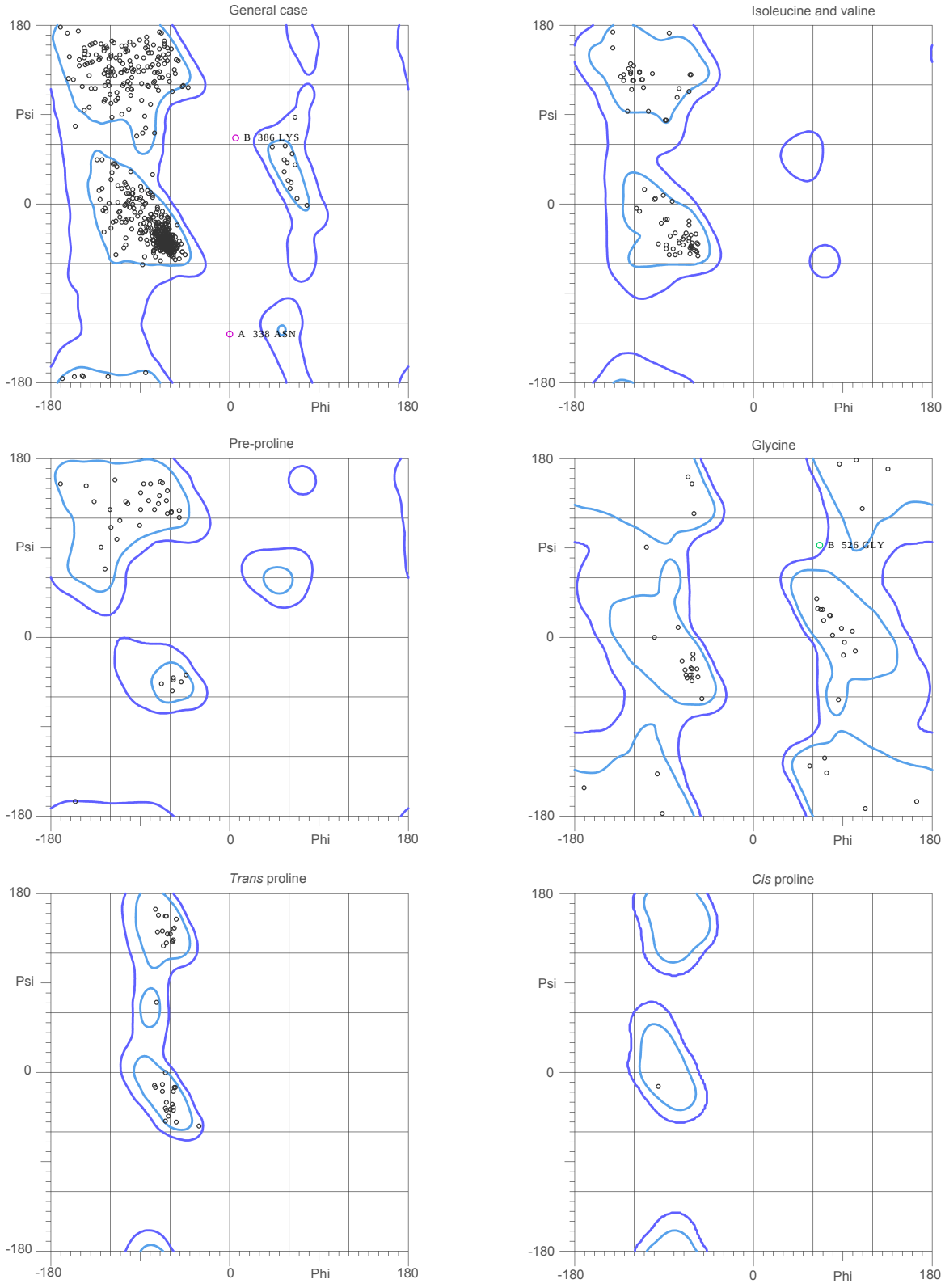
MolProbity Ramachandran analysis

6lzg.H.pdb, model 1



98.5% (776/788) of all residues were in favored (98%) regions.
99.6% (785/788) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):

A 338 ASN (0.0, -131.6)
B 386 LYS (6.8, 67.5)
B 526 GLY (67.9, 93.8)

Lovell, Davis, et al. Proteins 50:437 (2003)