
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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* Bernd Ihmels, Max-Planck-Institut fuer biophysikalische Chemie *

* Biocomputation Group, AG 061 *

* Am Fassberg, 37077 Goettingen *

* Tel: +49+551-201-1762 Fax: +49+551-201-1467 *

* Email: bihmels1@gwdg.de *
