



OpenWorm

Building a digital organism

Matteo Cantarelli

OpenWorm Co-founder

matteo@openworm.org

@tarelli

OCNS 2015 - Prague

OpenWorm's Goal

Long term

Full scale **simulation** of **C. elegans**

Medium term

Accurately predict **WormBehavior** database
using data-driven, cell-by-cell **3D**
neuromechanical model

Szigeti et al, - Front. Comp. Neuro., 2014

Why the *C. elegans*?

- Wide range of behaviours
 - Locomotion, feeding, reproduction
 - Learning, mating, social behaviours
- How does it do it?
 - 1mm long
 - 302 neurons
 - 956 cells in total
- The first organism to have its genome **fully sequenced**
 - 100 million base pairs
 - 3 billion in mouse and humans

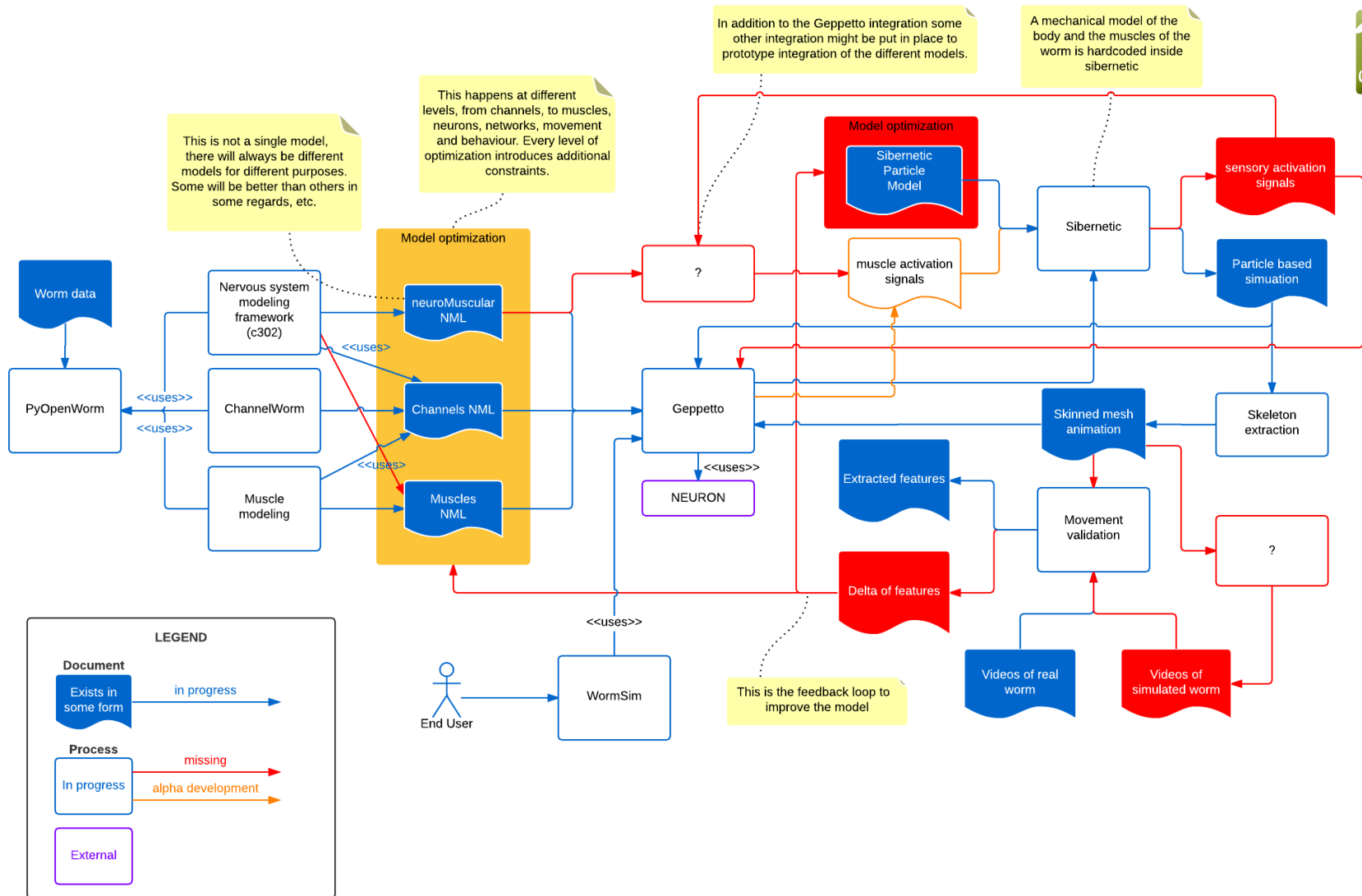


OpenWorm open science community

- Independent international **open science** community
 - 9 **core** members
 - 20 **active members** & 40 **additional contributors** across 14 **countries**
- Networked science
 - >35 **Git repositories**
 - ~350 **mailing list** members
 - Streamed 40+ online meetings last year on YouTube
- Collaborative open source construction
 - Every line of code committed to GitHub

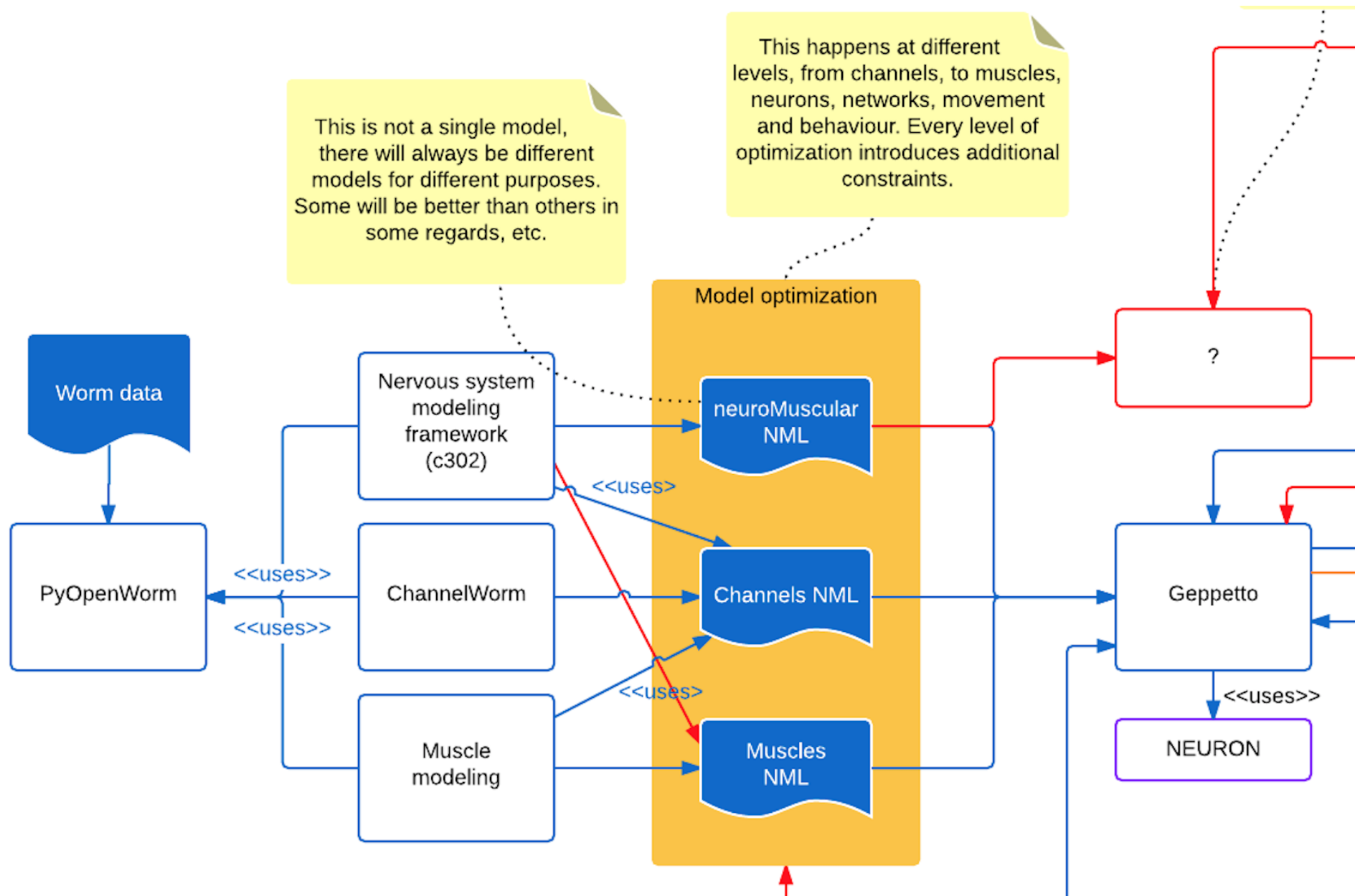
Where are we?





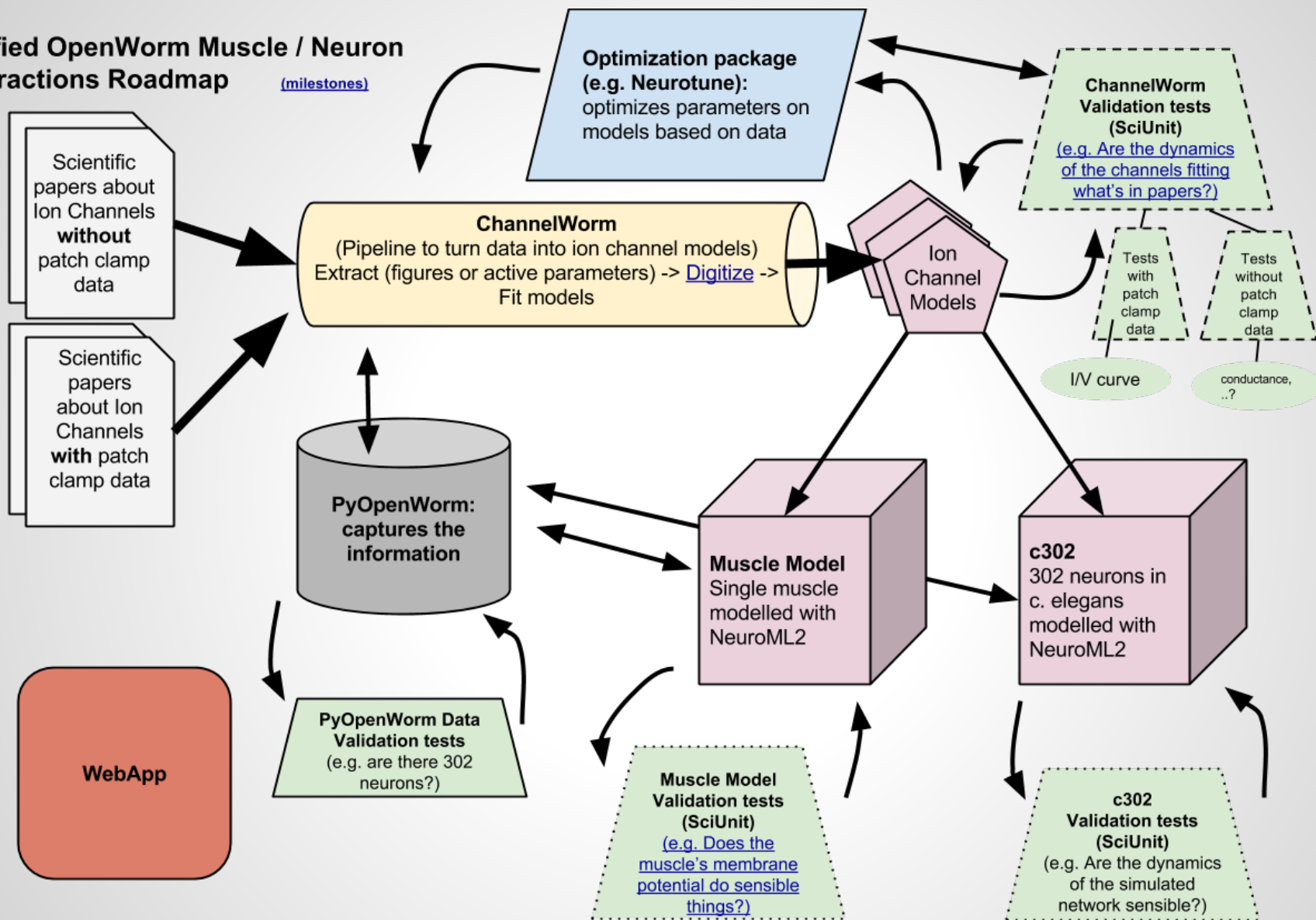
Modelling the channels and the cells



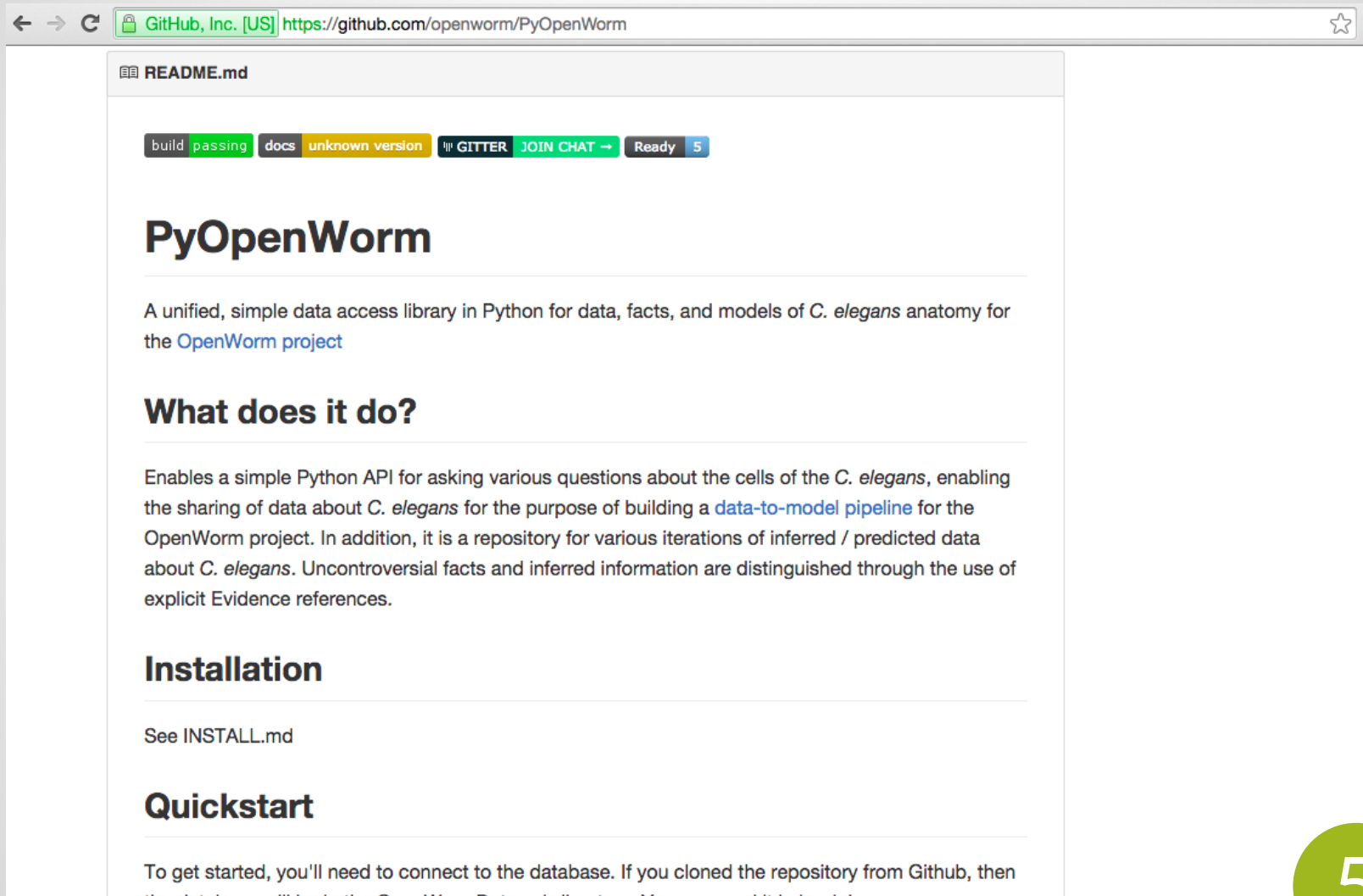


Unified OpenWorm Muscle / Neuron Interactions Roadmap



[\(milestones\)](#)




PyOpenWorm



The screenshot shows the GitHub repository page for PyOpenWorm. The browser address bar displays the URL <https://github.com/openworm/PyOpenWorm>. The repository name is "PyOpenWorm". The README file is selected, showing a header with status badges: "build passing", "docs unknown version", "GITTER", "JOIN CHAT", and "Ready 5". The main content of the README includes the title "PyOpenWorm", a description of the library, and sections for "What does it do?", "Installation", and "Quickstart".

← → ↻  GitHub, Inc. [US] <https://github.com/openworm/PyOpenWorm> 

 README.md

build passing docs unknown version GITTER JOIN CHAT → Ready 5

PyOpenWorm

A unified, simple data access library in Python for data, facts, and models of *C. elegans* anatomy for the [OpenWorm project](#)

What does it do?

Enables a simple Python API for asking various questions about the cells of the *C. elegans*, enabling the sharing of data about *C. elegans* for the purpose of building a [data-to-model pipeline](#) for the OpenWorm project. In addition, it is a repository for various iterations of inferred / predicted data about *C. elegans*. Uncontroversial facts and inferred information are distinguished through the use of explicit Evidence references.

Installation

See INSTALL.md

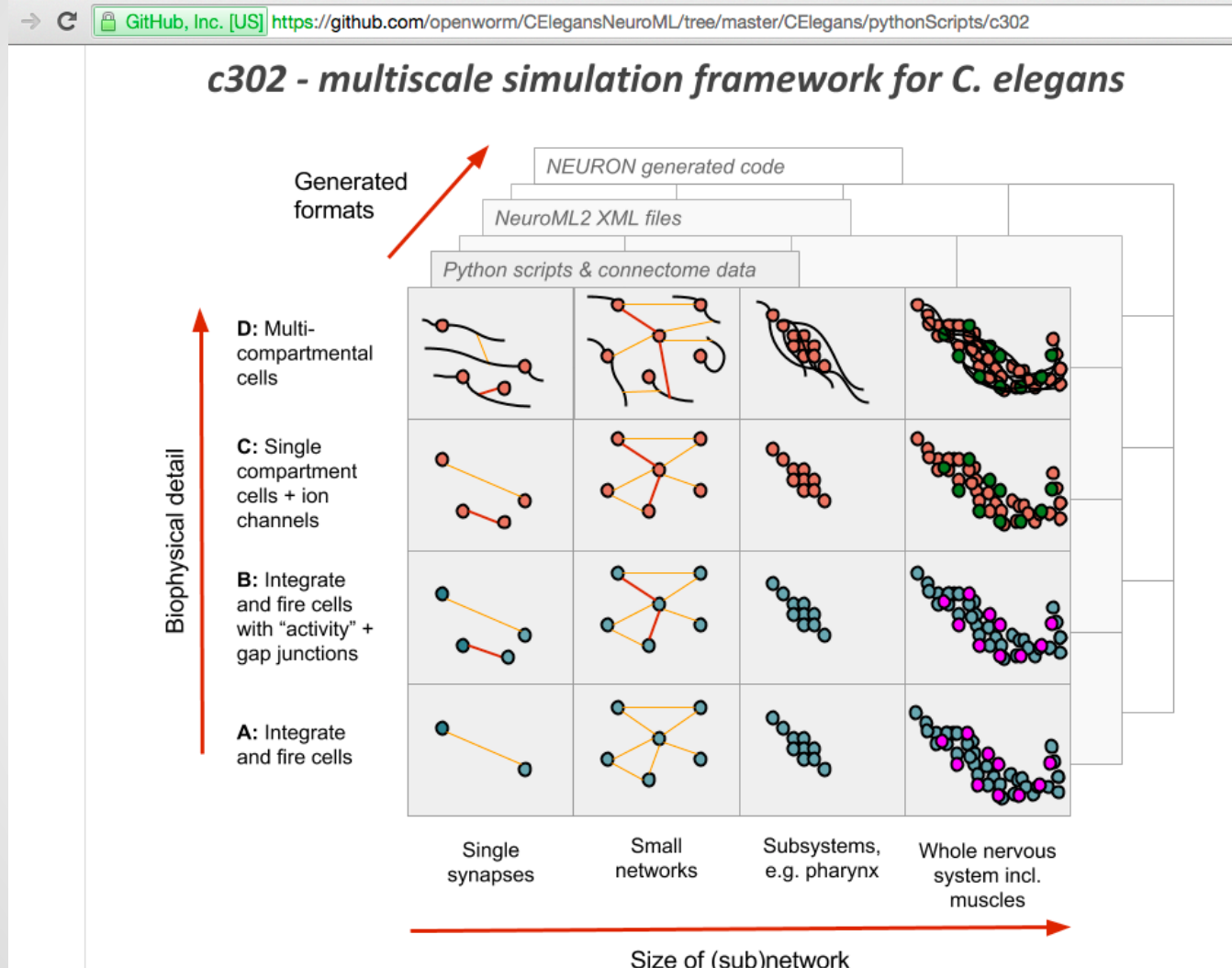
Quickstart

To get started, you'll need to connect to the database. If you cloned the repository from Github, then the database will be in the `OpenWormData` subdirectory. You can use it to build data...

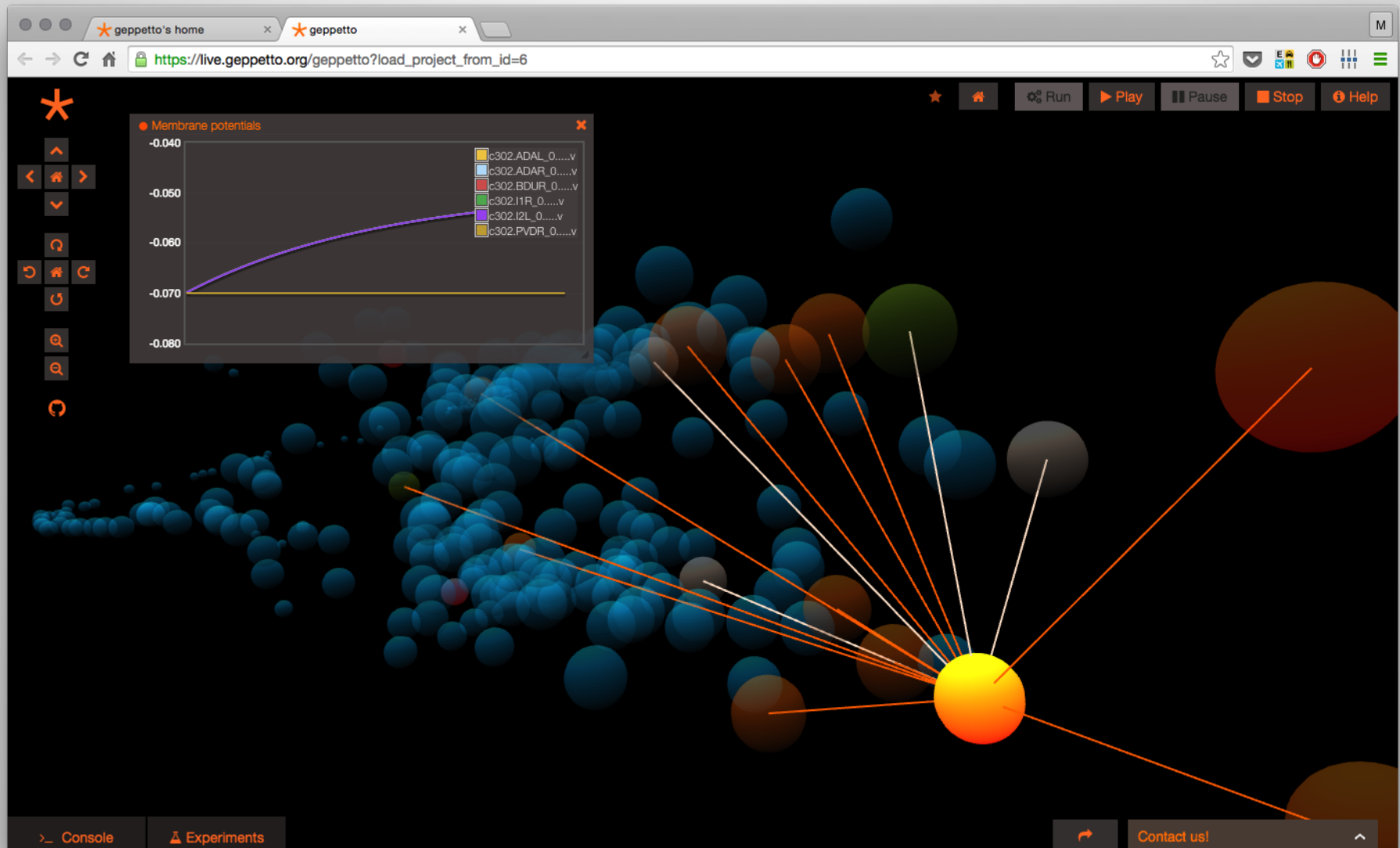
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<http://github.com/openworm/pyopenworm>

c302 model generation framework



c302 generated network




<https://github.com/openworm/CElegansNeuroML/tree/master/CElegans/pythonScripts/c302>

Model of a C. elegans muscle

→ C GitHub, Inc. [US] https://github.com/openworm/muscle_model

build passing

Open Worm muscle model

 GITTER [JOIN CHAT →](#)

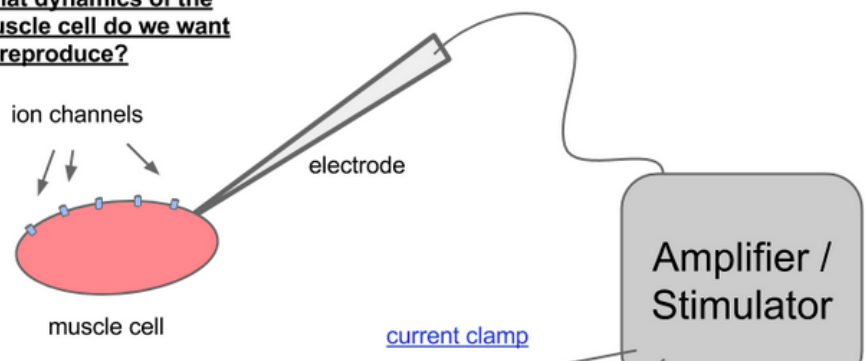
Authors: Mike Vella, Alex Dibert, Padraig Gleeson, Rayner Lucas email:mv333@cam.ac.uk

If you contribute to the project please add your name to the Authors field

Introduction

This repository contains several different subprojects all related to the construction of a biophysically-detailed model of the dynamic properties of electrical excitation of the body wall muscle of the c. elegans.

What dynamics of the muscle cell do we want to reproduce?



ion channels

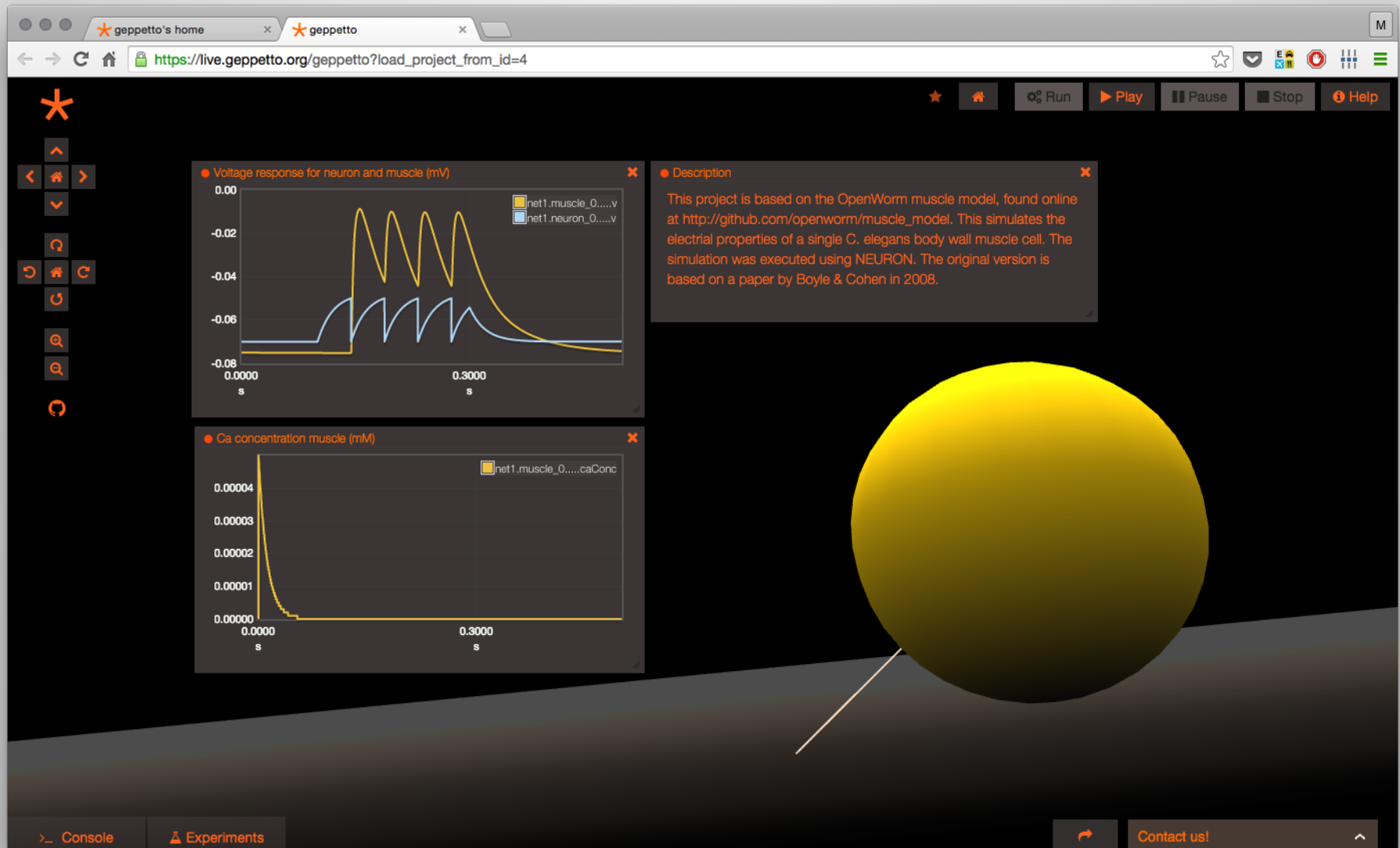
electrode

muscle cell

Amplifier / Stimulator

[current clamp](#)

Model of a C. elegans muscle



http://github.com/openworm/muscle_model

ChannelWorm

ChannelWorm

docs latest  GITTER JOIN CHAT → Ready 5 build passing

The aim of the **ChannelWorm** is to integrate information and tools related to modeling ion channels in *C. elegans* for the [OpenWorm Project](#).

Objectives

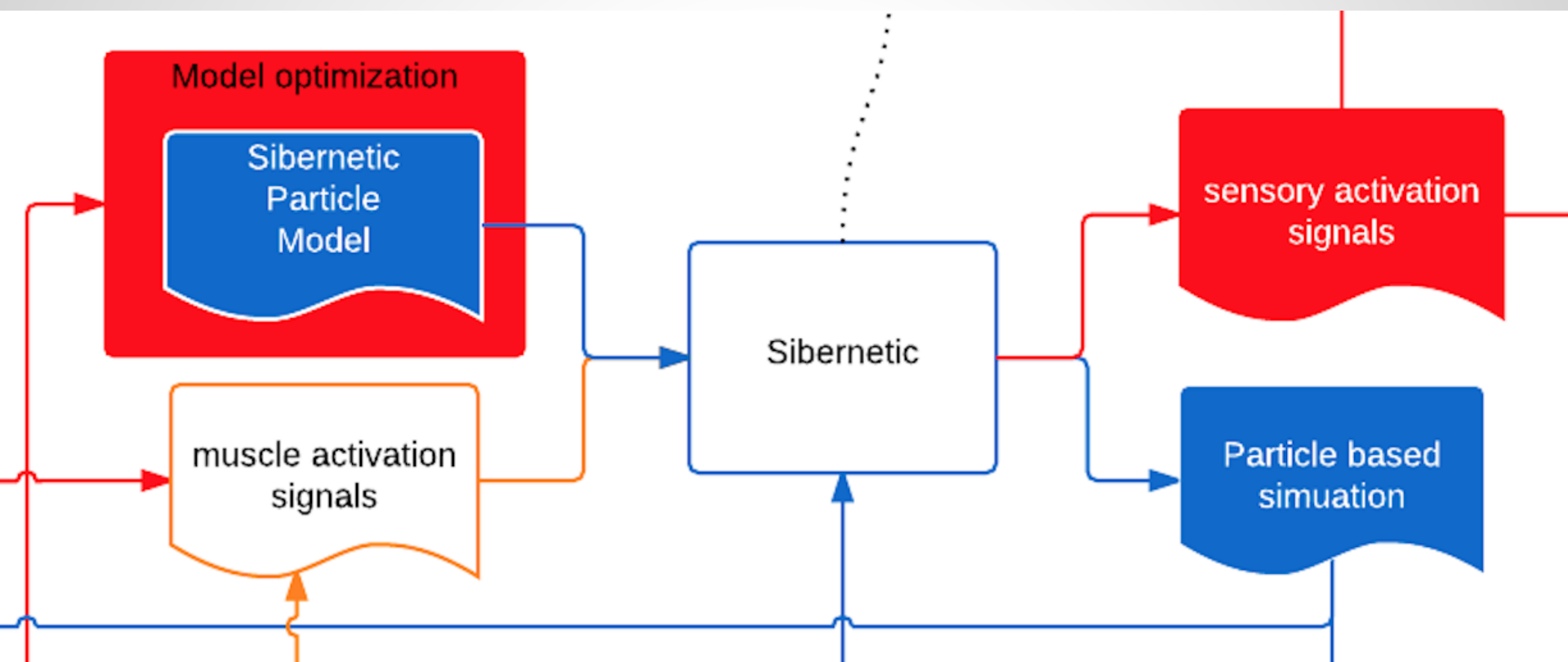
- **Information Management**
 - Integrate and structure data related to ion channels in *C. elegans*, from genotype to phenotype
 - Develop APIs for accessing data
 - Keep data up-to-date
- **Ion channel modeling**
 - Build Hodgkin-Huxley models for ion channels based on experimental patch clamp studies
 - Estimate kinetics and build models for ion channels with no patch clamp data available (based on homologous channel types)
 - Create verification & validation tests to prove matching of the models with experimental data
- **Setting up a simulation environment**
 - Simulate the computational analysis phase of a patch clamp experiment
 - Simulate and run customized versions of ion channel(s) in cell(s)
 - Check if the simulation fits the biological boundaries

Some use cases

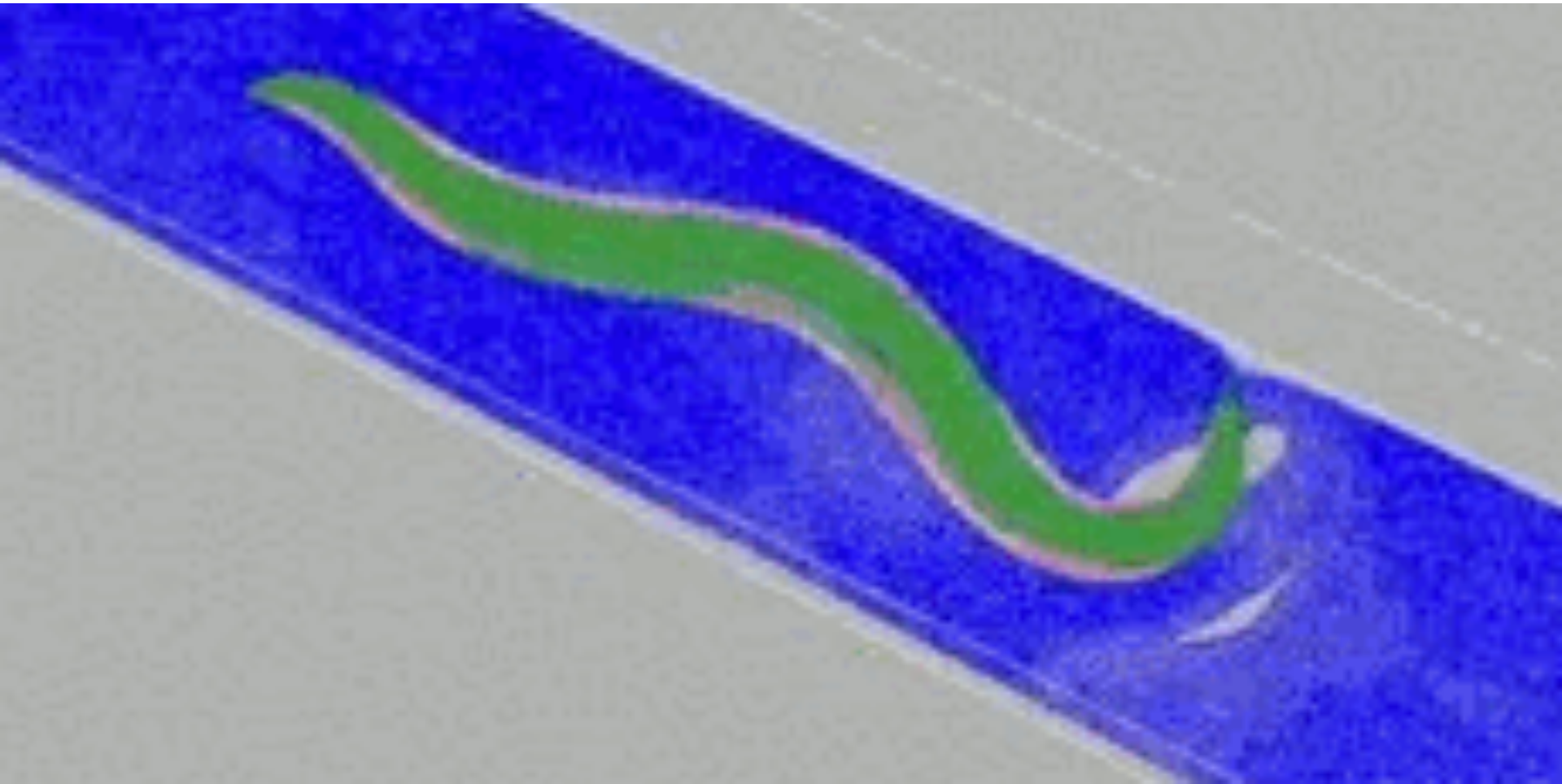
<https://github.com/VahidGh/ChannelWorm/>

Modelling the body





SIBERNETIC



3

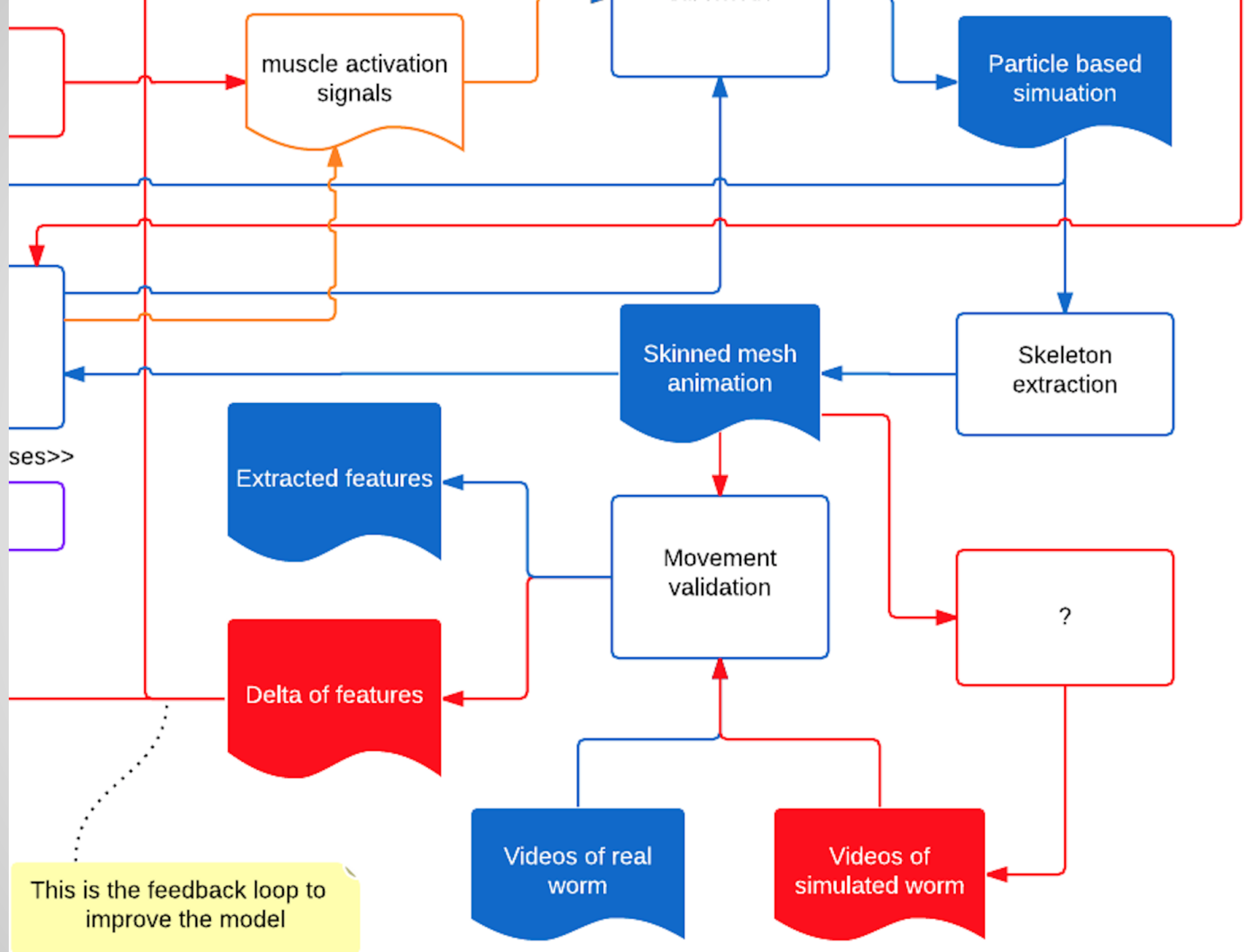
<http://github.com/openworm/sibernetik>

Palyanov et al., 2015, - In Revision

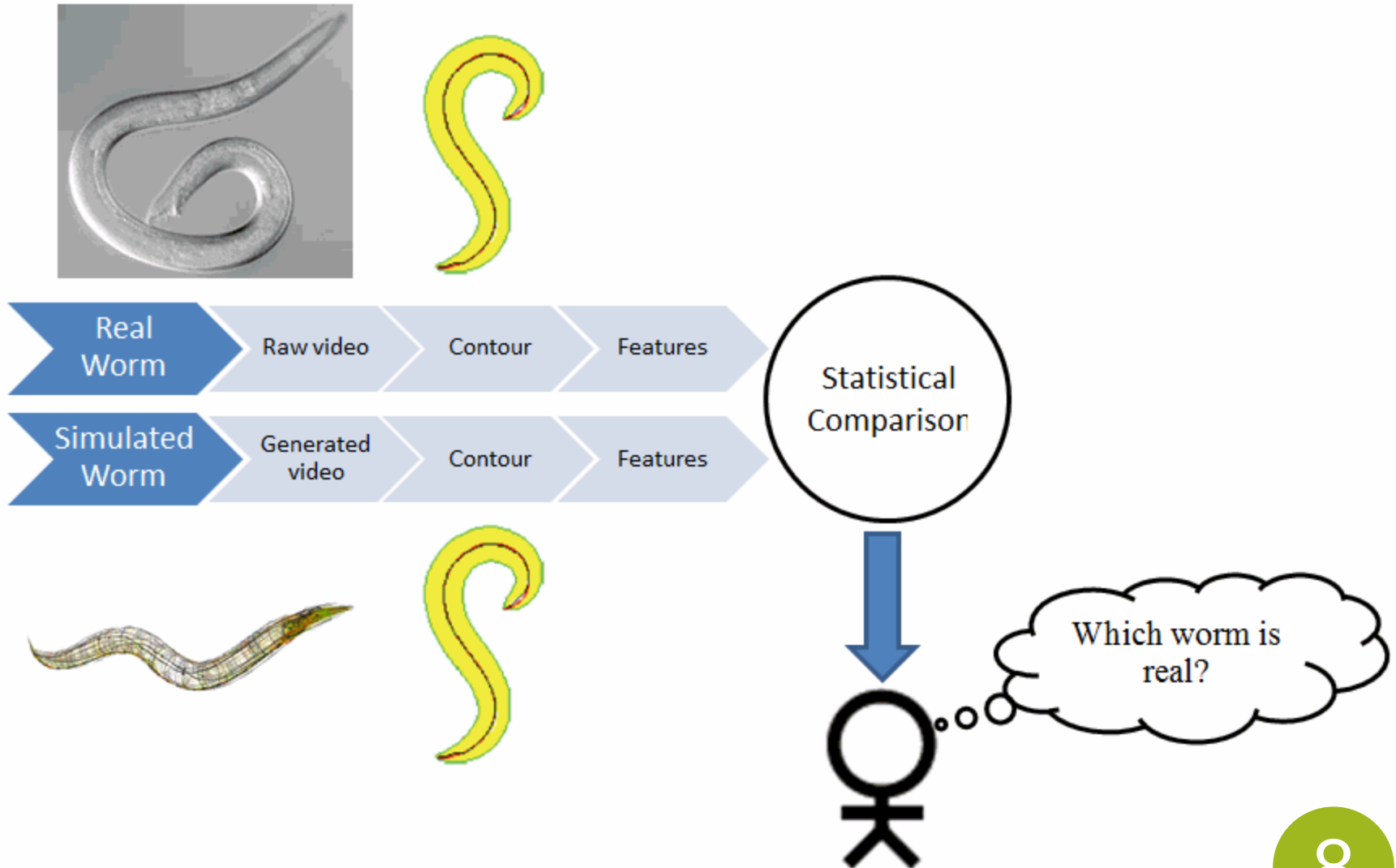


Movement Validation



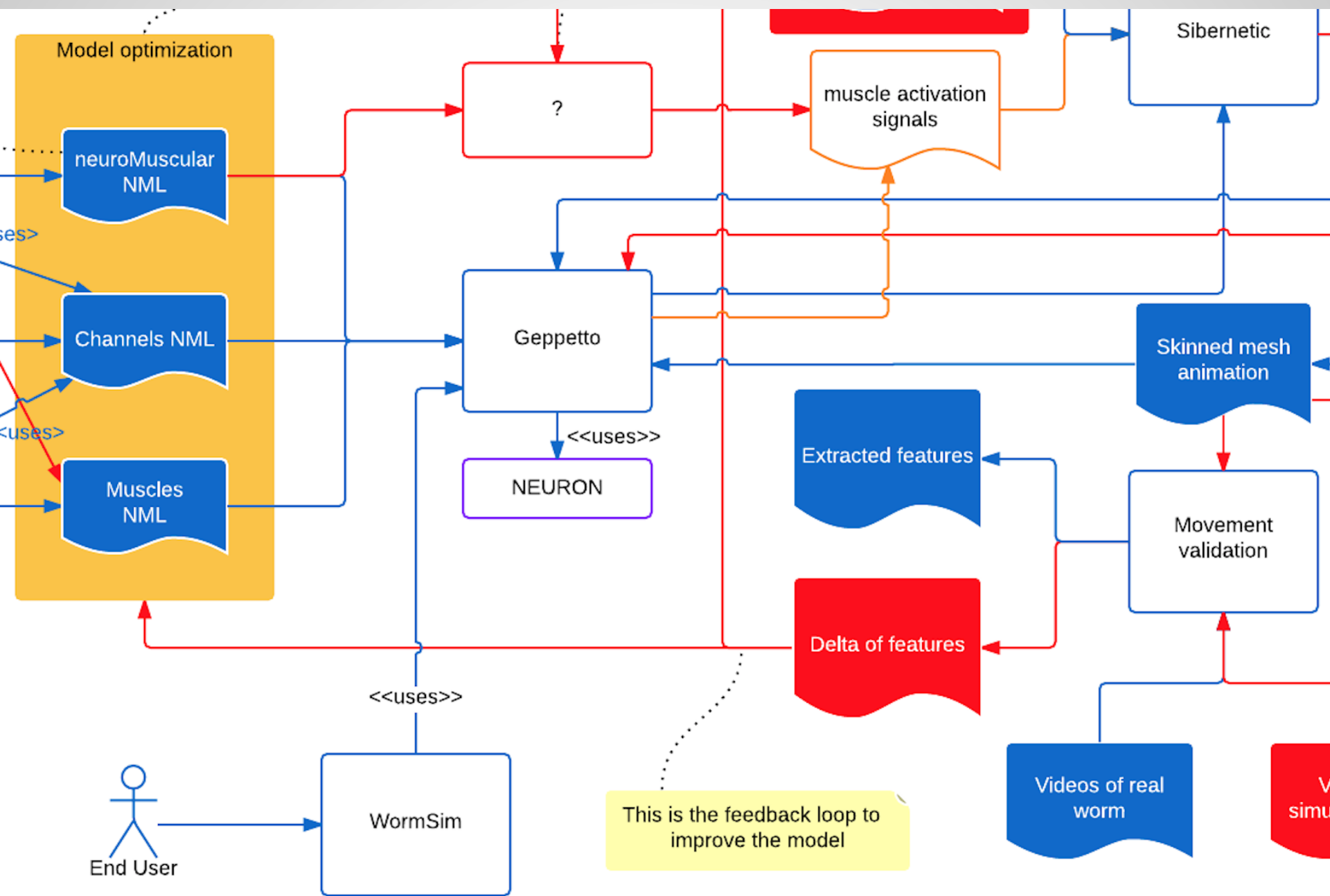


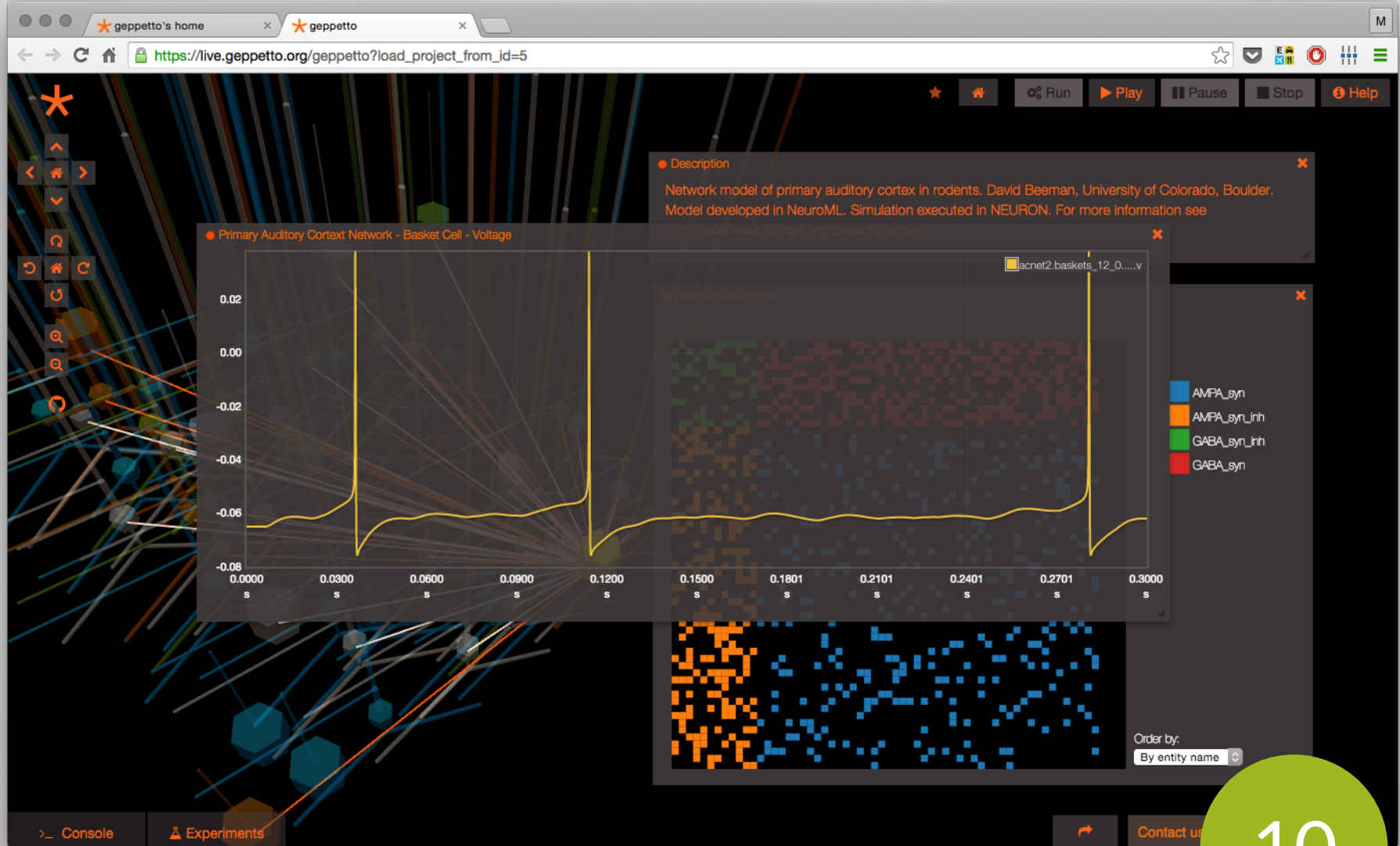
Movement validation



Model integration





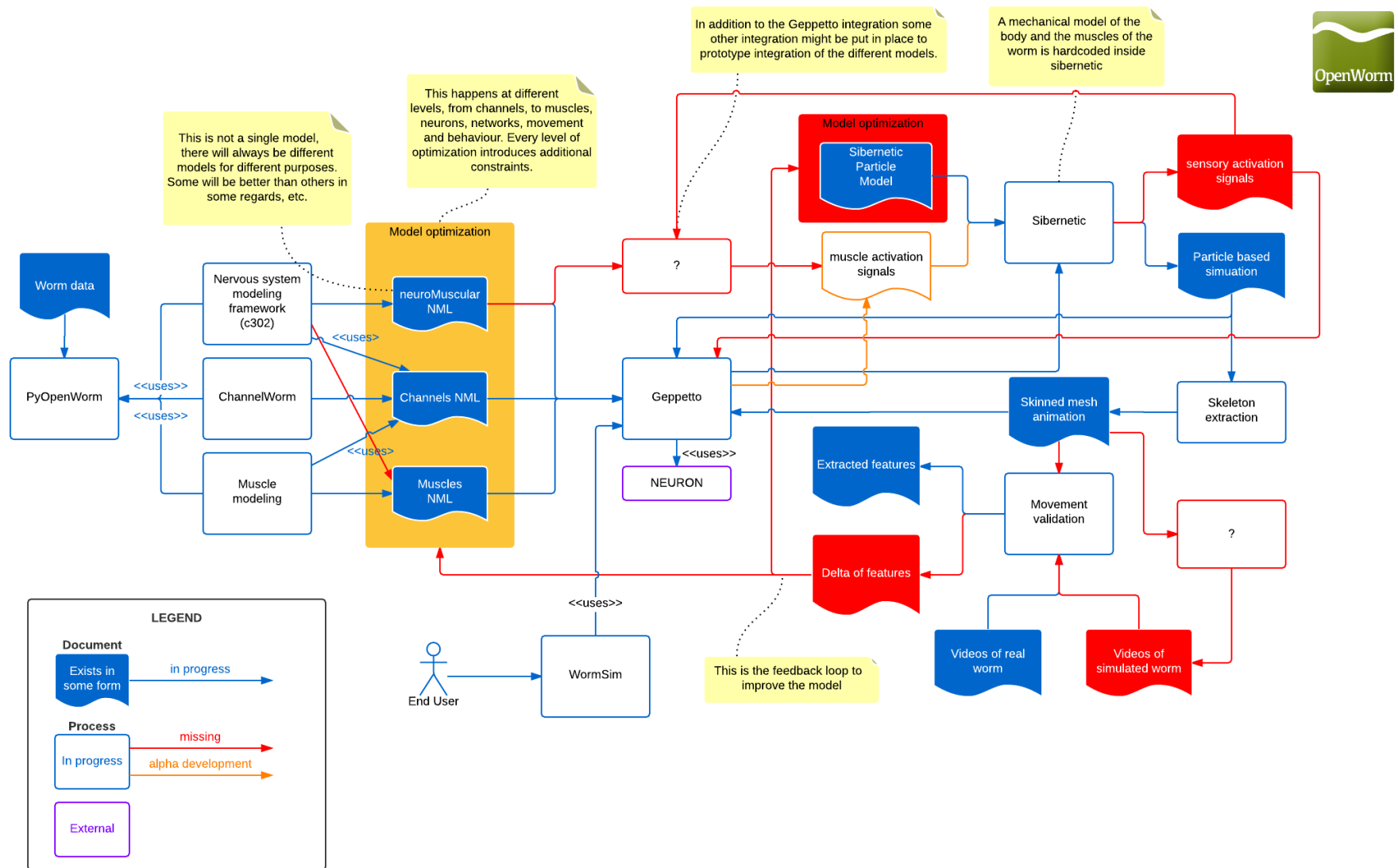


WormSim



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STARTER**



Recap

- OpenWorm is collaboratively building a *C. elegans* simulation in an open science manner
- The project is using a database of recorded worm behavior as its ground truth, in addition to reported cellular activity and observed anatomy
- Mechanical and neuronal simulations are currently used
- WormSim will let users interact with the different models and ultimately integrate them together

How can I contribute?

- Get in touch

info@openworm.org

<http://openworm.org>

@openworm

- Fork any of the repositories and issue a pull request
 - Join the conversations on GitHub



wellcome trust



Caltech



Massachusetts
Institute of
Technology



SALK INSTITUTE
FOR BIOLOGICAL STUDIES

UC San Diego



UNIVERSITY OF LEEDS

Core team



Our Superheroes.

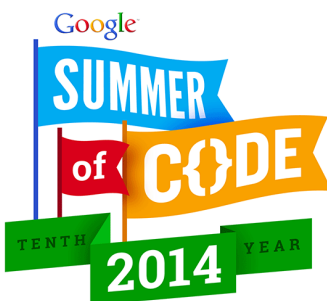
Andrey Palyanov	Balazs Szigeti	Giovanni Idili	Jim Hokanson	Matteo Cantarelli	Michael Currie
Padraig Gleeson	Sergey Khayrulin	Stephen Larson			

Contributors



Helping us with amazing support. Thanks!

Abdullah Haroon	Adrian Quintana	Alexander Dibert	Alexander N	Allie Love	Andrew Leifer
Andrew Papadopoli	Ari Richman	Bóris Marin	Bradley Alicia	Charles Cooper	Christian Grove
Chris Jensen	Christina Lee	Christopher Perez	Credentiality	Crystin Slade	Dan Knudsen
Dan Kruchinin	David Dalrymple	Deanne Taylor	Dmitriy Shabanov	Dmitar Shterionov	Dominic Denicola
Domenico Luciani	Ellen McGowan	Eviatar Yemini	Gaston Gentile	Gleb Kuznetsov	Jay Coggan
Jesus Martinez	Joe Bowen	Johannes Rieke	John Hurliman	John White	Jordan Boyle
Jinze Yu	Marius Bulbas	Mariusz Sasinski	Mark Watts	Martin Madaras	Matt Olson
Mike Vella	Mei Zhen	Nathan Daly	Netta Cohen	Pedro Tabacof	Peter McCluskey
Petr Baudis	Phil McElmurray	Rayner Lucas	Richard Gordon	Rich Stoner	Rick Gerkin
Steven Cook	Steve McGrew	Timothy Busbice	Travis Jacobs	Vanessa Adelmann	Vahid Ghayoomi





Thanks for the attention!

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