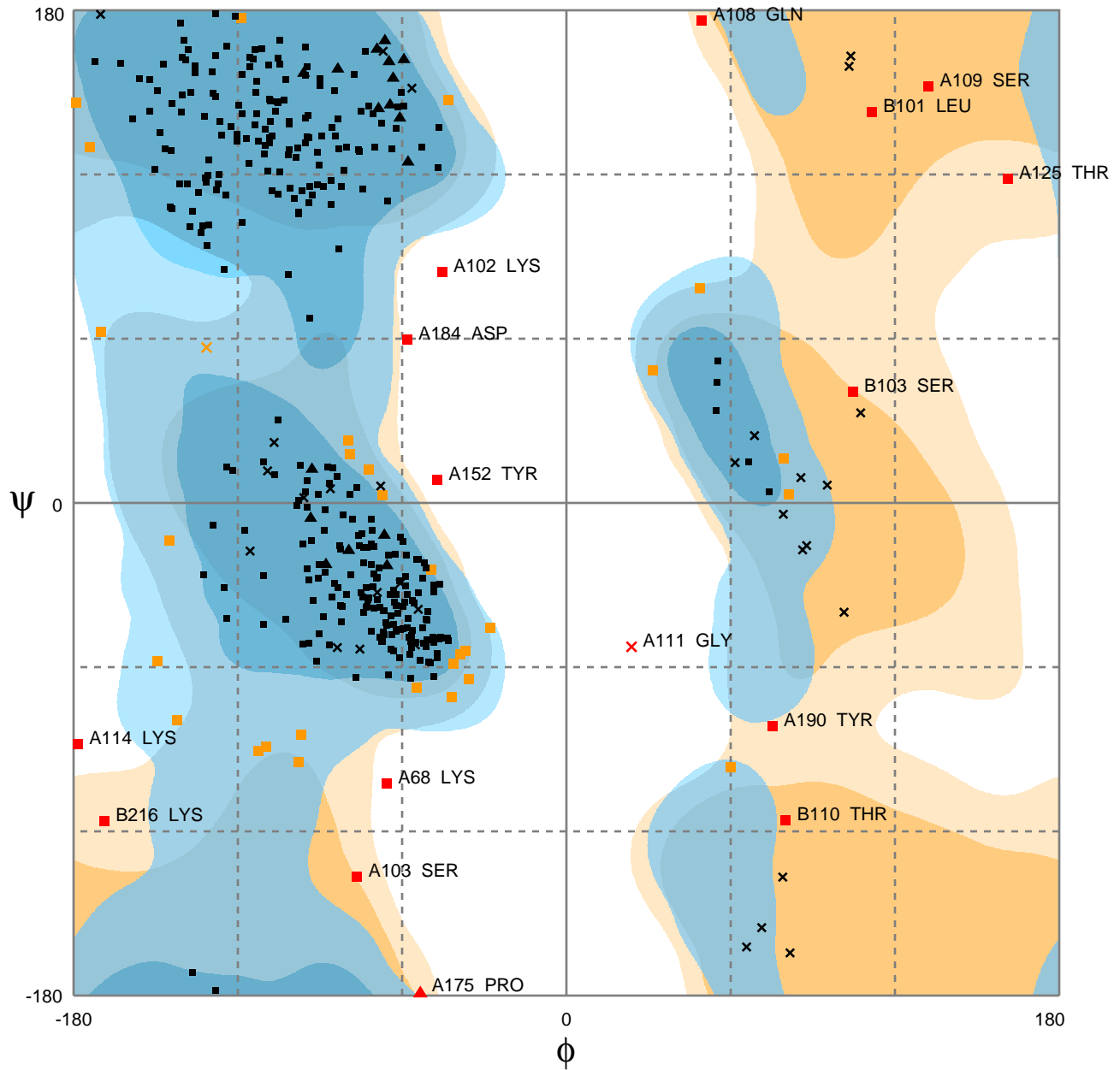


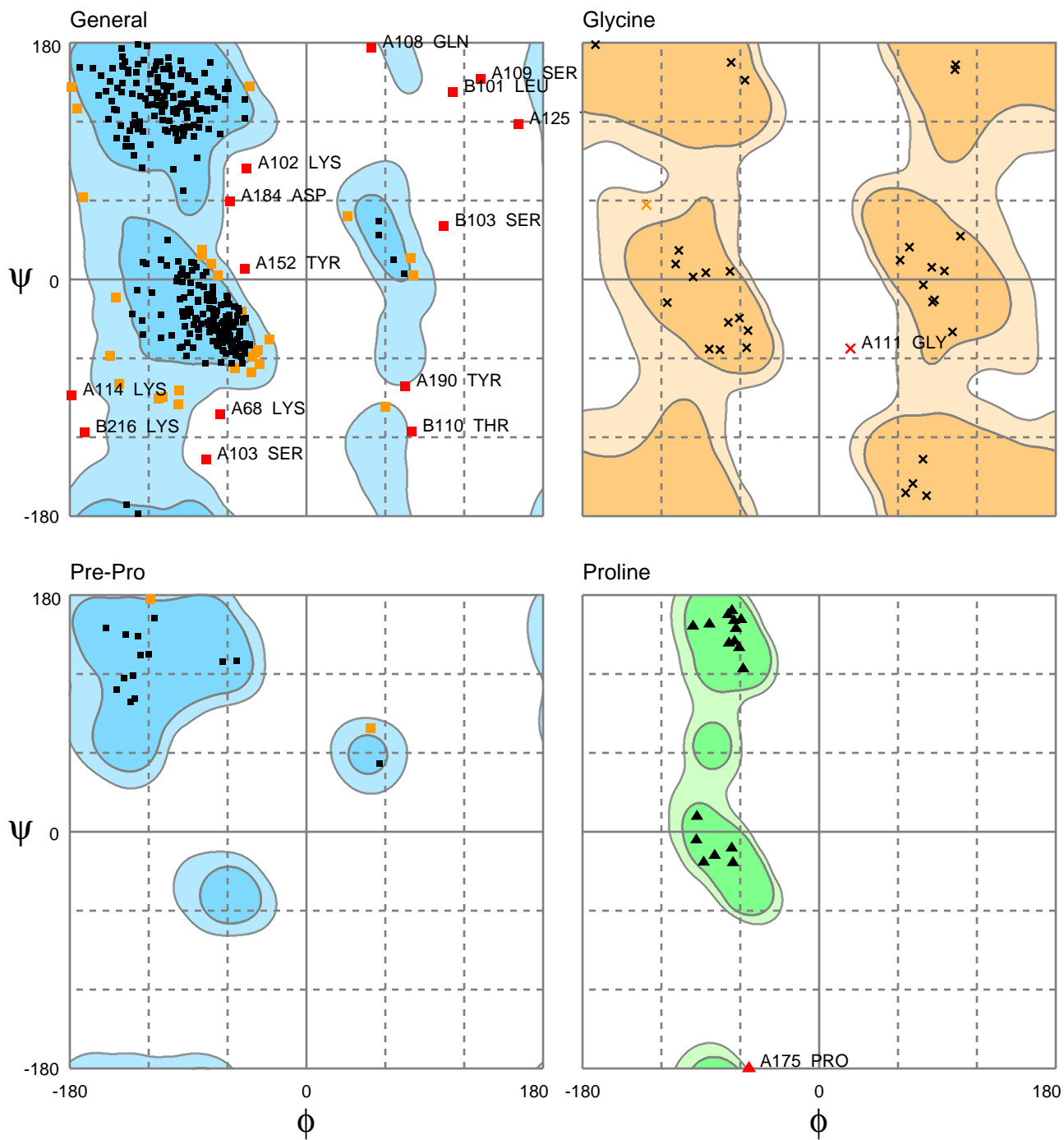
1HMP.pdb



Number of residues in favoured region (~98.0% expected) : 371 (89.0%)
 Number of residues in allowed region (~2.0% expected) : 30 (7.2%)
 Number of residues in outlier region : 16 (3.8%)

RAMPAGE by Paul de Bakker and Simon Lovell available at <http://www-cryst.bioc.cam.ac.uk/rampage/>

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