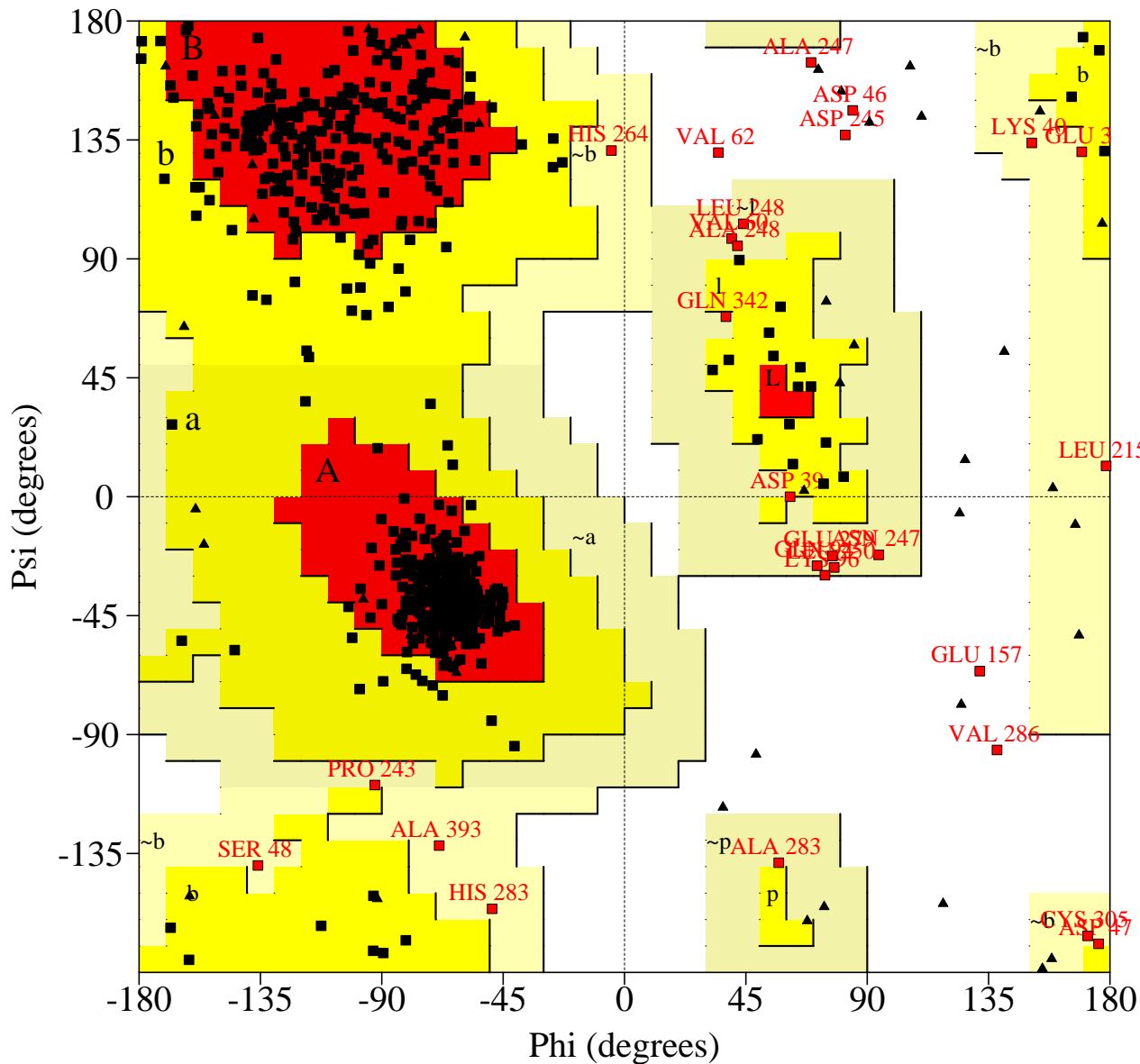


Ramachandran Plot



Plot statistics

| | | |
|------------------------------------------------------|-----|--------|
| Residues in most favoured regions [A,B,L] | 656 | 86.9% |
| Residues in additional allowed regions [a,b,l,p] | 73 | 9.7% |
| Residues in generously allowed regions [-a,-b,-l,-p] | 20 | 2.6% |
| Residues in disallowed regions | 6 | 0.8% |
| ----- | | |
| Number of non-glycine and non-proline residues | 755 | 100.0% |
| Number of end-residues (excl. Gly and Pro) | 4 | |
| Number of glycine residues (shown as triangles) | 68 | |
| Number of proline residues | 40 | |
| ----- | | |
| Total number of residues | 867 | |

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.