Quick Guide for Using NeMo-TMS

Compile mod files

Compile the .mod files inside the following folders:

'Model_Generation\Jarsky_files\lib_mech'
'Model_Generation\Aberra_files\lib_mech'

Note that you only need to do this once on each computer. Skip this step if you have already compiled the mod files. For more information on mod files and how to use them, see: https://www.neuron.yale.edu/phpBB/viewtopic.php?t=3263

Model generation

Generate the model by following the instructions for each model type:

Jarsky CA1 pyramidal cell: Change your MATLAB directory to 'Model_Generation' folder and run Jarsky_model() in MATLAB. The prompt asks for the input morphology file (.swc). You can use one of the sample morphology files provided in ('Model_Generation\morphos\'cell_#.swc).

Aberra cortical pyramidal cell: Change your MATLAB directory to 'Model_Generation' folder and run **Aberra_L5_model ()** or **Aberra_L23_model ()** in MATLAB.

The generated model is placed inside 'Models\model_name' folder. All the next steps are model-specific and therefore operate within this folder.

Note: Currently this step is not supported in macOS. However, models can be generated on other computers and copied to the macOS computer. In this case, the mod files for the generated model ('Models\model_name\lib_mech') should be recompiled in the new computer.

Electric field type

If you are interested in using a spatially uniform electric field, skip to 'TMS waveform generation'.

If you are interested in using a spatially realistic electric field, Simulate the spatial distribution of the electric fields in SimNIBS. Refer to the following resources for more details:

Online SimNIBS tutorial: https://simnibs.github.io/simnibs/build/html/tutorial/tutorial.html

Translational FEM models: https://zenodo.org/record/4009465

High-resolution human head FEM model: https://zenodo.org/record/5209082

Export neuron segment location

Export the neuron segment coordinates by running 'Code\NEURON\save_locations.hoc' in NEURON environment.

Tip: run NEURON scripts by double-clicking on the file on Windows or running the command **nrniv save_locations.hoc** in the terminal on macOS and Linux.

Tip: For the Aberra L2/3 model, the stack size needs to increase for the NEURON to run. Running 'save_locations.ps1' on Windows PowerShell, or 'save_locations.sh' on macOS/Linux does this automatically.

Realistic electric field coupling

Change your MATLAB directory to 'Code\E-Field_Coupling', run couple_gui(); in MATLAB, and follow the prompts.

TMS waveform generation

Change your MATLAB directory to 'Code\TMS_Waveform', run TMS_Waveform(); in MATLAB, and follow the prompts.

NEURON simulation

Select NEURON simulation parameters by running 'Code\NEURON\GUI_params.hoc' in NEURON environment and following the prompts. Then, start the simulation by running 'Code\NEURON\TMS_script.hoc' in NEURON environment. The window will automatically close upon successful completion of the simulation.

Tip: run NEURON scripts by double-clicking on the file on Windows or running the command **nrniv file_name.hoc** in the terminal on macOS and Linux.

Tip: For the Aberra L2/3 model, the stack size needs to increase for the NEURON to run. Running 'TMS_script.ps1' on Windows PowerShell, or 'TMS_script.sh' on macOS/Linux does this automatically.

Calcium simulation

Change your MATLAB directory to 'Code\Calcium', run run('calcium_simulation_setup.mlapp'); in MATLAB, enter the location of UG4 software and the parameters for calcium simulation. Then click on the Convert NEURON files button to convert the files to a format compatible with UG4. Afterward, click on the Generate script, Save script, and Run script buttons respectively and wait for the simulation to finish.

Visualization

Change your MATLAB directory to 'Code\Visualization', execute run('visualization.mlapp'); in MATLAB. Click on File and select Open Voltage Data and Open Calcium Data to visualize the 3D voltage and calcium results across time respectively.

Full tutorial

For more information refer to the full PDF tutorial.