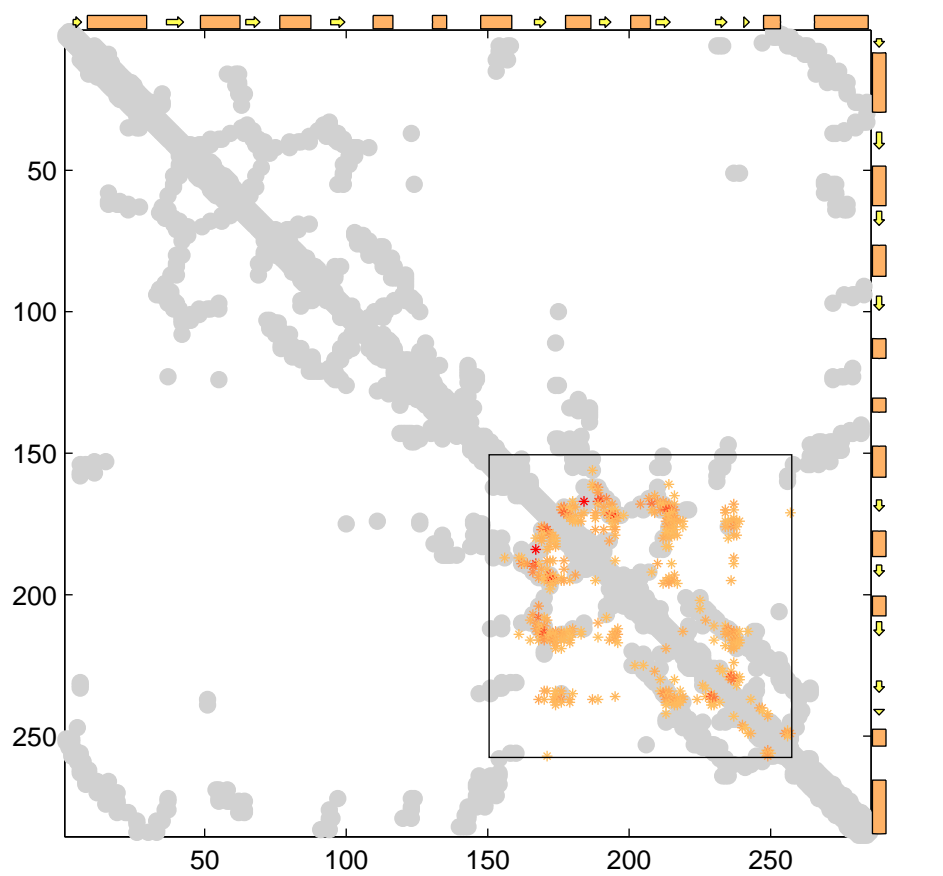


1a4i_A, Residues less than 5.00 Angstroms apart



Legend:

200 Evolutionary Couplings shown

- * Evolutionary Coupling (Max Score)
- * Evolutionary Coupling (Min Score)

- Known Structure Contact (from PDB structure)
- Alpha Helix (T.M. Span)
- Alpha Helix (Other)
- Beta Strand