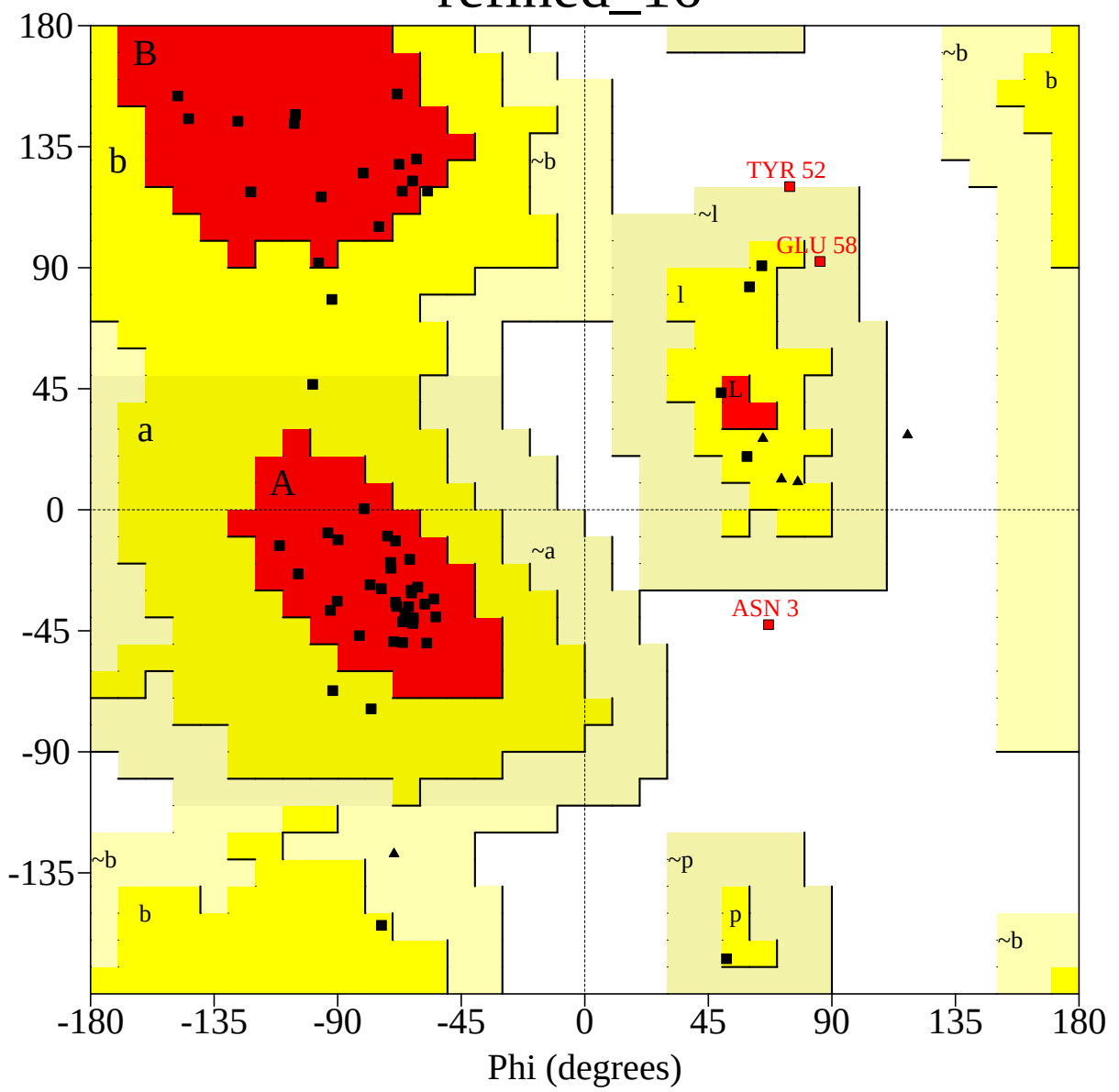
PROCHECK

Ramachandran Plot

refined_16



Plot statistics

Residues in most favoured regions [A,B,L]	43	76.8%
Residues in additional allowed regions [a,b,l,p]	10	17.9%
Residues in generously allowed regions [~a,~b,~l,~p]	1	1.8%
Residues in disallowed regions	2	3.6%
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Number of non-glycine and non-proline residues	56	100.0%
Number of end-residues (excl. Gly and Pro)	2	
Number of glycine residues (shown as triangles)	5	
Number of proline residues	4	

Total number of residues	67	

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.

refined_16_01.ps