

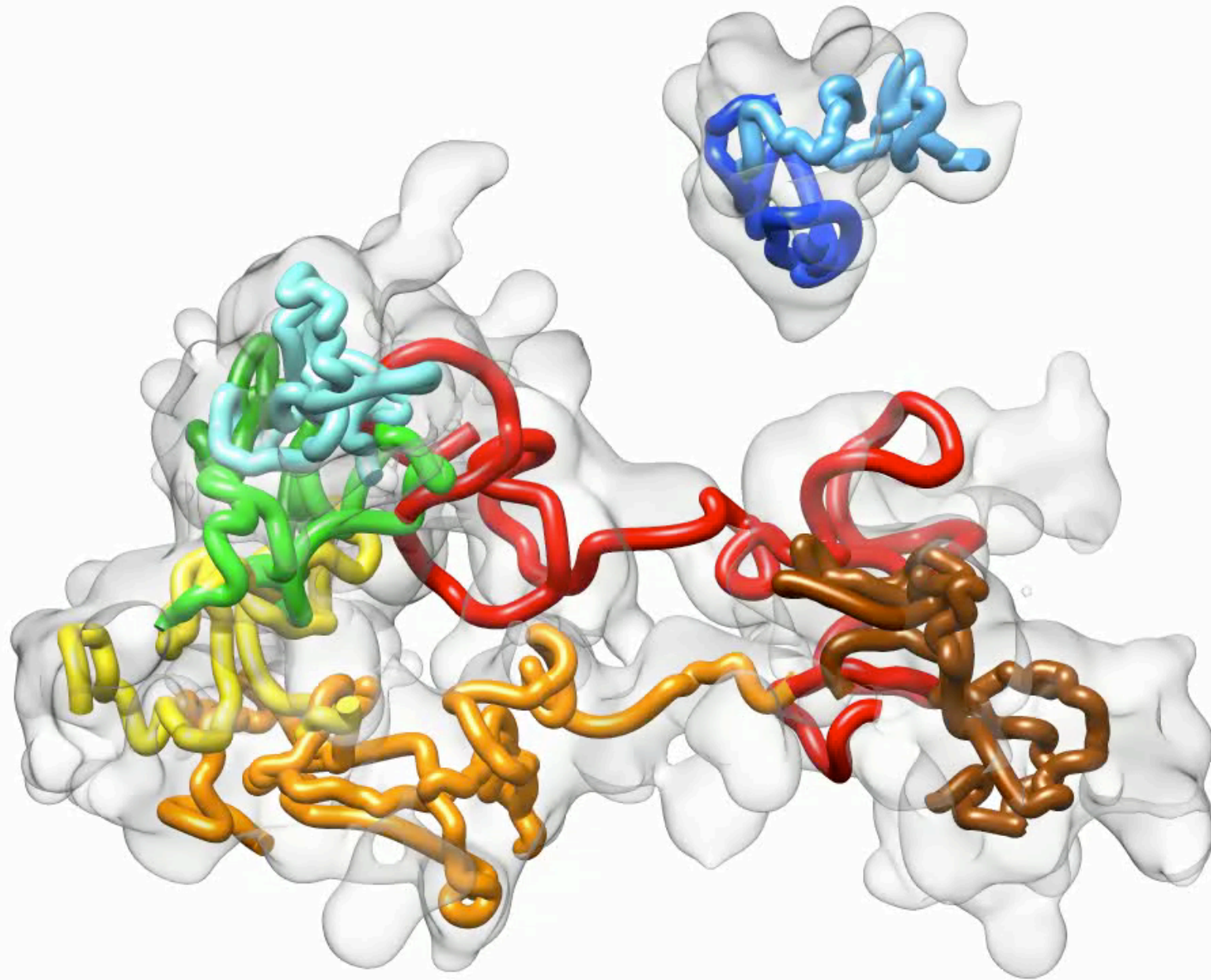
3DAROC18

Summary day #4

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Structural Genomics Group (CNAG-CRG)

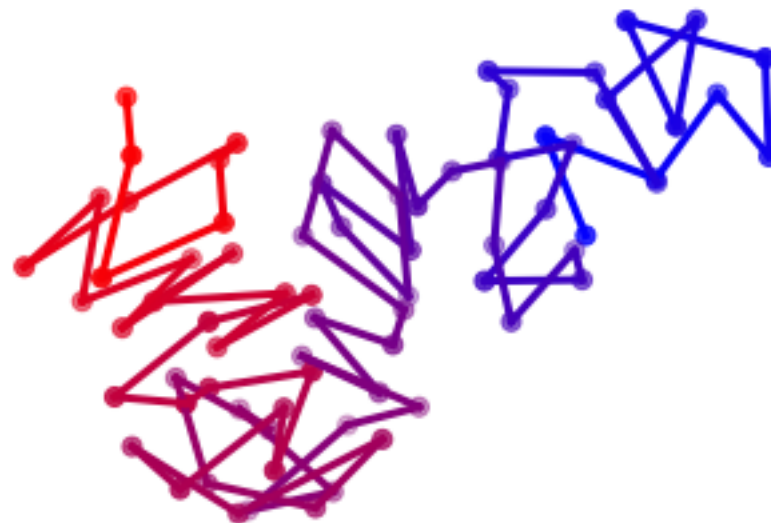
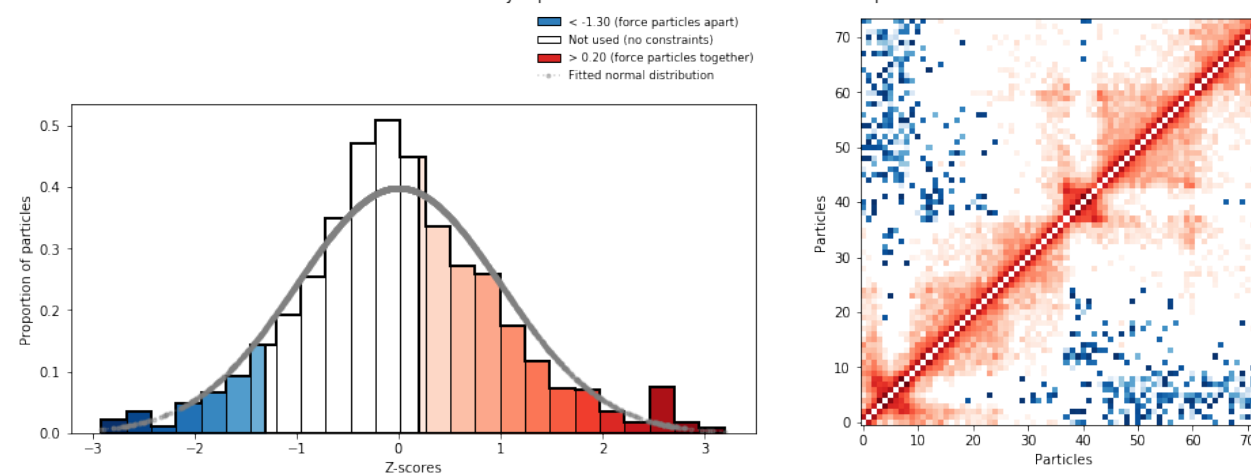
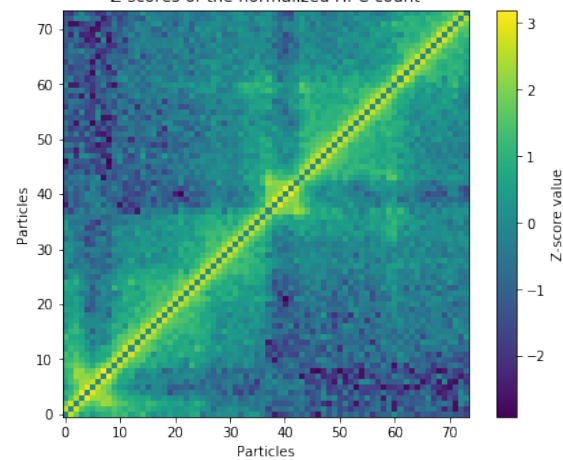
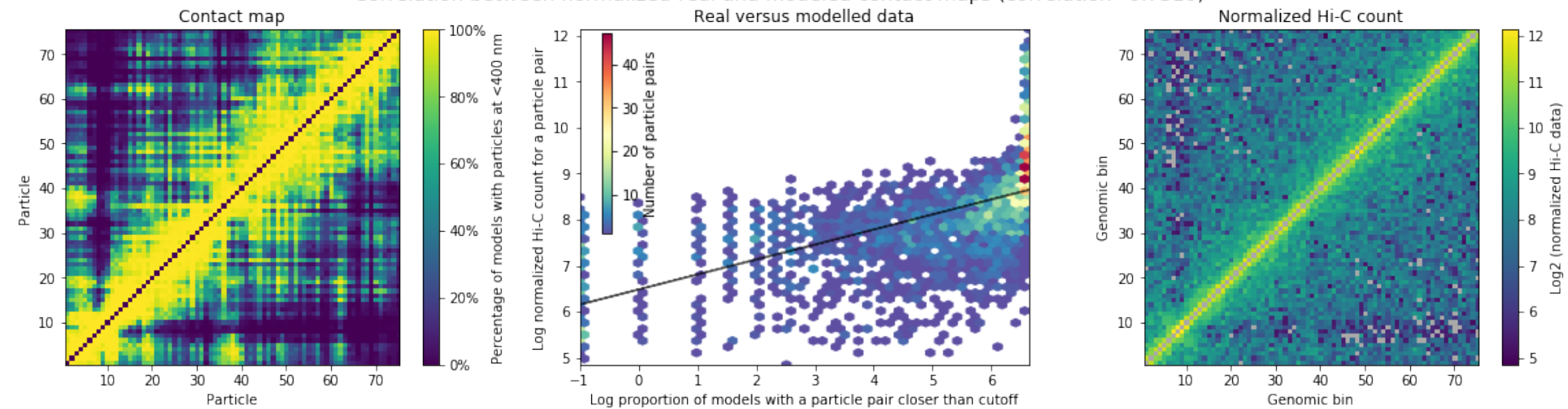


Integrating data types



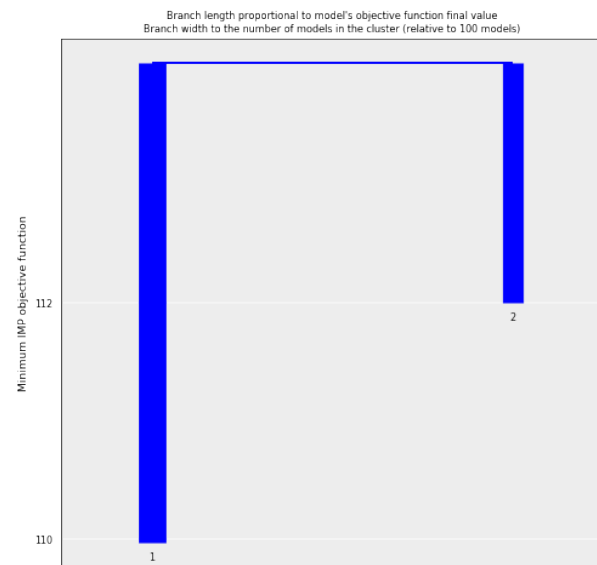
MuG - VRE



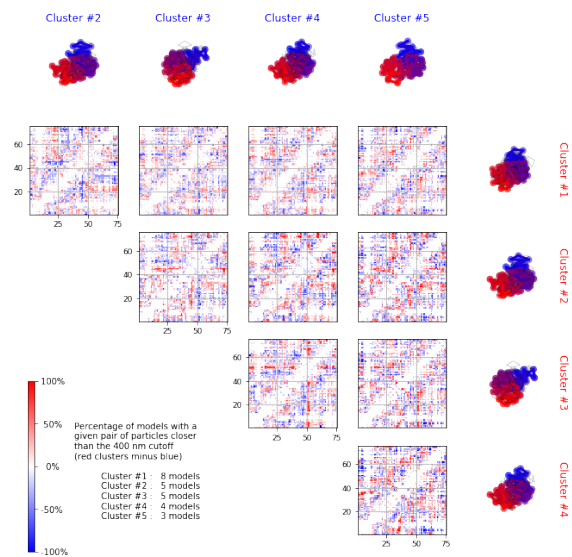


Model analysis

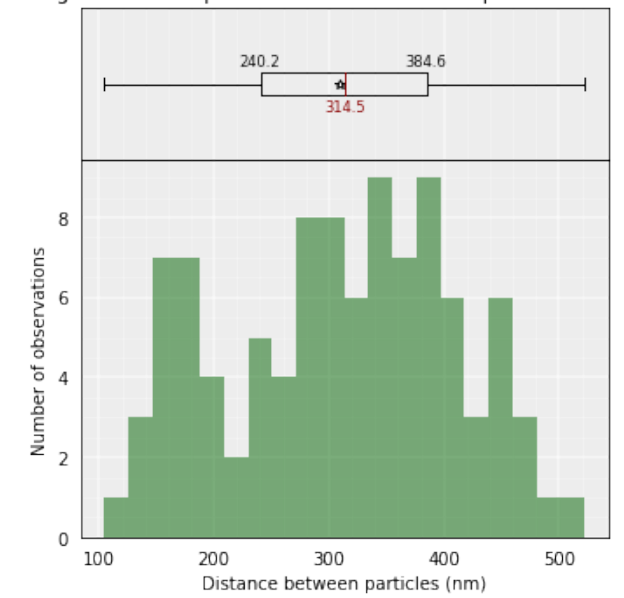
Dendrogram of clusters of 3D models



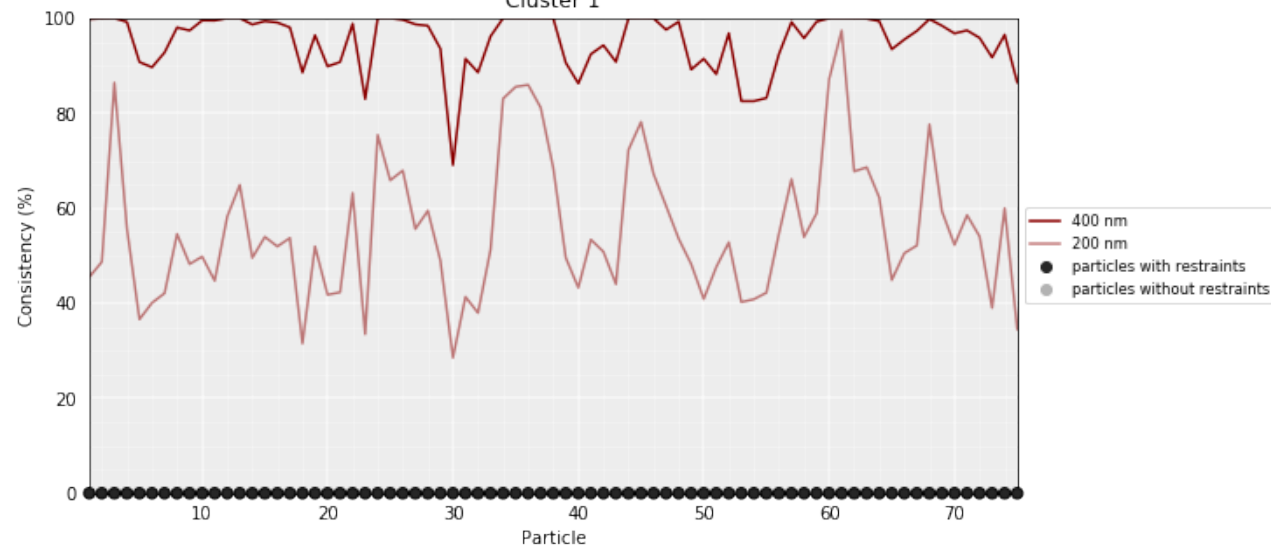
Deconvolution analysis for the 5 top clusters (cutoff=400 nm)



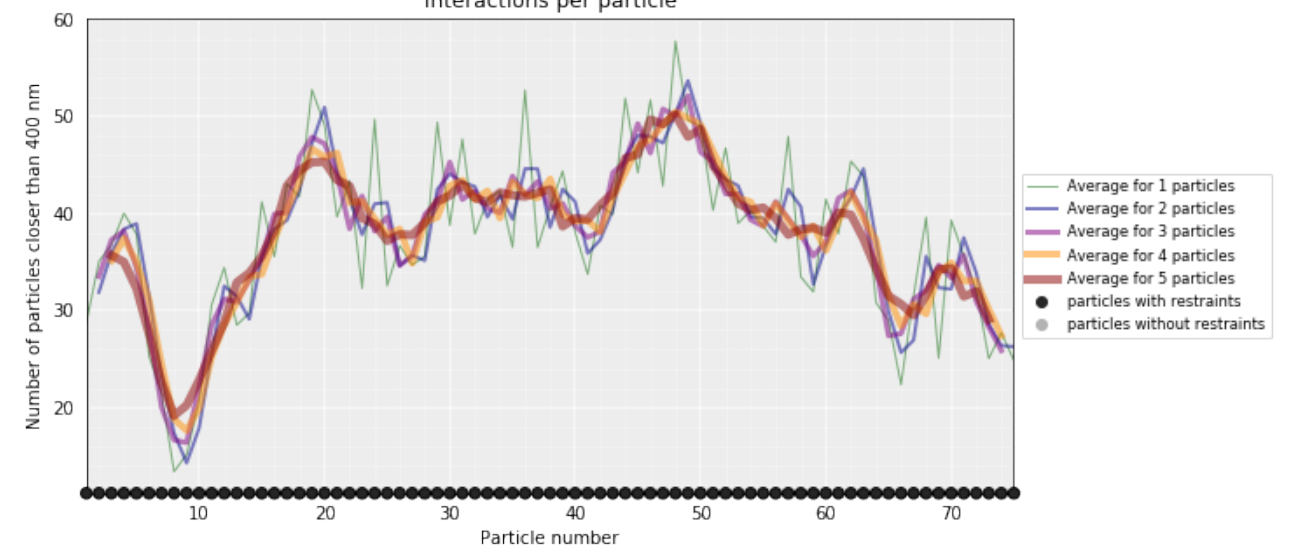
Histogram and boxplot of distances between particles 30 and 50



Cluster 1



Interactions per particle



How comfortable are you with...

- modeling genomes?
- analyzing the 3D genomes?
- reading the output of the genome analysis?

