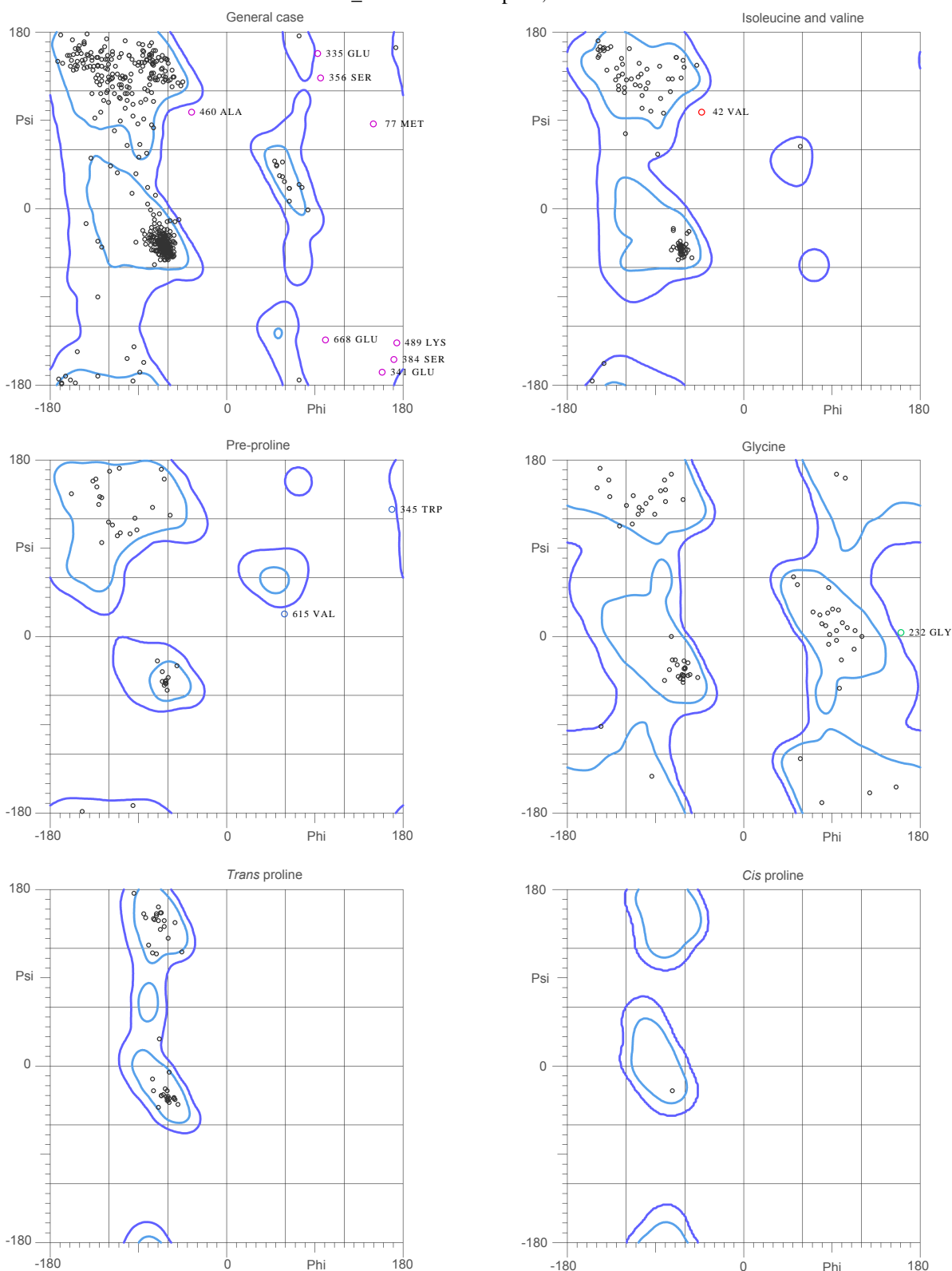


# MolProbity Ramachandran analysis

1aco\_.B99990036.pdb, model 1



93.1% (699/751) of all residues were in favored (98%) regions.  
98.4% (739/751) of all residues were in allowed (>99.8%) regions.

There were 12 outliers ( $\phi$ ,  $\psi$ ):

42 VAL (-43.7, 99.7)  
77 MET (150.5, 87.9)  
232 GLY (162.0, 4.6)  
335 GLU (93.6, 159.7)  
341 GLU (159.5, -167.6)  
345 TRP (169.7, 130.7)  
356 SER (96.5, 134.6)  
384 SER (171.6, -154.0)  
460 ALA (-36.4, 99.7)  
489 LYS (174.4, -137.5)  
615 VAL (59.6, 23.7)  
668 GLU (101.0, -134.1)