

THEORETICAL *and* COMPUTATIONAL
BIOPHYSICS GROUP

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Cube Plugin, Version 1.1

This plugin reads the atom positions and volumetric data sets in the plain text "cube" format as created by the **Gaussian program package**. The plugin supports cube files with single grid (e.g. densities, laplacian, electrostatic potential) and orbital cube files with multiple orbitals.

There are other programs that can create "cube" files natively or convert native data to the "cube" files. This plugin can only read those files, that comply with the following description:

```
C  WRITE A FORMATTED CUBEFILE VERY SIMILAR TO THOSE CREATED BY
C  THE GAUSSIAN PROGRAM OR THE CUBEGEN UTILITY.
C  THE FORMAT IS AS FOLLOWS (LAST CHECKED AGAINST GAUSSIAN 98):
C
C  LINE   FORMAT   CONTENTS
C  =====
C  1      A         TITLE
C  2      A         DESCRIPTION OF PROPERTY STORED IN CUBEFILE
C  3      I5,3F12.6  #ATOMS, X-,Y-,Z-COORDINATES OF ORIGIN
C  4-6    I5,3F12.6  #GRIDPOINTS, INCREMENT VECTOR
C  #ATOMS LINES OF ATOM COORDINATES:
C  ...    I5,4F12.6  ATOM NUMBER, CHARGE, X-,Y-,Z-COORDINATE
C  REST:  6E13.5    CUBE DATA (WITH Z INCREMENT MOVING FASTEST, THEN
C                   Y AND THEN X)
C
C  FOR ORBITAL CUBE FILES, #ATOMS WILL BE < 0 AND THERE WILL BE ONE
C  ADDITIONAL LINE AFTER THE FINAL ATOM GIVING THE NUMBER OF ORBITALS
C  AND THEIR RESPECTIVE NUMBERS. ALSO THE ORBITAL NUMBER WILL BE
C  THE FASTEST MOVING INCREMENT.
C
C  ALL COORDINATES ARE GIVEN IN ATOMIC UNITS.
```

Notes:

Coordinate Rotation / Periodic Display

VMD currently can only display periodic images of the unit cell correctly, if the first cell vector is collinear with the x-axis and the second cell vector in the xy-plane. Upon loading of the cube file, the coordinates and the grid vectors are rotated accordingly. Unfortunately, that may make the atomic positions incompatible to some other coordinate files containing matching coordinates.

Read-only

This plugin is only used to read files. VMD does not currently support writing this file type.

Memory Usage / Large Cube Files

Due to the way cube files with multiple grids (e.g. orbitals) are written, the plugin will temporarily hold all gridded data in memory to get a **huge** performance increase. So loading a single orbital data set from a large cube file with many orbitals may need a machine with a lot of available memory and/or swap space. Also the plugin uses some features internally, which may not work for files larger than 2GB on 32-bit systems.

Atom Masses

The plugin assigns masses to the atoms according to the natural isotope distribution of the elements (see e.g. <http://www.webelements.com/>). If an element could not be recognized, a mass of 0 is assigned. To set the mass for those elements or assign a different mass for the other atoms, you have to use to create a selection with **atomselect** and then use the selection subcommands to set the mass.

Binary Cube Files

Unformatted binary cube files, as they were created by versions before Gaussian 98 are not supported by this plugin.

Atomic Positions in Angstrom

The atomic positions and the grid vectors in a cube file are **always** in atomic units. There is a comment in the Gaussian documentation, that describes, how you can provide a custom grid in angstrom in the **input** of a gaussian job.

Incompatibilities

If VMD cannot read your cube file, please check the above notices. If your cube file has a negative number of atoms, but is missing the orbitals line, you may still be able to read it, by changing the sign of the number of atoms. But most likely the positions and the grid will be given in angstrom then (see the comment above), so all distances will be too short by a factor of about 1.9.