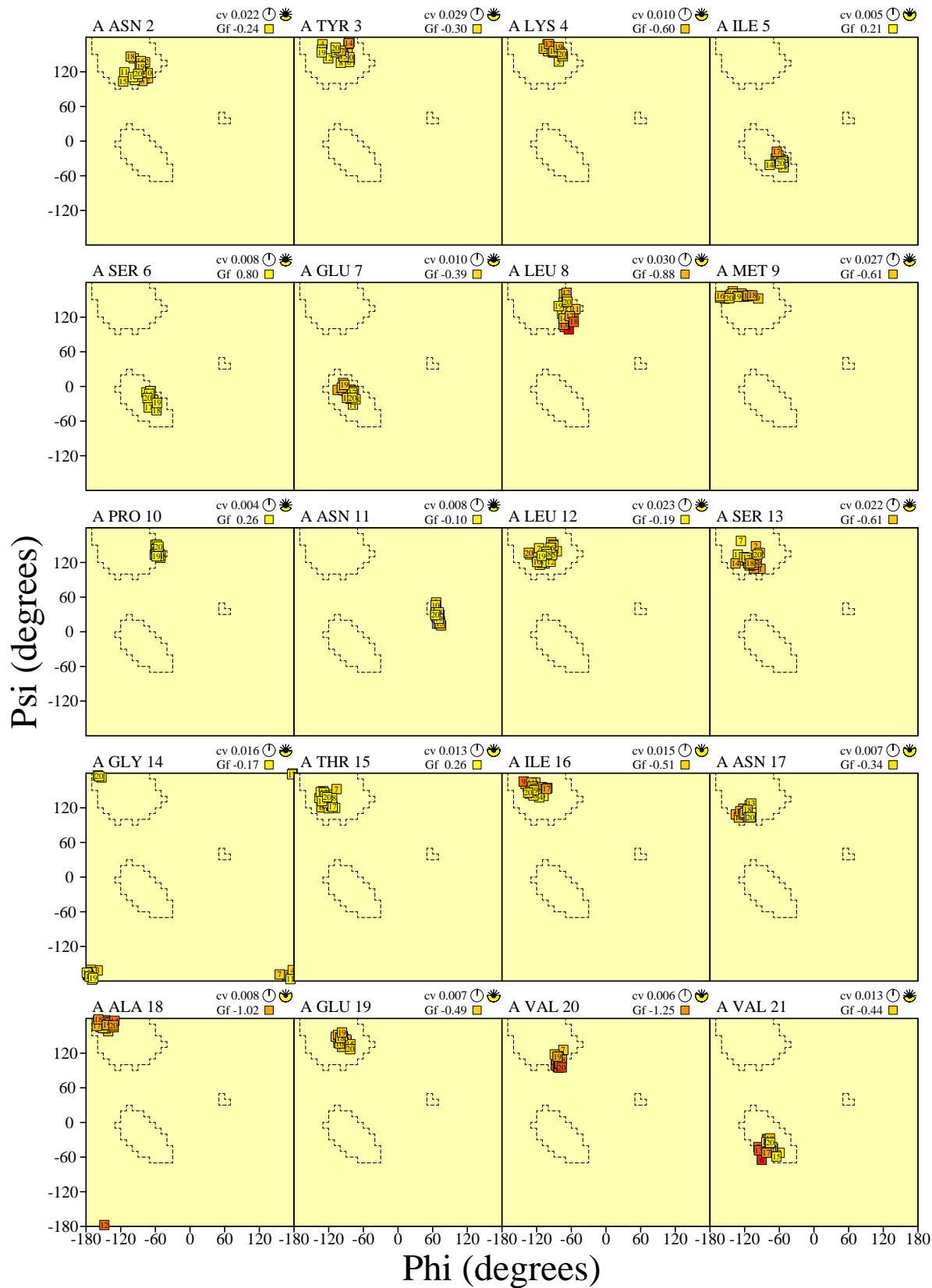


Ensemble Ramachandran plots

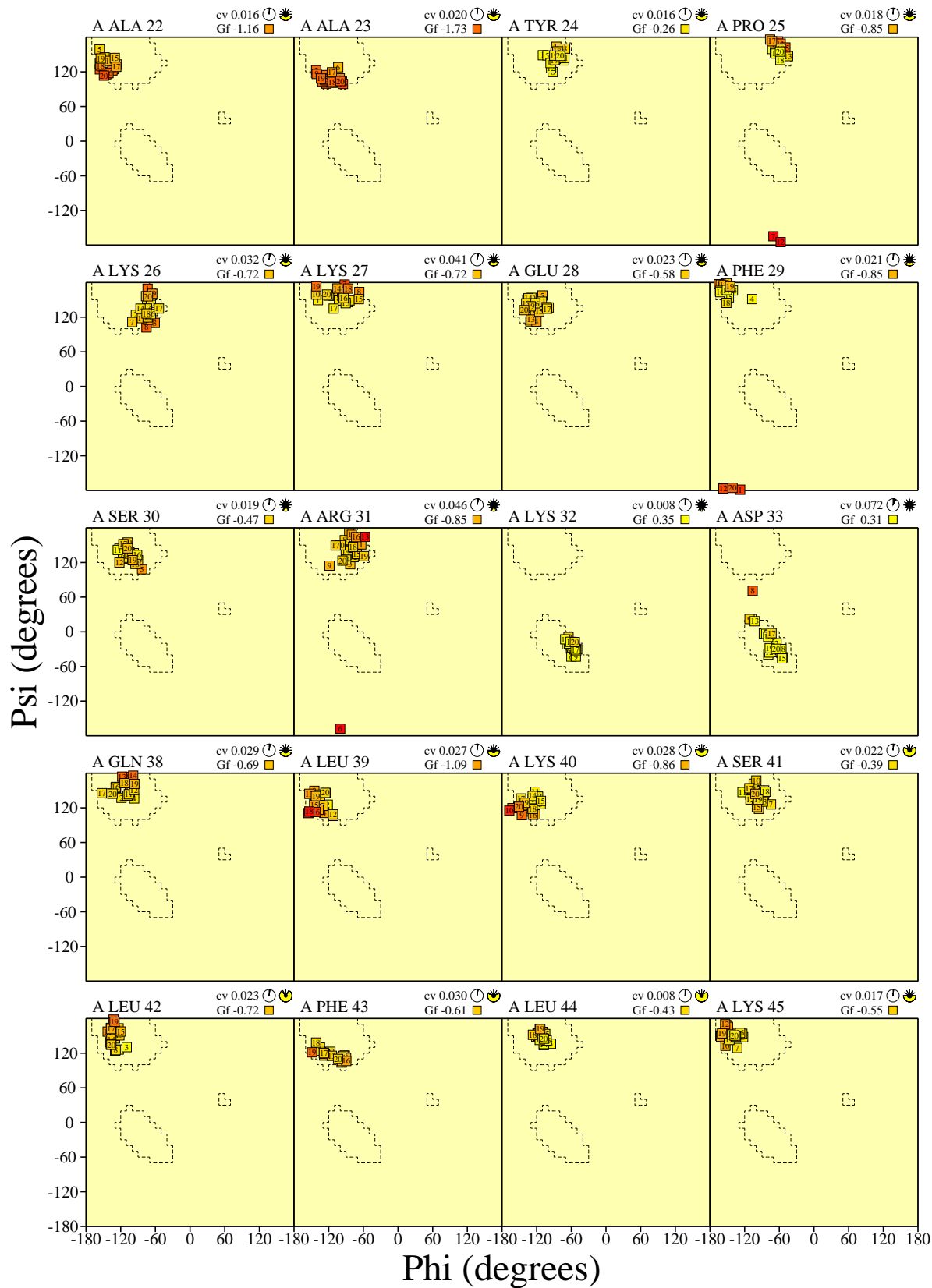
MRR110B_NMR_em_bcr3 (20 models)**



cv = Circular Variance (low values signify high clustering of the data points). ☀ Accessible ☾ Buried
 Gf = Average G-factor for the residue (the higher the value the more favourable the conformations) based on analysis of high-res. Xstal structures
 Data points coloured according to G-factor: Favourable Unfavourable

Ensemble Ramachandran plots

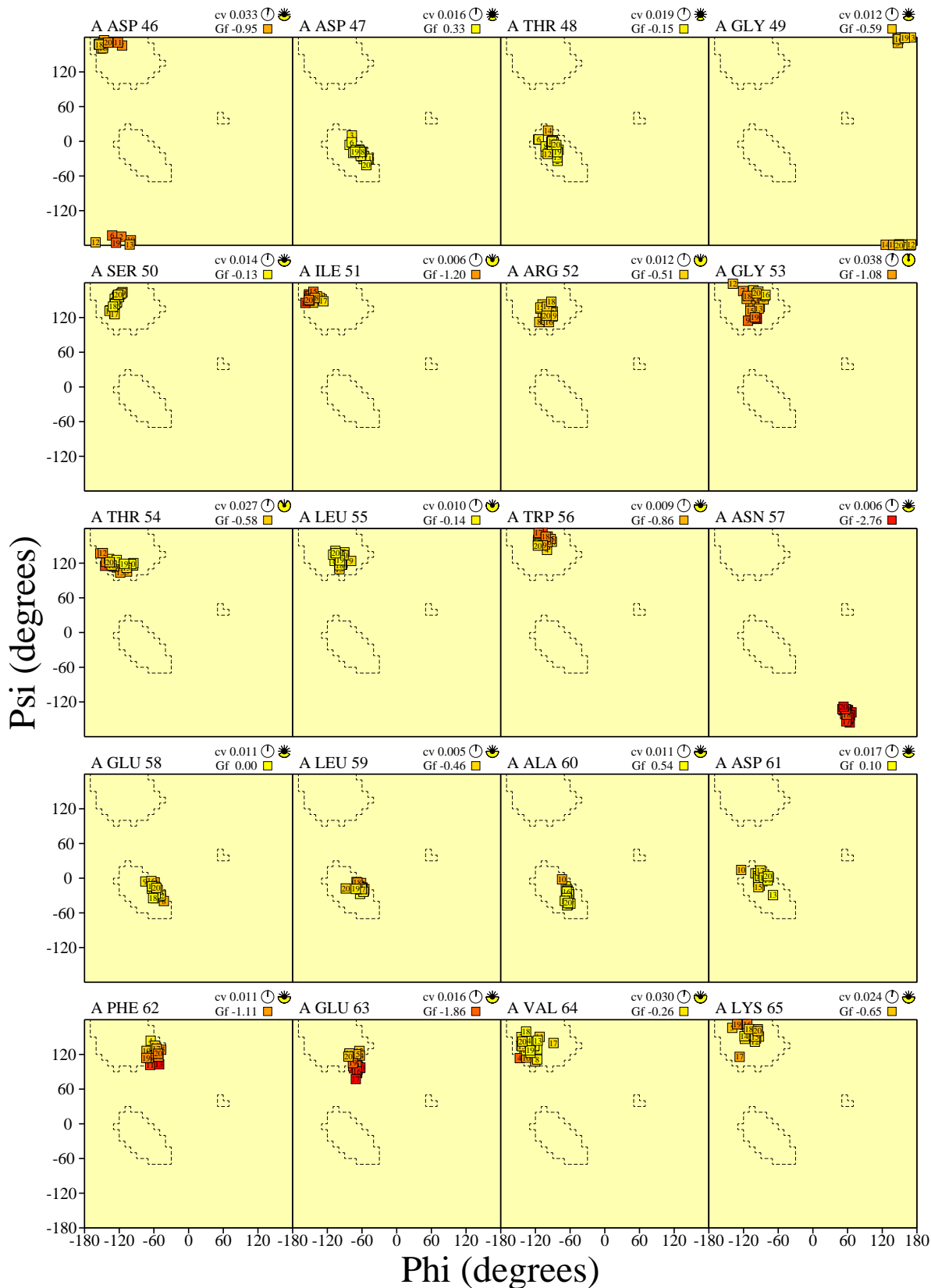
MRR110B_NMR_em_bcr3 (20 models)**



cv = Circular Variance (low values signify high clustering of the data points). * Accessible (circle) Buried (star)
 Gf = Average G-factor for the residue (the higher the value the more favourable the conformations) based on analysis of high-res. Xstal structures
 Data points coloured according to G-factor: Favourable (yellow) Unfavourable (red)

Ensemble Ramachandran plots

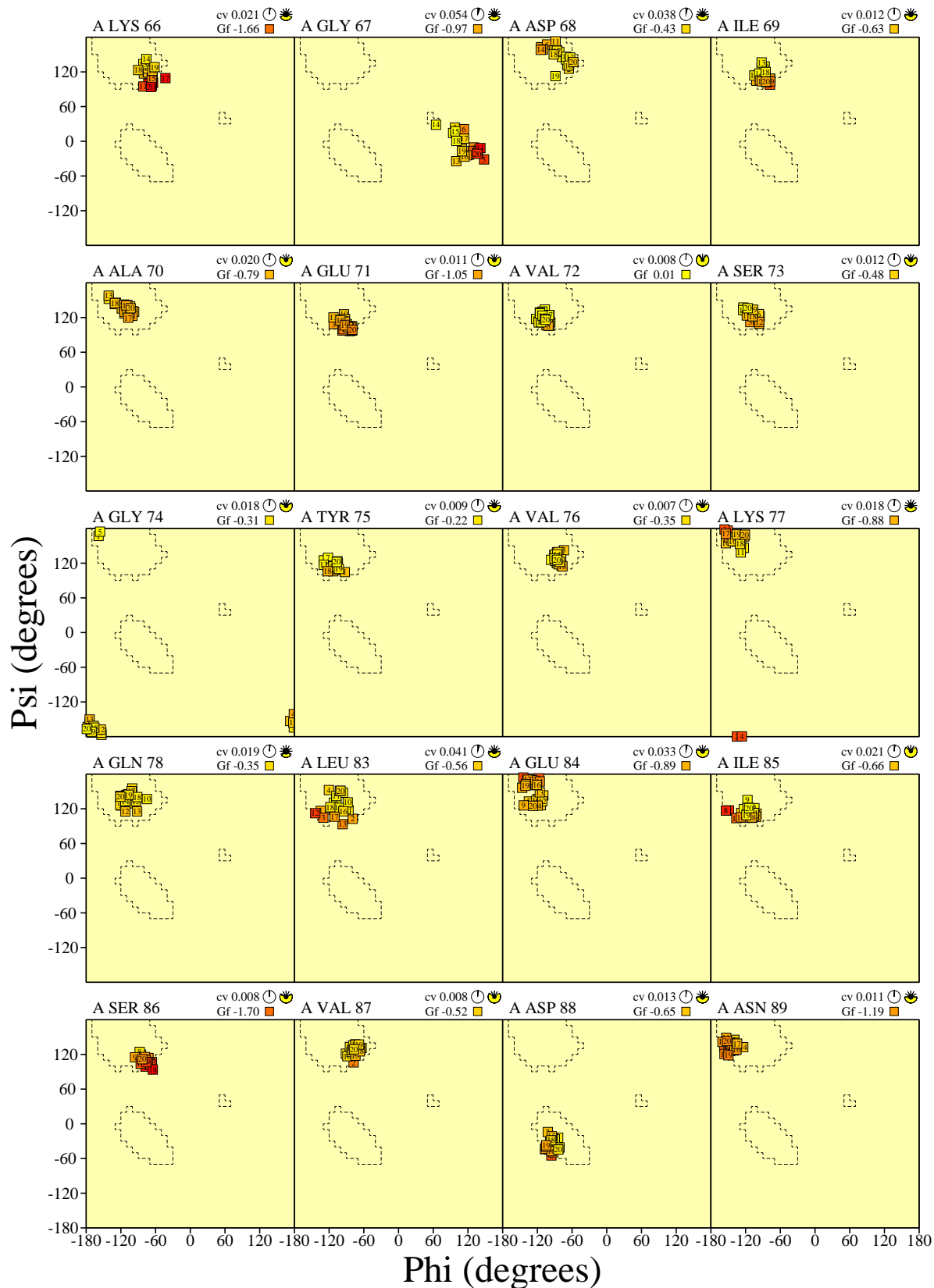
MRR110B_NMR_em_bcr3 (20 models)**



cv = Circular Variance (low values signify high clustering of the data points). * Accessible ☾ Buried
 Gf = Average G-factor for the residue (the higher the value the more favourable the conformations) based on analysis of high-res. Xstal structures
 Data points coloured according to G-factor: Favourable Unfavourable

Ensemble Ramachandran plots

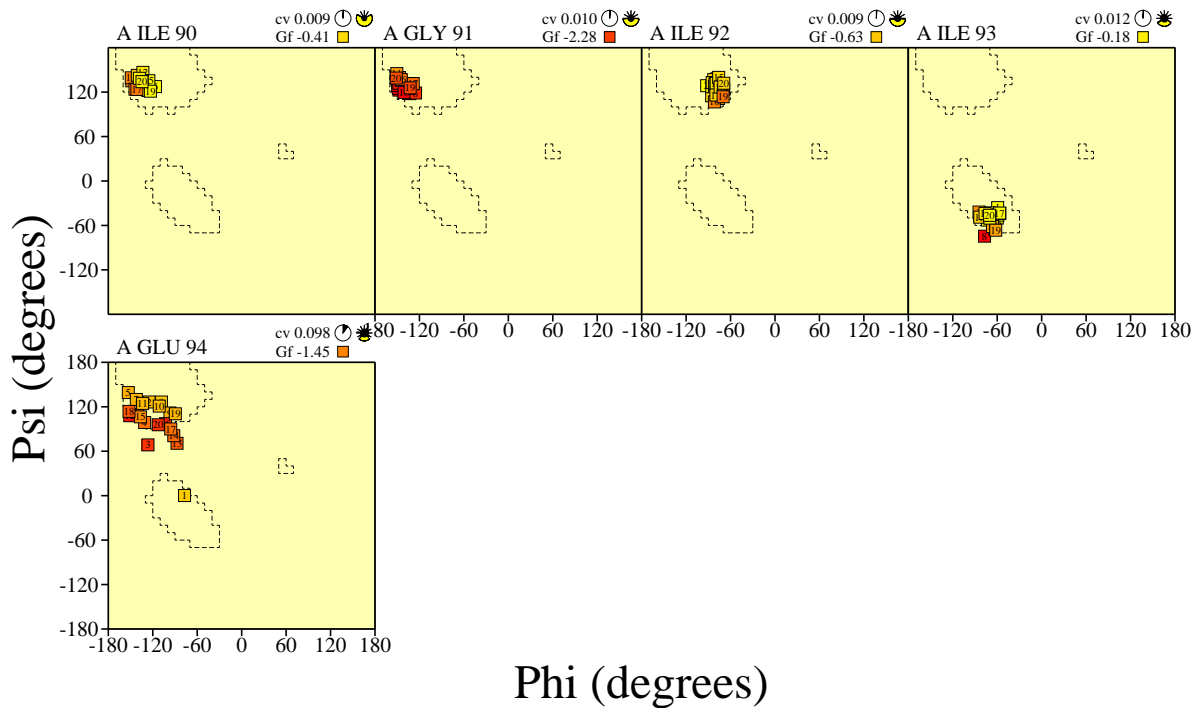
MRR110B_NMR_em_bcr3 (20 models)**



cv = Circular Variance (low values signify high clustering of the data points). ☀ Accessible ☾ Buried
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Ensemble Ramachandran plots

MRR110B_NMR_em_bcr3 (20 models)**



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