 1



90.6% (1179/1302) of all residues were in favored (90%) regions.

96.2% (1251/1302) of all residues were in allowed (49.8%) regions.

These were 49 outliers (49.8%):

121 MET (166.9, -162.6)
151 LEU (106.6, -96.9)
169 ASN (106.6, -106.5)
171 GLN (106.6, -96.9)
178 LYS (161.1, -149.1)
180 ASN (161.1, -149.1)
181 ASN (161.1, -149.1)
182 ASN (161.1, -149.1)
183 ASN (161.1, -149.1)
184 ASN (161.1, -149.1)
185 ASN (161.1, -149.1)
186 ASN (161.1, -149.1)
187 ASN (161.1, -149.1)
188 ASN (161.1, -149.1)
189 ASN (161.1, -149.1)
190 ASN (161.1, -149.1)
191 ASN (161.1, -149.1)
192 ASN (161.1, -149.1)
193 ASN (161.1, -149.1)
194 ASN (161.1, -149.1)
195 ASN (161.1, -149.1)
196 ASN (161.1, -149.1)
197 ASN (161.1, -149.1)
198 ASN (161.1, -149.1)
199 ASN (161.1, -149.1)
200 ASN (161.1, -149.1)

504 CYS (144.9, -144.3)

505 PRO (159.3, -159.3)

506 ILE (149.9, -149.1)

507 VAL (159.3, -159.3)

508 ALA (159.3, -159.3)

509 LYS (159.3, -159.3)

510 LYS (159.3, -159.3)

511 LYS (159.3, -159.3)

512 LYS (159.3, -159.3)

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