DiPDE: A simulator for population density modeling

http://alleninstitute.github.io/dipde/

HBP CodeJam Workshop #7, January 11, 2016 Shrigley Hall Hotel, Manchester, England

Nicholas Cain Scientist 1

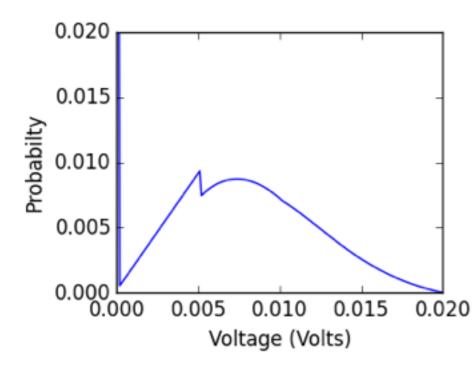


DiPDE (dipde) is a simulation platform for numerically solving the time evolution of coupled networks of neuronal populations. Instead of solving the subthreshold dynamics of individual model leaky-integrate-and-fire (LIF) neurons, dipde models the voltage distribution of a population of neurons with a single population density equation.

In this way, dipde can facilitate the fast exploration of mesoscale (population-level) network topologies, where large populations of neurons are treated as homogeneous with random fine-scale connectivity.

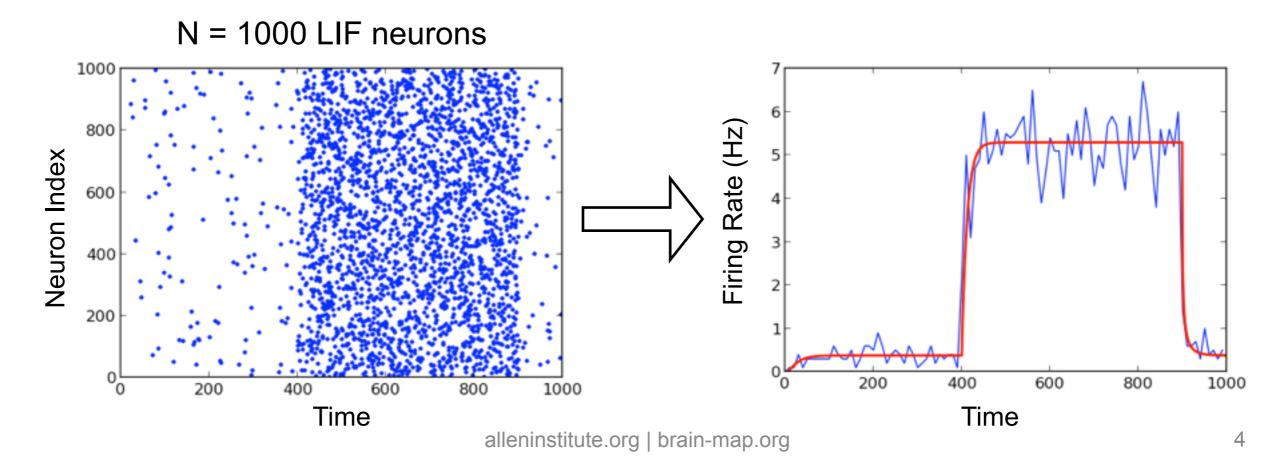
Goal: Provide a fast, flexible, python-based population statistic simulator for coarse-scale modeling

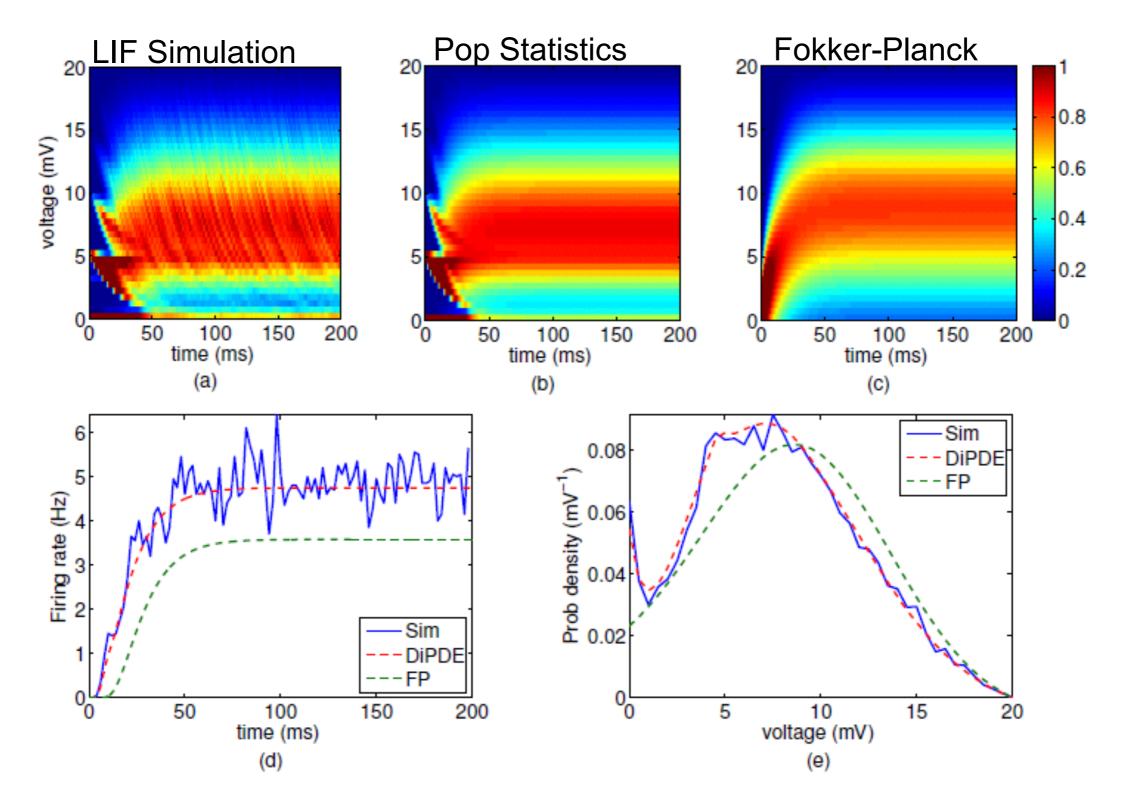
- Solve coupled population density equations (instantaneous synapses)
- Absorbing boundary condition at threshold
- Same mean firing rate as LIF population as N increases
- Exact when representing, ex., firing rate code





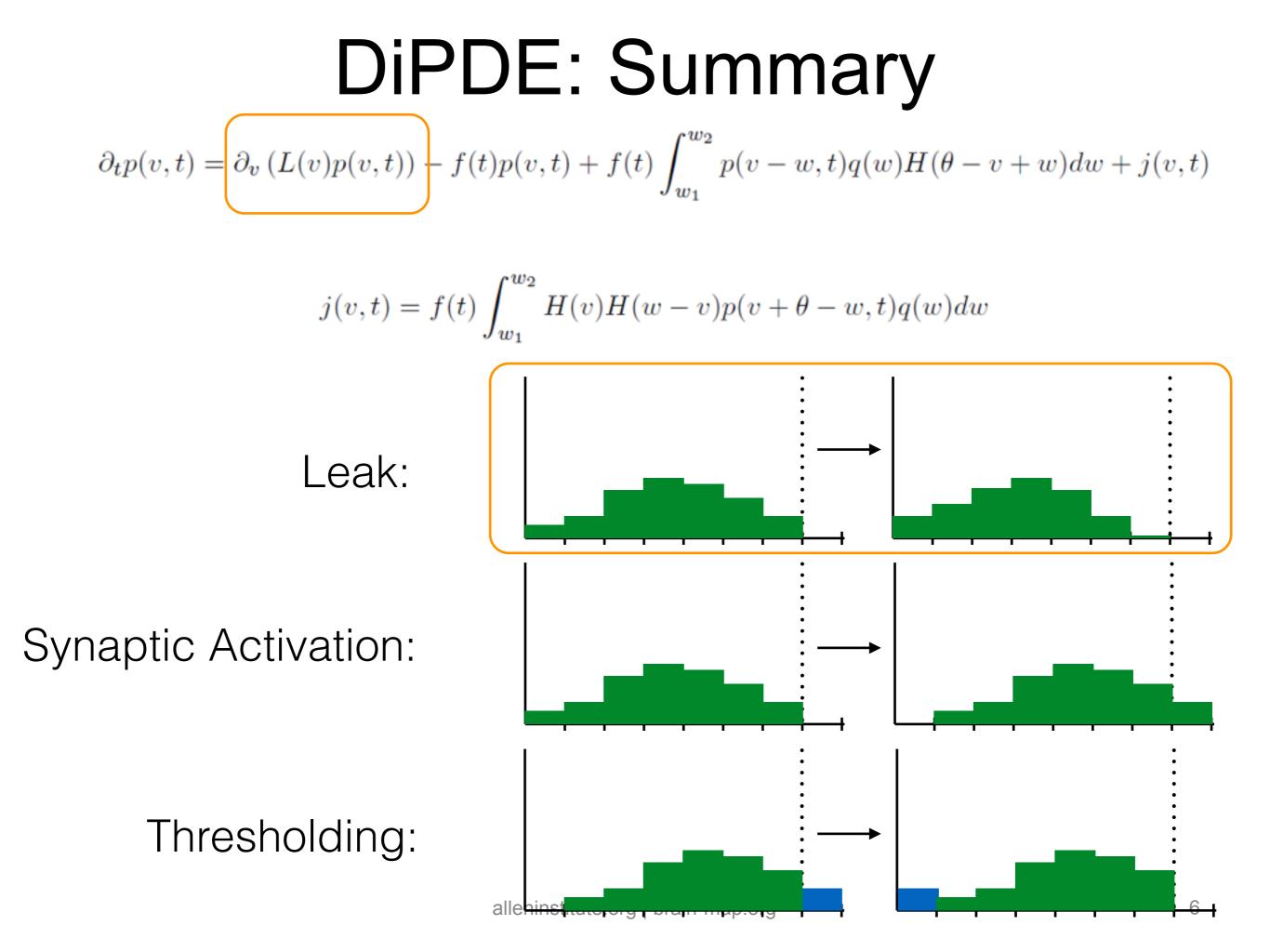
- Approximate mean firing rate of a LIF population
- Essentially a coupled PDE solver:
 - Boundary condition of source provides drive for target
 - Coupling through synaptic weight/delay distribution
- Allows fast:
 - Stability analysis
 - Stimulus/network topology/parameter exploration
 - Sensitivity analysis

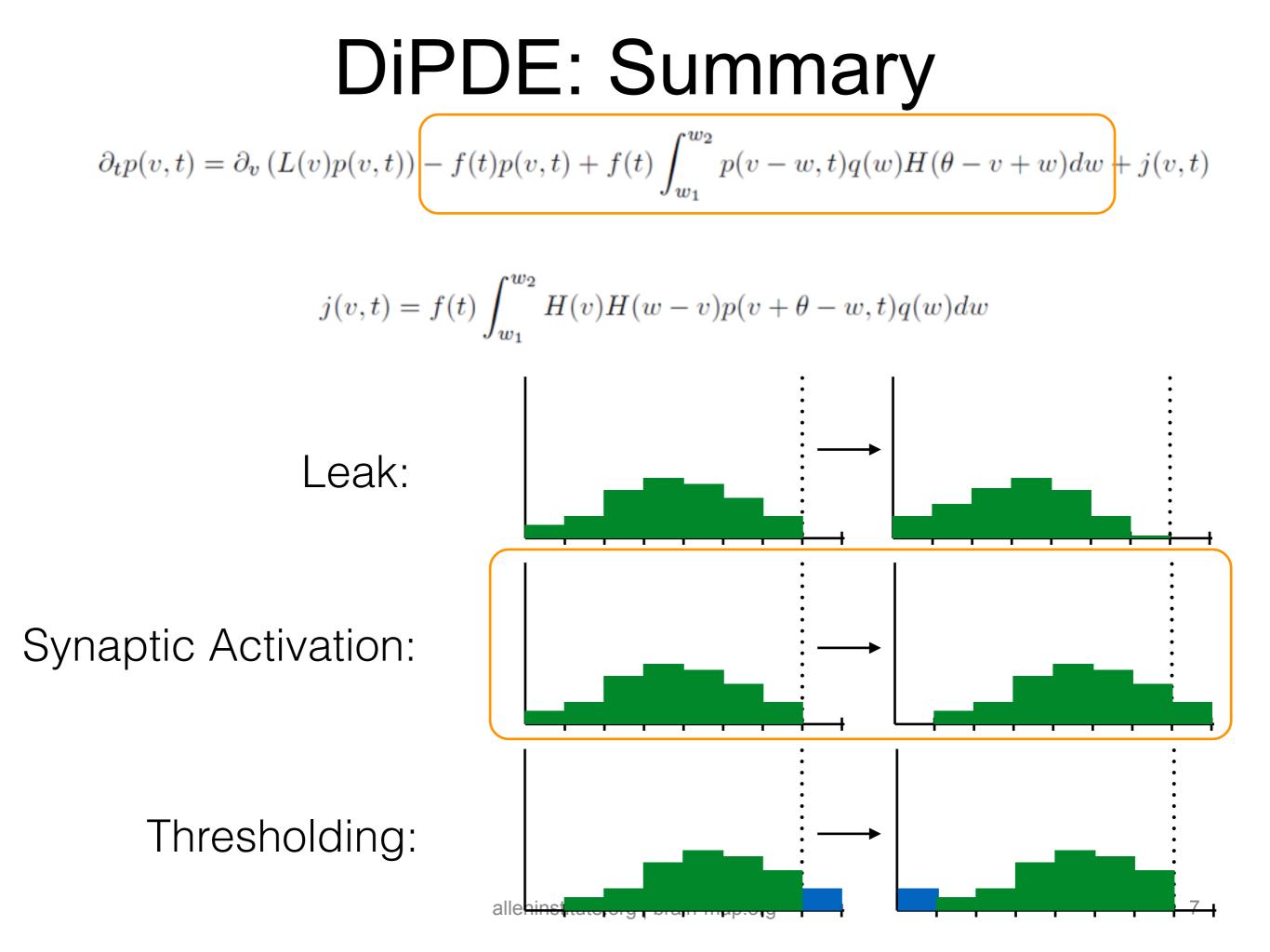




Iyer et al. PLoS Comp. Biol. 2013

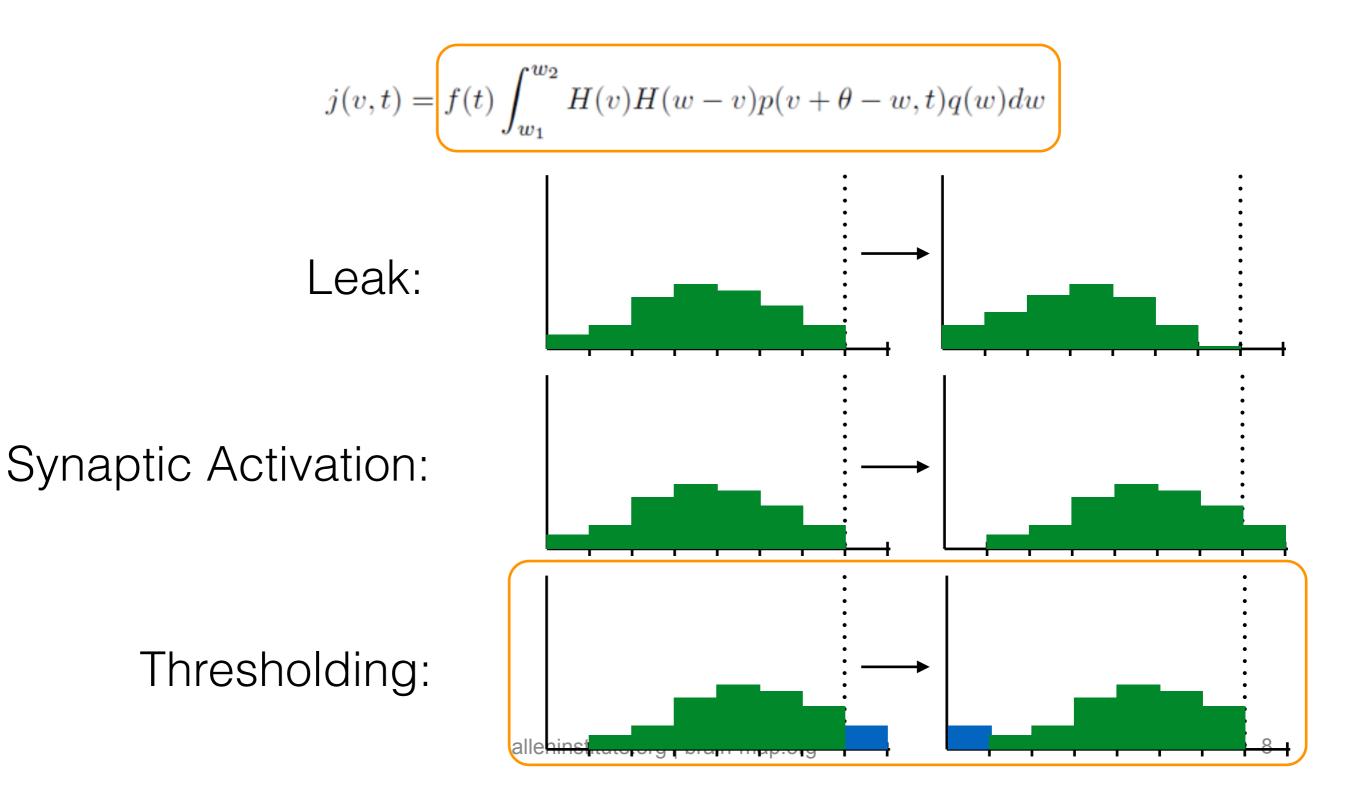
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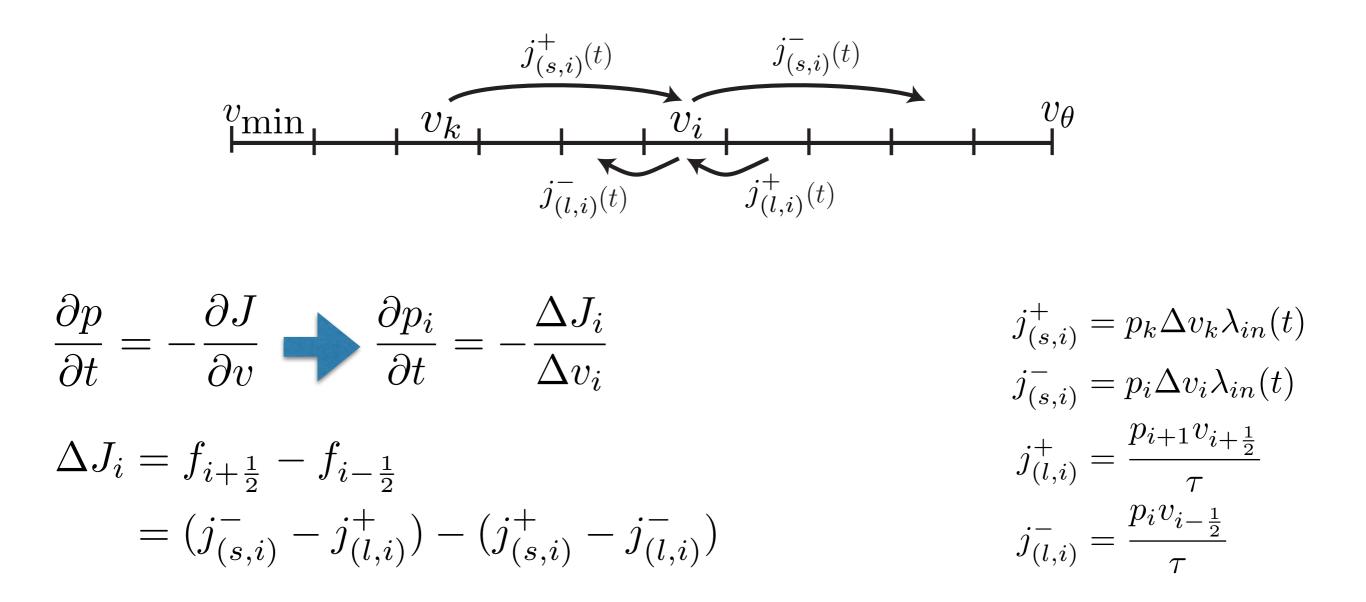
DiPDE: Summary

$$\partial_t p(v,t) = \partial_v \left(L(v) p(v,t) \right) - f(t) p(v,t) + f(t) \int_{w_1}^{w_2} p(v-w,t) q(w) H(\theta-v+w) dw + j(v,t)$$

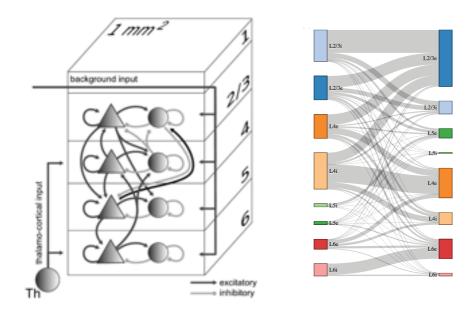


DiPDE: Summary

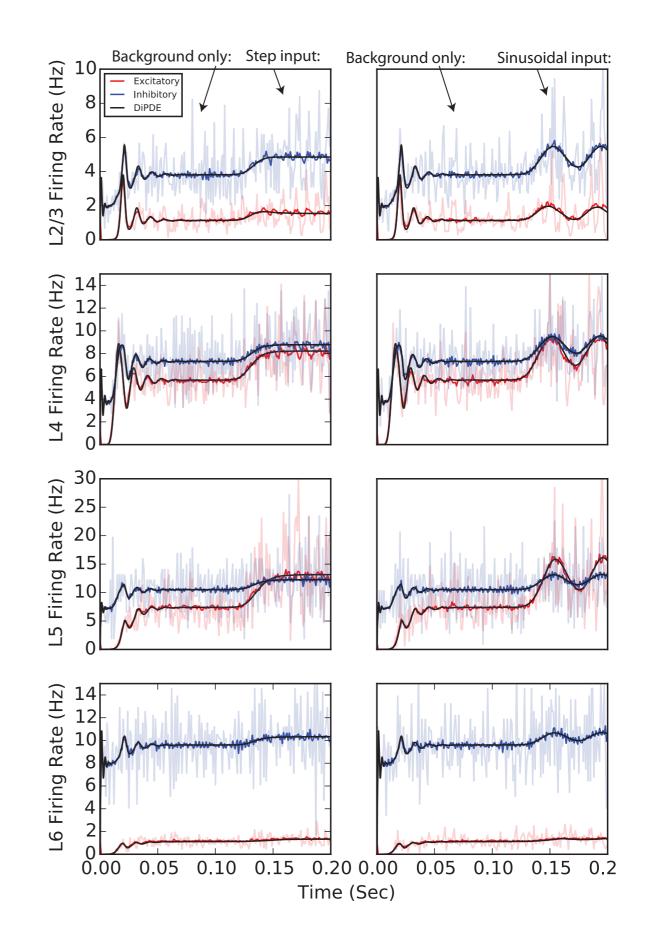
- Finite-volume method on the bounded interval
 - Assumes piecewise constant (over dt) recurrent drive
 - Evolve as a continuous time Markov chain



- DiPDE well-approximates a simplified cortical column
- Modified version of Potjans and Diesmann (2014)



- Plotted: averaged results from 100 LIF simulations (NEST)
- ~30 second run-time



Current Release:

- Current release: 0.1.0
 - <u>http://alleninstitute.github.io/dipde/</u>
 - 100% code coverage on tests
 - Pure python; only need numpy/scipy/sympy
- Tutorial features 5 simple examples:
 - <u>https://goo.gl/kZj2XN</u>

Documented Features:

- Populations:
 - External: Strings, sympy-functions, python functions
 - Internal: Variable dv/dt/time-step accuracy
- Connections:
 - Synaptic weight distributions
 - Discrete transmission delay distributions
- Simulation: run/pause/continue

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CONTENTS	WELCOME TO DIPO	E'S DOCUMENTATION			
User Guide	DAME (SING) is a similar	intice station for numerically polyton th	time qualities of counted automatic of	neuronal populations. Instant	and and and size
Quick Start Install Using Pip Regulared Dependencies	DPDE (dpdx) is a simulation platform for numerically solving the time evolution of coupled networks of neuronal populations. Instead of solving subthreshold dynamics of individual model leaky-integrate-and-fire (LIF) neurons, dpde models the voltage distribution of a population framma a single population dimative equation. In this way, dpde can facilitate the fast expiration of measurable (population-level) network topologies, wh				
Packages	large populations of neur	rons are treated as homogeneous with nar	don fine-scale connectivity.		
Examples (dipde.examples)	neurons. Beginning with	approach in computational neuroscience the work of Knight and Sirovich [3] (see age probability distribution receiving syna	also [2]), the approach typically formula	stes a partial integra-differe	ritial equation
Singlepop Singlepop	follow from the assumpt	tion of a leaky integrate-and fire model. of leaky integrate-and-fire neurons with i	We implement a numerical scheme for	computing the time evolution	
Singlepop (recurrent) Excitatory/Inhibitory	For a Quick Start, you o	an begin with the Examples Odpde.exam	ples). For a more complicated example,	check out the Cartical Cal	umn Model,
Singlepop (sine)	population example base	el on the Petjans and Diesmann (4) leaky	integrate and fire cortical column model	L	
Singlepop (exponential)	Details on the classes us	ed to define a dipde simulation can be fou	nd in Core functionality (dipole.internals)		
Core functionality (diple internals)	CONTENTS:				
External Population Internal Population	 User Guide 				
Connection	 Quick Start Required D 	Install Using Pa	6		
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	DIPDE TEAM				
	DIPDE is developed by the Modeling, Analysis and Theory group at the Alien Institute, with contributions from Nicholas Cain, Ram Iyer, Vilas Wi Michael Buice, Tim Files, Keth Godfrey, David Feng, and Sonten Wihalas.				
	INDICES AND TABLES				
	Index Module Index				
	Search Page				
	CITATIONS				
	[1] Knight, N.W., Hanin, D., & Sinovich, L. (1996) Dynamical models of interacting neuron populations in visual contex. Symposium on Robotics an				
	Catemetrics; Computational Engineering in Systems Application: 1–5. [2] Omuntag, A., Kright, B.W., B.Sinwich, L. (2005) On the Simulation of Large Populations of Neurons. Journal of Computational Neuroscience 8: 51–65.				
		A simple and stable numerical solution fi			
	[4] Patjans T.C., & Diesmann, H. (2014) The cell-type specific cortical microcircuit: relating structure and activity in a full scale spiking network model. Garderal Carter 24: 785–806.				
	DIPDE MANUSCRIPTS AND POSTERS				
	 Can, N., Isen, R., Koch, C., & Mihales, S. (2013) The computational properties of a simplified cortical column model. In Properation. (6) Can, N., Fisa, T., Manon, V., Iyer, R., Koch, C., & Mihalas, S. (2014) Simulations of the statistics of fring activity of neuronal populations in the Cano, N., Fisa, T., Manon, V., Iyer, R., Koch, C., & Mihalas, S. (2014) Simulations of the statistics of fring activity of neuronal populations in the Cano, N., Fisa, T., Manon, V., Iyer, R., Koch, C., & Mihalas, S. (2014) Simulations of the statistics of fring activity of neuronal populations in the Cano, N., Fisa, T., Manon, V., Iyer, R., Koch, C., & Mihalas, S. (2014) Simulations of the statistics of fring activity of neuronal populations in the Cano, N., Fisa, T., Manon, V., Iyer, R., Koch, C., & Mihalas, S. (2014) Simulations of the statistics of fring activity of neuronal populations in the Cano, N., Fisa, T., Manon, V., Iyer, R., Koch, C., & Mihalas, S. (2014) Simulations of the statistics of fring activity of neuronal populations in the Cano, N., Fisa, T., Manon, V., Iyer, R., Koch, C., & Mihalas, S. (2014) Simulations of the statistics of fring activity of neuronal populations in the Cano, S., Statistics, S. (2014) Simulations of the statistics of fring activity of neuronal populations of the statistics of fring activity of neuronal populations in the statistics of fring activity of neuronal populations of the statistics of fring activity of neuronal populations of the statistics of fring activity of neuronal populations of the statistics of the st				
	entire mouse brain, Program No. 100.02/0018. 2013 Neuroscience Heeting Planner, Washington, DC: Society for Neuroscience, 2014. Online. [7] Iger, R., Honen, V., Bulce, M., Koch, C., & Hhalins, S. (2015). The Influence of Spragtic Weight Distribution on Neuronal Population Dynamics.				
	PLIS Computational Biology, 9(10), e1093148. doi: 10.1371/jburnal.pdbi.2003148 [8] Iyer, R., Cain, N., & Mihalas, S. (2014). Dynamics of excitatory-inhibitory neuronal networks with exponential synaptic weight. Cosyne Abstract 2014. See Lake Circ ViSA.				
	CITING DIPDE				
	Website: © 2015 Allen Institute for Brain Science. DIPDE Simulator (Internet), Available from: https://pithub.com/AllenInstitute/dipole.				
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Example: singlepop.ipynb

In [12]: import matplotlib.pyplot as plt
from dipde.internals.internalpopulation import InternalPopulation
from dipde.internals.externalpopulation import ExternalPopulation
from dipde.internals.connection import Simulation
from dipde.internals.connection import Connection as Connection
%matplotlib inline

In [13]: # Settings: t0 = 0. dt = .0001 dv = .0001 tf = .1 tol = 1e-14 verbose = False

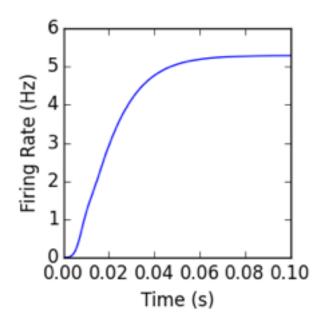
```
In [14]: # Create and run simulation:
bl = ExternalPopulation('100', record=True)
i1 = InternalPopulation(v_min=0, v_max=.02, dv=dv, tol=tol)
bl_i1 = Connection(b1, i1, 1, weights=[.005], probs=[1.], delay=0.0)
simulation = Simulation([b1, i1], [b1_i1], verbose=verbose)
simulation.run()
```

```
In [15]: # Visualize:
    i1 = simulation.population_list[1]
    plt.figure(figsize=(3,3))
    plt.plot(i1.t_record, i1.firing_rate_record)
    plt.xlim([0,tf])
    plt.ylim(ymin=0)
    plt.xlabel('Time (s)')
    plt.ylabel('Firing Rate (Hz)')
```

class Simulation(object):

def run(self):

```
for p in self.population_list:
    p.initialize()
for c in self.connection_list:
    c.initialize()
while self.t < self.tf:
    self.t += self.dt
    for p in self.population_list:
        p.update()
    for c in self.connection_list:
        c.update()</pre>
```

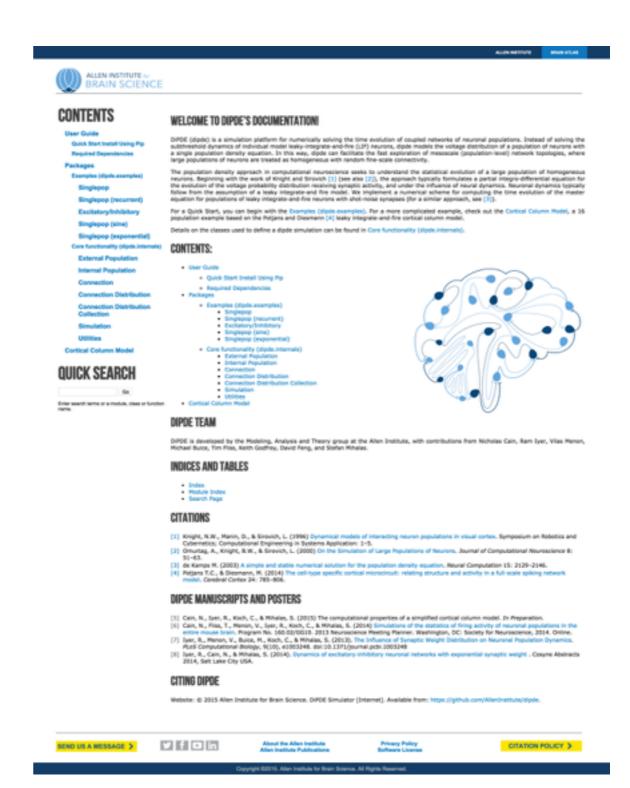


Under Development:

- Next release: 0.2.0 (March 2016)
 - https://github.com/nicain/dipde_dev

Features: (completed, debugged, in-progress)

- Distributions of:
 - Synaptic weights
 - Transmission delays
 - Membrane time-constants
- Simple Serialization (JSON)
- Flexible Interface (extend populations)
- ZMQ server/client inputs/outputs
- Run/pause/marshal/unmarshal/continue
- Callbacks on critical functions
- Logging, profiling
- Adapter to NEST/Brian/PyNN
- Algorithmic improvements:
 - Sparse storage
 - (2x-10x) speed up
- Prototype distributed version
- NWB export interface



Code Example: Distributions

 In 0.2.0, the following are all equivalent ways of specifying a connection distribution:

```
c = Connection(source, target, 1, weights=.005)
c = Connection(source, target, 1, weights=(.005, 1.))
c = Connection(source, target, 1, weights=sps.rv_discrete(values=(.005, 1.)))
c = Connection(source, target, 1, weights=sps.expon(0,.005))
c = Connection(source, target, 1, weights=(sps.expon(0,.005), 201))
c = Connection(source, target, 1, weights={'distribution': 'exponential', 'lambda':.005})
```

Code Example: Interface

• Basic interface to create (firing rate) populations

```
class PopulationInterface(object):
    '''Abstract Base Class for source populations'''
    def initialize(self):
        '''Override with behavior that sets an initial value'''
       self.set_curr_firing_rate(None)
    def update(self):
        '''Override with behavior that gets called once per time step'''
       logger.debug('GID(%s) Firing rate: %s' % (self.gid, self.curr_firing_rate))
    def set_curr_firing_rate(self, curr_firing_rate):
        '''Call to make "curr_firing_rate" visible to other populations.
        Typically invoked once at initialization, and once in update'''
       self._curr_firing_rate = curr_firing_rate
    @property
    def t(self): return self.simulation.t
    @property
    def dt(self): return self.simulation.dt
    @property
    def gid(self): return self.simulation.gid_dict[self]
    @property
    def curr_firing_rate(self): return self._curr_firing_rate
    @propertv
    def source_connection_list(self): return [c for c in self.simulation.connection_list if c.target == self]
    @property
    def source_firing_rate_dict(self):
       return dict((c.source.gid, self.simulation.get_curr_firing_rate(c.source.gid)) for c in self.source_connection_list)
```

Code Example: ZMQ REQ/REP Servers

• Callable that can be used as the firing_rate arg of an ExternalPopulation

```
class RequestFiringRate(object):
   def __init__(self, port):
        self.port = port
        self.socket = context.socket(zmg.REQ)
        self.socket.connect("tcp://localhost:%s" % port)
   def __call__(self, t):
        self.socket.send('%s' % t)
        message = self.socket.recv_multipart()
        return float(message[0])
class ReplyFiringRateServer(object):
   def __init__(self, port, reply_function):
        self.port = port
        self.reply_function = reply_function
        self.socket = context.socket(zmq.REP)
        self.socket.bind("tcp://*:%s" % self.port)
   def run(self):
        while True:
            message = self.socket.recv()
            if message == 'SHUTDOWN':
                break
            requested_t = float(message)
            self.socket.send_multipart([b"%s" % self.reply_function(requested_t)])
        self.socket.send('DOWN')
```

Code Example: NEST Adapter

Construct an analogous NEST simulation from a dipde simulation

```
def get_kernel(dt=.0001, tf=.1, seed=None, number_of_processors=2, verbose=True):
    if seed is None: seed = np.random.randint(1,100000)
    import nest as kernel
    kernel.ResetKernel()
    kernel.SetKernelStatus({"local_num_threads": number_of_processors})
    N_vp = kernel.GetKernelStatus(['total_num_virtual_procs'])[0]
    kernel.SetKernelStatus({'grng_seed' : seed+N_vp})
    kernel.SetKernelStatus({'rng_seeds' : range(seed+N_vp+1, seed+2*N_vp+1)})
    kernel.SetKernelStatus({"resolution": dt*1000, "print_time": verbose})
    return kernel
class PoissonPopulation(object):
    def __init__(self, name, firing_rate, number_of_neurons, kernel, start=0.):
        self.name = name
        self.firing_rate = firing_rate
        self.number_of_neurons = number_of_neurons
        self.gids = kernel.Create("poisson_generator", number_of_neurons, params={"rate": float(firing_rate),
                                                                                  'start':float(start)/.001})
class IAFPSCDeltaPopulation(object):
    def __init__(self, name, number_of_neurons, kernel, tau_refrac=0.):
        self.name = name
        if tau_refrac == 0.: tau_refrac = kernel.GetKernelStatus("resolution")/1000
                              "V_reset"
                                                                            : 10..
        curr_neuron_params= {
                                            : 0.,
                                                                "tau_m"
                                "C_m"
                                            : 250..
                                                                "V_th"
                                                                            : 15..
                                           : tau_refrac*1000, "V_m"
                                "t_ref"
                                                                            : 0..
                                "E L"
                                            : 0.1
        self.gids = kernel.Create("iaf_psc_delta", number_of_neurons, params=curr_neuron_params)
```

Goals for This CodeJam:

Meet as many new people and technologies as I can
 Have fun writing code with all of you

- 3. Get feedback on the technical approaches I am taking
- 4. Help anyone who is interested to learn more about dipde
- 5. Work in interfacing any/all AIBS code and data formats with community standards. (I do more than just dipde)
- 6. Work on model construction tools, and description formats/adapters
- 7. Interface dipde with any relevant visualization tools
- 8. Get help prioritizing features for the future dipde

THANKS!

THANK YOU

We wish to thank the Allen Institute for Brain Science founders, Paul G. Allen and Jody Allen, for their vision, encouragement and support.

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