





PMV: <http://www.scripps.edu/~sanner/python/>

Coot: <http://www.yasbl.york.ac.uk/~emsl/ey/coot/>



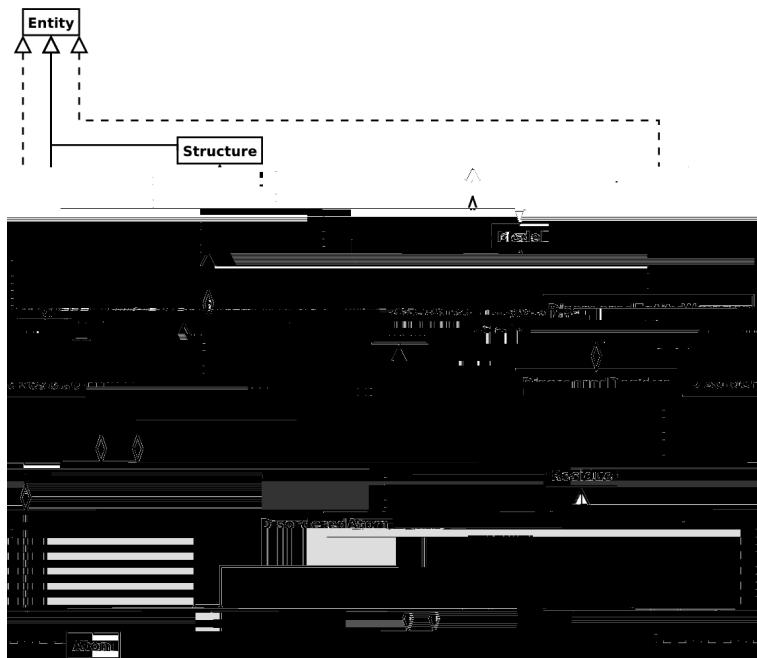
**How do I keep a local copy of the PDB up-to-date?**

This can also be done using the PDBList object. One simply creates a PDBList object (specifying the directory where the local copy of the PDB is present) and calls the update\_pdb method:

```
pl =PDBList(pdb='/data/pdb')
pl.update_pdb()
```

One can of course make a weekly cronjob out of this to keep the local copy automatic-







**How is disorder handled?**

This is one of the strong points of Bio.PDB. It can handle both disordered atoms and point mutations (ie. a Gly and an Ala residue in the same position).

**What about B factors?**

Well, yes! Bio.PDB supports isotropic and anisotropic B factors, and also deals with  
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**How do I measure angles?**







