

## Refmac: Final Results Summary



Values for a good fitted 3D map.

R factor ~ 0.3,

Rms BondLength ~ 0.02.

Save Data

	Initial	Final
R factor	0.4597	0.4509
Rms BondLength	0.0275	0.0227
Rms BondAngle	2.8824	2.6286
Rms ChirVolume	0.1845	0.1674