

Modelling the Population Level and Beyond

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University of Leeds

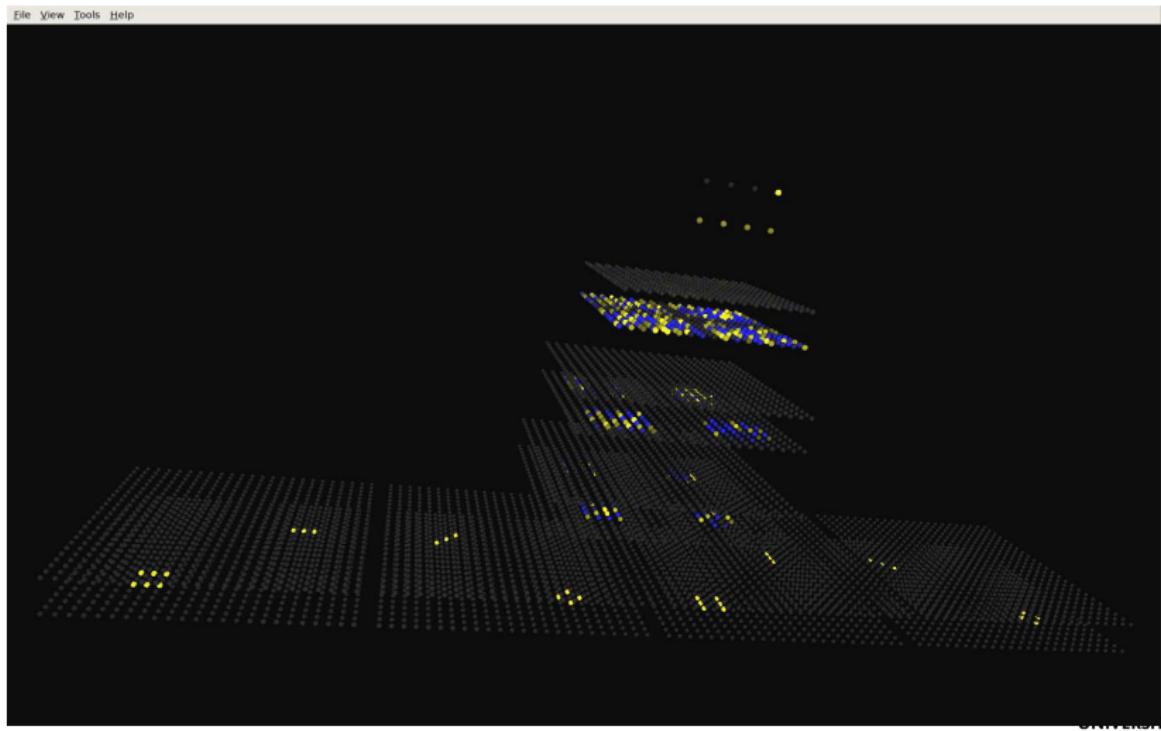
14th March, 2012

Funding for the Neurosciences

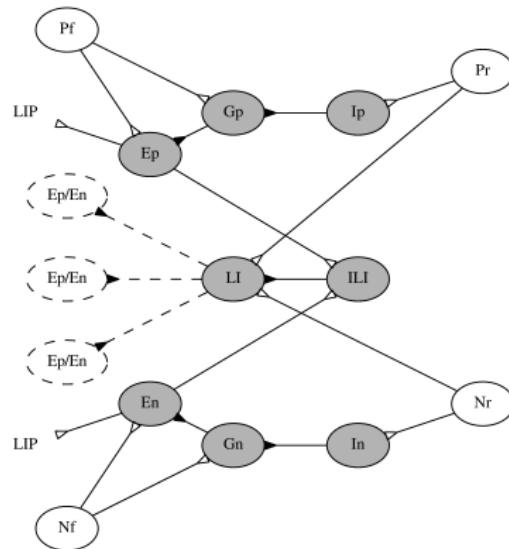
Three drivers:

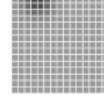
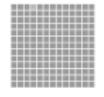
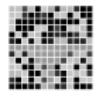
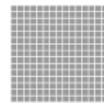
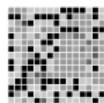
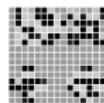
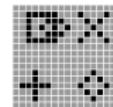
- Health related
- Brain-inspired technology
- Understanding brain/mi(i)nd

Spatial Structure



Local Circuit

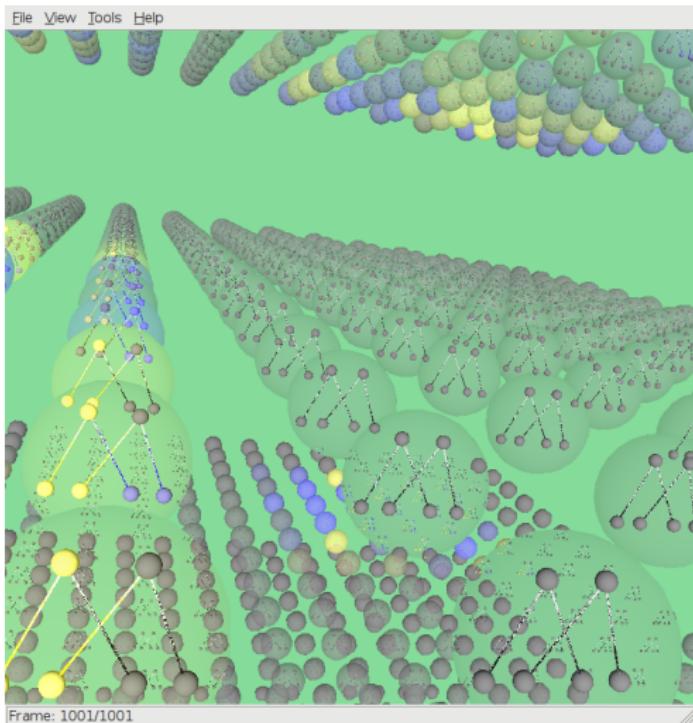




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Circuits



Outline

1 Model of Visual Attention

2 CLAMVis

- CLAMVis XML Structure

3 MIIND

CLAMVis Top Level Elements

```
<CLAMVisProject name="XMLTest" xsi:noNamespaceSchemaLocation="CLAMVisProject.xsd">
  <ProjectDescription> Description of Project </ProjectDescription>
  +<LayerDescriptions></LayerDescriptions>
  +<Networks></Networks>
  +<CircuitDescriptions></CircuitDescriptions>
  +<DynamicNetworks></DynamicNetworks>
  +<Mappers></Mappers>
  +<Simulations></Simulations>
</CLAMVisProject>
```

Layer Descriptions and Networks

```

<LayerDescriptions>
  <LayerDescription name="XML_V1" x_pixels="32" y_pixels="32" features="4"/>
  <LayerDescription name="XML_V2" x_pixels="30" y_pixels="30" x_field="3" y_field="3"/>
  <LayerDescription name="XML_V4" x_pixels="28" y_pixels="28" x_field="3" y_field="3"/>
  <LayerDescription name="XML_PIT" x_pixels="26" y_pixels="26" x_field="3" y_field="3"/>
  <LayerDescription name="XML_AIT" x_pixels="1" y_pixels="1" features="6" x_field="26"
    y_field="26"/>
  <LayerDescription name="LIP" x_pixels="32" y_pixels="32"/>
</LayerDescriptions>
<Networks>
- <Network name="ANN_FWD" filename="xml_all_fwd.net">
  <Layers>
    <Layer>XML_V1</Layer>
    <Layer>XML_V2</Layer>
    <Layer>XML_V4</Layer>
    <Layer>XML_PIT</Layer>
    <Layer>XML_AIT</Layer>
  </Layers>
  <LinkRelation>DenseOverlapLinkRelation</LinkRelation>
  <TrainingParameters bias="0.01" momentum="0.0" sigma="0.25" stepsize="0.2"
    train_threshold="false" train_threshold_value="1e-5" init="true"
    training_set="xml_all.trainingset" noise_level="0.005" noise_strength="0.5"/>
</Network>
+ <Network name="ANN_REV" filename="xml_all_rev.net" reverse="ANN_FWD" type="Hebbian">
  </Network>
</Networks>

```

Circuit Descriptions

```
<CircuitDescriptions>
  -<CircuitDescription name="perceptron" external="e_p" number_of_nodes="6">
    <CircuitNodeRole name="e_p" type="exc" x_pos="3.0" y_pos="0.0" z_pos="-2.0"/>
    <CircuitNodeRole name="i_p" type="inh" x_pos="-3.0" y_pos="0.0" z_pos="-2.0"/>
    <CircuitNodeRole name="e_n" type="exc" x_pos="1.0" y_pos="0.0" z_pos="-2.0"/>
    <CircuitNodeRole name="i_n" type="inh" x_pos="-1.0" y_pos="0.0" z_pos="-2.0"/>
    <CircuitNodeRole name="P_OUT" type="exc" x_pos="2.5" y_pos="0.0" z_pos="2.0"
      isOutput="true"/>
    <CircuitNodeRole name="N_OUT" type="exc" x_pos="-2.5" y_pos="0.0" z_pos="2.0"
      isOutput="true" isNegative="true"/>
    <Connection from="i_p" to="N_OUT" weight="-2.0"/>
    <Connection from="e_p" to="P_OUT" weight="2.0"/>
    <Connection from="i_n" to="P_OUT" weight="-2.0"/>
    <Connection from="e_n" to="N_OUT" weight="2.0"/>
  </CircuitDescription>
  -<CircuitDescription name="lip_circuit" external="lipNode" number_of_nodes="1">
    <CircuitNodeRole name="lipNode" type="exc" x_pos="0.0" y_pos="0.0" z_pos="0.0"
      isOutput="true"/>
  </CircuitDescription>
</CircuitDescriptions>
```



Dynamic Networks

```
<DynamicNetworks>
  -<Parameters>
    <WilsonCowanParameter name="exc" time_membrane="20e-3" rate_maximum="1.0"
      noise="1.0"/>
    <WilsonCowanParameter name="inh" inhibitory="true" time_membrane="10e-3"
      rate_maximum="1.0" noise="1.0"/>
  </Parameters>
  -<DynamicNetwork name="fwd" ANN="ANN_FWD" exc_param="exc" inh_param="inh">
    <CircuitCreator type="Perceptron" circuitDescription="perceptron"/>
    -<Layers>
      <Layer>XML_V1</Layer>
      <Layer>XML_V2</Layer>
      <Layer>XML_V4</Layer>
      <Layer>XML_PIT</Layer>
      <Layer>XML_AIT</Layer>
    </Layers>
  </DynamicNetwork>
</DynamicNetworks>
<Mappers>
  -<Mapper type="GaussianLayerNodeMapper" from_network="fwd" to_network="disinhibition"
    from_layer="1" to_layer="0" sigma="0.1" strength="1.0">
    <Mapping from_id="P_OUT" to_id="i_gat_p" from_feature="0" to_feature="0"/>
    <Mapping from_id="P_OUT" to_id="e_dis_p" from_feature="0" to_feature="0"/>
    <Mapping from_id="N_OUT" to_id="i_gat_n" from_feature="0" to_feature="0"/>
    <Mapping from_id="N_OUT" to_id="e_dis_n" from_feature="0" to_feature="0"/>
  </Mapper>
</Mappers>
```

Simulations

```
<Simulations>
--<Simulation name="Simulation">
  <SimulationDescription>Description of Simulation</SimulationDescription>
  --<SimulationPatterns>
    <SimulationPattern network="fwd">0 1 2 3</SimulationPattern>
    <SimulationPattern network="rev" time="0.5">3</SimulationPattern>
  </SimulationPatterns>
  <SimulationRunParameter max_iterations="100000000" start_time="0.0" end_time="1.0"
    report_time="1e-1" update_time="1e-2" network_step_time="1e-3" feedback_rate="0.35"
    feedback_time="0.5" progress_file_name="progress.txt"
    output_file_name="simulation_results.root"/>
</Simulation>
</Simulations>
```



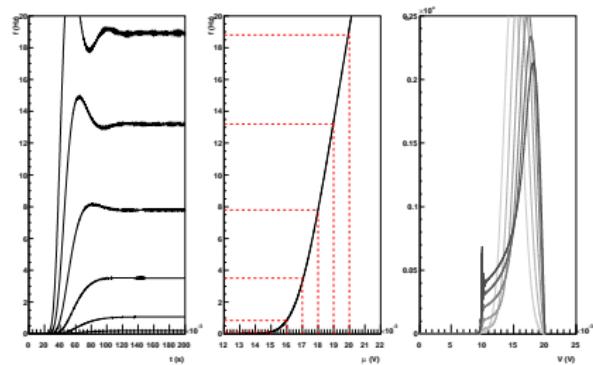
Is This Neuroscience?



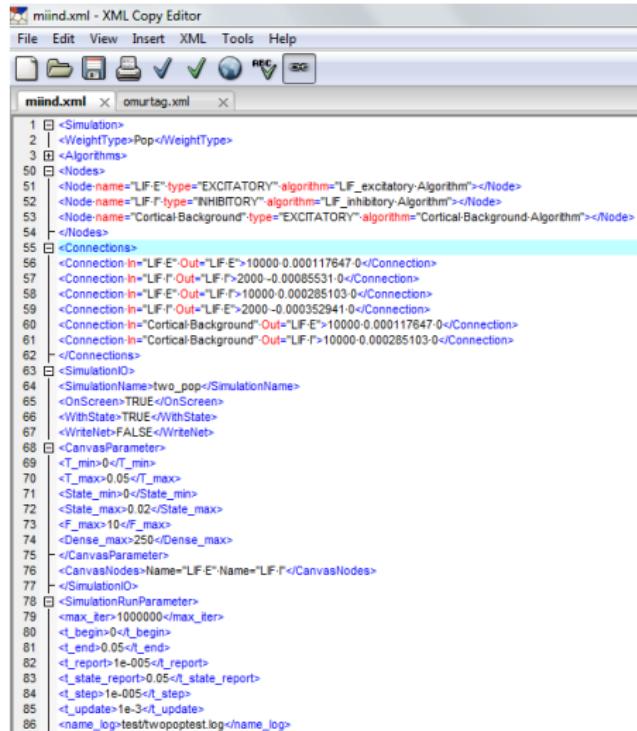
Modelling Populations: MIIND

Sophisticated Population Dynamics

- Wilson-Cowan Dynamics
- Population Density method
 - Describes large populations of leaky-integrate-and-fire neurons
 - Fokker-Planck but better; no diffusion limit
 - $\rho(v)dv$: fraction of population with *membrane potential* in $[v, v + dv]$
 - Balance excitation-inhibition



Short Demo



The screenshot shows a window titled "miind.xml - XML Copy Editor". The menu bar includes File, Edit, View, Insert, XML, Tools, and Help. The toolbar contains icons for file operations like Open, Save, Print, and XML validation. The main area displays the XML code for a simulation named "two_pop". The code defines nodes (LIF-E and LIF-I), algorithms (excitatory and inhibitory), connections between them, and a simulation run parameter. The XML uses namespaces and attributes to specify parameters like connection weights and algorithm names.

```

1 <?xml version="1.0" encoding="UTF-8"?>
2 <Simulation>
3   <WeightType>Pop</WeightType>
4   <Algorithms>
50     <Nodes>
51       <Node name="LIF-E" type="EXCITATORY" algorithm="LIF_excitatory_Algorithm"></Node>
52       <Node name="LIF-I" type="INHIBITORY" algorithm="LIF_inhibitory_Algorithm"></Node>
53       <Node name="Cortical/Background" type="EXCITATORY" algorithm="Cortical Background Algorithm"></Node>
54     </Nodes>
55     <Connections>
56       <Connection in="LIF-E" Out="LIF E">10000.000117647-0</Connection>
57       <Connection in="LIF-I" Out="LIF I">2000-0.00085531-0</Connection>
58       <Connection in="LIF-E" Out="LIF I">10000.000285103-0</Connection>
59       <Connection in="LIF-I" Out="LIF E">2000-0.000352941-0</Connection>
60       <Connection in="Cortical/Background" Out="LIF E">10000-0.000117647-0</Connection>
61       <Connection in="Cortical/Background" Out="LIF I">10000-0.000285103-0</Connection>
62     </Connections>
63   <SimulationIO>
64     <SimulationName>two_pop</SimulationName>
65     <OnScreen>TRUE</OnScreen>
66     <WithState>TRUE</WithState>
67     <WriteNet>FALSE</WriteNet>
68     <CanvasParameter>
69       <T_min>0</T_min>
70       <T_max>0.05</T_max>
71       <State_min>0</State_min>
72       <State_max>0.02</State_max>
73       <r_max>10</r_max>
74       <Dense_max>250</Dense_max>
75     </CanvasParameter>
76     <CanvasNodes>Name="LIF-E" Name="LIF-I"</CanvasNodes>
77   </SimulationIO>
78   <SimulationRunParameter>
79     <max_iter>1000000</max_iter>
80     <l_begin></l_begin>
81     <l_end>0.05</l_end>
82     <l_report>1e-005</l_report>
83     <l_state_report>0.05</l_state_report>
84     <l_step>1e-005</l_step>
85     <l_update>1e-3</l_update>
86     <name_log>test\wopoptest.log</name_log>

```

Conclusions

- Many 'cognitive' models use stereotypic coding patterns which can easily be captured in XML
- The population is a natural bridge between 'basic' neuroscience and more higher level models
- The population level scales well, although it may not be appropriate in every situation
- Much replication effort can be avoided by extending the NeuroML, NineML domains
- Very much in the interest of the neurosciences, given the drivers for funding

Status

- MIIND: public; XML available (slightly brittle). Documentation, patchy, but improving: <http://miind.sf.net>
- Soon: generic 1D neural model solver (not just LIF: QIF as well)
- Aim: Generic 2D population solver
 - Izhikevich
 - adaptive exponential
 - synapses
- Working on cloud implementation, web interface, tutorial
- ClamVis: Dave's project. Not really public but check out:
<http://stacker.me.uk/~daveh/NetSimDocs/projectxmlformat.html>

Acknowledgement

- Dave Harrison: modelling, XML stuff
- Robert Cannon
- Padraig Gleesson, Angus Silver