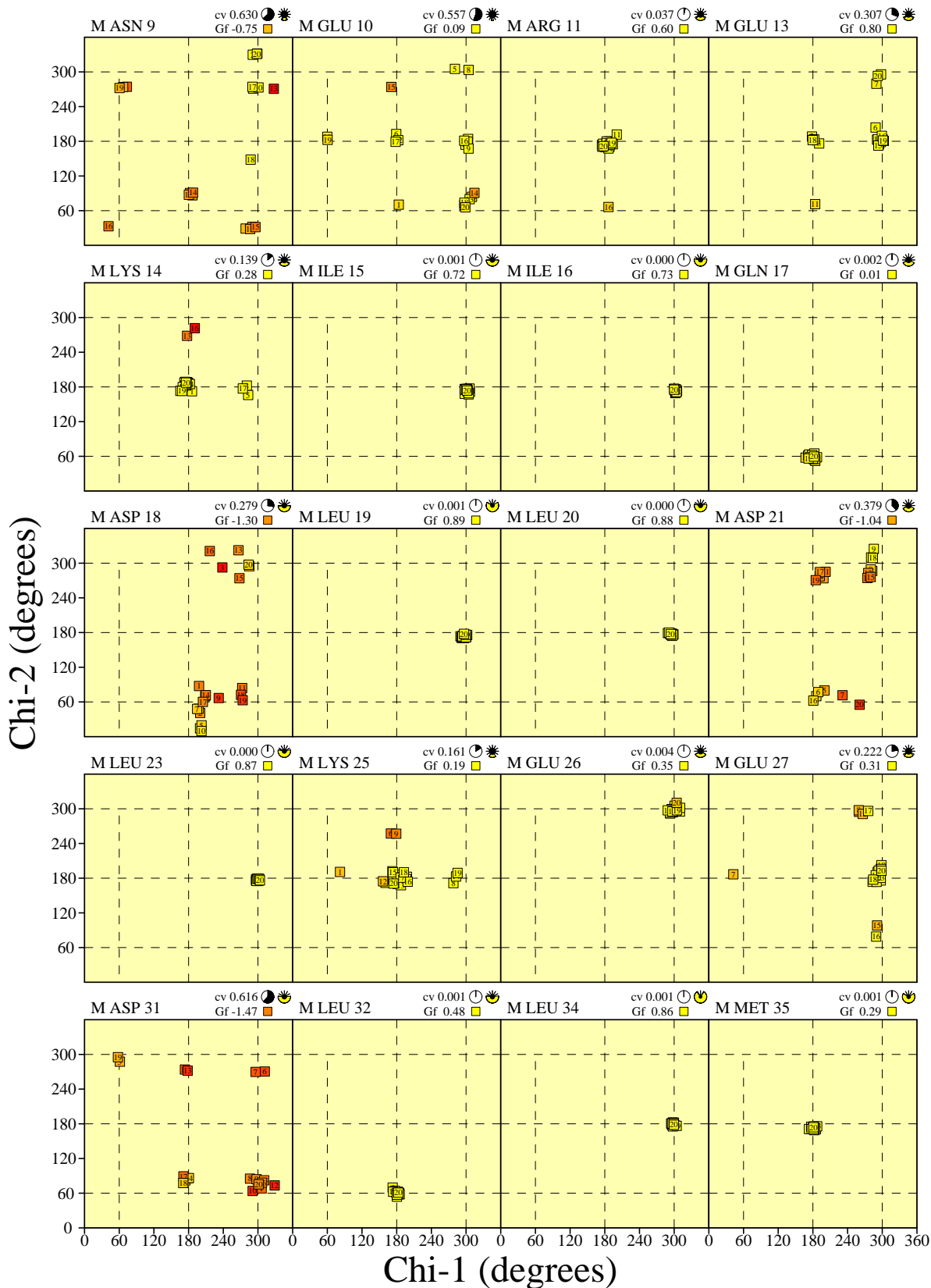


# Ensemble chi1-chi2 plots

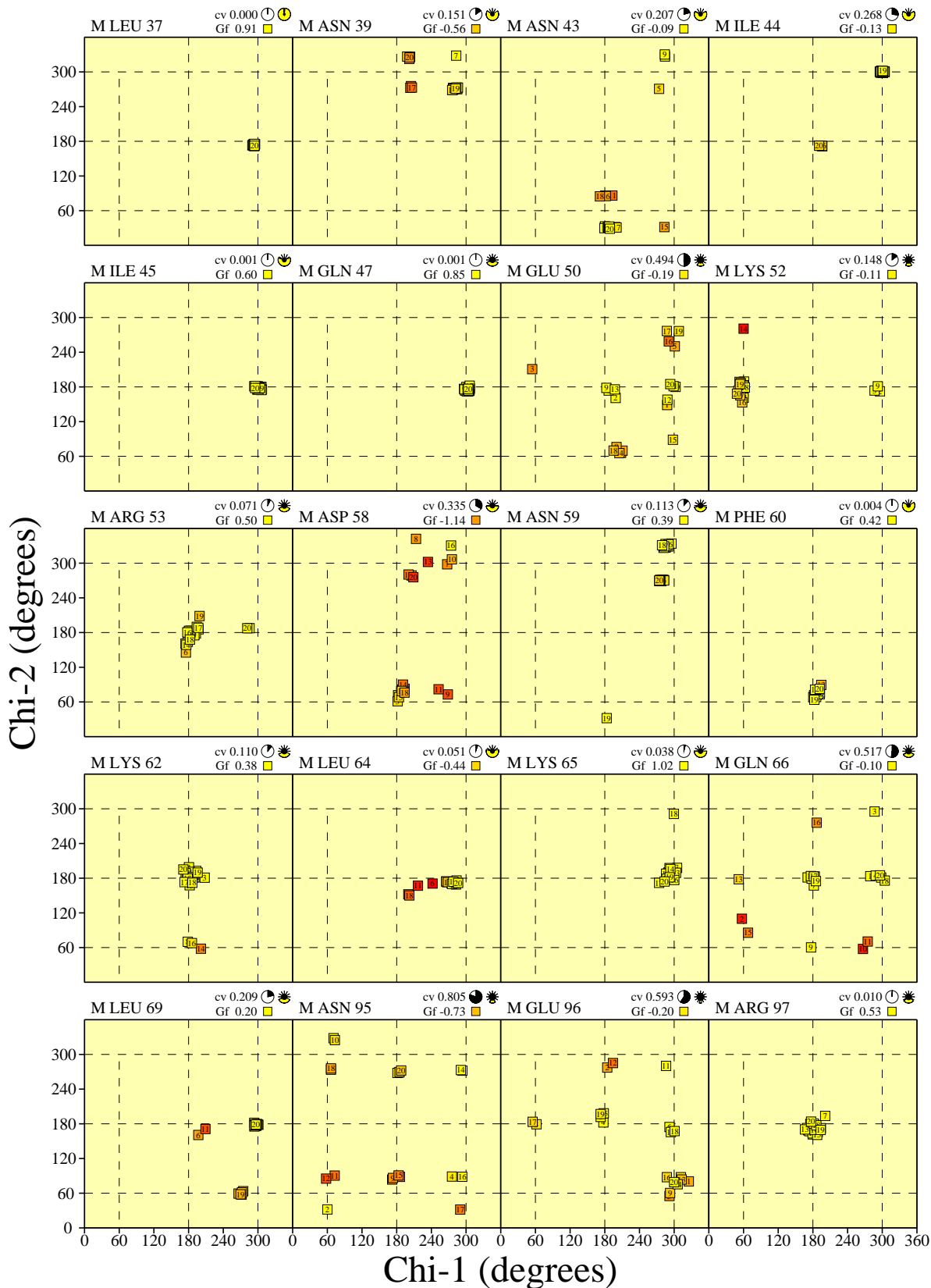
## CSR4\_NMR\_em\_bcr3 (20 models)\*\*



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

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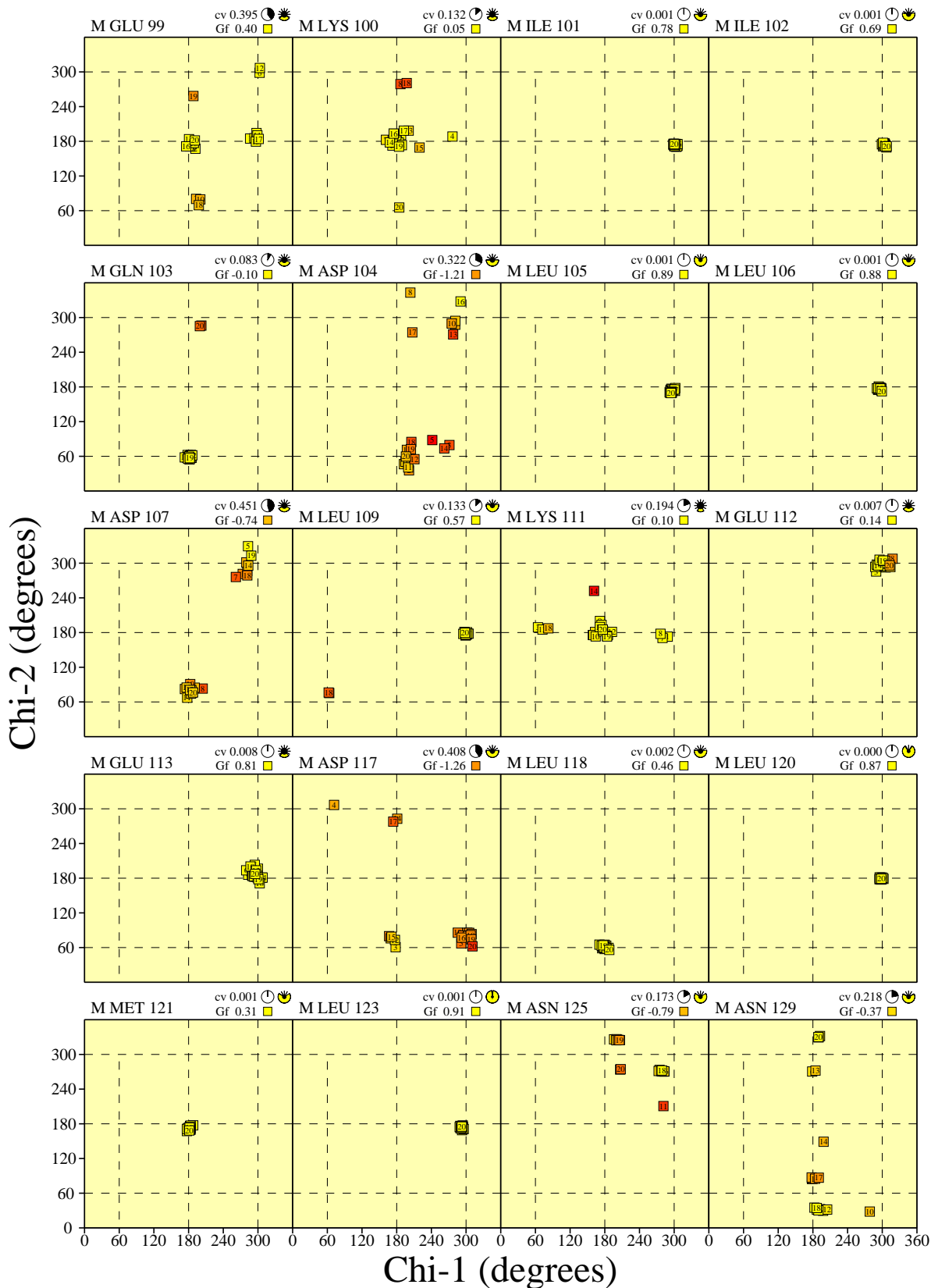
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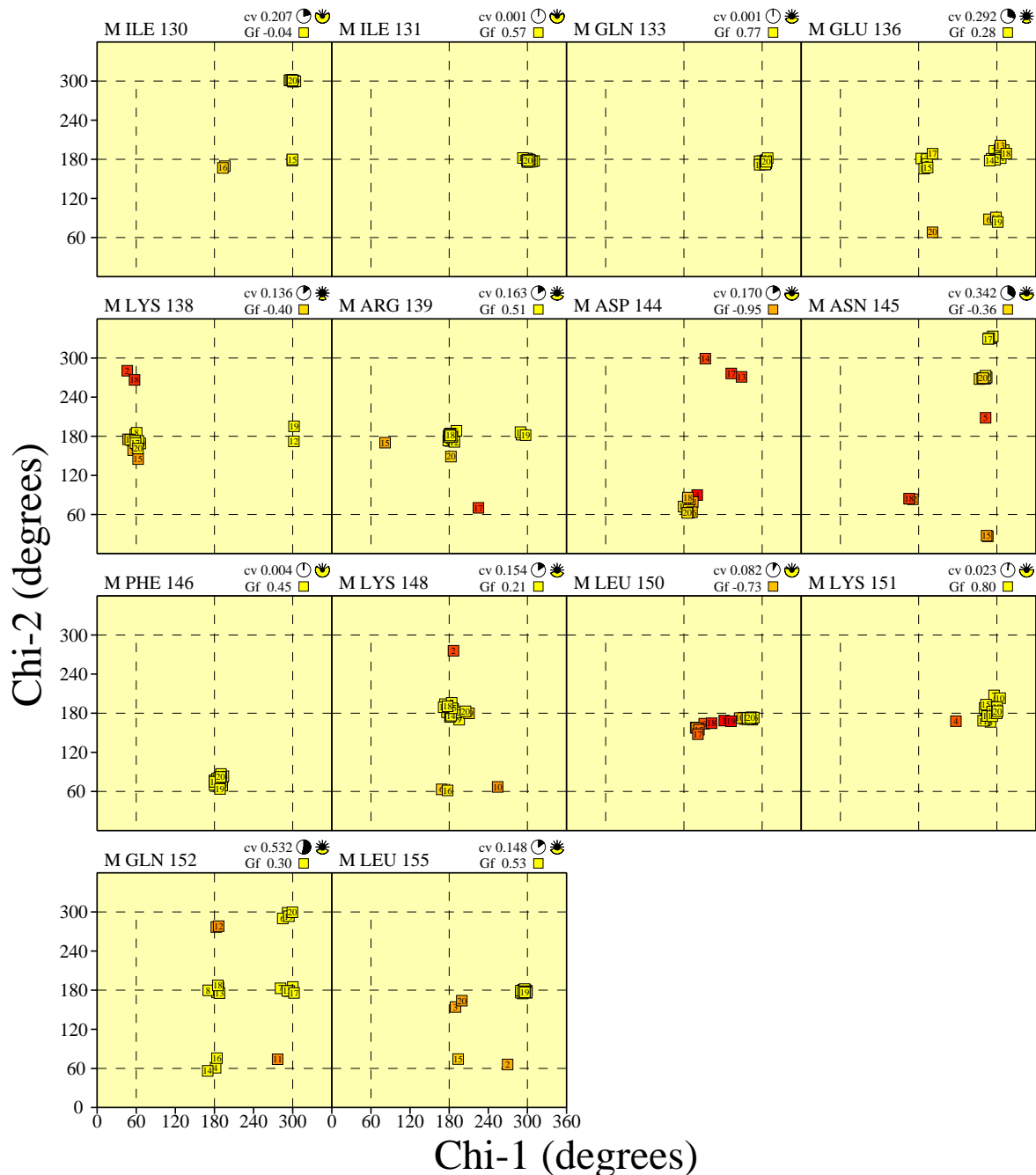
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