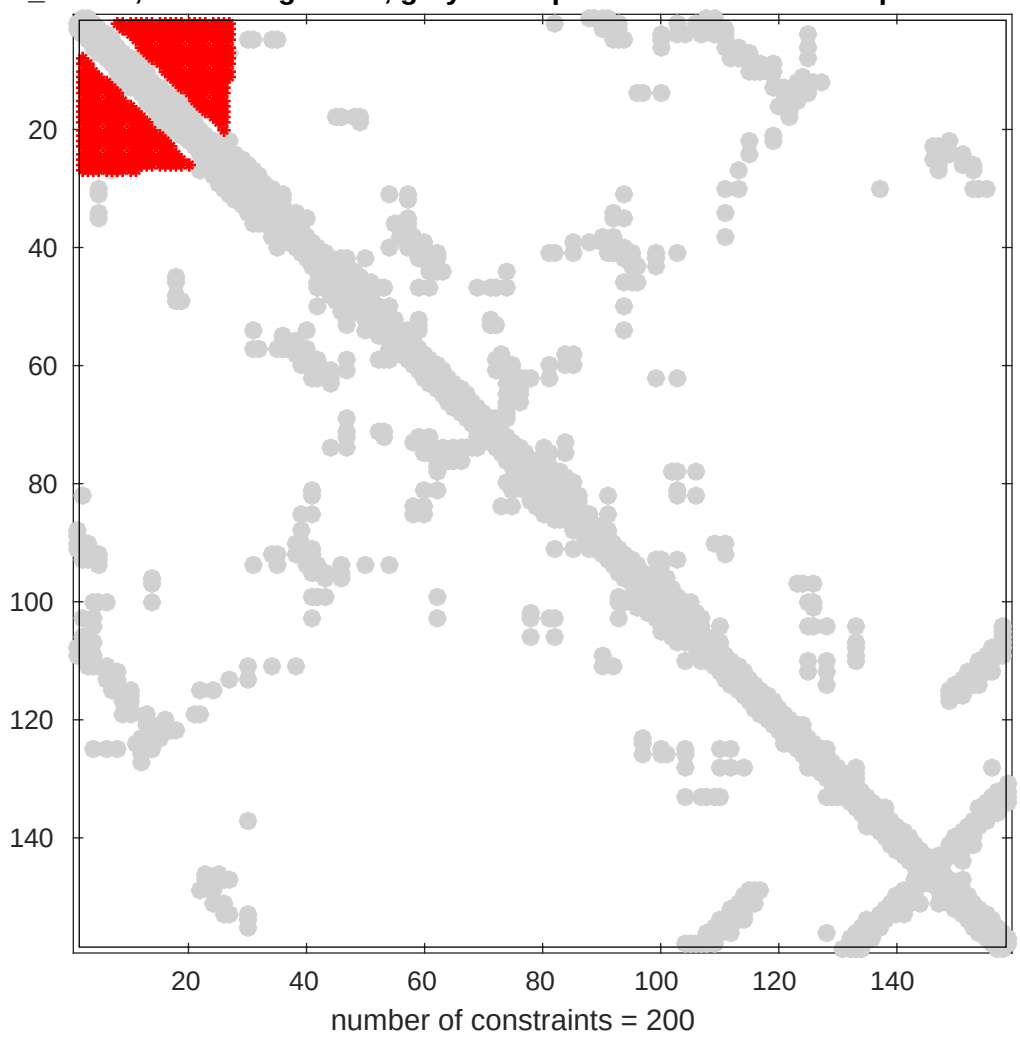DYR_ECOLI, red e3 alignment, grey 1RX2.pdb residues < 5.00 A apart