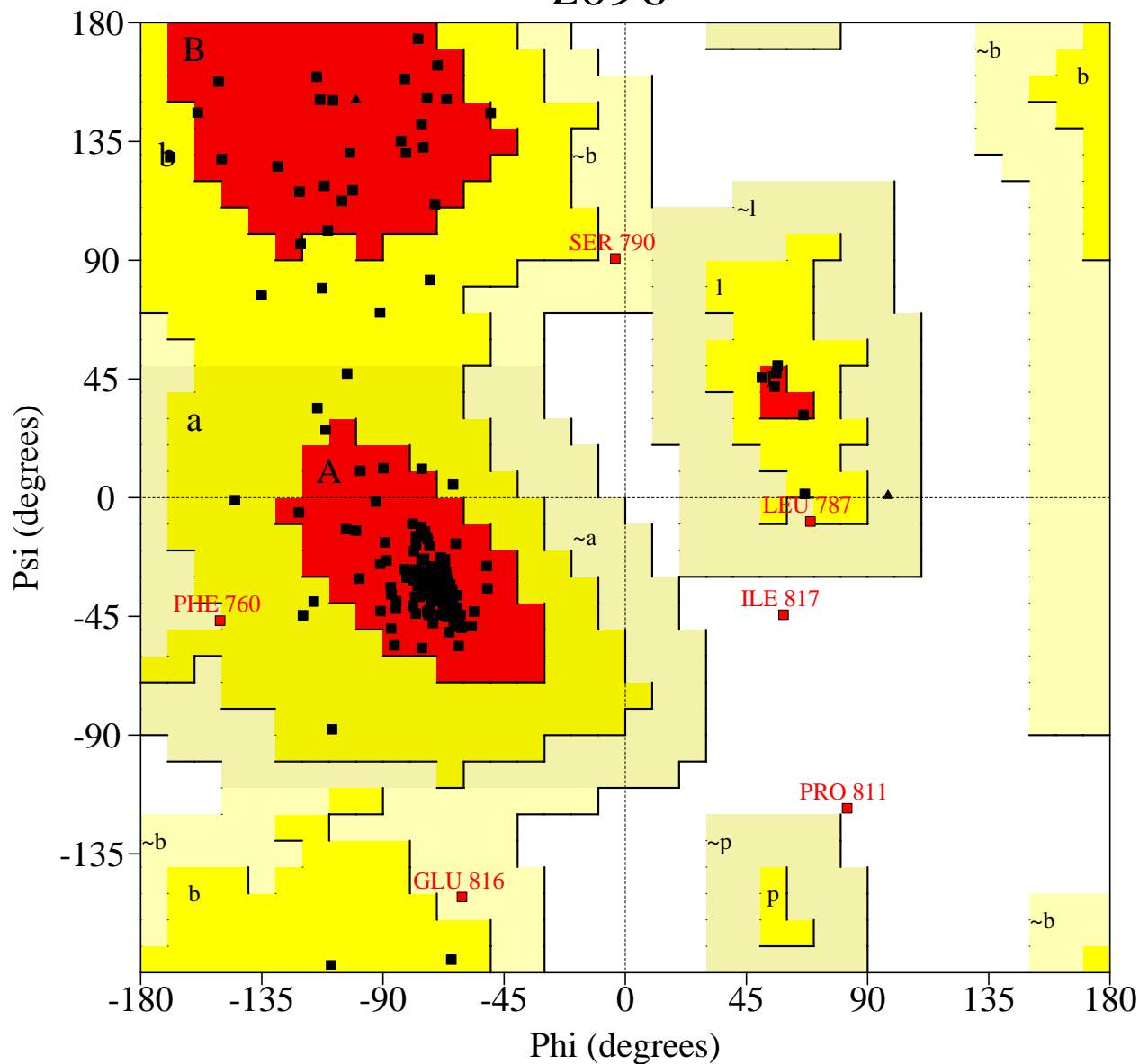


Ramachandran Plot

z096



Plot statistics

Residues in most favoured regions [A,B,L]	150	86.2%
Residues in additional allowed regions [a,b,l,p]	19	10.9%
Residues in generously allowed regions [-a,-b,-l,-p]	4	2.3%
Residues in disallowed regions	1	0.6%

Number of non-glycine and non-proline residues	174	100.0%
Number of end-residues (excl. Gly and Pro)	10	
Number of glycine residues (shown as triangles)	4	
Number of proline residues	8	

Total number of residues	196	

Based on an analysis of 118 structures of resolution of at least 2.0 Angstroms and R-factor no greater than 20%, a good quality model would be expected to have over 90% in the most favoured regions.