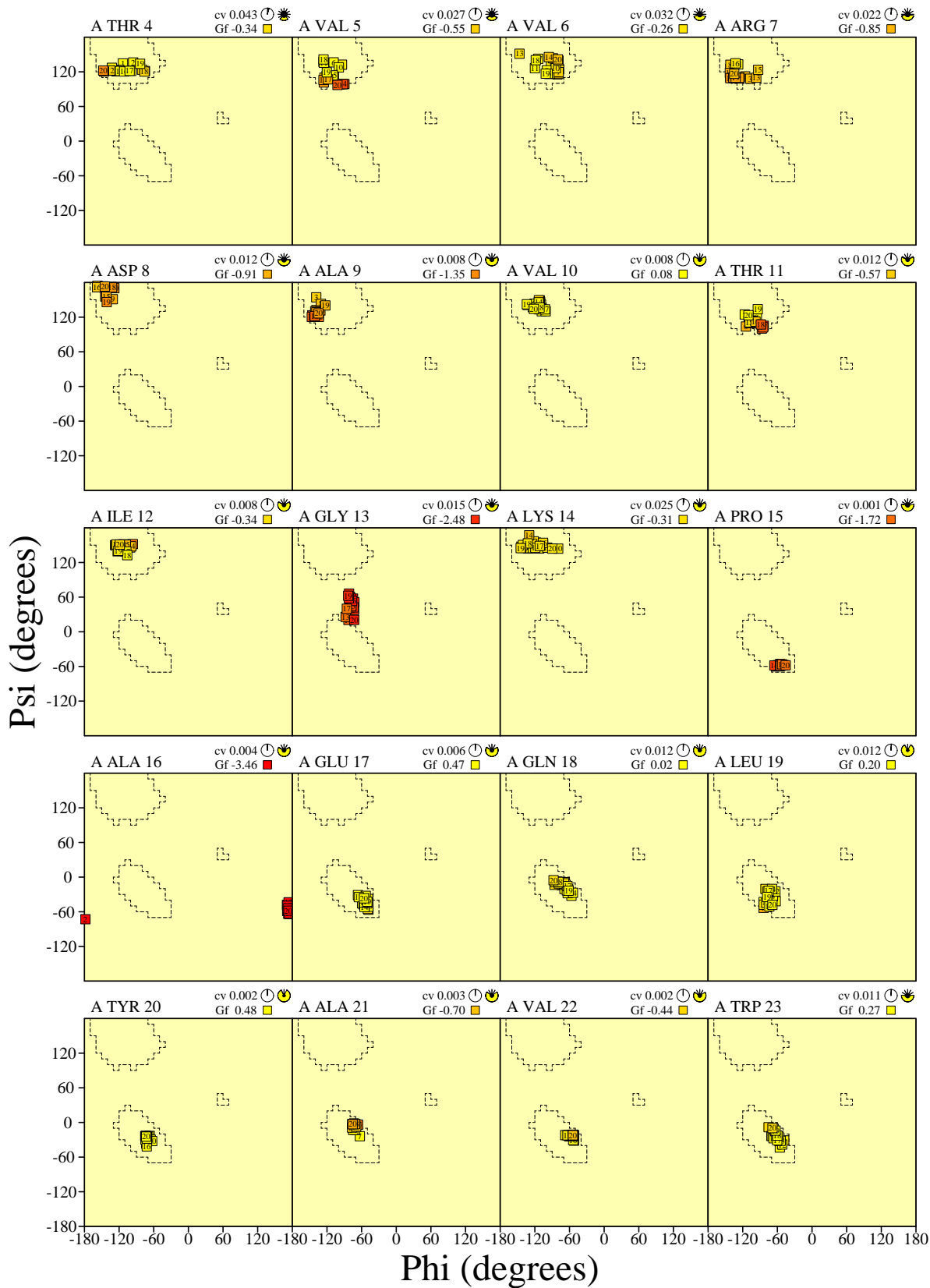


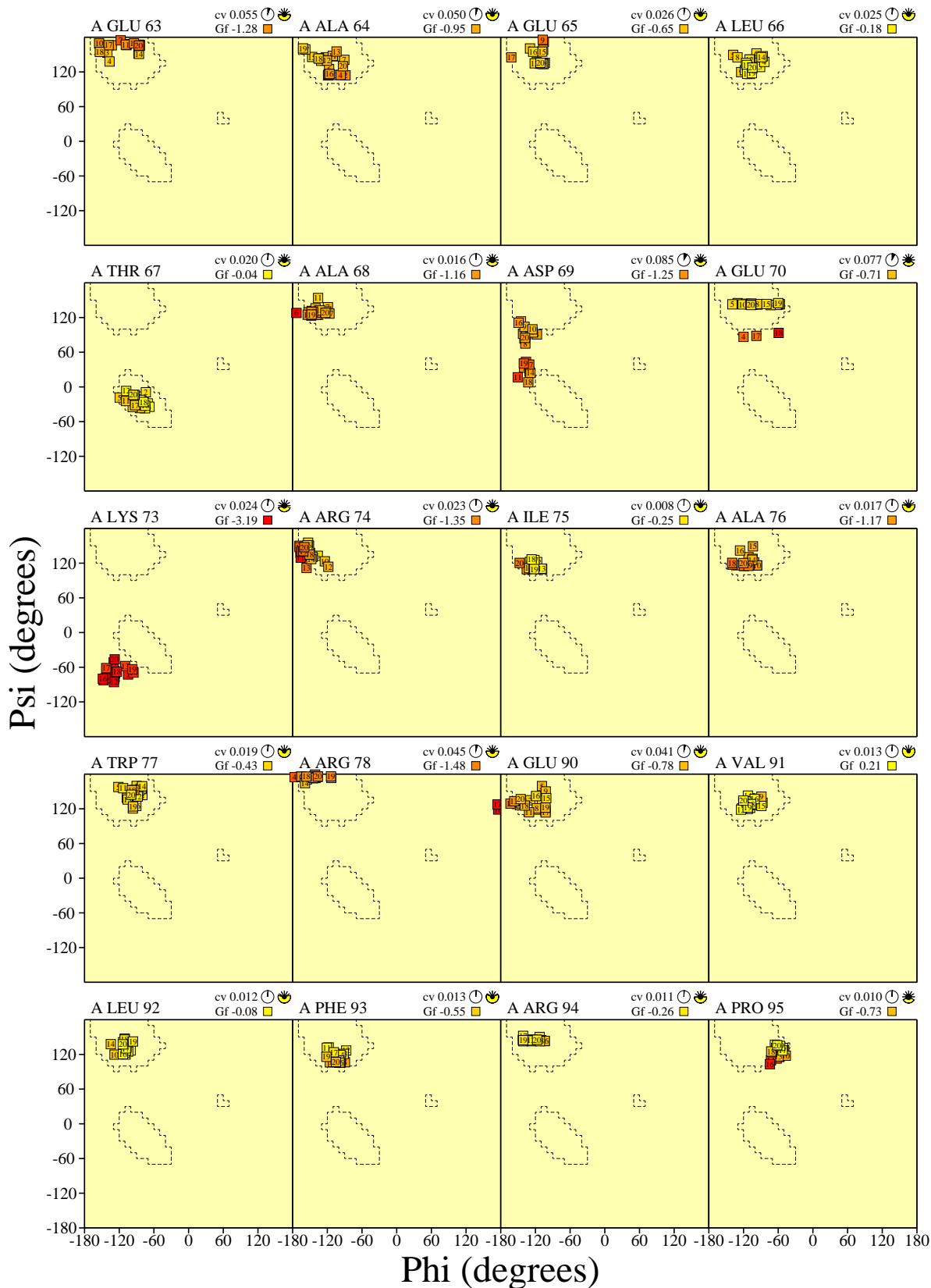
# Ensemble Ramachandran plots DRR147D\_NMR\_em\_bcr3 (20 models)\*\*



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

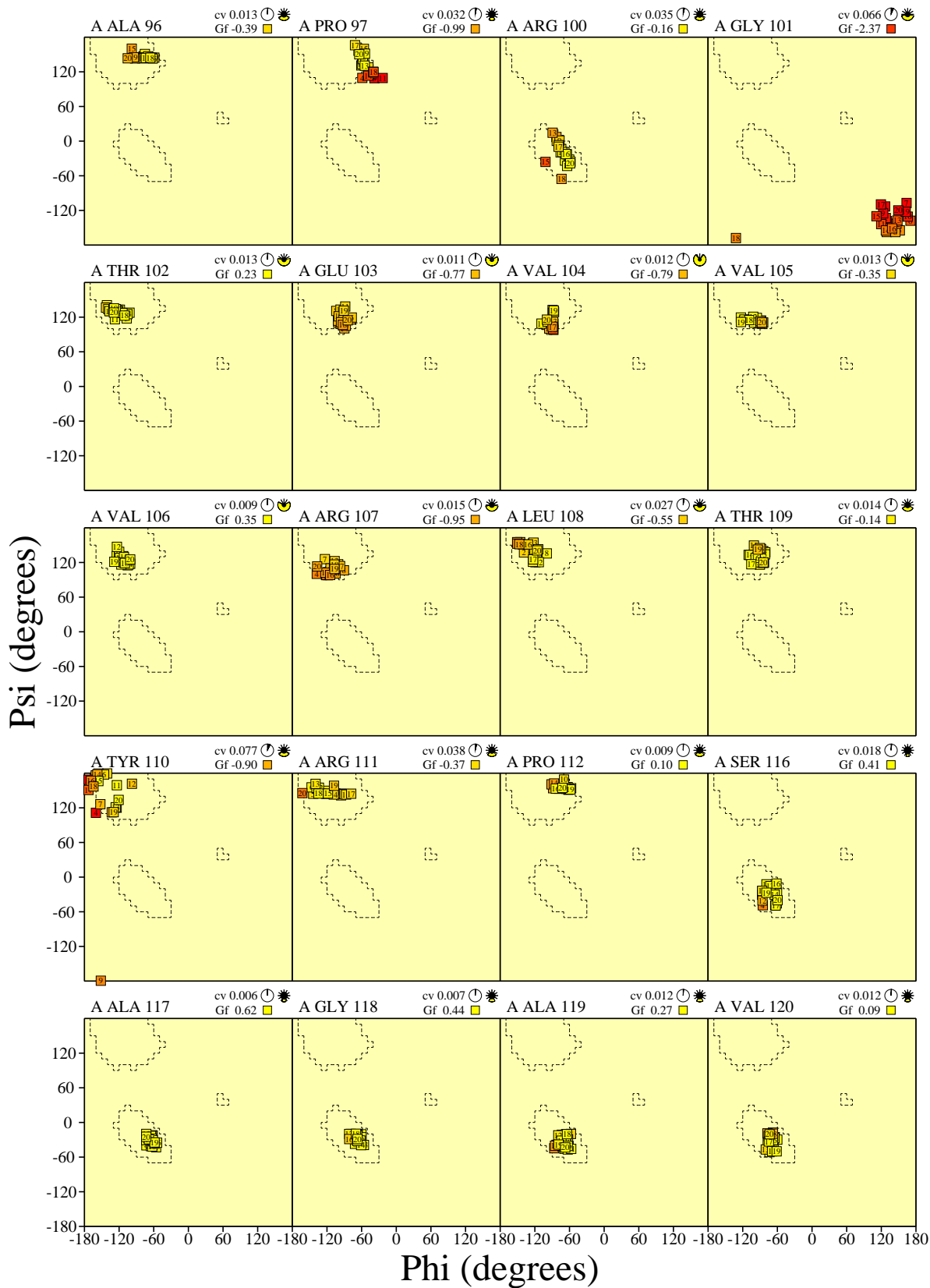
## DRR147D\_NMR\_em\_bcr3 (20 models)\*\*



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

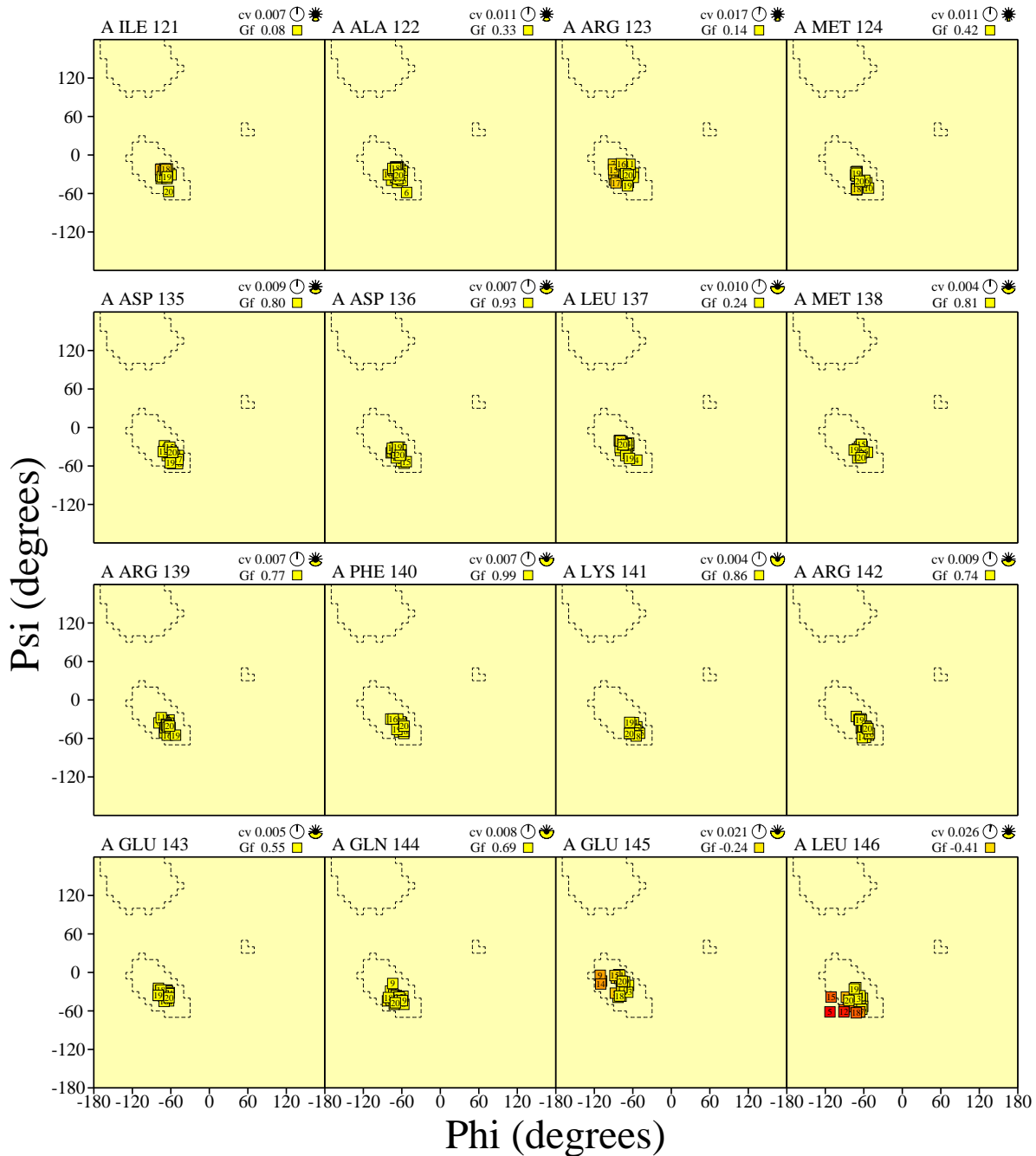
# Ensemble Ramachandran plots

## DRR147D\_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 DRR147D\_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