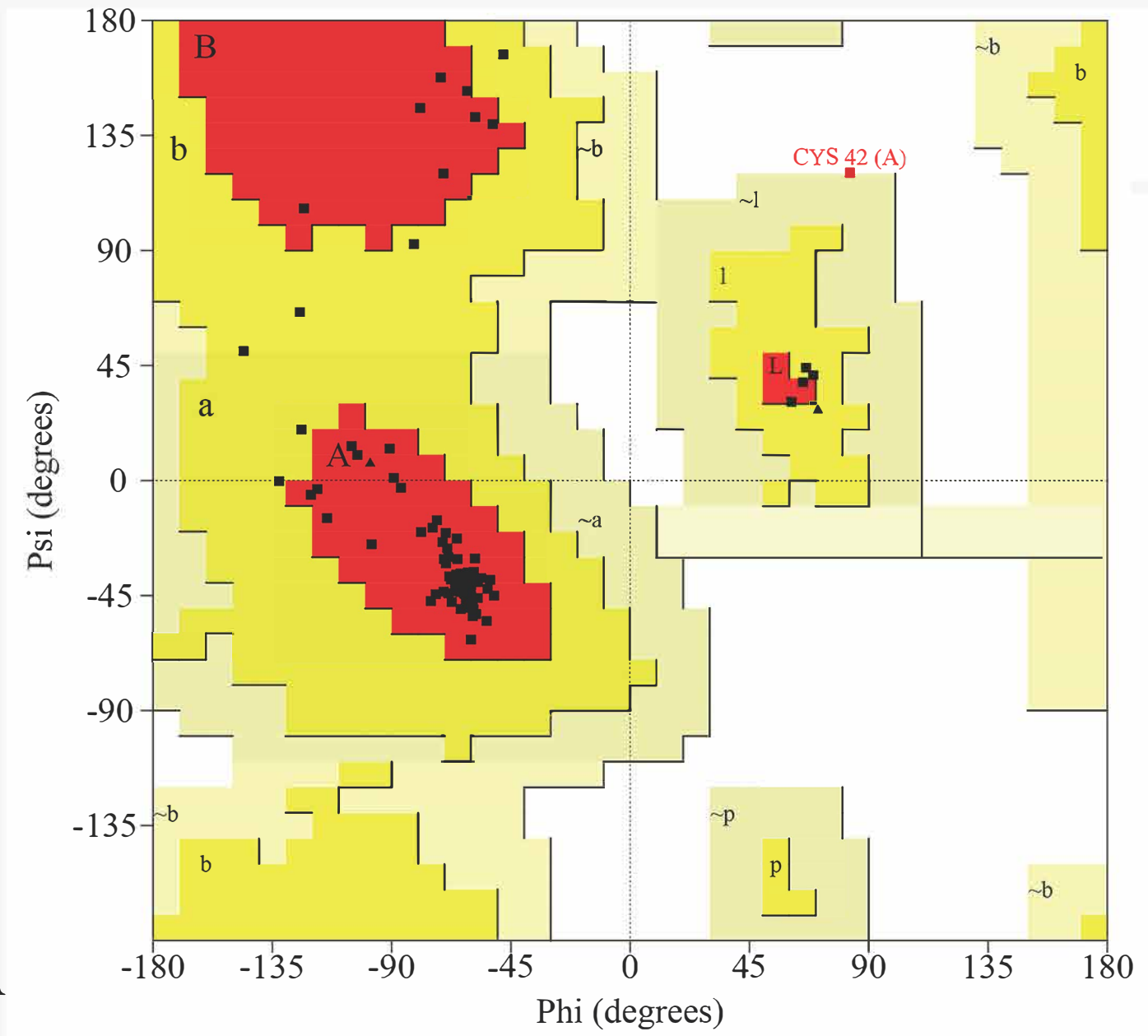
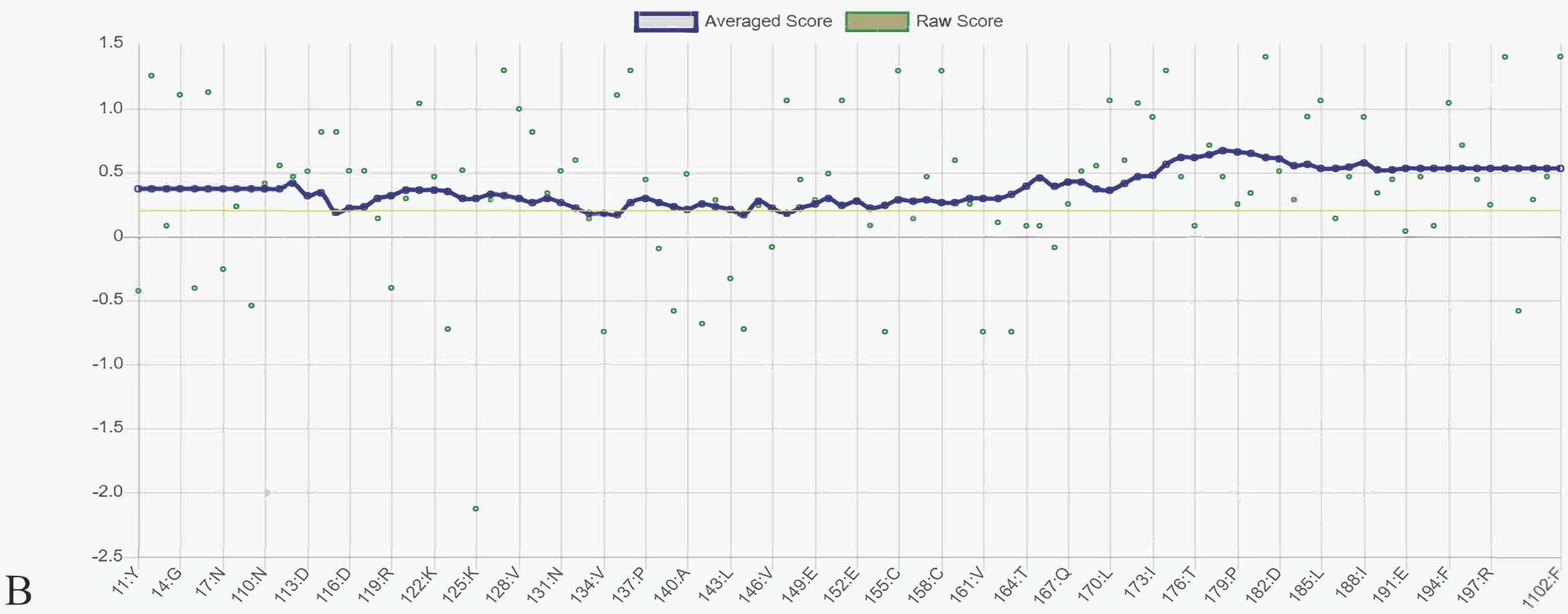


Plot statistics

Residues in most favoured regions [A,B,L]	84	91.3%
Residues in additional allowed regions [a,b,l,p]	7	7.6%
Residues in generously allowed regions [~a,~b,~l,~p]	0	0.0%
Residues in disallowed regions	1	1.1%
	---	------
Number of non-glycine and non-proline residues	92	100.0%
Number of end-residues (excl. Gly and Pro)	2	
Number of glycine residues (shown as triangles)	2	
Number of proline residues	6	

Total number of residues	102	



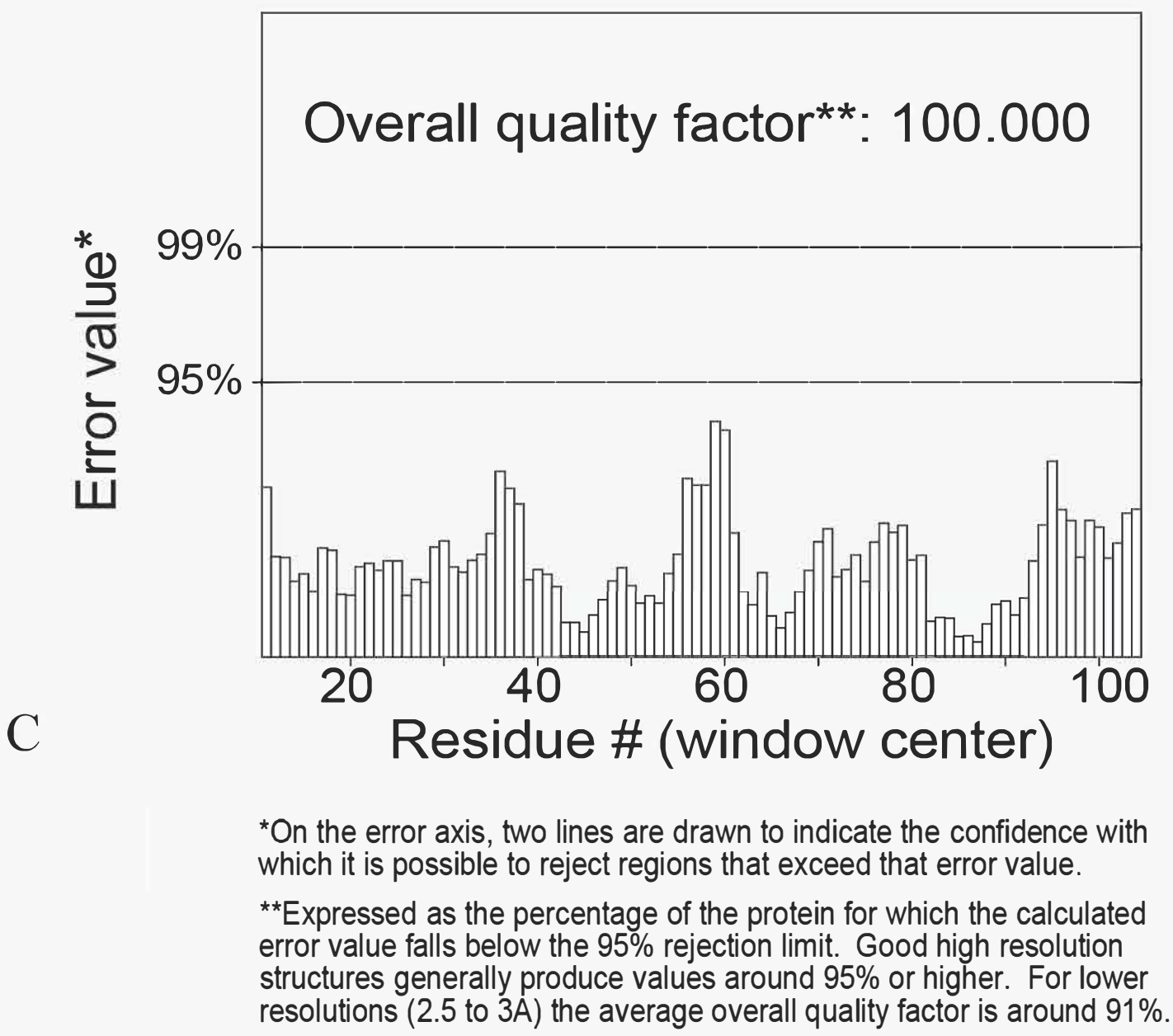


Figure S1. Quality evaluation of the SnitCSP4 model. (**A**) Ramachandran plot; (**B**) Verify-3D; (**C**) ERRAT.