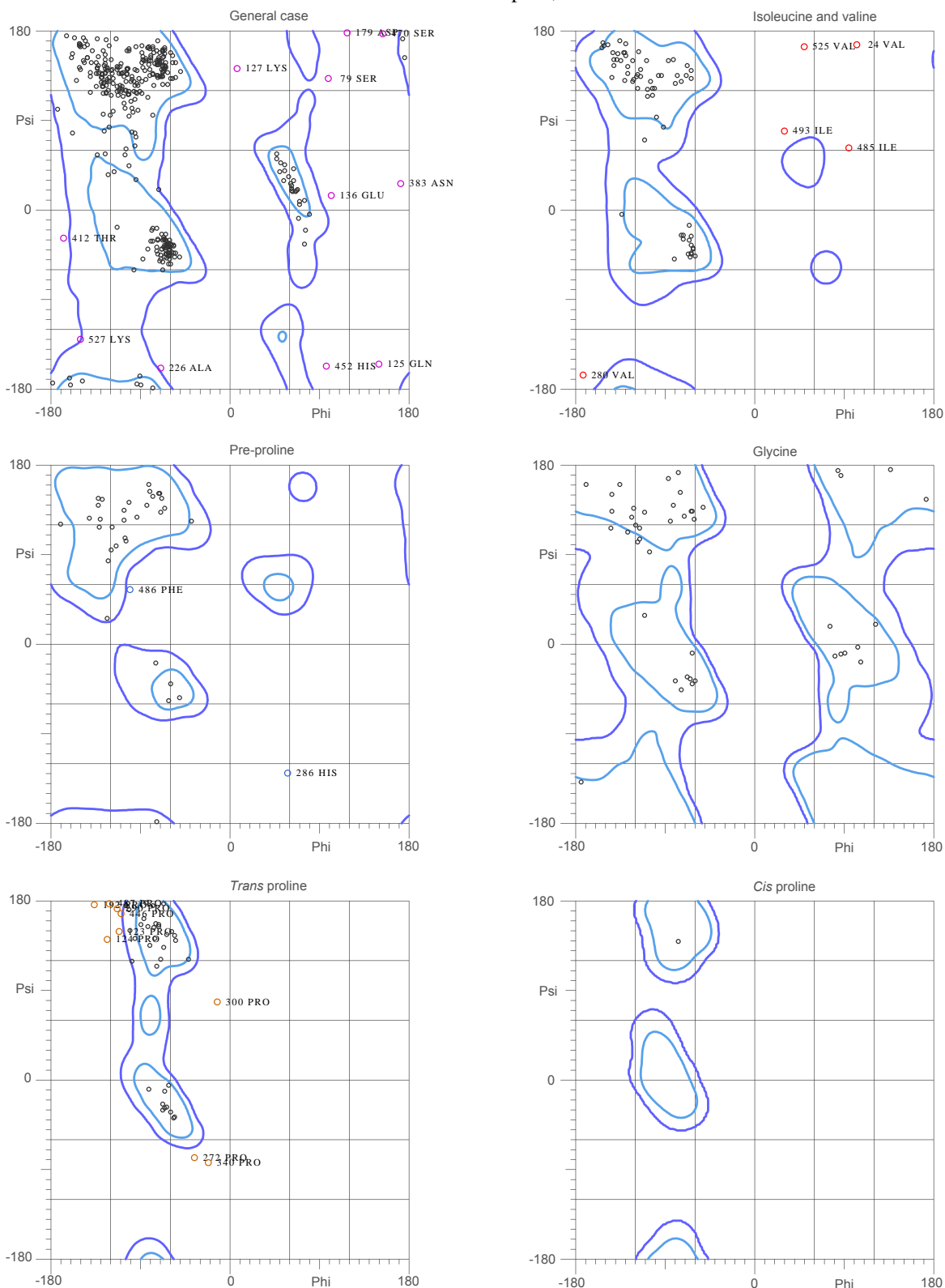


# MolProbity Ramachandran analysis

1aozA.B99990001.pdb, model 1



88.0% (182/208) of all residues were in favored (Fav) regions.  
10.7% (22/208) of all residues were in allowed (All) regions.

Residue name (chain ID, residue number, Phi, Psi)

24 VAL	(1aozA, 187, 7)
113 SER	(1aozA, 113, 3)
114 SER	(1aozA, 114, 3)
115 PRO	(1aozA, 115, 2)
116 SER	(1aozA, 116, 3)
117 SER	(1aozA, 117, 3)
118 SER	(1aozA, 118, 3)
119 ASP	(1aozA, 119, 3)
120 PRO	(1aozA, 120, 3)
121 SER	(1aozA, 121, 3)
122 SER	(1aozA, 122, 3)
123 SER	(1aozA, 123, 3)
124 SER	(1aozA, 124, 3)
125 GLN	(1aozA, 125, 3)
126 PRO	(1aozA, 126, 3)
127 LYS	(1aozA, 127, 3)
128 SER	(1aozA, 128, 3)
129 SER	(1aozA, 129, 3)
130 SER	(1aozA, 130, 3)
131 SER	(1aozA, 131, 3)
132 SER	(1aozA, 132, 3)
133 SER	(1aozA, 133, 3)
134 SER	(1aozA, 134, 3)
135 SER	(1aozA, 135, 3)
136 GLU	(1aozA, 136, 3)
137 SER	(1aozA, 137, 3)
138 SER	(1aozA, 138, 3)
139 SER	(1aozA, 139, 3)
140 SER	(1aozA, 140, 3)
141 SER	(1aozA, 141, 3)
142 SER	(1aozA, 142, 3)
143 SER	(1aozA, 143, 3)
144 SER	(1aozA, 144, 3)
145 SER	(1aozA, 145, 3)
146 SER	(1aozA, 146, 3)
147 SER	(1aozA, 147, 3)
148 SER	(1aozA, 148, 3)
149 SER	(1aozA, 149, 3)
150 SER	(1aozA, 150, 3)
151 SER	(1aozA, 151, 3)
152 SER	(1aozA, 152, 3)
153 SER	(1aozA, 153, 3)
154 SER	(1aozA, 154, 3)
155 SER	(1aozA, 155, 3)
156 SER	(1aozA, 156, 3)
157 SER	(1aozA, 157, 3)
158 SER	(1aozA, 158, 3)
159 SER	(1aozA, 159, 3)
160 SER	(1aozA, 160, 3)
161 SER	(1aozA, 161, 3)
162 SER	(1aozA, 162, 3)
163 SER	(1aozA, 163, 3)
164 SER	(1aozA, 164, 3)
165 SER	(1aozA, 165, 3)
166 SER	(1aozA, 166, 3)
167 SER	(1aozA, 167, 3)
168 SER	(1aozA, 168, 3)
169 SER	(1aozA, 169, 3)
170 SER	(1aozA, 170, 3)
171 SER	(1aozA, 171, 3)
172 SER	(1aozA, 172, 3)
173 SER	(1aozA, 173, 3)
174 SER	(1aozA, 174, 3)
175 SER	(1aozA, 175, 3)
176 SER	(1aozA, 176, 3)
177 SER	(1aozA, 177, 3)
178 SER	(1aozA, 178, 3)
179 ASP	(1aozA, 179, 3)
180 SER	(1aozA, 180, 3)