## **NAME**

TRANSGEOM - Transfers optimized geometry from a model system to a production run input

## **SYNOPSIS**

```
Transgeom [ -d? (?:0~3) ] [ -h ] [ -q ] [ -atmp ] [ -atmtypp ] [ -chgp ] [ -coordp ] [ -layerp ] [
-movep ] [ -t tolerance ] filename
```

# **DESCRIPTION**

This program extracts the atomic setup information from calculations of a model system and prepares a production run input

## **OPTIONS**

Command line option specifications are processed from left to right and may be specified more than once. If conflicting options are specified, later specifications override earlier ones.

<b>-d?</b> (?:0~3)	Turn on debug printing. The printing level can be controlled by a given number. The larger the number, the more information will be printed when the program is running.
-h	
help	Print full TRANSGEOM documentation via perldoc. Cannot be used with other options.
<b>-q</b>	Run in quiet mode and do not print progress messages.
-atmp	If set, use the element information from the production ONIOM input file. Default uses model system input file.
-atmtypp	If set, use the atom type information from the production ONIOM input file. Default uses model system input file.
-chgp	If set, use the partial charges information from the production ONIOM input file. Default uses model system input file.
-coordp	If set, use the coordinate information from the production ONIOM input file. Default uses model system input file.
-movep	If set, use the move flag information from the production ONIOM input file. Default uses model system input file.
-layerp	If set, use the layer setup information from the production ONIOM input file. Default uses model system input file.
-t tolerance	Tolerance to decide if an atom has been moved during the optimization. Default is 0.00001;

filename

Format:
---------

Modelgjf ONIOM input file name for model system Modelonb

ONIOM ONB file for model system

Productiongjf ONIOM input file name for production run system

ONIOM ONB file for production run system Productiononb

Productioninput New ONIOM input file for production run system (Def

These files can be specified in any order.

The parameter filename must be the last command line argument when running transgeom.

## **EXAMPLES**

Called without any parameters, TRANSGEOM will display usage information. If -h or --help is passed, then the full TRANSGEOM documentation is displayed via perldoc.

transgeom foo.in

TRANSGEOM reads foo.in, then creates a production run input. Please note that there is no flag needed before setup file name in command line. Sample foo.in

ModelgjfONIOMmodelinput.gjfModelonbONIOMmodelinput.onbProductiongjfONIOMproductioninput.gjfProductiononbONIOMproductioninput.onb

Productioninput ONIOMproductioninputwithmodelgeom.gjf

## **NOTES**

The model system is not necessary smaller than the production system. A full protein model can be used as a model system, and a reduced size model for the same protein can be used as the production system. TRANSGEOM will extract corresponding portion of the geometry from the full size model and build a partial model.

# **VERSION**

1.1

# **AUTHOR**

Peng Tao, <tao.21@osu.edu>

# **COPYRIGHT**

Copyright (c) 2009~2010 by Peng Tao