

Finding intermodel H-bonds
Finding intramodel H-bonds
Constraints relaxed by 0.4 angstroms and 20 degrees
Models used:
#0 complexwildtypedrug1.pdb

H-bonds (donor, acceptor, hydrogen, D..A dist, D-H..A dist):
LYS 257 NZ LIG 1.het 0 LYS 257 HZ3 2.622 1.772
LYS 736 NZ LIG 1.het 0 LYS 736 HZ3 2.852 2.013