

An introduction to NineML

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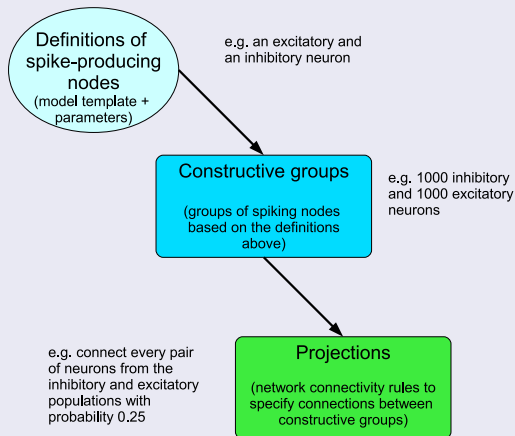


Background

- NineML is a domain-specific language for describing neuronal network models.
- Joint effort of developers of neuroscience simulator software and simulator-independent description language initiatives (NeuroML, PyNN).
- The NineML effort is initiated and coordinated by the International Neuroinformatics Coordinating Facility (INCF).

Model structure of NineML

Model



A model description consists of:

- Neuronal dynamics template
- List of parameter values
- Groups of neurons—populations, layers, etc.
- Connectivity patterns—projections, connection probabilities, etc.

Spike-producing node definition

Example

The user starts out by choosing a model template:

Model template: `IaF_gL`
(supplied by model library)

$$C_m \frac{dV}{dt} = -g_L V + \sum_i I_i$$

$$\text{if } V > \theta : \begin{cases} \text{while } t < t_{spike} + t_{refr} \\ V = V_{rest} \end{cases}$$

Having chosen a model template, the user can then provide parameter values:

Parameter description
(supplied by user)

$$C_m = 1\mu F \quad g_L = 0.5mS$$

$$\theta = 20mV \quad V_{rest} = -60mV$$

$$t_{refr} = 0.1ms$$

Spike-producing node definition

Example

Model template: `IaF_gL`

(supplied by model library)

$$\textcircled{C_m} \frac{dV}{dt} = -g_L V + \sum_i I_i$$

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Spike-producing node definition

Example

```
<spikingNode name="E">  
  
  <definition>  
    <type>laF_gL</type>  
    <link><url>http://www.nineml.org/1.0/laF_gL</url></link>  
  </definition>  
  
  <parameters>  
    <membraneCapacitance name="C_m">...  
      <value>1</value>  
      <unit>uF</unit>  
    </membraneCapacitance>  
    <threshold>...</threshold>  
    <refractoryTime>...</refractoryTime>  
    <resetPotential>...</resetPotential>  
  </parameters>
```

Spike-producing node definition

Example

```
<spikingNode name="E">
```

```
  <definition>
```

```
    <type>laF_gL</type>
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```
    <link><url>http://www.nineml.org/1.0/laF_gL</url></link>
```

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  </definition>
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```
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    <membraneCapacitance name="C_m">...
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```
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```
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```
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  </parameters>
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Spike-producing node definition

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  </parameters>
```


User Layer and Abstraction Layer

User Layer

Parameter description

(supplied by user)

$$C_m = 1\mu F \quad g_L = 0.5mS$$

$$\theta = 20mV \quad V_{rest} = -60mV$$

$$t_{refr} = 0.1ms$$

Abstraction Layer

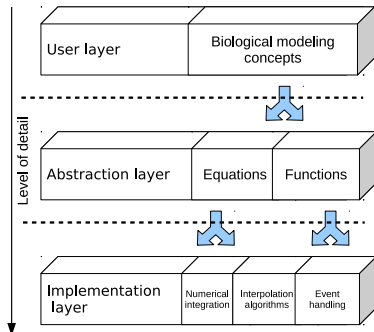
Model template: I_{aF_gL}

(supplied by model library)

$$C_m \frac{dV}{dt} = -g_L V + \sum_i I_i$$

$$\text{if } V > \theta : \begin{cases} \text{while } t < t_{spike} + t_{refr} \\ V = V_{rest} \end{cases}$$

The Layer View



Parameters for model **IaF_gL**

$$C_m = 1\mu F \quad g_L = 0.5mS$$

$$\theta = 20mV \quad V_{rest} = -60mV$$

$$t_{refr} = 0.1ms$$

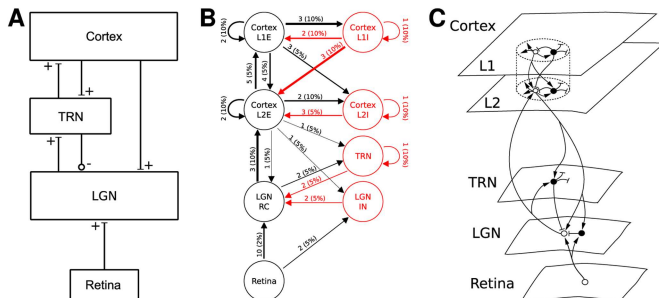
Model template: **IaF_gL**

$$C_m \frac{dV}{dt} = -g_L V + \sum_i I_i$$

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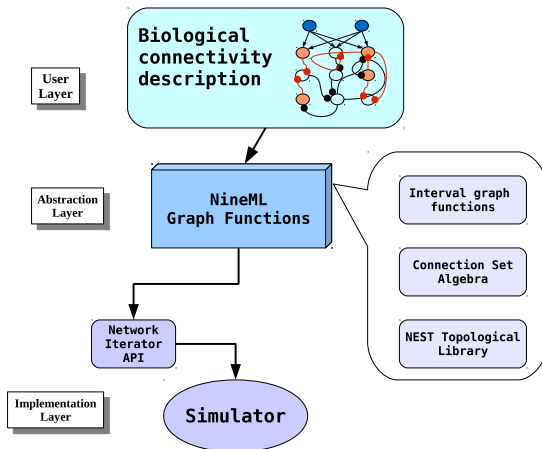
- In the user layer, we characterize biological concepts by means of a model name and a set of parameter values.
- In the abstraction layer, we define parameterized equations that characterize a particular class of models.

Representations of Network Connectivity

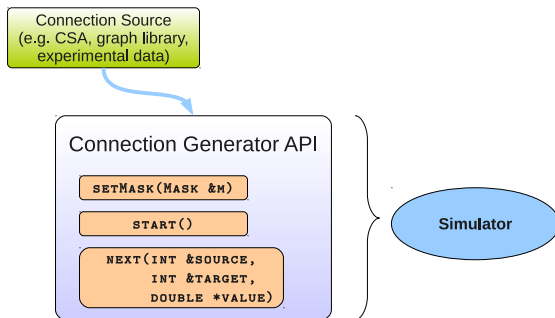


Nordlie, E et al. *Towards reproducible descriptions of neuronal network models.*, PLoS Comput Biol (2009).

NineML Connectivity Overview



Connection Generator Interface



Masks and Interval Sets

```
class Mask
{
  IntervalSet sources;
  IntervalSet targets;
}
```

```
class IntervalSet
{
  std::vector<ClosedInterval> ivals;
}
```

Network Connectivity Example

**Create index intervals for excitatory,
inhibitory and all cells**

```
binding e = Interval.closed_interval (0, 599)
```

```
binding i = Interval.closed_interval (600, 899)
```

```
binding a = Interval.union (e, i)
```

Network Connectivity Example

Create geometry function g and metric d

```
binding g = CSA.random2d (900)
```

```
binding d = CSA.euclidMetric2d (g)
```

Network Connectivity Example

**Excitatory and inhibitory gaussian value sets
(gaussian of the distance for every index pair)**

```
binding g_e = CSA.gaussian (0.1, 0.3) (d)
```

```
binding g_i = CSA.gaussian (0.2, 0.3) (d)
```


Network Connectivity Example

Create connection-sets with gaussian dependent random masks, gaussian dependent conductance and distance dependent delay

```
binding c_e = CSA.cset (CSA.random (g_e), g_e, d)
```

```
binding c_i = CSA.cset (CSA.random (g_i), -g_i, d)
```

Network Connectivity Example

Combine excitatory and inhibitory connectivity into one network using cartesian and multiset sum operators

```
binding c = CSA.sum (CSA.cartesian (e, a) (c_e),  
                    CSA.cartesian (i, a) (c_i))
```

Network Connectivity Example

Create index intervals for excitatory, inhibitory and all cells

```
binding e = Interval.closed_interval (0, 599)
binding i = Interval.closed_interval (600, 899)
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Excitatory and inhibitory gaussian value sets
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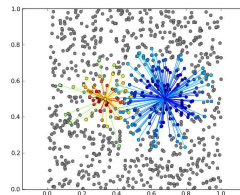
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Combine excitatory and inhibitory connectivity into
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Home page

`http://software.incf.org/software/nineml`

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NineML

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NineML: APIs and documentation

Purpose
 NineML is a simulator independent language with the aim of providing an unambiguous description of neuronal network models for efficient model sharing and reusability.
 This language has emerged from a joint effort of experts in the fields of computational neuroscience, simulator development and simulator-independent language initiatives ([NeuroML](#), [PyNN](#)), grouped in the [INCf Multiscale Modeling Task Force](#).
 This effort was initiated and is coordinated and supported by the International Neuroinformatics Coordinating Facility (INCf), as part of the

Diagram: A central box labeled 'NineML' is connected by double-headed arrows to three surrounding boxes: 'NEURON' (top), 'NEST' (right), and 'NeuroML /LEMS' (bottom).

Topics
[Large scale modeling](#)
[Computational neuroscience](#)

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Why the Need for Layers?

Let us consider a non-layered solution, e.g. PYNN or NEUROML:

A non-layered solution for model standardization

Standard model library

IaF_gL

$$\begin{aligned}C_m &= 1\mu F & g_L &= 0.5mS \\ \theta &= 20mV & V_{rest} &= -60mV \\ t_{refr} &= 0.1ms\end{aligned}$$

NEURON

NEST

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QIaF

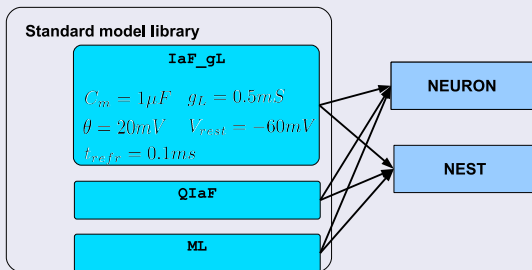
NEURON

NEST

Why the Need for Layers?

Let us consider a non-layered solution, e.g. PYNN or NEUROML:

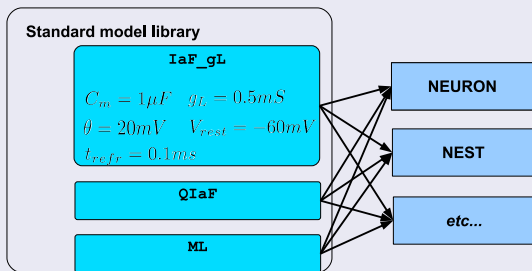
A non-layered solution for model standardization



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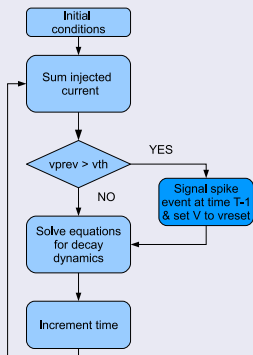
Problems with the non-layered approach:

- Potential duplication of implementation effort.
- Superficial model equivalence.
- Insufficiently flexible for development of new models.

Why the Need for Layers?

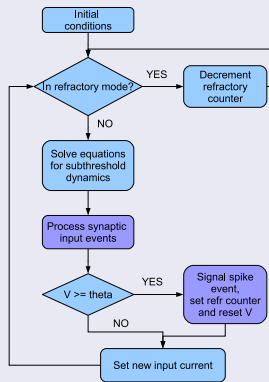
The problem of porting models between simulators:

Generalized Integrate-and-Fire



Pillow, JW, Paninski, L., et al. (2005): Prediction and Decoding of Retinal Ganglion Cell Responses with a Probabilistic Spiking Model. *Journal of Neuroscience* 25:11003-11013.

NEST extension module



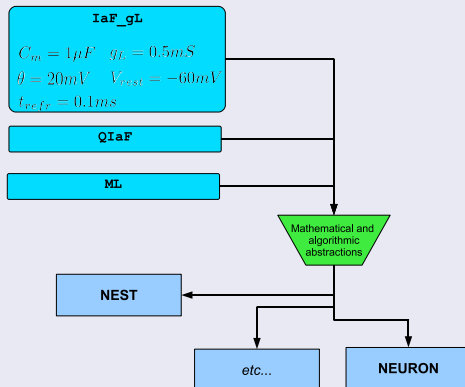
Writing an Extension Module with NEST.

http://www.nest-initiative.org/index.php/Writing_an_Extension_Module

Why the Need for Layers?

The answer: flexibility, ease of porting models, establishing true model equivalence.

A layered solution for model standardization



We've Got Layers, Now What?

- What should the abstraction layer for networks of integrate-and-fire models look like?

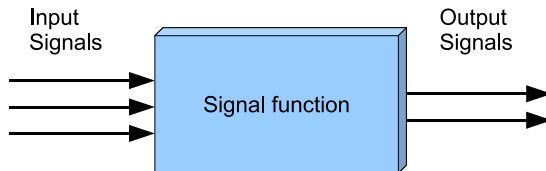
We've Got Layers, Now What?

- What should the abstraction layer for networks of integrate-and-fire models look like?
- Such an abstraction layer must:
 - provide enough flexibility for a variety of existing models
 - capture mathematical and algorithmic model properties required to achieve reproducibility of results
 - provide extensibility for future model development

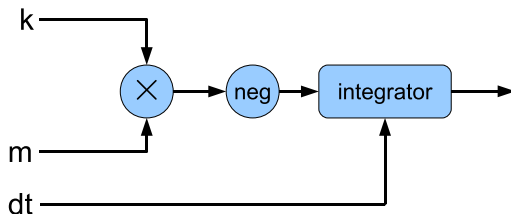
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- What should the abstraction layer for networks of integrate-and-fire models look like?
- Such an abstraction layer must:
 - provide enough flexibility for a variety of existing models
 - capture mathematical and algorithmic model properties required to achieve reproducibility of results
 - provide extensibility for future model development
- Proposal: an abstraction layer that can represent:
 - event handling
 - state transitions
 - differential and algebraic equations
 - other equation types: chemical reaction equations, etc.

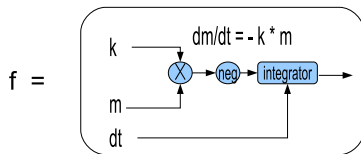
Signal Functions



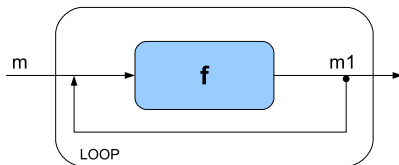
$$dm/dt = -k * m$$



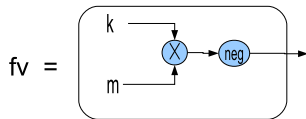
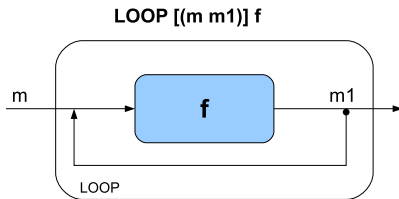
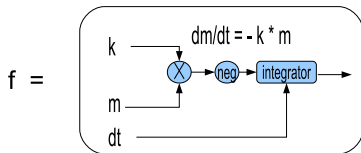
Signal Function Combinators



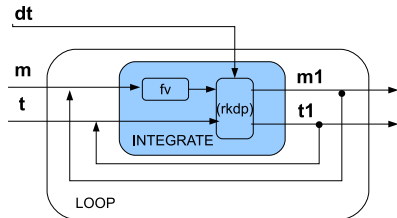
LOOP [(m m1)] f



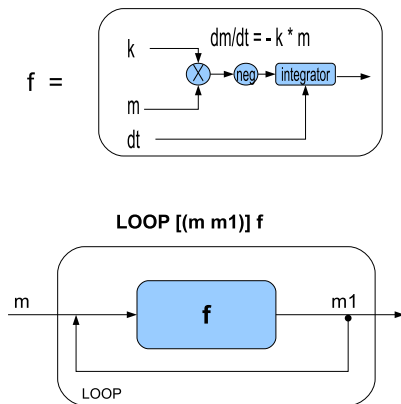
Signal Function Combinators



LOOP [(m m1) (t t1)]
INTEGRATE rkdp [(m m1) (t t1) dt] fv



Signal Function Combinators



Advantages of signal function combinators:

- precise control over the structure of the abstraction layer
- much flexibility in building the mathematical description of a model
- efficient executable code from model specification

Signal Function Combinators: Example 1

```

DEFINE dv (v,w, Istim , gl , vl , v1 , v2 , gca , vca , gk , vk , c)
  (Istim + (gl * (vl - v)) +
    ica(v,gca,vca,v1,v2) +
    ik(v,w,gk,vk)) / c

```

```

DEFINE dw (w,v, phi , v3 , v4)
  lamw (v, phi , v3 , v4) * (winf (v,v3,v4) - w)

```

```

DEFINE Morris-Lecar
  LOOP [(v nv) (w nw) (t nt)]
    SEQ
      INTEGRATE [(v nv) (t nt) tstep]
        SENSE [v w Istim gl vl v1 v2 gca vca gk vk c]
        PURE (dv) dv
      INTEGRATE [(w nw) (t nt) tstep]
        SENSE (w v phi v3 v4)
        PURE (dw) dw

```

Signal Function Combinators: Example 1

```

DEFINE dv (v,w, Istim , gl , vl , v1 , v2 , gca , vca , gk , vk , c)
  ( Istim + ( gl * ( vl - v ) ) +
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    ik ( v , w , gk , vk ) ) / c

```

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DEFINE dw (w,v, phi , v3 , v4)
  lamw ( v , phi , v3 , v4 ) * ( winf ( v , v3 , v4 ) - w )

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    PURE (dv) dv

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  INTEGRATE [(w nw) (t nt) tstep]
    SENSE (w v phi v3 v4)
    PURE (dw) dw

```

Signal Function Combinators: Example 1

```

DEFINE dv (v,w, lstim , gl , vl , v1 , v2 , gca , vca , gk , vk , c)
  ( lstim + ( gl * ( vl - v ) ) +
    ica ( v , gca , vca , v1 , v2 ) +
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```

```

DEFINE dw (w,v, phi , v3 , v4)
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```

```

INTEGRATE [(w nw) (t nt) tstep]
  SENSE (w v phi v3 v4)
  PURE (dw) dw

```

Signal Function Combinators: Example 2

```
DEFINE Hansel:98
  LOOP [(v nv) (t nt)]
    RSWITCH [spike! recur]
    SEQ
      INTEGRATE rkdp [(v nv) (t nt) tstep]
      SENSE [v t spikes Istim c vl gl gsyn ve tau1 tau2]
      PURE [dv] dv
      ACTUATE [spike!]
      SENSE [nv nt theta]
      PURE [spike!] spike?
    ACTUATE [v recur]
    SENSE [Vrest]
    PURE/list [recur] refr
```

Signal Function Combinators: Example 2

```
DEFINE Hansel:98  
  LOOP [(v nv) (t nt)]
```

```
    RSWITCH [spike! recur]
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```
      SEQ  
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Signal Function Combinators: Example 2

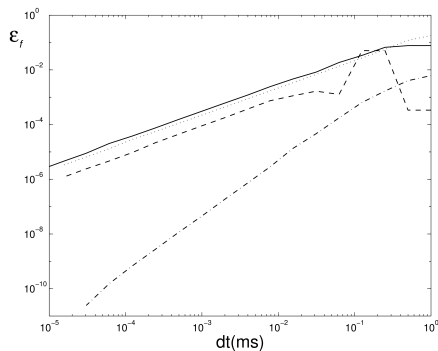
```
DEFINE Isyn (v,t,spikes,gsyn,ve,tau1 tau2)
  (gsyn - (v - ve)) *
  (vector-fold Isynf 0.0 spikes)
```

```
DEFINE dv (v,t,spikes,lstim,c,vl,gl,gsyn,ve,tau1,tau2)
  (((neg gl) * (- v vl)) + Isyn (v,t,spikes,gsyn,ve,tau1,tau2)) / c
```

```
DEFINE refr (Vrest)
  (list Vrest 1)
```

```
DEFINE spike? (v,t,theta)
  if (>= v theta)
  then t
  else undefined-signal
```

Numerical Issues



- Standard Euler
- Standard Euler + interpolation
- - - RK2
- . - . RK2 + interpolation

$$\epsilon_f = \left| \frac{\hat{f} - f}{f} \right|$$

Hansel D, Mato G, Meunier C, Neltner L (1998): On numerical simulations of integrate-and-fire neural networks. *Neural Computation* 10:467–483.