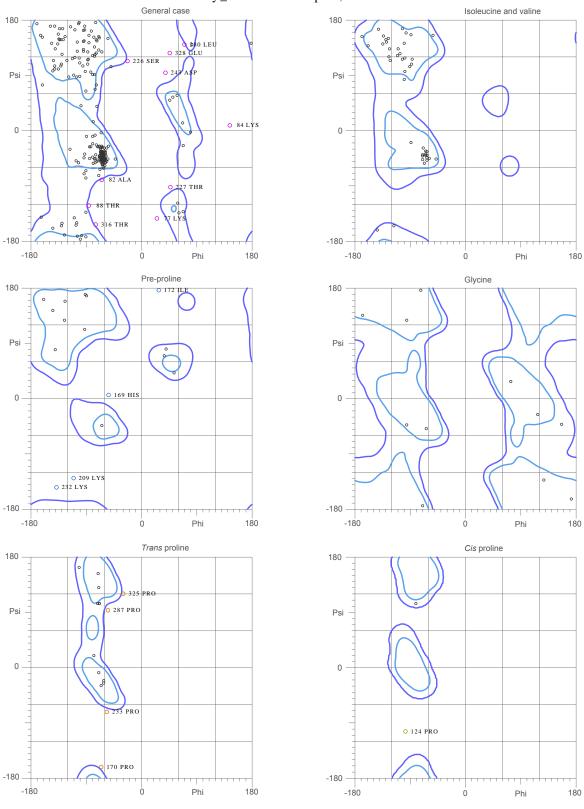
MolProbity Ramachandran analysis





\$2,1% (276/336) of all residues were in favored (98%) regions.

4,3% (317/336) of all residutes (ph.; 4,3% (317/336) of all residutes; (ph.; 4) of 12 kg (25,143k) sequence 19 outliers (ph.; 4-143k) sequence 24 kg (25,143k) sequence 24