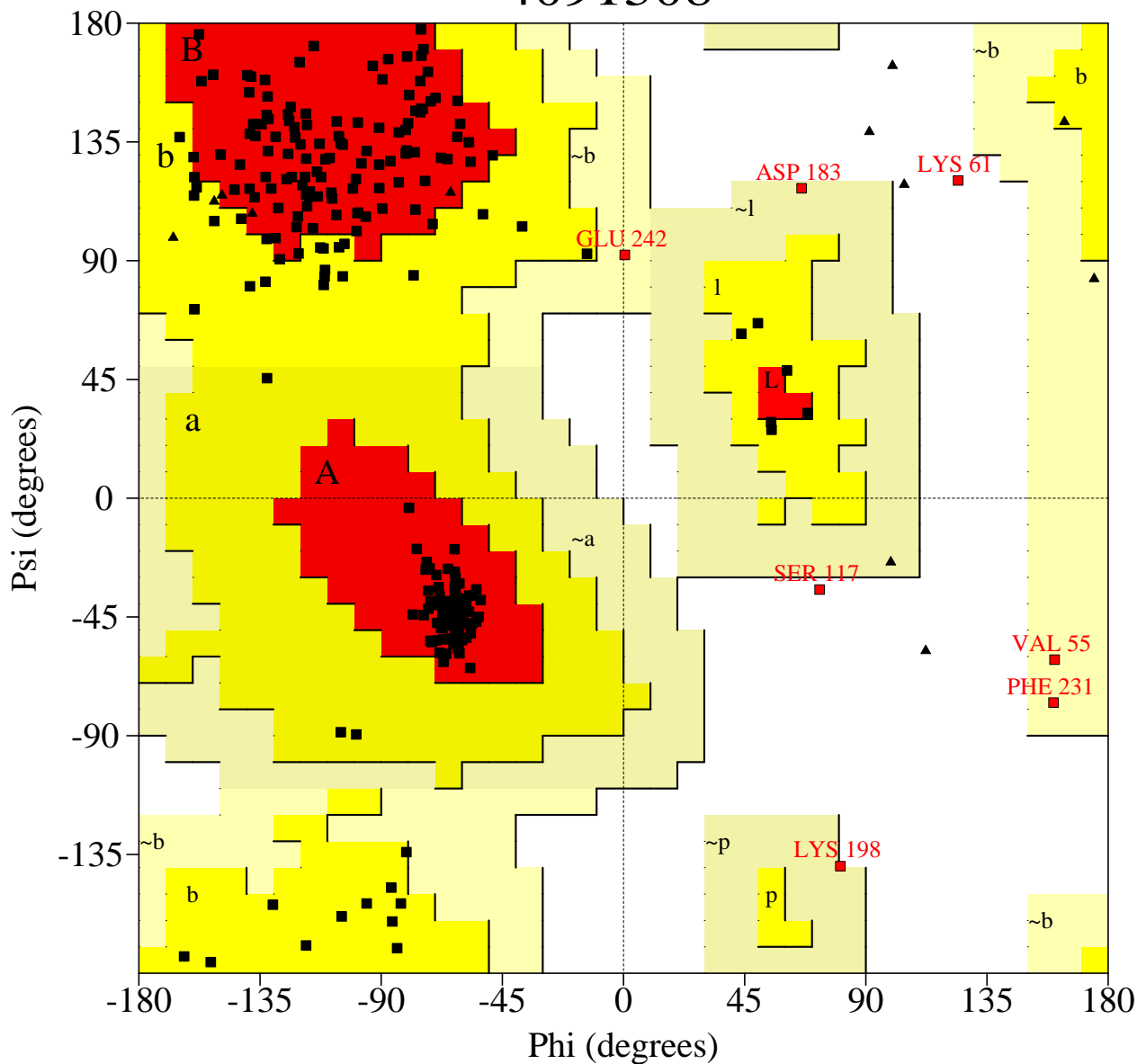


Ramachandran Plot

4091308



Plot statistics

| | | |
|--|-----|--------|
| Residues in most favoured regions [A,B,L] | 200 | 81.0% |
| Residues in additional allowed regions [a,b,l,p] | 40 | 16.2% |
| Residues in generously allowed regions [~a,~b,~l,~p] | 4 | 1.6% |
| Residues in disallowed regions | 3 | 1.2% |
| ----- | | |
| Number of non-glycine and non-proline residues | 247 | 100.0% |
| Number of end-residues (excl. Gly and Pro) | 2 | |
| Number of glycine residues (shown as triangles) | 14 | |
| Number of proline residues | 12 | |
| ----- | | |
| Total number of residues | 275 | |

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.