

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

H-bonds (donor, acceptor, hydrogen, D..A dist, D-H..A dist):
HIS 112 N GLY 108 O HIS 112 HN 3.058 2.090
SER 147 N THR 143 O SER 147 HN 2.814 1.843
LYS 844 NZ SER 841 O LYS 844 HZ2 3.508 2.573