Report 3

Computational Neuroscience

Computer Assignment 3

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```
In [1]: %matplotlib notebook import torch import numpy as np

In [2]: from cnsproject.network.neural_populations import LIFPopulation from cnsproject.plotting.plotting import plotting from cnsproject.utils import step_function, two_way_step_function, random_step_function from cnsproject.network.monitors import Monitor from cnsproject.network.connections import DenseConnection
```

Global Variables

time parameter shows how often (seconds*scale/dt) we want to run our neuron. dt means with what resolution (scale) we want our seconds move forward. I used 10 neurons for start and split them into two part which first part representing the excitatory population with 0.8*N number of neurons and the second part that is representing the inhibitory population has 0.2*N neurons.

```
In [3]: time = 1500
    scale = 100
    dt = 1
    neuron_size = 10
    shape1 = (int(neuron_size*0.8),)
    shape2 = (int(neuron_size*0.2),)
```

Walkthrough

The default implemented model is a fully connected and homogeneous population.

First we need to specify the input current.

```
In [4]: I1 = step_function(time = time, I_value = 3, scale = scale)
I2 = I1
```

Then two LIF Population created withe shapes that we talked about. np1 population is the excitatory population and np2 is the inhibitory population.

Here we creat connection between the two population.

For monitoring the 2 population, two Monitor created.

```
In [7]: monitor1 = Monitor(pn1, state_variables=["s", "u"])
    monitor1.set_time_steps(time, dt)
    monitor1.reset_state_variables()

monitor2 = Monitor(pn2, state_variables=["s", "u"])
    monitor2.set_time_steps(time, dt)
    monitor2.reset_state_variables()
```

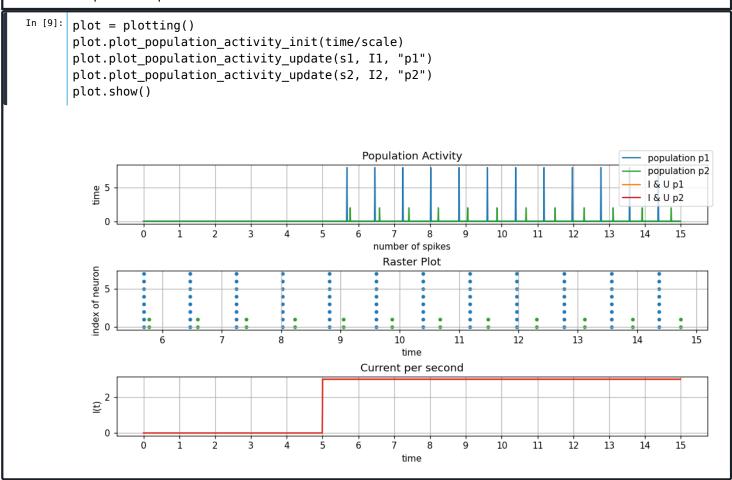
In this step first one population starts and will forward but before the next population forwarding, we need to compute the current that effects the other populations. This will continue till the end.

```
In [8]: I_pop=0
for i in range(len(II)):
    pn1.forward(II[i]-I_pop)
    I_pop = con1.compute()
    pn2.forward(I2[i]-I_pop)
    I_pop = con2.compute()
    monitor1.record()
    monitor2.record()

s1 = torch.transpose(monitor1.get("s")*1, 0, 1)
s2 = torch.transpose(monitor2.get("s")*1, 0, 1)

//home/aref/Desktop/projects/UT/Computational_Neuroscience/package/cnsproject/network/neural_populations.py:190: UserWarning:
    To copy construct from a tensor, it is recommended to use sourceTensor.clone().detach() or sourceTensor.clone().detach().req
    uires_grad_(True), rather than torch.tensor(sourceTensor).
    self.dt = torch.tensor(self.dt)
```

In the end lets plot the output.

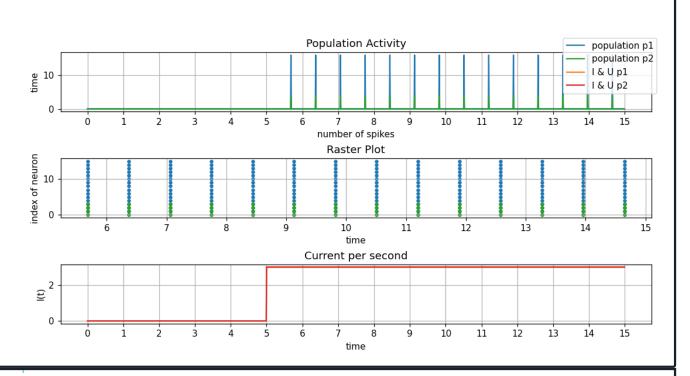


Population Behavior

Isolate Populations

It's better to see the population activity when there is no connection between them.

```
In [24]:
       time = 1500
       scale = 100
       dt = 1
       neuron_size = 20
       shape1 = (int(neuron_size*0.8),)
       shape2 = (int(neuron size*0.2),)
       I1 = step_function(time = time, I_value = 3, scale = scale)
       I2 = I1
       pn1 = LIFPopulation(
               shape = shape1, spike_trace = True, additive_spike_trace = True, tau_s = 10, tra
       ce scale = 1.,
               is_inhibitory = False, learning = False, R = 10, C = 10, threshold = -55
       pn1.dt = dt
       pn2 = LIFPopulation(
               shape = shape2, spike_trace = True, additive_spike_trace = True, tau_s = 10, tra
       ce_scale = 1.,
               is_inhibitory = False, learning = False, R = 10, C = 10, threshold = -55
       pn2.dt = dt
       monitor1 = Monitor(pn1, state_variables=["s", "u"])
       monitor1.set time steps(time, dt)
       monitor1.reset_state_variables()
       monitor2 = Monitor(pn2, state_variables=["s", "u"])
       monitor2.set_time_steps(time, dt)
       monitor2.reset_state_variables()
       for i in range(len(I1)):
           pn1.forward(I1[i])
           pn2.forward(I2[i])
           monitor1.record()
           monitor2.record()
       s1 = torch.transpose(monitor1.get("s")*1, 0, 1)
       s2 = torch.transpose(monitor2.get("s")*1, 0, 1)
       plot = plotting()
       plot.plot_population_activity_init(time/scale)
       plot.plot_population_activity_update(s1, I1, "p1")
       plot.plot_population_activity_update(s2, I2, "p2")
       plot.show()
```

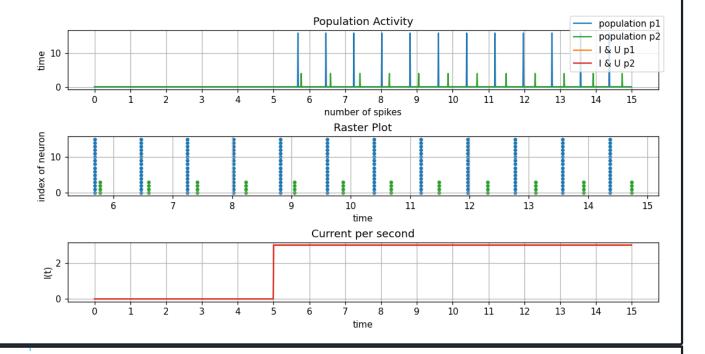


Connected Populations

In this part we have the two same populations from previous part, but the only difference is they are connected. the second population is inhibitory. As we can see the effect of connection is showing that the populations have distance and they are getting farther becaus I used a parameter ($trace_scale$) to achieve the additive behavior.

```
In [26]:
       time = 1500
       scale = 100
       dt = 1
       neuron_size = 20
       shape1 = (int(neuron_size*0.8),)
       shape2 = (int(neuron size*0.2),)
       I1 = step_function(time = time, I_value = 3, scale = scale)
       I2 = I1
       pn1 = LIFPopulation(
               shape = shape1, spike_trace = True, additive_spike_trace = True, tau_s = 10, tra
       ce scale = 1.,
               is_inhibitory = False, learning = False, R = 10, C = 10, threshold = -55
       pn1.dt = dt
       pn2 = LIFPopulation(
               shape = shape2, spike_trace = True, additive_spike_trace = True, tau_s = 10, tra
       ce_scale = 1.,
               is_inhibitory = False, learning = False, R = 10, C = 10, threshold = -55
       pn2.dt = dt
       con1 = DenseConnection(
               pre = pn1, post = pn2, lr = None, weight decay = 0.0,
               J = 2, tau_s = 10, trace_scale = 1., dt = dt
       con1.dt = dt
       con2 = DenseConnection(
               pre = pn2, post = pn1, lr = None, weight_decay = 0.0,
               J = 2, tau_s = 10, trace_scale = 1., dt = dt
       con2.dt = dt
       monitor1 = Monitor(pn1, state_variables=["s", "u"])
       monitor1.set time steps(time, dt)
       monitor1.reset_state_variables()
       monitor2 = Monitor(pn2, state_variables=["s", "u"])
       monitor2.set_time_steps(time, dt)
       monitor2.reset_state_variables()
       I pop=0
       for i in range(len(I1)):
           pn1.forward(I1[i]-I_pop)
           I pop = conl.compute()
           pn2.forward(I2[i]-I_pop)
           I_pop = con2.compute()
           monitor1.record()
           monitor2.record()
       s1 = torch.transpose(monitor1.get("s")*1, 0, 1)
       s2 = torch.transpose(monitor2.get("s")*1, 0, 1)
       plot = plotting()
       plot.plot_population_activity_init(time/scale)
       plot.plot_population_activity_update(s1, I1, "p1")
       plot.plot_population_activity_update(s2, I2, "p2")
       plot.show()
```

/home/aref/Desktop/projects/UT/Computational_Neuroscience/package/cnsproject/network/neural_populations.py:190: UserWarning:
To copy construct from a tensor, it is recommended to use sourceTensor.clone().detach() or sourceTensor.clone().detach().req
uires_grad_(True), rather than torch.tensor(sourceTensor).
 self.dt = torch.tensor(self.dt)



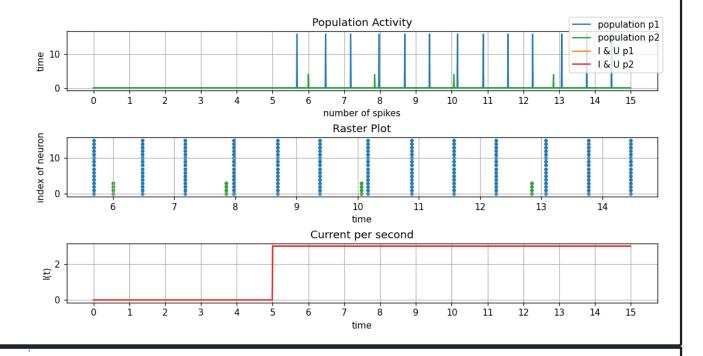
Affect of tau_s

In this part the effect of tau_s is clear. This parameter affects in the decaying, and increasing it will cause the decay to become zero.

```
In [12]: time = 1500
       scale = 100
       dt = 1
       neuron_size = 20
       shape1 = (int(neuron_size*0.8),)
       shape2 = (int(neuron_size*0.2),)
       I1 = step_function(time = time, I_value = 3, scale = scale)
       I2 = I1
       pn1 = LIFPopulation(
               shape = shape1, spike_trace = True, additive_spike_trace = True, tau_s = 10, tra
       ce_scale = 1.,
               is_inhibitory = False, learning = False, R = 10, C = 10, threshold = -55
       pn1.dt = dt
       pn2 = LIFPopulation(
               shape = shape2, spike_trace = True, additive_spike_trace = True, tau_s = 10, tra
       ce_scale = 1.,
               is_inhibitory = False, learning = False, R = 10, C = 10, threshold = -55
       pn2.dt = dt
```

```
con1 = DenseConnection(
        pre = pn1, post = pn2, lr = None, weight_decay = 0.0,
        J = 2, tau_s = 50, trace_scale = 1., dt = dt
con1.dt = dt
con2 = DenseConnection(
        pre = pn2, post = pn1, lr = None, weight_decay = 0.0,
        J = 2, tau s = 10, trace scale = 1., dt = dt
   )
con2.dt = dt
monitor1 = Monitor(pn1, state_variables=["s", "u"])
monitor1.set_time_steps(time, dt)
monitor1.reset_state_variables()
monitor2 = Monitor(pn2, state_variables=["s", "u"])
monitor2.set_time_steps(time, dt)
monitor2.reset_state_variables()
I pop=0
for i in range(len(I1)):
   pn1.forward(I1[i]-I_pop)
   I_pop = con1.compute()
   pn2.forward(I2[i]-I_pop)
   I_pop = con2.compute()
   monitor1.record()
   monitor2.record()
s1 = torch.transpose(monitor1.get("s")*1, 0, 1)
s2 = torch.transpose(monitor2.get("s")*1, 0, 1)
plot = plotting()
plot.plot_population_activity_init(time/scale)
plot.plot_population_activity_update(s1, I1, "p1")
plot.plot_population_activity_update(s2, I2, "p2")
plot.show()
```

/home/aref/Desktop/projects/UT/Computational_Neuroscience/package/cnsproject/network/neural_populations.py:190: UserWarning:
To copy construct from a tensor, it is recommended to use sourceTensor.clone().detach() or sourceTensor.clone().detach().req
uires_grad_(True), rather than torch.tensor(sourceTensor).
 self.dt = torch.tensor(self.dt)

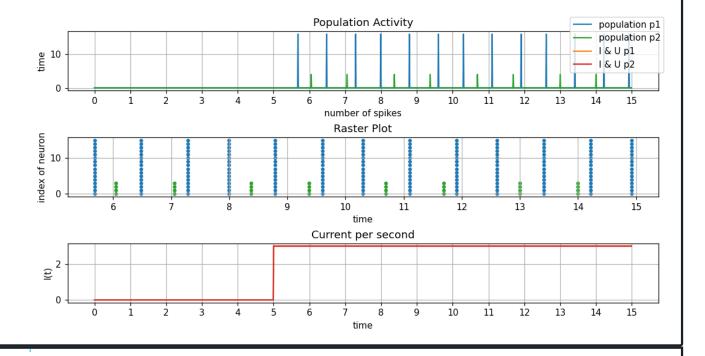


Affect of $trace_scale$

As I said before, $trace_scale$ will change the additivity of the trace. By increasing this parameter, the adaptation of these neurons will become easier.

```
In [15]:
       time = 1500
       scale = 100
       dt = 1
       neuron_size = 20
       shape1 = (int(neuron_size*0.8),)
       shape2 = (int(neuron size*0.2),)
       I1 = step_function(time = time, I_value = 3, scale = scale)
       I2 = I1
       pn1 = LIFPopulation(
               shape = shape1, spike_trace = True, additive_spike_trace = True, tau_s = 10, tra
       ce scale = 1.,
               is_inhibitory = False, learning = False, R = 10, C = 10, threshold = -55
       pn1.dt = dt
       pn2 = LIFPopulation(
               shape = shape2, spike_trace = True, additive_spike_trace = True, tau_s = 10, tra
       ce_scale = 1.,
               is_inhibitory = False, learning = False, R = 10, C = 10, threshold = -55
       pn2.dt = dt
       con1 = DenseConnection(
               pre = pn1, post = pn2, lr = None, weight decay = 0.0,
               J = 2, tau_s = 10, trace_scale = 3, dt = dt
       con1.dt = dt
       con2 = DenseConnection(
               pre = pn2, post = pn1, lr = None, weight_decay = 0.0,
               J = 2, tau_s = 10, trace_scale = 1., dt = dt
       con2.dt = dt
       monitor1 = Monitor(pn1, state_variables=["s", "u"])
       monitor1.set time steps(time, dt)
       monitor1.reset_state_variables()
       monitor2 = Monitor(pn2, state_variables=["s", "u"])
       monitor2.set_time_steps(time, dt)
       monitor2.reset_state_variables()
       I pop=0
       for i in range(len(I1)):
           pn1.forward(I1[i]-I_pop)
           I pop = conl.compute()
           pn2.forward(I2[i]-I_pop)
           I_pop = con2.compute()
           monitor1.record()
           monitor2.record()
       s1 = torch.transpose(monitor1.get("s")*1, 0, 1)
       s2 = torch.transpose(monitor2.get("s")*1, 0, 1)
       plot = plotting()
       plot.plot_population_activity_init(time/scale)
       plot.plot_population_activity_update(s1, I1, "p1")
       plot.plot_population_activity_update(s2, I2, "p2")
       plot.show()
```

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 self.dt = torch.tensor(self.dt)

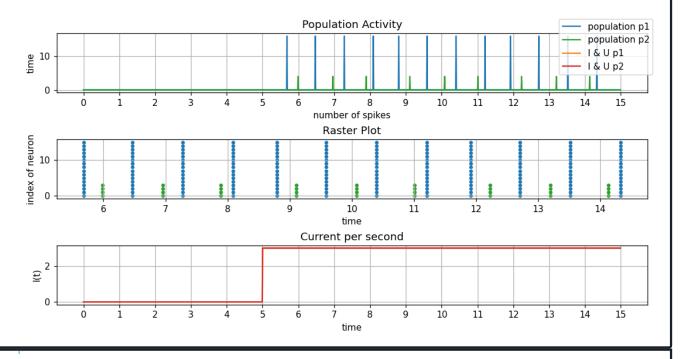


Affect of J

The J parameter has a linear relationship with the weights of connection between each pair of neurons. By increasing this parameter the effect of the other population will increase. As we can see the distance between the populations gets longer.

```
In [16]:
       time = 1500
       scale = 100
       dt = 1
       neuron_size = 20
       shape1 = (int(neuron_size*0.8),)
       shape2 = (int(neuron size*0.2),)
       I1 = step_function(time = time, I_value = 3, scale = scale)
       I2 = I1
       pn1 = LIFPopulation(
               shape = shape1, spike_trace = True, additive_spike_trace = True, tau_s = 10, tra
       ce scale = 1.,
               is inhibitory = False, learning = False, R = 10, C = 10, threshold = -55
       pn1.dt = dt
       pn2 = LIFPopulation(
               shape = shape2, spike_trace = True, additive_spike_trace = True, tau_s = 10, tra
       ce_scale = 1.,
               is_inhibitory = False, learning = False, R = 10, C = 10, threshold = -55
       pn2.dt = dt
       con1 = DenseConnection(
               pre = pn1, post = pn2, lr = None, weight decay = 0.0,
               J = 5, tau_s = 10, trace_scale = 1, dt = dt
       con1.dt = dt
       con2 = DenseConnection(
               pre = pn2, post = pn1, lr = None, weight_decay = 0.0,
               J = 2, tau_s = 10, trace_scale = 1., dt = dt
       con2.dt = dt
       monitor1 = Monitor(pn1, state_variables=["s", "u"])
       monitor1.set time steps(time, dt)
       monitor1.reset_state_variables()
       monitor2 = Monitor(pn2, state_variables=["s", "u"])
       monitor2.set_time_steps(time, dt)
       monitor2.reset_state_variables()
       I pop=0
       for i in range(len(I1)):
           pn1.forward(I1[i]-I_pop)
           I pop = conl.compute()
           pn2.forward(I2[i]-I_pop)
           I_pop = con2.compute()
           monitor1.record()
           monitor2.record()
       s1 = torch.transpose(monitor1.get("s")*1, 0, 1)
       s2 = torch.transpose(monitor2.get("s")*1, 0, 1)
       plot = plotting()
       plot.plot_population_activity_init(time/scale)
       plot.plot_population_activity_update(s1, I1, "p1")
       plot.plot_population_activity_update(s2, I2, "p2")
       plot.show()
```

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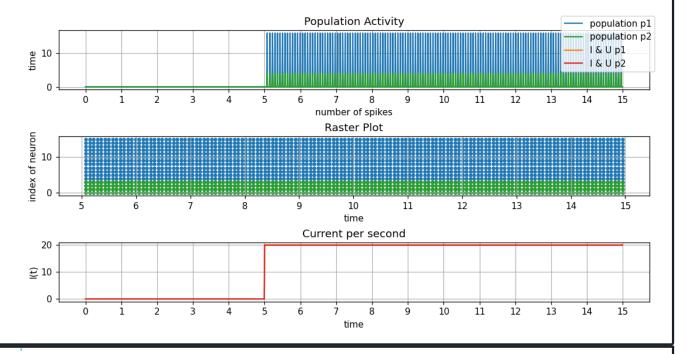


Affect of excitatory

If we change the nature (excitatory and inhibitory) of a population, as we expected, they amplify each other.

```
In [23]:
       time = 1500
       scale = 100
       dt = 1
       neuron_size = 20
       shape1 = (int(neuron_size*0.8),)
       shape2 = (int(neuron size*0.2),)
       I1 = step_function(time = time, I_value = 20, scale = scale)
       I2 = I1
       pn1 = LIFPopulation(
               shape = shape1, spike trace = True, additive spike trace = True, tau s = 10, tra
       ce scale = 1.,
               is_inhibitory = False, learning = False, R = 10, C = 10, threshold = -55
       pn1.dt = dt
       pn2 = LIFPopulation(
               shape = shape2, spike_trace = True, additive_spike_trace = True, tau_s = 10, tra
       ce_scale = 1.,
               is_inhibitory = False, learning = False, R = 10, C = 10, threshold = -55
       pn2.dt = dt
       con1 = DenseConnection(
               pre = pn1, post = pn2, lr = None, weight decay = 0.0,
               J = 2, tau_s = 10, trace_scale = 1, dt = dt
       con1.dt = dt
       con2 = DenseConnection(
               pre = pn2, post = pn1, lr = None, weight_decay = 0.0,
               J = 2, tau_s = 10, trace_scale = 1., dt = dt
       con2.dt = dt
       monitor1 = Monitor(pn1, state_variables=["s", "u"])
       monitor1.set time steps(time, dt)
       monitor1.reset_state_variables()
       monitor2 = Monitor(pn2, state_variables=["s", "u"])
       monitor2.set_time_steps(time, dt)
       monitor2.reset state variables()
       I pop=0
       for i in range(len(I1)):
           pn1.forward(I1[i]+I_pop)
           I pop = conl.compute()
           pn2.forward(I2[i]+I_pop)
           I_pop = con2.compute()
           monitor1.record()
           monitor2.record()
       s1 = torch.transpose(monitor1.get("s")*1, 0, 1)
       s2 = torch.transpose(monitor2.get("s")*1, 0, 1)
       plot = plotting()
       plot.plot_population_activity_init(time/scale)
       plot.plot_population_activity_update(s1, I1, "p1")
       plot.plot_population_activity_update(s2, I2, "p2")
       plot.show()
```

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 self.dt = torch.tensor(self.dt)

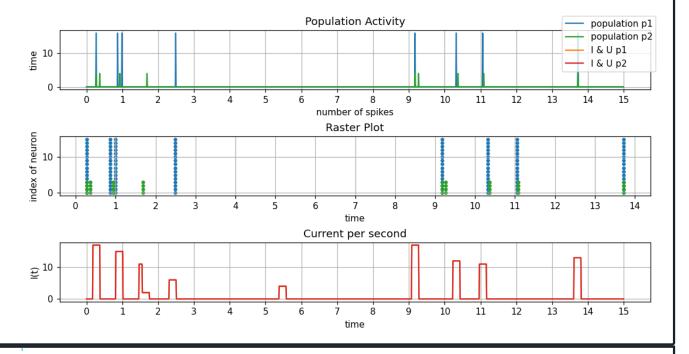


Random Input

If we change the input current to a random function, we can observe some behavior like below.

```
In [20]:
       time = 1500
       scale = 100
       dt = 1
       neuron_size = 20
       shape1 = (int(neuron_size*0.8),)
       shape2 = (int(neuron size*0.2),)
       I1 = random_step_function(time = time, I_value = 20, scale = scale)
       I2 = I1
       pn1 = LIFPopulation(
               shape = shape1, spike_trace = True, additive_spike_trace = True, tau_s = 10, tra
       ce scale = 1.,
               is_inhibitory = False, learning = False, R = 10, C = 10, threshold = -55
       pn1.dt = dt
       pn2 = LIFPopulation(
               shape = shape2, spike_trace = True, additive_spike_trace = True, tau_s = 10, tra
       ce_scale = 1.,
               is_inhibitory = False, learning = False, R = 10, C = 10, threshold = -55
       pn2.dt = dt
       con1 = DenseConnection(
               pre = pn1, post = pn2, lr = None, weight decay = 0.0,
               J = 2, tau_s = 10, trace_scale = 1, dt = dt
       con1.dt = dt
       con2 = DenseConnection(
               pre = pn2, post = pn1, lr = None, weight_decay = 0.0,
               J = 2, tau_s = 10, trace_scale = 1., dt = dt
       con2.dt = dt
       monitor1 = Monitor(pn1, state_variables=["s", "u"])
       monitor1.set time steps(time, dt)
       monitor1.reset_state_variables()
       monitor2 = Monitor(pn2, state_variables=["s", "u"])
       monitor2.set_time_steps(time, dt)
       monitor2.reset_state_variables()
       I pop=0
       for i in range(len(I1)):
           pn1.forward(I1[i]-I_pop)
           I pop = conl.compute()
           pn2.forward(I2[i]-I_pop)
           I_pop = con2.compute()
           monitor1.record()
           monitor2.record()
       s1 = torch.transpose(monitor1.get("s")*1, 0, 1)
       s2 = torch.transpose(monitor2.get("s")*1, 0, 1)
       plot = plotting()
       plot.plot_population_activity_init(time/scale)
       plot.plot_population_activity_update(s1, I1, "p1")
       plot.plot_population_activity_update(s2, I2, "p2")
       plot.show()
```

/home/aref/Desktop/projects/UT/Computational_Neuroscience/package/cnsproject/network/neural_populations.py:190: UserWarning:
To copy construct from a tensor, it is recommended to use sourceTensor.clone().detach() or sourceTensor.clone().detach().req
uires_grad_(True), rather than torch.tensor(sourceTensor).
 self.dt = torch.tensor(self.dt)



Two Random Current

If we change the second population's current, the population is no longer Homogeneous and became hetergeneous. Lets have a berief and short review of this change.

```
In [28]:
       time = 1500
       scale = 100
       dt = 1
       neuron_size = 20
       shape1 = (int(neuron_size*0.8),)
       shape2 = (int(neuron size*0.2),)
       I1 = random_step_function(time = time, I_value = 20, scale = scale)
       I2 = random step function(time = time, I value = 20, scale = scale)
       pn1 = LIFPopulation(
               shape = shape1, spike_trace = True, additive_spike_trace = True, tau_s = 10, tra
       ce scale = 1.,
               is inhibitory = False, learning = False, R = 10, C = 10, threshold = -55
       pn1.dt = dt
       pn2 = LIFPopulation(
               shape = shape2, spike_trace = True, additive_spike_trace = True, tau_s = 10, tra
       ce_scale = 1.,
               is_inhibitory = False, learning = False, R = 10, C = 10, threshold = -55
       pn2.dt = dt
       con1 = DenseConnection(
               pre = pn1, post = pn2, lr = None, weight decay = 0.0,
               J = 2, tau_s = 10, trace_scale = 1, dt = dt
       con1.dt = dt
       con2 = DenseConnection(
               pre = pn2, post = pn1, lr = None, weight_decay = 0.0,
               J = 2, tau_s = 10, trace_scale = 1., dt = dt
       con2.dt = dt
       monitor1 = Monitor(pn1, state_variables=["s", "u"])
       monitor1.set time steps(time, dt)
       monitor1.reset_state_variables()
       monitor2 = Monitor(pn2, state_variables=["s", "u"])
       monitor2.set_time_steps(time, dt)
       monitor2.reset_state_variables()
       I pop=0
       for i in range(len(I1)):
           pn1.forward(I1[i]-I_pop)
           I pop = conl.compute()
           pn2.forward(I2[i]-I_pop)
           I_pop = con2.compute()
           monitor1.record()
           monitor2.record()
       s1 = torch.transpose(monitor1.get("s")*1, 0, 1)
       s2 = torch.transpose(monitor2.get("s")*1, 0, 1)
       plot = plotting()
       plot.plot_population_activity_init(time/scale)
       plot.plot_population_activity_update(s1, I1, "p1")
       plot.plot_population_activity_update(s2, I2, "p2")
       plot.show()
```

