

# Logic Modeling

## Requirements

- numpy
- cython
- pypar

Can download files from github at <https://github.com/LoLab-VU/LogicalModel>  
Models directory contains 5 models.

```
Models/core_iron_6variables_3states.txt
Models/final_continuous_model_21_nodes.txt
Models/func-example.txt
Models/IRP2_overexp_functions_1.txt
Models/IRP2_overexp_functions_2.txt
Models/core_iron_6variables_3states.txt
```

## Example Usage

For this example we will use the core\_iron\_6variables\_3states.txt model. The main program is called main\_attractor\_synch\_cython.py. The option are

```
usage: main_attractor_synch_cython.py [-h] [-n NSTATES] [-s START] [-e END]usage:
main_attractor_synch_cython.py [-h] [-n NSTATES] [-s START] [-e END]
                                [-m MODEL] [-v VERBOSE] [-p PARALLEL]

optional arguments:
  -h, --help                show this help message and exit
  -n NSTATES, --nstates NSTATES
                            provide a number of states
  -s START, --start START
                            starting string to convert to base Nstates
  -e END, --end END         ending string to convert to based Nstates
  -m MODEL, --model MODEL
                            model to run simulation, assumes file to end in .txt
  -v VERBOSE, --verbose VERBOSE
                            if you want verbose updates (use with single
                                processor)
  -p PARALLEL, --parallel PARALLEL
                            run in parallel, use 0 or 1
```

The only required arguments are the number of states (-n) and the model file (-f)

```
python main_attractor_synch_cython.py -n 3 -m Model/core_iron_6variables_3states.txt
```

```
pinojc@LoLab-760:~/Projects/LogicalModel$ python main_attractor_synch_cython.py -n 3
-m Models/core_iron_6variables_3states.txt
Running on single CPU
Started
Computed 729 samples 0.0034 minutes
Attractors ['002211', '111111', '011111']
Frequencies [266, 29, 434]
Total 729
```

## Running on multiple processors/across nodes

To run on a multi core or cluster, you need to comment the main() (line 105) and uncomment all the multiprocessing lines below that. In python it is just the two lines that contain "", line 111 and 191.

Then you run using mpirun

```
mpirun -n #proc /path/to/python main_attractor_synch_cython.py -n 3 -p 1
```

```
pinojc@LoLab-760:~/Projects/LogicalModel$ mpirun -np 4 python
main_attractor_synch_cython.py -n 3 -m Models/core_iron_6variables_3states.txt -p 1
Processor 3 initialised on node LoLab-760
main_attractor_synch_cython.py:54: UserWarning: Shared object file
'core_iron_6variables_3states.so' already exists: moving on.
  warnings.warn(warning_string + 'moving on.')
Processor 1 initialised on node LoLab-760
Processor 0 initialised on node LoLab-760
samplesize = 728
split up into 72 segments
Processor 2 initialised on node LoLab-760
Computed 729 samples in 0.0012 minutes
Attractors ['002211', '111111', '011111']
Frequencies [266, 29, 434]
Total 729
```