Subject: Routine to read files in pdb-format?
Posted by Bernd Ihmels on Tue, 23 Jan 1996 08:00:00 GMT
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Has anyone ever written a routine to read the coordinates of a molecule from
a file in the common pdb-format and plot this molecule as a spacefill-plot,
i.e. to plot the atoms of the molecule as balls with the van der Waals
radius?

Any help will be appreciated.

Bernd
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