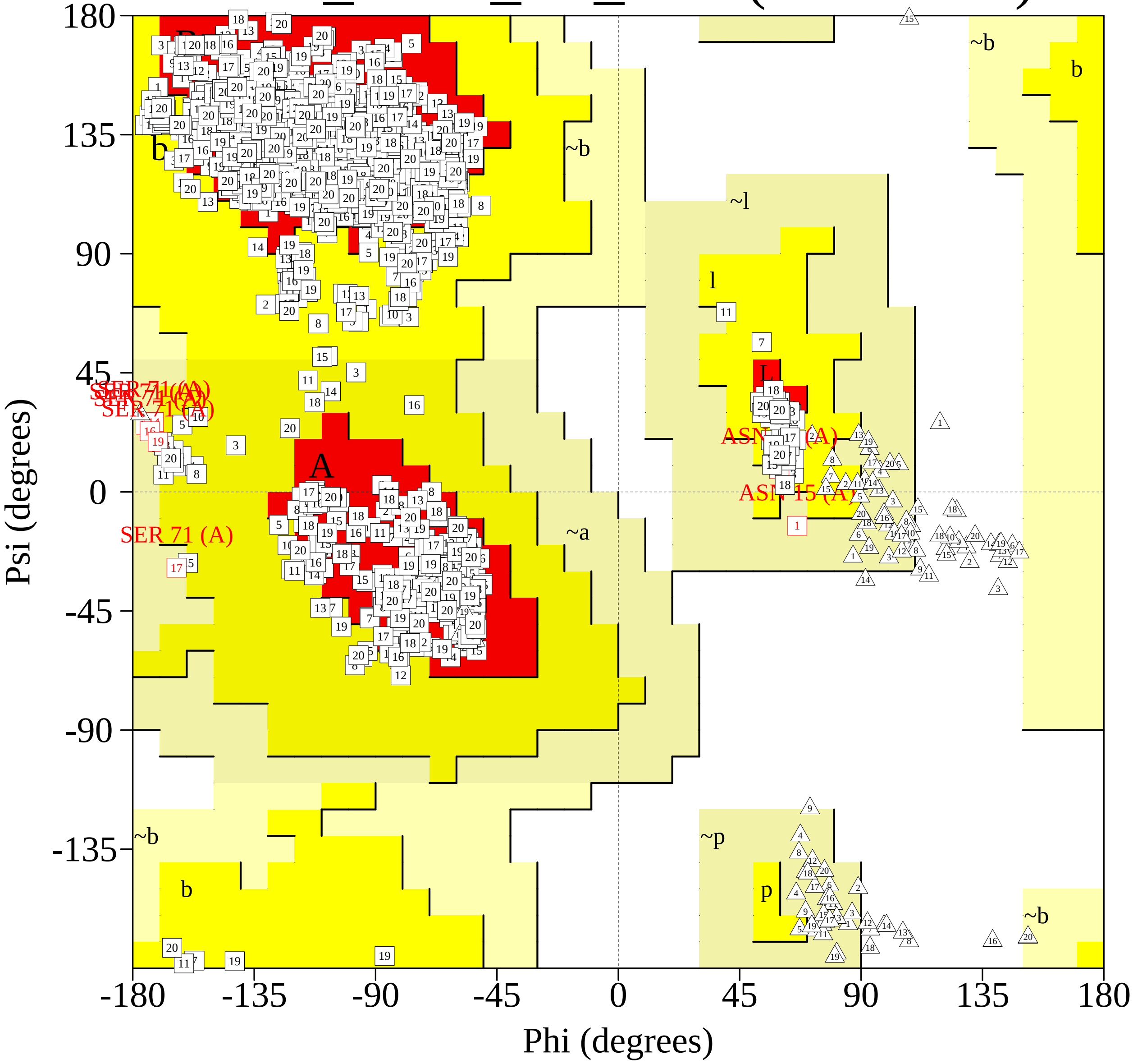
PROCHECK-NMR

Ramachandran Plot

LKR112_NMR_em_bcr3 (20 models)**



Plot statistics

Residues in most favoured regions [A,B,L]	1196	85.4%
Residues in additional allowed regions [a,b,l,p]	197	14.1%
Residues in generously allowed regions [~a,~b,~l,~p]	7	0.5%
Residues in disallowed regions	0	0.0%
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Number of non-glycine and non-proline residues	1400	100.0%
Number of end-residues (excl. Gly and Pro)	0	
Number of glycine residues (shown as triangles)	120	
Number of proline residues	60	

Total number of residues	1580	

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.
Model numbers shown inside each data point.

LKR112_NMR_em_bcr3_01_ramachand.ps

** Selected residues only