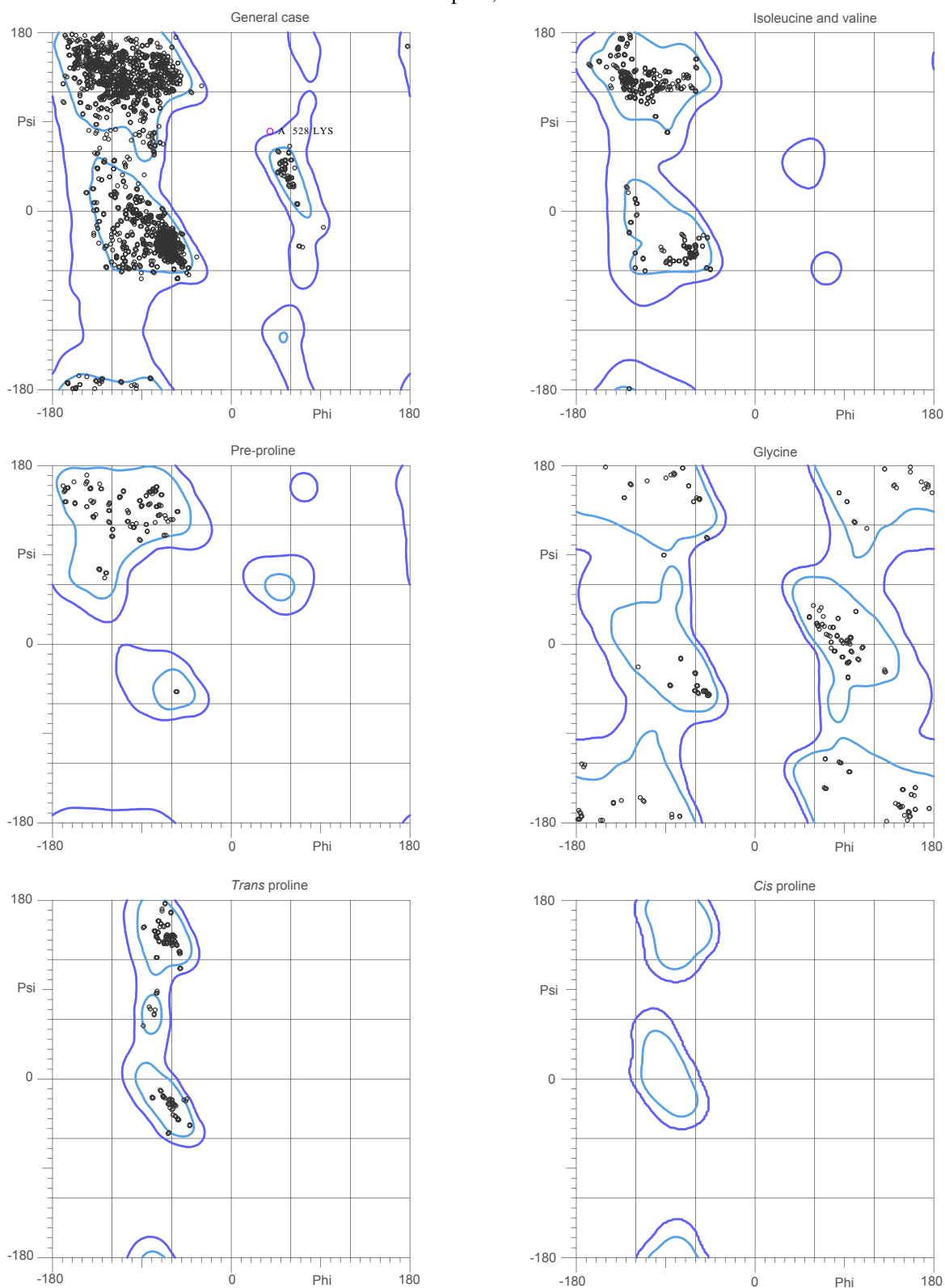


# MolProbity Ramachandran analysis

6zhd.H.pdb, model 1



96.3% (3175/3296) of all residues were in favored (98%) regions.

100.0% (3295/3296) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

A 528 LYS (39.2, 81.6)