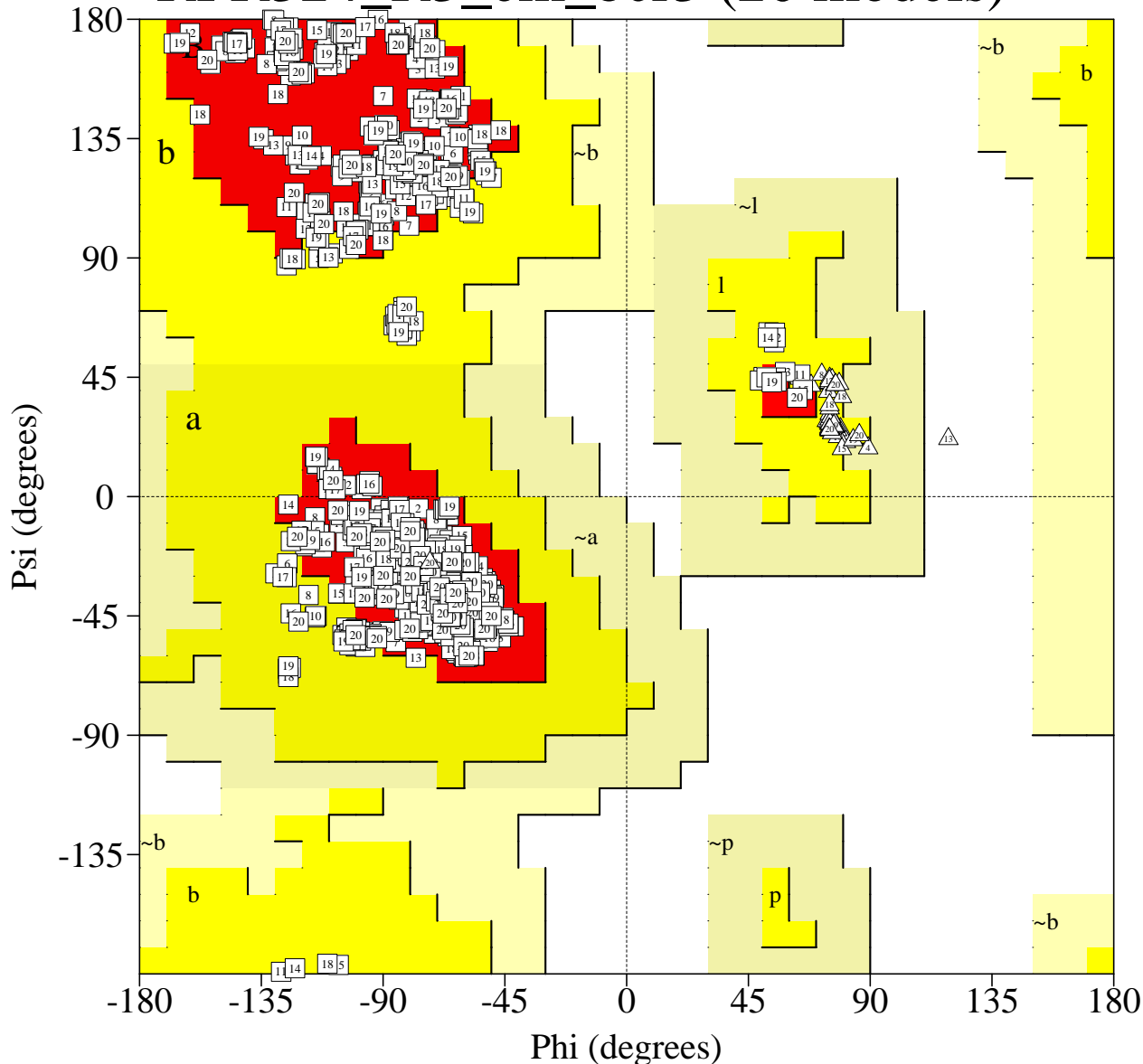


# Ramachandran Plot

## RPR324\_R3\_em\_bcr3 (20 models)\*\*



### Plot statistics

|  |      |        |
|--|------|--------|
| Residues in most favoured regions [A,B,L]            | 1621 | 94.2%  |
| Residues in additional allowed regions [a,b,l,p]     | 99   | 5.8%   |
| Residues in generously allowed regions [-a,-b,-l,-p] | 0    | 0.0%   |
| Residues in disallowed regions                       | 0    | 0.0%   |
|  | ---- | -----  |
| Number of non-glycine and non-proline residues       | 1720 | 100.0% |
| Number of end-residues (excl. Gly and Pro)           | 0    |        |
| Number of glycine residues (shown as triangles)      | 80   |        |
| Number of proline residues                           | 60   |        |
|  | ---- |        |
| Total number of residues                             | 1860 |        |

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.