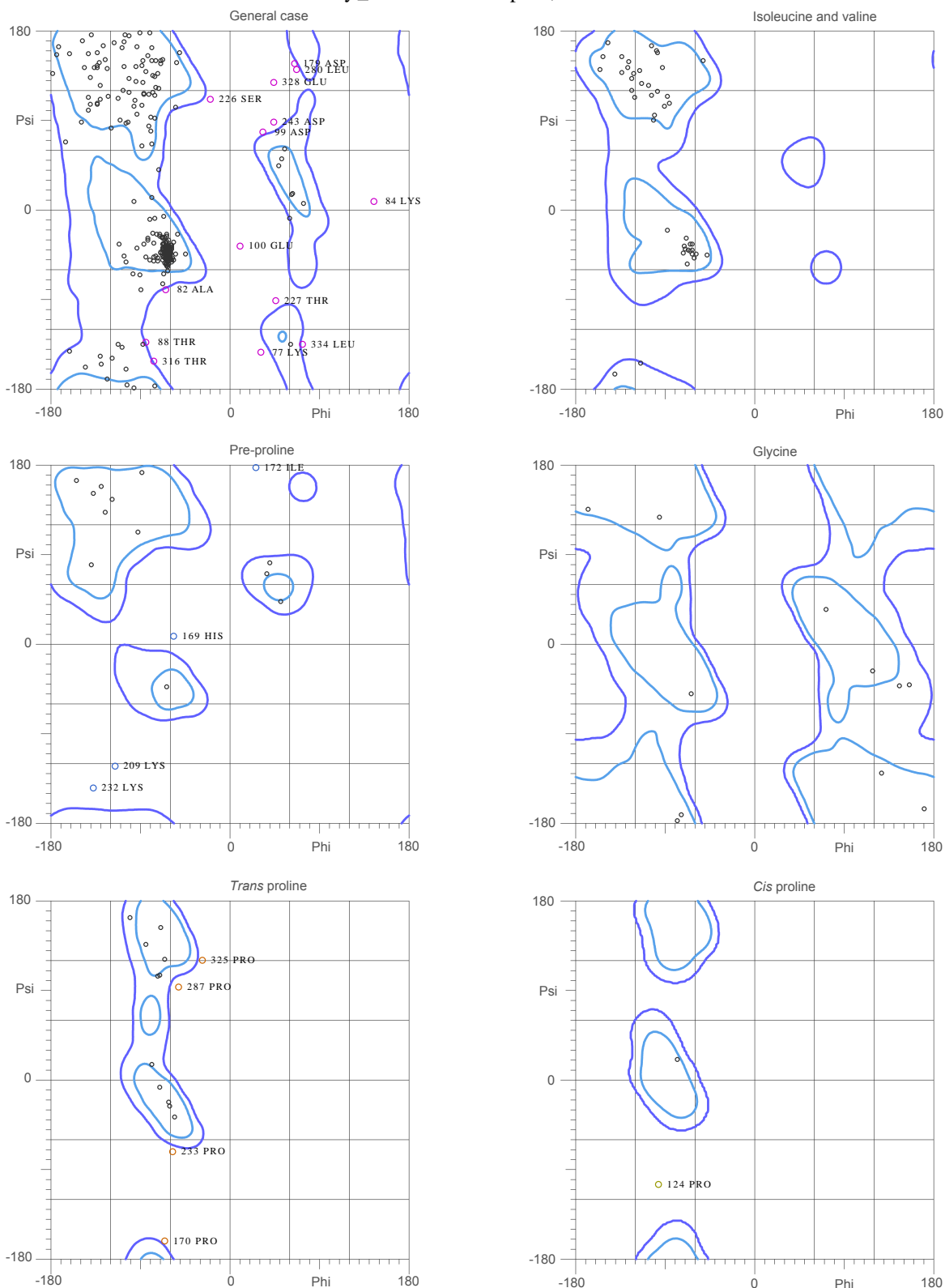


# MolProbity Ramachandran analysis

1a8y\_.B99990024.pdb, model 1



82.7% (278/336) of all residues were in Allowed (99%) regions.  
16.2% (53/336) of all residues were in allowed (95.9%) regions.

There were 31 outliers (out):

179 ASP (15.1, -143.7)  
280 LEU (14.1, -86.9)  
226 SER (14.1, 9.1)  
243 ASP (14.1, -151.3)  
99 ASP (14.1, 79.3)  
100 GLU (14.1, -106.7)  
100 PRO (15.1, -100.3)  
82 ALA (15.1, 8.1)  
170 PRO (14.1, -143.7)  
172 ILE (14.1, -174.9)  
169 HIS (14.1, -143.7)  
209 LYS (14.1, -143.7)  
232 LYS (14.1, -143.7)  
325 PRO (14.1, -143.7)  
287 PRO (14.1, -143.7)  
233 PRO (14.1, -143.7)  
170 PRO (14.1, -143.7)  
124 PRO (14.1, -143.7)  
179 ASP (14.1, -143.7)  
280 LEU (14.1, -143.7)  
226 SER (14.1, -143.7)  
243 ASP (14.1, -143.7)  
99 ASP (14.1, -143.7)  
100 GLU (14.1, -143.7)  
100 PRO (14.1, -143.7)  
82 ALA (14.1, -143.7)  
227 THR (14.1, -143.7)  
334 LEU (14.1, -143.7)  
77 LYS (14.1, -143.7)  
88 THR (14.1, -143.7)  
316 THR (14.1, -143.7)