Spike-based machine learning with GeNN

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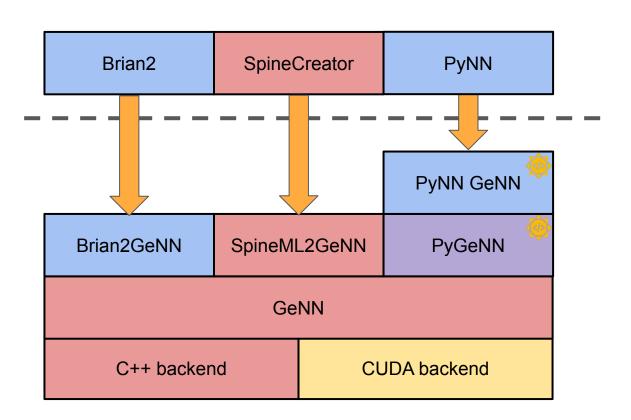


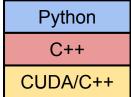


GeNN

- Cross-platform C++ library for generating optimised CUDA code for GPU accelerated SNN simulations.
- Can also generate C++ code for testing on computers without GPUs (e.g. here with your laptops!)
- Hopefully you've learnt all about it in our talks earlier in the week!
- All GeNN features are now available from Python for easier interoperability with other ML and Computational Neuroscience tools

The GeNN ecosystem





Installation

CUDA on Linux

- Each version of CUDA only supports a subset of GCC versions so if you have a very old or very bleeding edge OS you may need to install an additional version of GCC.
- Installing CUDA via the NVIDIA proprietary packages tends to work best if your OS is supported.
- Ensure that the CUDA_PATH environment variable is set

CUDA on Windows

- CUDA is nicely integrated into Visual Studio and provided graphical debugging and profiling tools
- Historically, because Visual Studio is frequently updated, compiler/CUDA version mismatches were more prevalent than on Linux but, as of CUDA 10 and Visual Studio 2017, this no longer appears to be the case!
- If installing from scratch we recommend:
 - CUDA 10.1
 - Visual Studio 2017

CUDA on Mac

- Sadly Apple hasn't built any machines with NVIDIA GPUs since 2014
- However, if you're lucky enough to have:
 - MacBook Pro (Retina, 15-inch, Late 2013)
 - MacBook Pro (Retina, 15-inch, Mid 2014)
 - Equivalent iMac models (probably not with you!)
- You may have a NVIDIA GPU that's usable with the current version of CUDA!
- Ensure that the CUDA_PATH environment variable is set





Windows notes

- PyGeNN requires at least Python 3.5 and Visual Studio 2015 installed
- I recommend using Anaconda for Python development on Windows
- You should install CUDA after Visual Studio so they are correctly integrated
- PyGeNN should be used from a command prompt with Python and Visual studio properly configured i.e. by activating conda within a Visual Studio "x64 Native Tools Command Prompt"

Installing PyGeNN from binary wheels

- Select a suitable wheel from the latest release available at https://github.com/genn-team/genn/releases
 For example, if you have a Linux system with Python 3.7, you would pick pygenn-0.2-cp37-cp37m-linux_x86_64.whl
 Note: the Mac OS X wheel are built for CUDA 9, all others for CUDA 10
- 2. Install the wheel using pip e.g. pip install pygenn-0.2-cp37-cp37m-linux x86 64.whl

Installation from source on Linux/Mac

- 1. Download latest release of GeNN from https://github.com/genn-team/genn/releases
- 2. Make sure you have swig installed
- 3. From GeNN directory, build as a dynamic library, directly into the PyGeNN directory using:

```
make DYNAMIC=1
LIBRARY_DIRECTORY=`pwd`/pygenn/genn_wrapper/
```

4. Build and install python module with setuptools using: python setup.py develop

Installation from source on Windows

- 1. Download latest release of GeNN from https://github.com/genn-team/genn/releases
- 2. Make sure you have swig installed
- 3. From GeNN directory, build as a dynamic library using: msbuild genn.sln /t:Build /p:Configuration=Release_DLL
- 4. Copy the newly built DLLs into pygenn using copy /Y lib\genn*Release_DLL.* pygenn\genn_wrapper
- 5. Build and install python module with setuptools using: python setup.py develop

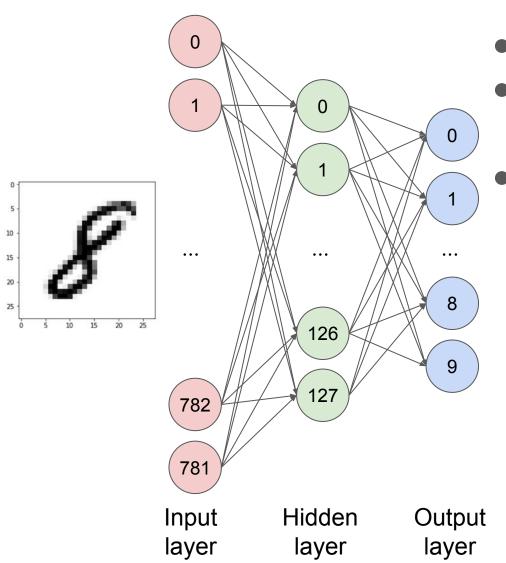
Tutorial repository

- Checkout from:
 - https://github.com/neworderofjamie/pygenn_ml_tutorial
- Contains
 - testing_images.npy and testing_labels.npy testing portion of MNIST dataset
 - weights_0_1.npy and weights_1_0.npy weights trained in Keras
 - tutorial_1.py and tutorial_2.py Code for this tutorial
- Test your installation with:

```
python tutorial_1.py
```

Tutorial: MNIST inference

Part 0: Training the ANN



- Not state-of-the-art!
- I have already done this part for you!
- Achieves 97.6%accuracy on MNIST

Part 1: Classifying a single image

- Basics of using PyGeNN
- 2. Building a spiking network based on ANN
- Recording spikes resulting from presenting single MNIST digit

```
import numpy as np
from os import path
```

Import some standard Python packages

Import some useful PyGeNN components

```
import numpy as np
from os import path
from pygenn.genn model import (create custom neuron class,
                            create custom current source class,
                            GeNNModel)
from pygenn.genn wrapper import NO DELAY
# Parameters
                                       Input required for neurons to spike
IF PARAMS = {"Vthr": 5.0} ←
TIMESTEP = 1.0
PRESENT TIMESTEPS = 100 ◀
                                       How long to present each digit
INPUT CURRENT SCALE = 1.0 / 100.0
                                       How to scale image intensity to
                                       input currents
```

```
# Custom GeNN models
# Very simple integrate-and-fire neuron model
if_model = create_custom_neuron_class(
    "if_model",
    param_names=["Vthr"],
    var_name_types=[("V", "scalar"), ("SpikeCount", "unsigned int")],
    sim_code="$(V) += $(Isyn) * DT;",
    reset_code="""
    $(V) = 0.0;
    $(SpikeCount)++;
    """,
    threshold_condition_code="$(V) >= $(Vthr)")
```

```
Custom GeNN models
# Very simple integrate-and-fire neuron model
if model = create custom neuron class(
    "if model",
    param names=["Vthr"],
    var_name_types=[("V", "scalar"), ("SpikeCount", "unsigned int")],
    sim code="$(V) += $(Isyn) * DT;",
    reset code="""
    $(V) = 0.0;
    $(SpikeCount)++;
    threshold condition code="$(V) >= $(Vthr)
```

Parameters are common across all neurons in population (layer)

State variables used to track per-neuron membrane voltage and spike count

```
# Custom GeNN models
# Very simple integrate-and-fire neuron model
if_model = create_custom_neuron_class(
    "if_model",
    param_names=["Vthr"],
    var_name_types=[("V", "scalar"), ("SpikeCount", "unsigned int")],
    sim_code="$(V) += $(Isyn) * DT;",
    reset_code="""
    $(V) = 0.0;
    $(SpikeCount)++;
    """,
    threshold condition code="$(V) >= $(Vthr)
Neuron simply integrates input
    current $(Isyn) every timestep
```

Neuron spikes when \$(V) goes above \$(Vthr)

```
# Custom GeNN models
# Very simple integrate-and-fire neuron model
if model = create custom neuron class(
    "if model",
    param names=["Vthr"],
    var_name_types=[("V", "scalar"), ("SpikeCount", "unsigned int")],
    sim code="$(V) += $(Isyn) * DT;",
    reset code="""
    $(V) = 0.0;
    $(SpikeCount)++;
    threshold condition code="$(V) >= $(Vthr)")
# Current source model which injects current with a magnitude specified by a state variable
cs model = create custom current source class(
    "cs model",
    var name types=[("magnitude", "scalar")],
    injection code="$(injectCurrent, $(magnitude));")
```

Current source 'injects' current specified by state variable each timestep

```
Build model
# Create GeNN model
                                         Create new network using
model = GeNNModel("float", "tutorial_1")
                                         single-precision by default and
model.dT = TIMESTEP
                                         generating code into tutorial 1
# Load weights
weights = []
                                         directory
while True:
   filename = "weights_%u_%u.npy" % (len(weights), len(weights) + 1)
   if path.exists(filename):
       weights.append(np.load(filename))
   else:
       break
```

Load any weights present in directory into list

```
# Build model
# Create GeNN model
model = GeNNModel("float", "spiking eval")
model.dT = TIMESTEP
# Load weights
weights = []
while True:
   filename = "weights %u %u.npy" % (len(weights), len(weights) + 1)
   if path.exists(filename):
       weights.append(np.load(filename))
   else:
       break
# Initial values to initialise all neurons
                                        Initial values for all IF neurons
# Create first neuron layer
neuron layers = [model.add neuron population("neuron0", weights[0].shape[0],
                                         if model, IF PARAMS, if init)]
# Create subsequent neuron layer
for i, w in enumerate(weights)
                            Create neuron population with a neuron for
   neuron layers.append(model
                            each of the first layer's input dimensions
```

```
# Build model
# Create GeNN model
model = GeNNModel("float", "spiking eval")
model.dT = TIMESTEP
# Load weights
weights = []
while True:
    filename = "weights %u %u.npy" % (len(weights), len(weights) + 1)
    if path.exists(filename):
       weights.append(np.load(filename))
    else:
       break
# Initial values to initialise all neurons to
if init = {"V": 0.0, "SpikeCount":0}
# Create first neuron layer
neuron layers = [model.add neuron population("neuron0", weights[0].shape[0],
                                            if model, IF PARAMS, if init)]
# Create subsequent neuron layer
for i, w in enumerate(weights):
    neuron layers.append(model.add neuron population("neuron%u" % (i + 1),
                                                   w.shape[1], if model,
                                                   IF PARAMS, if init))
Create neuron populations matching subsequent layer's output
dimensions
```

Name of synapse population

Dense matrix with individual state variables (weights) for each synapse

http://genn-team.github.io/genn/document ation/4/html/d5/d39/subsect34.html

```
# Create synaptic connections between layers
for i, (pre, post, w) in enumerate(zip(neuron_layers[:-1], neuron_layers[1:], weights)):
    model.add_synapse_population(
        "synapse%u" % i, "DENSE_INDIVIDUALG", NO_DELAY,
        pre, post,
        "StaticPulse" {}, {"g": w.flatten()}, {}, {},
        "DeltaCurr", {}, {}, {})
have no parameters
```

Use built-in static synapse model

Initialise the weight of each synapse to the pre-trained weights

```
# Create synaptic connections between layers
for i, (pre, post, w) in enumerate(zip(neuron_layers[:-1], neuron_layers[1:], weights)):
    model.add_synapse_population(
        "synapse%u" % i, "DENSE_INDIVIDUALG", NO_DELAY,
        pre, post,
        "StaticPulse", {}, {"g": w.flatten()}, {},
        "DeltaCurr" ({}), {})
```

Use built-in delta postsynaptic model

This model has no parameters or variables



```
# Create synaptic connections between layers
for i, (pre, post, w) in enumerate(zip(neuron layers[:-1], neuron layers[1:], weights)):
   model.add synapse population(
       "synapse%u" % i, "DENSE INDIVI
                                   Add current source to provide
       pre, post,
       "StaticPulse", {}, {"g": w.fla input into first layer (neuron0)
       "DeltaCurr", {}, {})
# Create current source to deliver input to first layers of neurons
current input = model.add current source("current input", cs model,
                                      "neuron0", {}, {"magnitude": 0.0})
                                        Uses GeNN to generate
# Build and load our model
model.build() 	◀
                                        simulation code for model
model.load() 
                                        Loads generated model into
                                         PyGeNN
```

```
# Simulate
# -----
# Load testing data
testing_images = np.load("testing_images.npy")
testing_labels = np.load("testing_labels.npy")

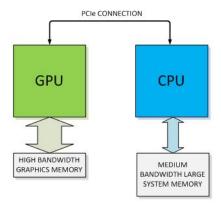
# Check dimensions match network
assert testing_images.shape[1] == weights[0].shape[0]
assert np.max(testing_labels) == (weights[1].shape[1] - 1)

# Set current input by scaling first image
```

Copy first image into memory view of current source magnitude

current_input.vars["magnitude"].view[:] = testing_images[0] * INPUT_CURRENT_SCALE

```
# Simulate
# Load testing data
testing images = np.load("testing_images.npy")
testing labels = np.load("testing labels.npy")
# Check dimensions match network
assert testing images.shape[1] == weights[0].shape[0]
assert np.max(testing labels) == (weights[1].shape[1] - 1)
# Set current input by scaling first image
current input.vars["magnitude"].view[:] = testing images[0] * INPUT CURRENT SCALE
# Upload
current input.push var to device("magnitude")
                   Upload this variable to GPU
```

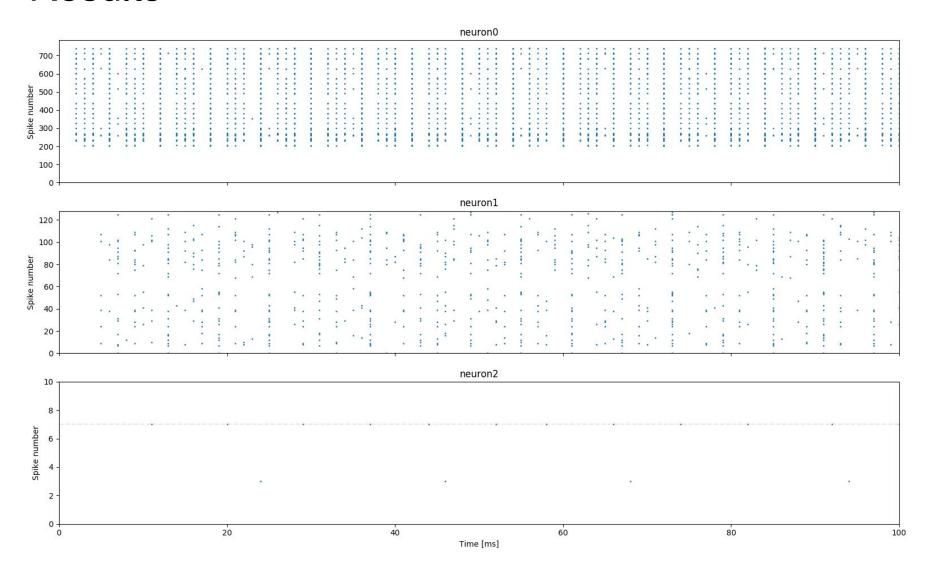


```
# Simulate
layer_spikes = [[] for _ in enumerate(neuron layers)]
while model.timestep < PRESENT TIMESTEPS:</pre>
   # Advance simulation
   model.step time() 	◄
                                                Advance simulation
   # Loop through neuron layers
                                                           Download spikes
   for i, 1 in enumerate(neuron_layers):
       # Download spikes
                                                           emitted this timestep
       1.pull current spikes from device()
                                                           from GPU
       # Add to data structure
       layer spikes[i].append(np.copy(1.current spikes))
                                                           Add spikes to data
                                                           structure
```

Time [ms]	0	0	1	1	2
Neuron index	1	14	13	100	56

```
# -----
# Plotting
import matplotlib.pyplot as plt
# Create a plot with axes for each
fig, axes = plt.subplots(len(neuron layers), sharex=True)
# Loop through axes and their corresponding neuron populations
for a, s, 1 in zip(axes, layer spikes, neuron layers):
    spike ids = np.concatenate(spk)
    spike times = np.concatenate([np.ones like(s) * i * TIMESTEP
                               for i, s in enumerate(spk)])
   # Plot spikes
   a.scatter(spike times, spike ids, s=1)
   # Set title, axis labels
   a.set title(1.name)
   a.set ylabel("Spike number")
   a.set xlim((0, PRESENT_TIMESTEPS * TIMESTEP))
    a.set ylim((0, 1.size))
# Add an x-axis label and translucent line showing the correct label
axes[-1].set xlabel("Time [ms]")
axes[-1].hlines(testing labels[0], xmin=0, xmax=PRESENT TIMESTEPS,
               linestyle="--", color="gray", alpha=0.2)
# Show plot
plt.show()
```

Results



Part 2: Evaluating entire dataset

- 1. Presenting entire MNIST testing set to network
- 2. Calculating inference performance

```
# Simulate
# -----
# Load testing data
testing_images = np.load("testing_images.npy")
testing_labels = np.load("testing_labels.npy")

# Check dimensions match network
assert testing_images.shape[1] == weights[0].shape[0]
assert np.max(testing_labels) == (weights[1].shape[1] - 1)

# Get views to efficiently access state variables
current_input_magnitude = current_input.vars["magnitude"].view
output_spike_count = neuron_layers[-1].vars["SpikeCount"].view
layer_voltages = [1.vars["V"].view for 1 in neuron_layers]
```

Cache memory views of required state variables

```
# Simulate
num correct = 0
while model.timestep < (PRESENT TIMESTEPS * testing images.shape[0]):</pre>
    # Calculate the timestep within the presentation
                                                             Divide timestep into
   timestep in example = model.timestep % PRESENT TIMESTEPS
    example = int(model.timestep // PRESENT TIMESTEPS) >
                                                             example index and
   # If this is the first timestep of presenting the example
                                                             timestep within
    if timestep in example == 0:
                                                             example
        current input magnitude[:] = testing images[example]
        current input.push var to device("magnitude")
        # Loop through all layers and their corresponding voltage views
        for 1, v in zip(neuron layers, layer voltages):
           # Manually 'reset' voltage
           v[:] = 0.0
           # Upload
           1.push var to device("V")
        # Zero spike count
        output spike count[:] = 0
        neuron layers[-1].push var to device("SpikeCount")
    # Advance simulation
   model.step time()
```

```
# Simulate
num correct = 0
while model.timestep < (PRESENT TIMESTEPS * testing images.shape[0]):</pre>
   # Calculate the timestep within the presentati
   timestep_in_example = model.timestep % PRESENT Copy image into memory
   example = int(model.timestep // PRESENT TIMEST
                                                 view and upload to GPU
   # If this is the first timestep of presenting the example
    if timestep in example == 0:
       current_input_magnitude[:] = testing_images[example] * INPUT CURRENT SCALE
       current input.push var to device("magnitude")
       # Loop through all layers and their corresponding voltage views
       for 1, v in zip(neuron layers, layer voltages):
           # Manually 'reset' voltage
           v[:] = 0.0 ←
                                             Set all neuron voltages to zero
           # Upload
           1.push var to device("V")
                                                       Upload to GPU
       # Zero spike count
       output spike count[:] = 0
       neuron layers[-1].push var to device("SpikeCount")
   # Advance simulation
   model.step time()
```

```
# Simulate
num correct = 0
while model.timestep < (PRESENT TIMESTEPS * testing images.shape[0]):</pre>
   # Calculate the timestep within the presentation
   timestep in example = model.timestep % PRESENT TIMESTEPS
   example = int(model.timestep // PRESENT TIMESTEPS)
   # If this is the first timestep of presenting the example
    if timestep in example == 0:
       current_input_magnitude[:] = testing_images[example] * INPUT_CURRENT_SCALE
       current input.push var to device("magnitude")
       # Loop through all layers and their corresponding voltage views
       for 1, v in zip(neuron layers, layer voltages):
           # Manually 'reset' voltage
                                                       Zero spike count for all
           v[:] = 0.0
                                                       output layer neurons
           # Upload
                                                       and upload to GPU
           1.push var to device("V")
       # Zero spike count
       output spike count[:] = 0
       neuron layers[-1].push var to device("SpikeCount")
   # Advance simulation
   Advance simulation
```

```
# If this is the LAST timestep of presenting the example
   if timestep in example == (PRESENT TIMESTEPS - 1):
       # Download spike count from last layer
       neuron layers[-1].pull var from device("SpikeCount")
       # Find which neuron spiked the most to get prediction
                                                             Download output
       predicted label = np.argmax(output spike count)
                                                             layer spike count
       true label = testing labels[example]
       print("\tExample=%u, true label=%u, predicted label=%u" % (example,
                                                                  true label,
                                                                  predicted label))
       if predicted label == true label:
           num correct += 1
print("Accuracy %f%%" % ((num correct / float(testing images.shape[0])) * 100.0))
```

Results

```
Example=9965, true label=3, predicted label=3
       Example=9966, true label=6, predicted label=6
       Example=9967, true label=8, predicted label=8
       Example=9968, true label=7, predicted label=7
       Example=9969, true label=1, predicted label=1
       Example=9970, true label=5, predicted label=5
       Example=9971, true label=2, predicted label=2
       Example=9972, true label=4, predicted label=4
        Example=9973, true label=9, predicted label=9
       Example=9974, true label=4, predicted label=4
        Example=9975, true label=3, predicted label=3
       Example=9976, true label=6, predicted label=6
       Example=9977, true label=4, predicted label=4
       Example=9978, true label=1, predicted label=1
       Example=9979, true label=7, predicted label=7
       Example=9980, true label=2, predicted label=2
       Example=9981, true label=6, predicted label=6
       Example=9982, true label=5, predicted label=5
       Example=9983, true label=0, predicted label=0
       Example=9984, true label=1, predicted label=1
        Example=9985, true label=2, predicted label=2
       Example=9986, true label=3, predicted label=3
       Example=9987, true label=4, predicted label=4
       Example=9988, true label=5, predicted label=5
       Example=9989, true label=6, predicted label=6
       Example=9990, true label=7, predicted label=7
       Example=9991, true label=8, predicted label=8
       Example=9992, true label=9, predicted label=9
       Example=9993, true label=0, predicted label=0
       Example=9994, true label=1, predicted label=1
       Example=9995, true label=2, predicted label=2
       Example=9996, true label=3, predicted label=3
       Example=9997, true label=4, predicted label=4
       Example=9998, true label=5, predicted label=5
       Example=9999, true label=6, predicted label=6
Accuracy 97.440000%
tensortlow) 1K421Ginf900801:~/offline train example$
```

Part 3: Play time!

- How does PRESENT_TIMESTEPS affect performance?
- Can you reduce the number of spikes while maintaining performance by modifying IF_PARAMS and INPUT_CURRENT_SCALE?
- Try training your own sequential model with dense layers,
 save the weights and see how this code performs

Thank you!

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