

■ ▲	General/Pre-Pro/Proline Favoured	□ ▲	General/Pre-Pro/Proline Allowed
×	Glycine Favoured	×	Glycine Allowed

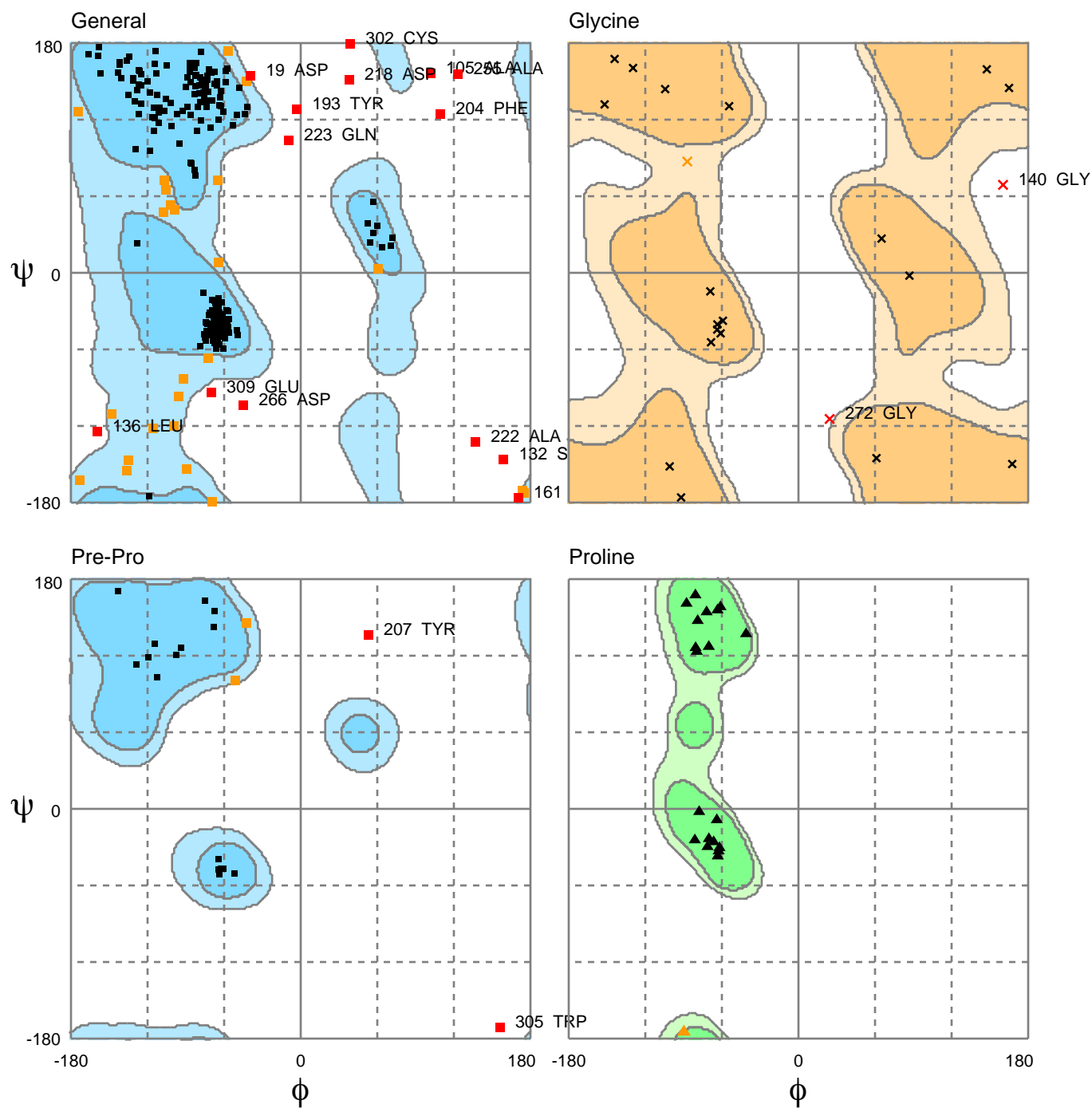
Number of residues in favoured region (~98.0% expected) : 303 (86.8%)

Number of residues in allowed region (~2.0% expected) : 28 (8.0%)

Number of residues in outlier region : 18 (5.2%)

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

Please cite: S.C. Lovell, I.W. Davis, W.B. Arendall III, P.I.W. de Bakker, J.M. Word, M.G. Prisant, J.S. Richardson & D.C. Richardson (2002)  
Structure validation by  $\alpha$  geometry:  $\phi/\psi$  and  $C\beta$  deviation. *Proteins: Structure, Function & Genetics*. **50**: 437-450



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