

Finding intermodel H-bonds
Finding intramodel H-bonds
Constraints relaxed by 0.4 angstroms and 20 degrees
Models used:
#0 complex14.pdb

H-bonds (donor, acceptor, hydrogen, D..A dist, D-H..A dist):

| | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-------------|-------|-----|
| ALA | 292 | N | GLY | 288 | O | no hydrogen | 2.874 | N/A |
| ASN | 296 | ND2 | LIG | 0 | 0 | no hydrogen | 2.836 | N/A |
| ASN | 296 | ND2 | ALA | 292 | O | no hydrogen | 2.720 | N/A |
| GLN | 773 | NE2 | ASN | 296 | OD1 | no hydrogen | 3.369 | N/A |
| GLY | 781 | N | PHE | 777 | O | no hydrogen | 2.983 | N/A |
| LYS | 826 | NZ | LIG | 0 | 0 | no hydrogen | 3.320 | N/A |
| SER | 831 | OG | LYS | 826 | O | no hydrogen | 3.522 | N/A |