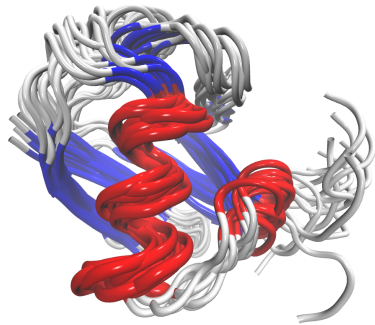
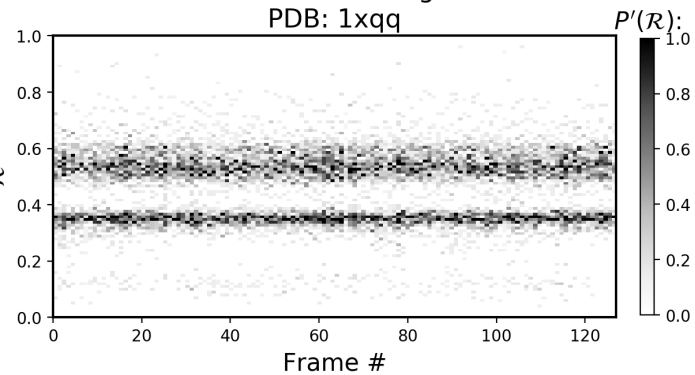


a.



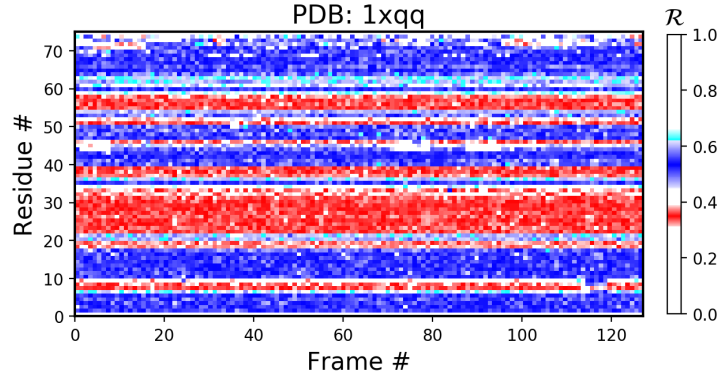
b.

Per-model \mathcal{R} -histogram
PDB: 1xqq



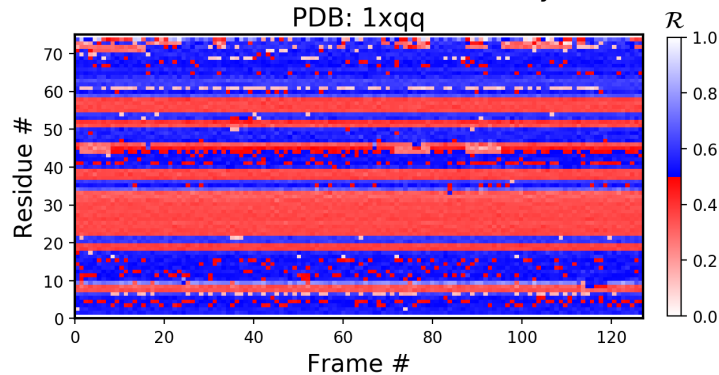
c.

Per-residue \mathcal{R} ; CMAP: SecondaryStructure
PDB: 1xqq



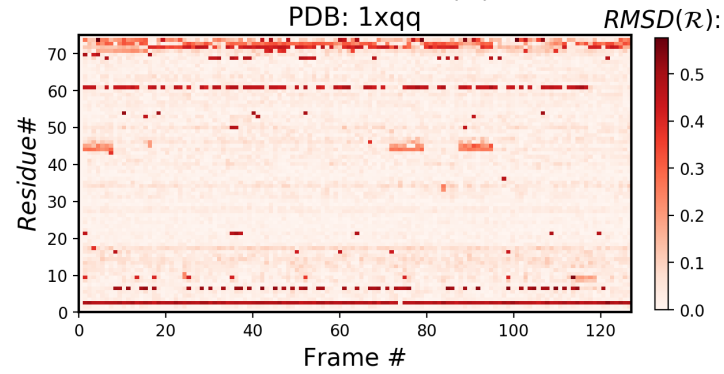
d.

Per-residue \mathcal{R} ; CMAP: Chirality
PDB: 1xqq



e.

Per-residue RMSD(\mathcal{R})
PDB: 1xqq



f.

Per-residue RMSF(\mathcal{R})
PDB: 1xqq

