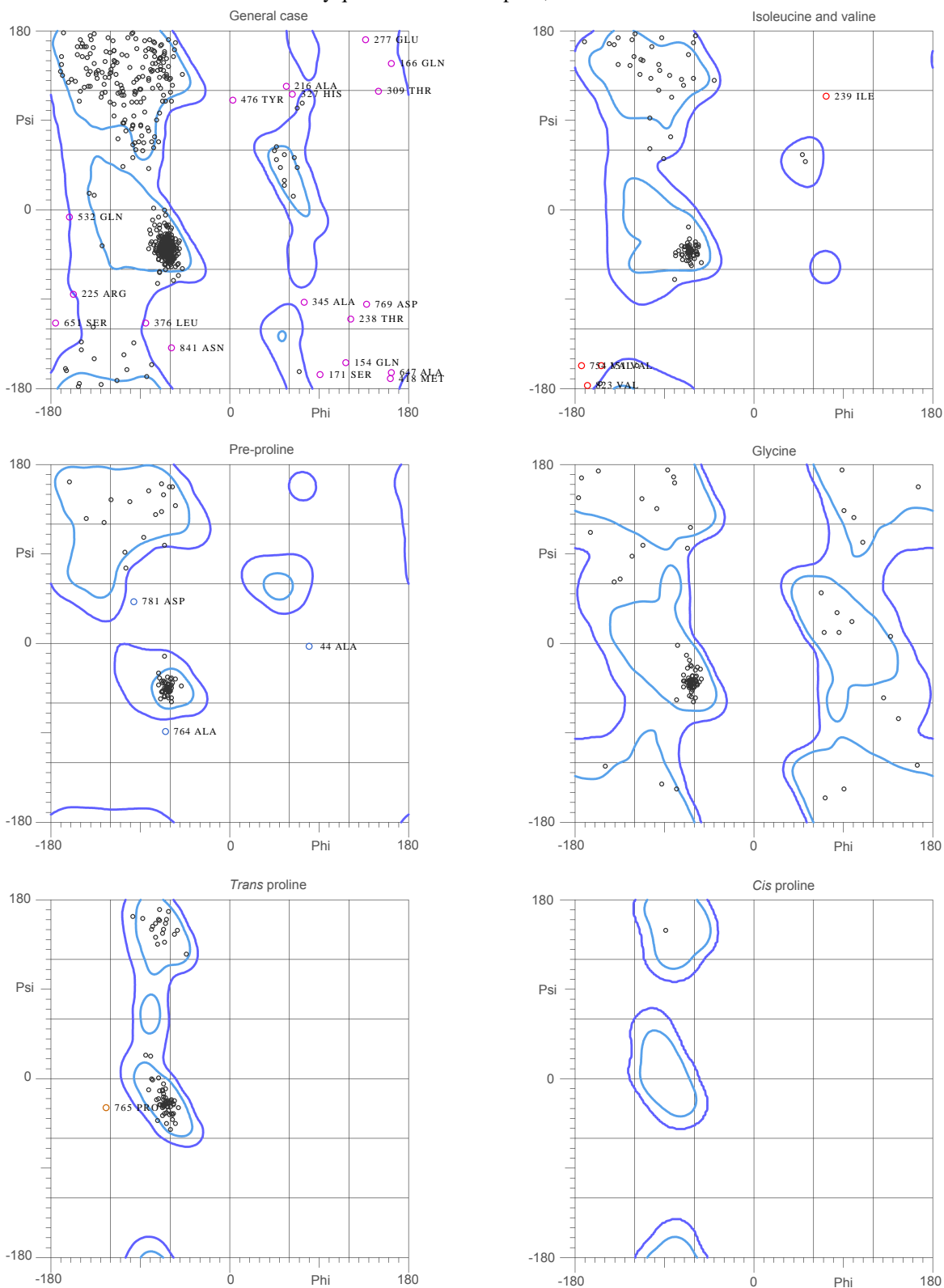


MolProbity Ramachandran analysis

1yq2.B99990011.pdb, model 1



95.7% (152/158) of all residues were in favored (FF) regions.
97.2% (155/158) of all residues were in allowed (FF+GA) regions.

There were 26 outliers (GA, GJ):

111 ALA (158.4, 1.5)
112 THR (158.4, 107.9)
113 GLN (171.4, 114.3)
114 SER (169.5, 108.3)
115 THR (161.1, 108.7)
116 ALA (157.1, 108.3)
117 THR (157.1, 108.3)
118 THR (157.1, 108.3)
119 THR (157.1, 108.3)
120 THR (157.1, 108.3)
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131 THR (157.1, 108.3)
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136 THR (157.1, 108.3)
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140 THR (157.1, 108.3)
141 THR (157.1, 108.3)
142 THR (157.1, 108.3)
143 THR (157.1, 108.3)
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146 THR (157.1, 108.3)
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