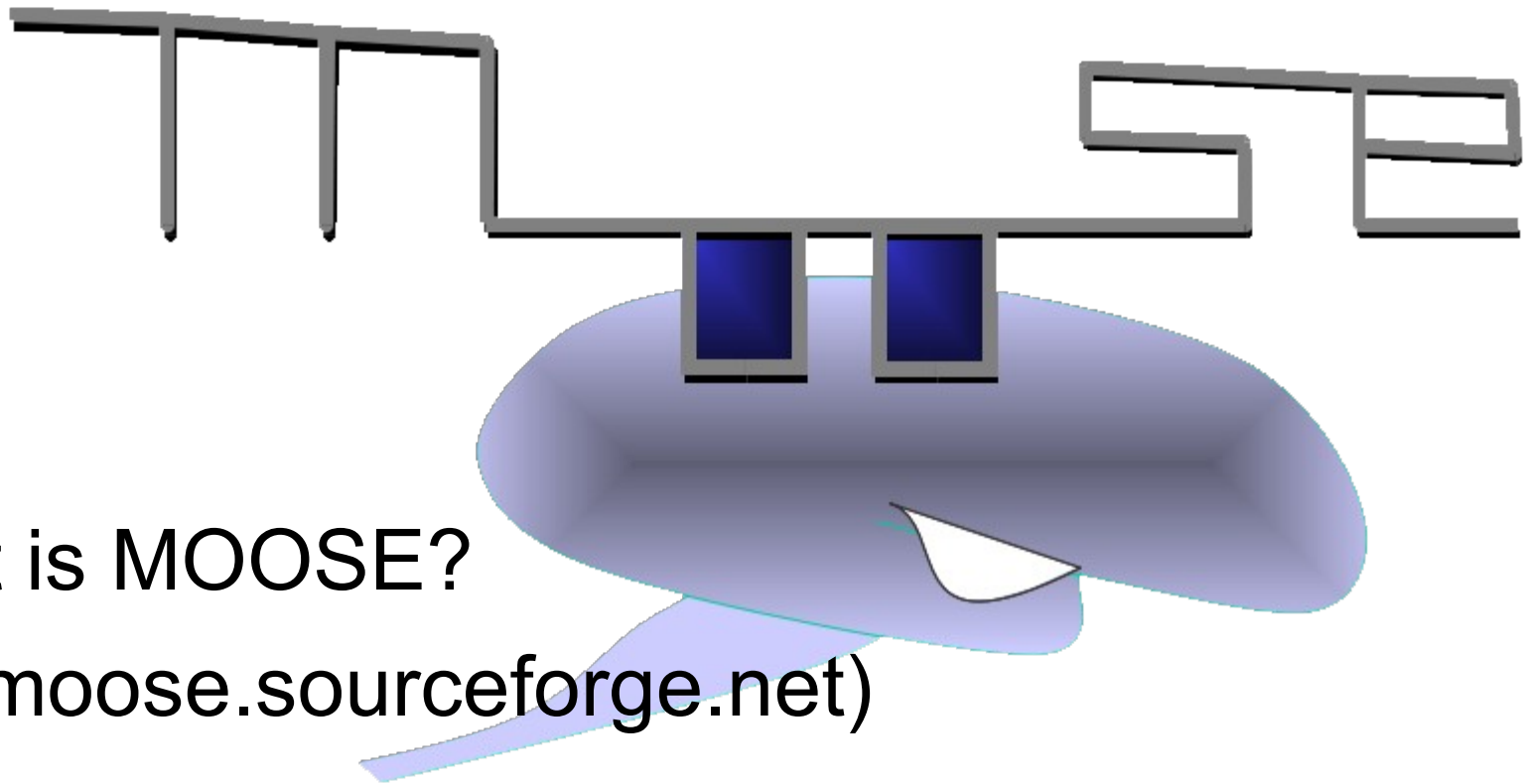


Experiences with Traub model in MOOSE

Subhasis Ray
NCBS, TIFR
Bangalore
INDIA

Introduction

- First a story ...



- What is MOOSE?
(<http://moose.sourceforge.net>)

Why Traub model?

- Highly detailed single cell models put together
- Possibly the most biologically realistic model of a cortical column at the time

Porting Traub 2005 model to moose

FORTRAN version

Almost impossible to read

Too many dialect specific material

Did not compile in the first place

Porting Traub 2005 model to moose

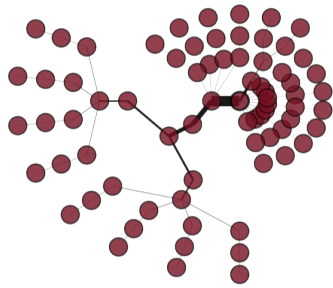
NEURON version

- + Closer to biological description and convention in GENESIS/MOOSE
- + Less code
- + More organized, numerics separated from model structure

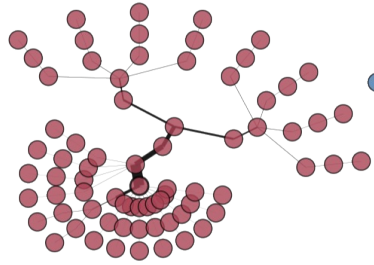
Easier to read.

Some of the cell morphologies

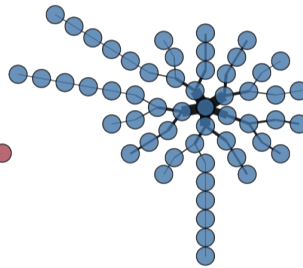
SupPyrRS



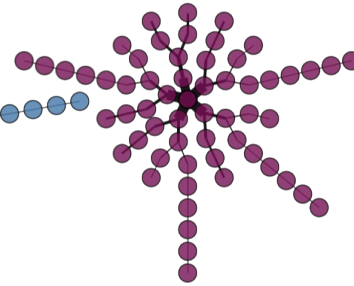
SupPyrFRB



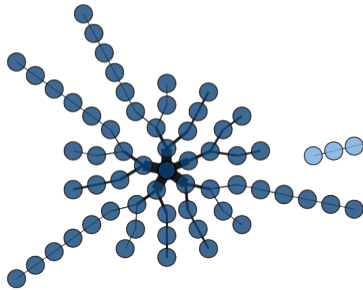
SupLTS



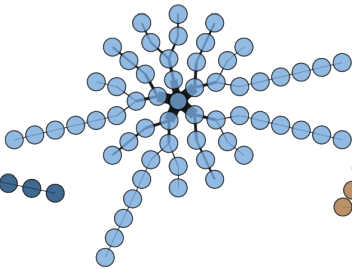
SpinyStellate



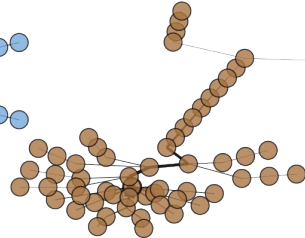
SupBasket



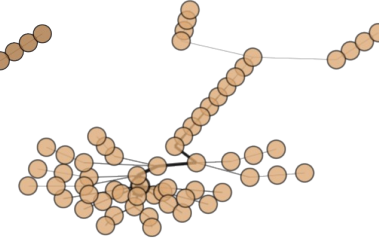
SupAxoaxonic



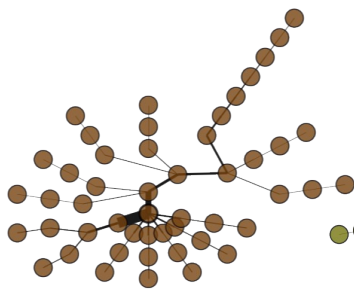
TuftedIB



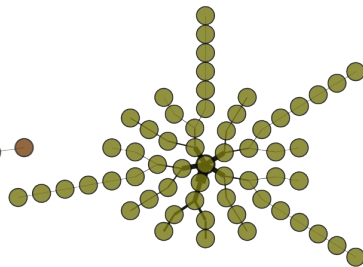
TuftedRS



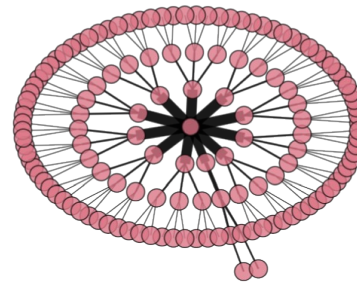
NontuftedRS



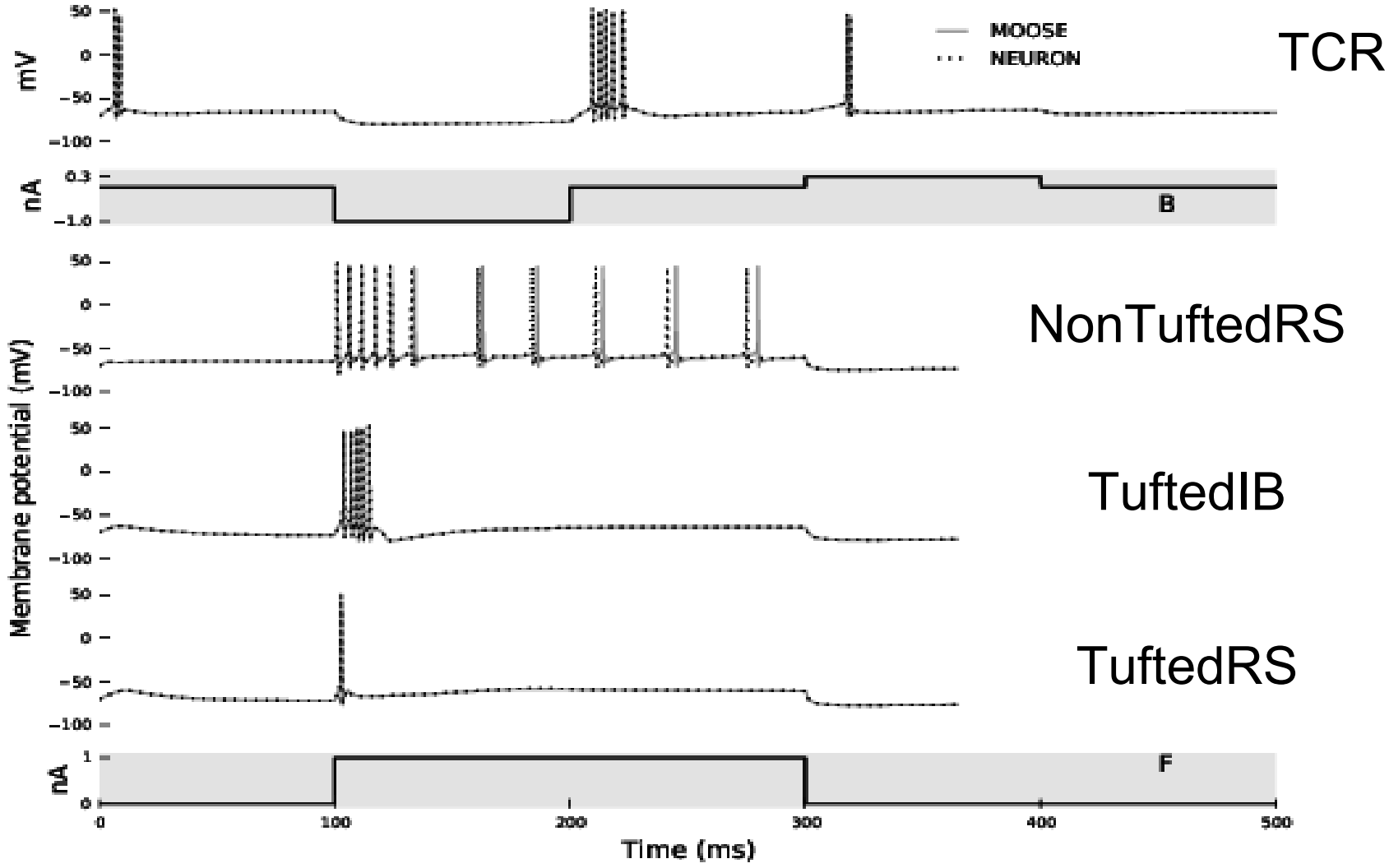
nRT

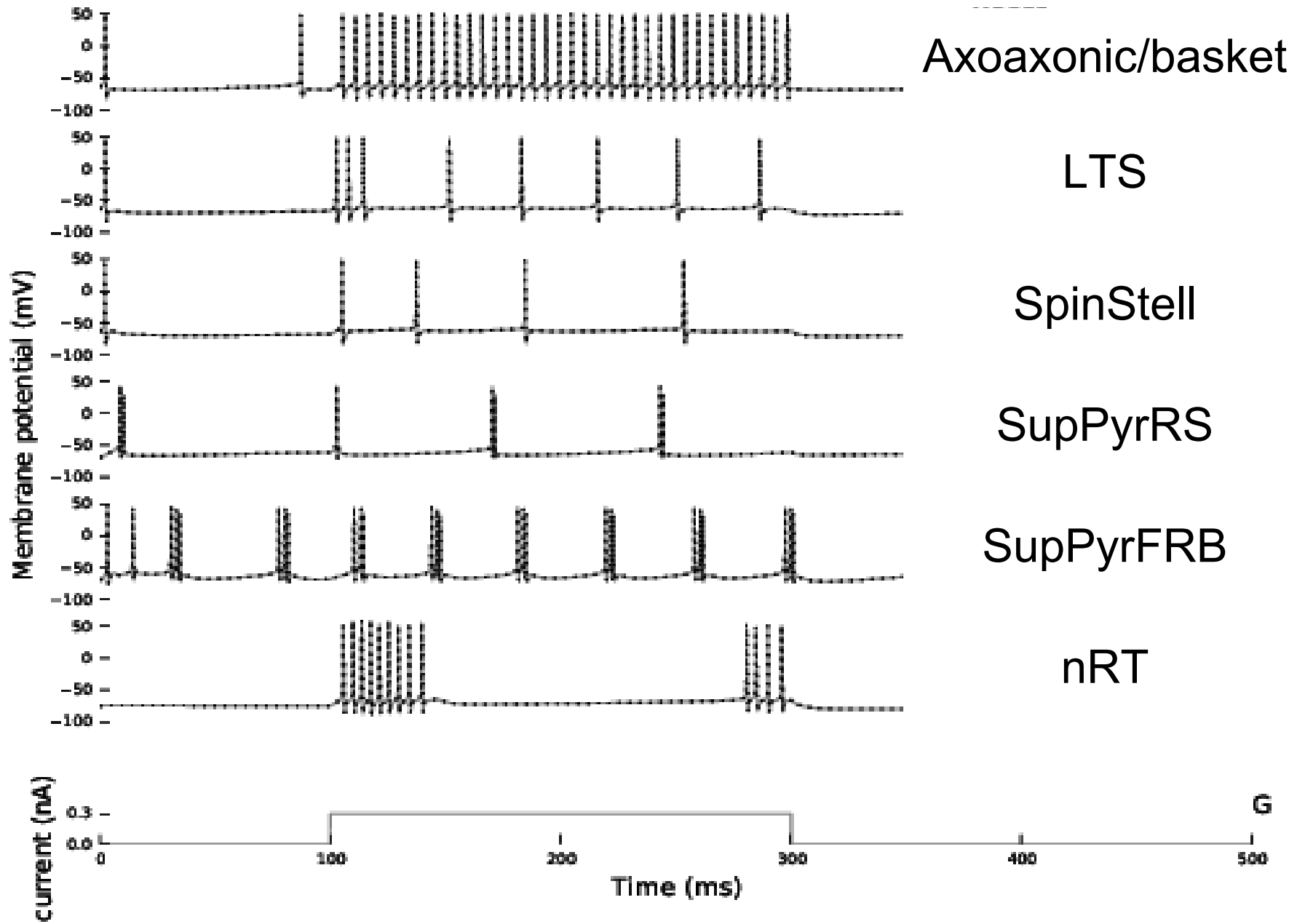


TCR



Comparison of single cell model behaviour





All single cell models are available with moose

<http://moose.sourceforge.net>

[moose/Demos/traub_2005](#)

Issues faced

- Non smooth functions for Ca^{2+} dependence, Mg^{2+} dependence – biological functions are usually smooth!

Issues faced

Units!

- The biggest source of problem (the world does not seem to have agreed upon SI units)!
- Ca^{2+} dynamics with arbitrary units

Issues faced

Initialization of parameters

What is m_{inf} , τ_m etc ? Is this reasonable?

Initial conditions ...

Huge spike at the start – does it affect in the long run?

Does this model make sense for longer simulations?

Different behaviour showing up in longer simulations

Plasticity will matter

Some of the lessons from the conversion process

- Incremental development
 - ... but this can be difficult if the model is not modular.
 - ... start with passive model
 - ... test single channel on single compartment
 - ... put multiple channels together (debugging can still be a combinatorial problem) in single compartment
 - ... test whole cells
- Minor differences in behaviour can come from major errors and vice versa

Some of the lessons from the conversion process

Manual conversion good for getting a hang of the model – but ultimately programmatic conversion is a must.

Looking at the model in multiple ways can find bugs.

**** Discussing it with somebody else helps!**

Can we be sure of our code?

Network behaviour can be highly variable depending on the exact connectivity.

- Comparison of exact connectivity important.
- Getting exact sequence of random numbers between simulators may be impossible.

Acknowledgements

Upi and MOOSE Team: Niraj, Harsha, Chaitanya

Lab13@ncbs

ncbs

DAE, EU-India Grid

NIGMS/ SBCNY (Systems Biology Centre of New York (5P50 GM071158-03))

Bristol Myers Squibb

Thank you

Questions?