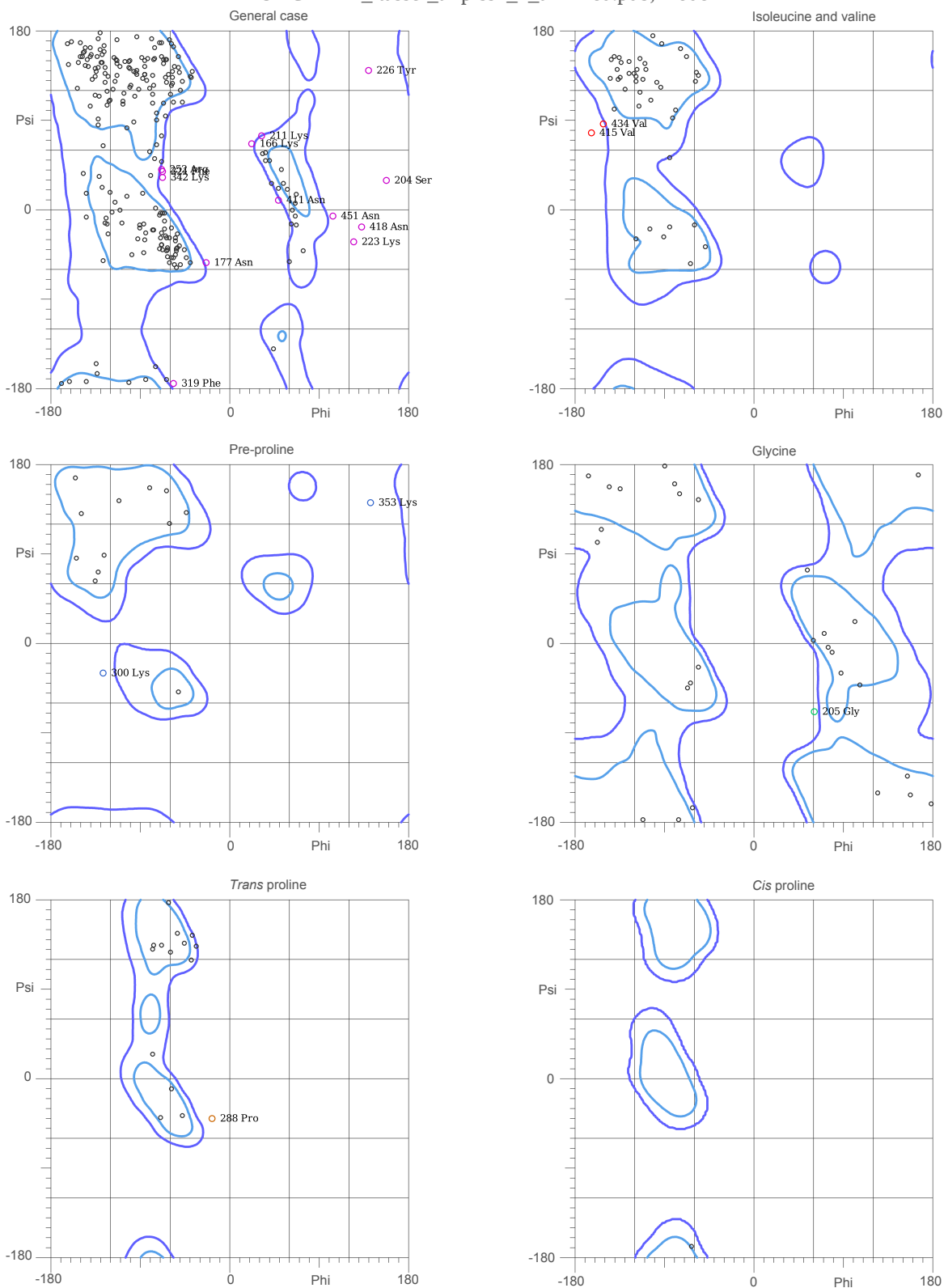


# MolProbity Ramachandran analysis

TRUNCATED\_itasser\_tmprss2\_1\_trimmed.pdb, model 1



78.6% (265/337) of all residues were in favored (98%) regions.

94.4% (318/337) of all residues were in allowed (>99.8%) regions.

418 Asn (133.7, -17.6)

434 Val (-152.0, 87.4)

451 Asn (104.7, -6.1)

There were 19 outliers (phi, psi):

|                        |                         |
|------------------------|-------------------------|
| 166 Lys (22.3, 67.8)   | 288 Pro (-18.4, -40.6)  |
| 177 Asn (-24.6, -53.4) | 300 Lys (-128.0, -30.2) |
| 204 Ser (158.1, 30.8)  | 319 Phe (-57.1, -175.8) |
| 205 Gly (61.0, -69.5)  | 321 Phe (-68.9, 39.0)   |
| 211 Lys (32.5, 75.5)   | 342 Lys (-68.9, 33.3)   |
| 223 Lys (125.9, -32.9) | 353 Lys (143.0, 142.6)  |
| 226 Tyr (140.9, 141.9) | 411 Asn (49.4, 10.1)    |
| 252 Arg (-69.9, 41.0)  | 415 Val (-164.3, 78.7)  |