# Documentation for networkDiffusion Code

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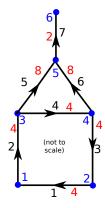
The code in this package will simulate diffusion on a network of 1D paths.

# Notes for myself describing code implementation

• Assume all diffusion is on purely 1D paths, so just need to keep track of where in the network the particle goes

Example network to use in code development:

blue = node numbersblack = edge numbersred = edge lengths



- Propagation in 1D interval with 2 absorbing boundaries:
  - Use either sine series expansion (long times) or image method expansion (short times)
  - When sampling from cumulative distribution, require convergence to SOLVE-TOLF in function value and SOLVETOLX in x value

## 1 Compilation Instructions

To compile and run the program, you will need the following:

- a compiler capable of handling Fortran 90. The code has been tested with the gfortran and compiler. The default compiler is gfortran.
- BLAS and LAPACK libraries installed in a place where the compiler knows to look for them
- Optionally: Matlab to visualize output data

The code has been tested on Ubuntu Linux.

To compile with gfortran, go into the source directory. Type make. To compile with any other compiler that can handle Fortran 90, type

#### make FC=compiler

substituting in the command you usually use to call the compiler.

If the compilation works properly, the executable chainBD.exe will appear in the main directory.

# 2 Usage Instructions

To run the program in the main directory, type:

./networkdiffusion.exe suffix > outputfile.out

Here, suffix can be any string up to 100 characters in length. The program reads in all input information from a file named param.suffix where, again, suffix is the command-line argument. If no argument is supplied, it will look for a file named param. If the desired parameter file does not exist, the program will exit with an error. You can supply multiple suffixes to read in multiple parameter files.

The parameters in the input file are given in the format "KEYWORD value" where the possible keywords and values are described in Section 4. Each keyword goes on a separate line. Any line that starts with "#" is treated as a comment and ignored. Any blank line is also ignored. The keywords in the parameter file are not case sensitive. For the most part, the order in which the keywords are given does not matter. All parameters have default values, so you need only specify keywords and values when you want to change something from the default.

## 3 Example for a Quick Start

### 3.1 Example 1

An example parameter file (param.example1) is provided. This will run a simulation of  $10^4$  particles propagating over a simple network, keeping track of the first passage time of each particle to each node. Run the example with

### ./networkdiffusion.exe example1

Use the script plotnetworkfpt.m to visualize the mean first passage time to each node on the network (where all particles start at node 1, shown in black).

The movie example1\_nodehit.avi shows the fraction of particles that have hit each node over time. Also made with the plotnetworkfpt.m script.



## 4 Keyword Index

The code will attempt to read parameters out of a file named param.suffix where "suffix" is the command line argument. If no command line arguments are supplied, it will look for a file named param. If multiple arguments are supplied, it will read multiple parameter files in sequence.

The parameter file should have one keyword per line and must end with a blank line. All blank lines and all lines beginning with # are ignored. For the most part, the order of the lines and the capitalization of the keywords does not matter. All keywords except ACTION are optional. The default values for each parameter are listed below. If a keyword is supplied, then values may or may not be needed as well. Again, the required and optional value types are listed below.

Keywords and multiple values are separated by spaces.

When reading the parameter file, lines longer than 500 characters will be truncated. To continue onto the next line, add "+++" at the end of the line to be continued. No individual keyword or value should be longer than 100 characters.

Floating point numbers can be formated as 1.0, 1.1D0, 10e - 1, -1.0E + 01, etc., where the exponential notation specifier must be D or E (case insensitive). Integer numbers can also be specified in exponential notation without decimal points (eg: 1000 or 1E3). Logical values can be specified as T, F, TRUE, FALSE, 1, or 0 (with 1 corresponding to true and 0 to false).

The length units used for the defaults are in nm and the energy units are in kT.

WARNING: documentation of keyword arguments is not finished. Look inside keys.f90 and readkey.f90 for the possible keywords

#### • ACTION

- value: 1 string of at most 20 characters; no default
- This keyword sets the overall calculation performed by the program (see Sec.??)
- Possible values are: GETFPT

### • OUTFILE

- value: 1 string, up to 100 characters; default: \*.out
- Output file for energy and coordinates of last bead.
- Any \* in the file name will be replaced by the command-line argument (suffix)

#### • RNGSEED

- 1 integer; default: 0
- seed for random number generator
- value of 0 will seed with system time in milliseconds
- value of -1 will use the last 5 characters in the suffix
- value of -2 will use the last 4 charactes in the suffix and the millisecond time
- other positive value: the seed is used directly for repeatable simulations (should be positive)

#### • SNAPSHOTS (not implemented yet)

- 1 optional integer, 1 optional string, 1 optional logical; defaults: 1, \*.snap.out, false
- Dump snapshots over the course of the calculation
- integer: how often to dump snapshots; string: snapshot file (\* is replaced with suffix); logical: append rather than rewriting the snapshot file

### • SNAPSHOTFILE (not implemented yet)

- value: 1 string; default: \*.snap.out
- File for dumping out snapshots. Can also be specified within SNAPSHOTS keyword.

#### VERBOSE

- value: 1 logical; default: false
- Print extra output. Not currently implemented