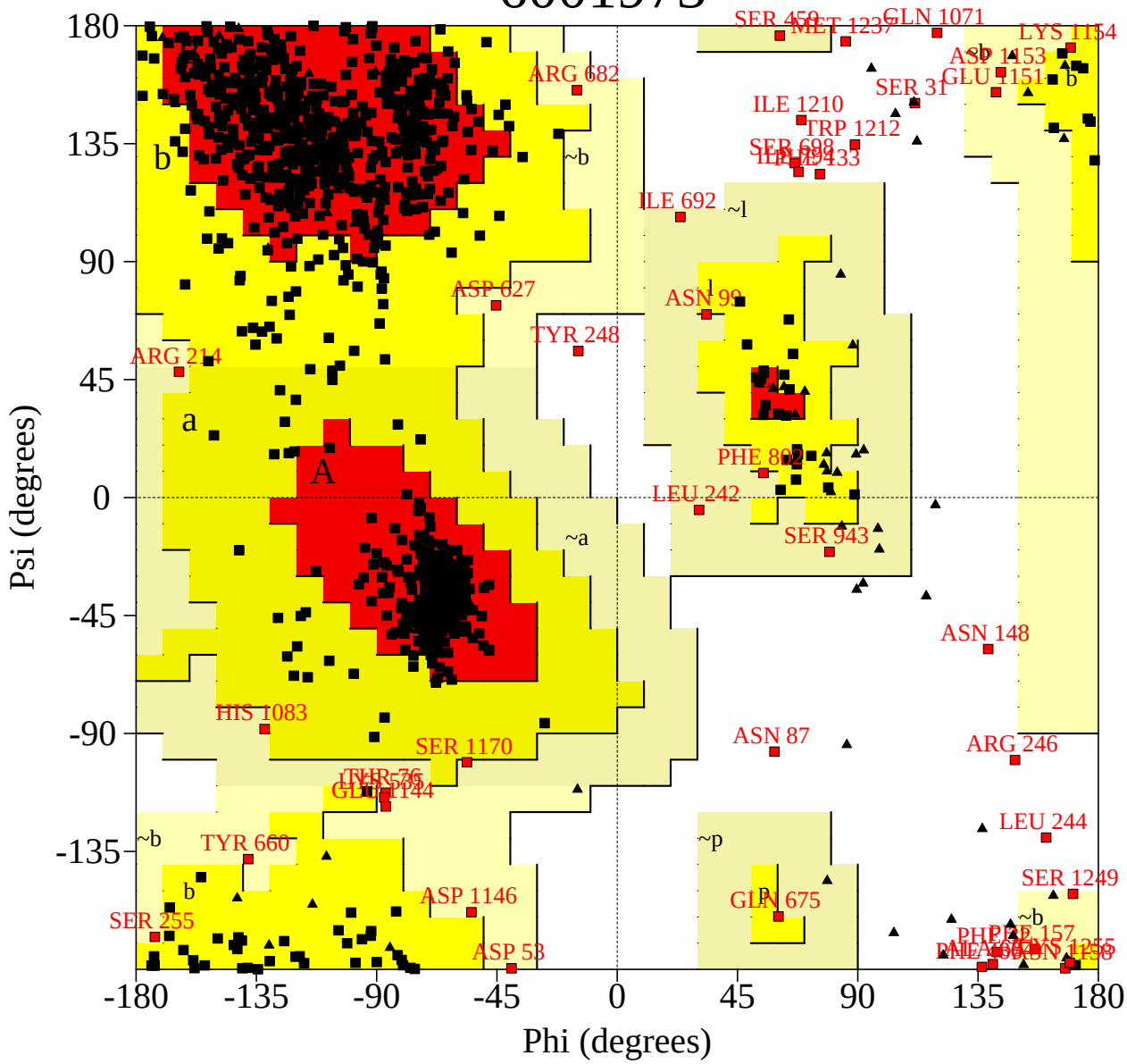
PROCHECK

Ramachandran Plot
6001973



Plot statistics

Residues in most favoured regions [A,B,L]	938	82.9%
Residues in additional allowed regions [a,b,l,p]	151	13.4%
Residues in generously allowed regions [~a,~b,~l,~p]	26	2.3%
Residues in disallowed regions	16	1.4%
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Number of non-glycine and non-proline residues	1131	100.0%
Number of end-residues (excl. Gly and Pro)	2	
Number of glycine residues (shown as triangles)	82	
Number of proline residues	58	

Total number of residues	1273	

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.

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