mm_trj2pdb V.1.1.0

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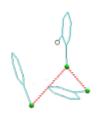
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mm_trj2pdb



1.1 Introduction

About this program:

· Program that creates PDB file from trajectory snapshot

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```
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```

License:

• GPL

1.2 How to use

Usage:

2 mm_trj2pdb

Install

2.1 Requirements

The application mm_trj2pdb requires the following external stuff:

```
• cmake >= 2.8
```

• gcc >= 4.8

2.2 How to install

2.2.1 Linux

```
* mkdir build && cd build

* cmake -DCMAKE_INSTALL_PREFIX=/usr -DCMAKE_BUILD_TYPE=Release ../

* make

* make install
```

2.2.2 Windows

```
* create project file using 'cmake'
* compile project
```

You may also download compiled executable file for Win_x86.

Install

Changelog

V.1.0.3 (2013-08-30)

• Bug fixes

V.1.0.1 (2013-07-27)

• initial release

Changelog 6

File Index

4.1 File List

Here is a list of all files with brief descriptions:

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src/coords.c															 					 			10)
src/main.c																								
src/messages.c																								
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File Documentation

5.1 src/add_main.c File Reference

```
#include <stdio.h>
#include "messages.h"
```

Functions

• int error_checking (const const char *input, const char *output)

function that checks errors in input variables

int print_message (const int quiet, FILE *std_output, const int log, FILE *f_log, const int mode, const char *str)

function that prints message in log and stdout

int set_defaults (char *input, int *log, char *output, int *quiet)

function that sets default values of variables

5.1.1 Function Documentation

```
5.1.1.1 int error_checking ( const const char * input, const char * output )
```

function that checks errors in input variables

```
* error_checking (input, output);
*
```

Parameters

aglinp	aglomerate file name
cell	massive of cell size
input	input file name
output	output file name

Returns

```
11 - error in 'input'
```

12 - error in 'output'

0 - exit without errors

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5.1.1.2 int print_message (const int quiet, FILE * std_output, const int log, FILE * f_log, const int mode, const char * str)

function that prints message in log and stdout

```
* print_message (quiet, stdout, log, f_log, 0, str);
```

Parameters

quiet	status of quiet-mode
std_output	stdout
log	status of log-mode
f_log	log file
mode	number of message in "messages.c"
str	additional text in message

Returns

0 - exit without errors

```
5.1.1.3 int set_defaults ( char * input, int * log, char * output, int * quiet )
```

function that sets default values of variables

```
* set_defaults (input, &log, output, &quiet);
```

Parameters

input	mask of trajectory files
log	status of log-mode
output	output file name
quiet	status of quiet-mode

Returns

0 - exit without errors

5.2 src/coords.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
```

Functions

• int reading_coords (const int mode, const char *filename, const int type_inter, const int *label_atom, const float *cell, int *num_mol, int *num_atoms, int *true_label_mol, int *label_mol, int *type_atoms, float *coords, char *ch_type_atoms)

function that reads coordinates from special file format

5.2.1 Function Documentation

5.2.1.1 int reading_coords (const int *mode*, const char * *filename*, const int *type_inter*, const int * *label_atom*, const float * *cell*, int * *num_mol*, int * *num_atoms*, int * *true_label_mol*, int * *label_mol*, int * *type_atoms*, float * *coords*, char * *ch_type_atoms*)

function that reads coordinates from special file format

```
* reading_coords (0, filename, type_inter, label_atom, cell, &num_mol, &num_atoms,

* true_label_mol, label_mol, type_atoms, coords, ch_type_atoms);

*
```

Parameters

mode of reading; '1' is statgen, '2' is envir or frad, '3' is agl
input file name
number of needed atoms (number of needed molecules)
massive of needed atom types (massive of needed molecules)
massive of cell size
number of molecules
number of atoms
massive of true numbers of molecule for atoms
massive of numbers of molecule for atoms
massive of atom types
massive of coordinates
massive of char atom types

Returns

- 1 file \$filename does not exist
- 2 unknown mode
- 0 exit without errors

Work blocks

```
reading file translation free memory
```

5.3 src/main.c File Reference

```
#include <math.h>
#include <stdio.h>
#include <stdlib.h>
#include <string.h>
#include "add_main.h"
#include "coords.h"
#include "messages.h"
#include "print_struct.h"
```

Functions

• int main (int argc, char *argv[])

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5.3.1 Function Documentation

5.3.1.1 int main (int argc, char * argv[])

Returns

- 1 error in error_checking
- 2 input file does not exist
- 3 memory error
- 4 unknown flag
- 0 exit without errors

5.4 src/messages.c File Reference

```
#include <stdio.h>
#include <time.h>
```

Functions

int message (const int log, const int mode, const char *text, FILE *output)
 function that prints messages to output

5.4.1 Function Documentation

5.4.1.1 int message (const int log, const int mode, const char * text, FILE * output)

function that prints messages to output

```
* message (log, mode, text, output);
*
```

Parameters

log	equal to 1 if print to logfile
mode	number of message
text	additional text
output	output file (may be stdout)

Returns

- 1 unknown mode
- 0 exit without errors

5.5 src/print_struct.c File Reference

```
#include <stdio.h>
```

Functions

• int print_structure (const char *output, const int num_needed_mol, const int *needed_mol, const int num_atoms, const int *label_mol, const char *ch_type_atoms, const float *coords)

function that prints structure to pdb file

5.5.1 Function Documentation

5.5.1.1 int print_structure (const char * output, const int num_needed_mol, const int * needed_mol, const int num_atoms, const int * label_mol, const char * ch_type_atoms, const float * coords)

function that prints structure to pdb file

Parameters

output	output file name
num_needed	number of needed molecules
mol	
needed_mol	massive of number of needed molecules
num_atoms	number of atoms
label_mol	massive of numbers of molecule for atoms
ch_type_atoms	massive of char atom types
coords	massive of coordinates

Returns

0 - exit without errors

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