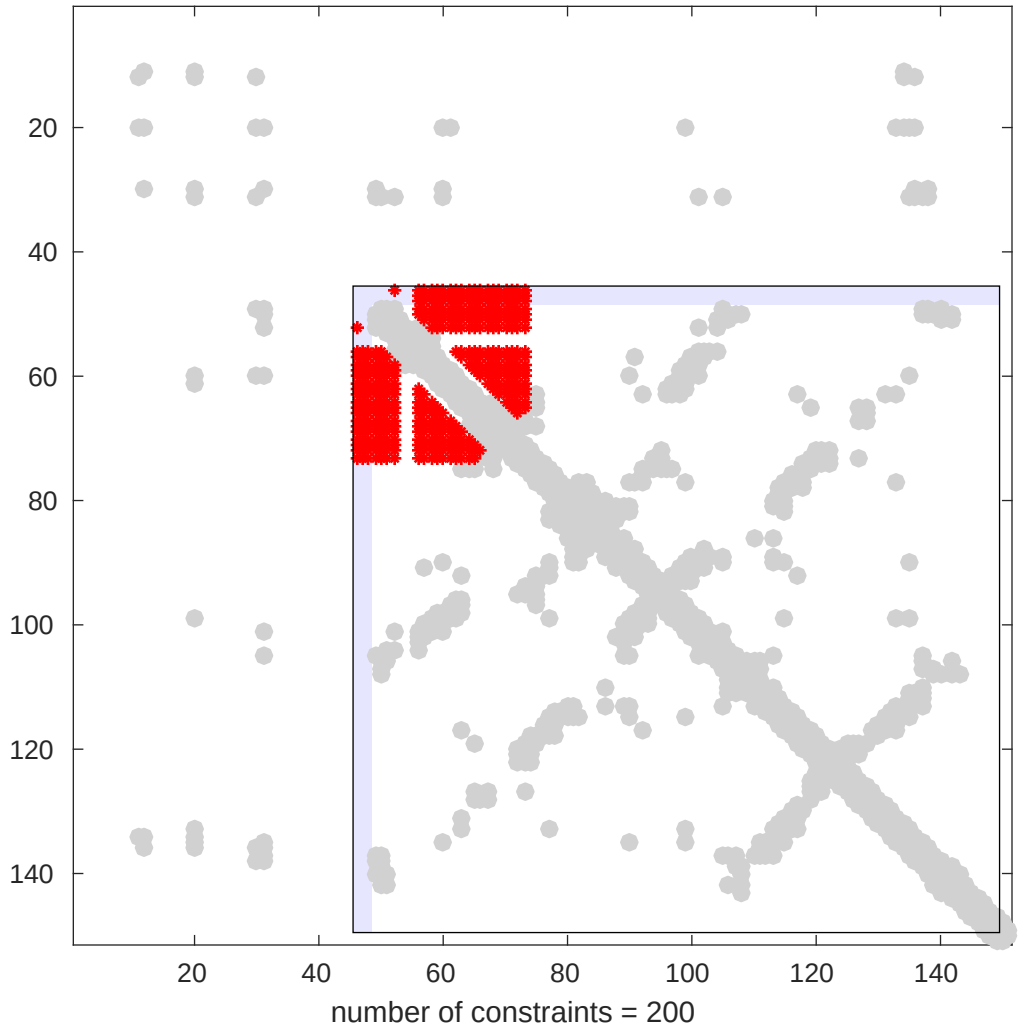CADH1_HUMAN, red e3 alignment, grey 2O72.pdb residues < 5.00 A apart