

Allowed overlap: 0.6
H-bond overlap reduction: 0.4
Ignore contacts between atoms separated by 4 bonds or less
Detect intra-residue contacts also

3 contacts

atom1	atom2	overlap	distance
GLN 900 CG	GLN 748 2HE2	0.795	2.085
GLN 900 CB	GLN 748 2HE2	0.736	2.144
GLN 900 CB	GLN 748 NE2	0.638	2.867