

# Pruning Multiple neurons at one play

March 30, 2017

Compute the performance of MAB methods of pruning Multiple neurons at one time  
MAP for choosing multi arms at one time

```
In [1]: import numpy as np
import time
import sys
from numpy import *
import matplotlib.pyplot as plt
from sklearn import metrics
%matplotlib inline
plt.rcParams['figure.figsize'] = (15, 6)
```

## 1 Load Bokeh

```
In [2]: from bokeh.layouts import row, gridplot
from bokeh.plotting import figure, output_notebook, show
from bokeh.models import Legend
TOOLS = 'box_zoom,box_select,crosshair,resize,reset,lasso_select,pan,save,poly_select,tap'
output_notebook()
```

## 2 Load the data

```
In [3]: X_train = np.load('./valley/X_train.npy')
y_train = np.load('./valley/y_train.npy')
X_test = np.load('./valley/X_test.npy')
y_test = np.load('./valley/y_test.npy')
X_deploy = np.load('./valley/X_deploy.npy')
y_deploy = np.load('./valley/y_deploy.npy')
print('Number of training examples',len(X_train))
print('Number of validation examples',len(X_test))
print('Number of testing examples',len(X_deploy))
```

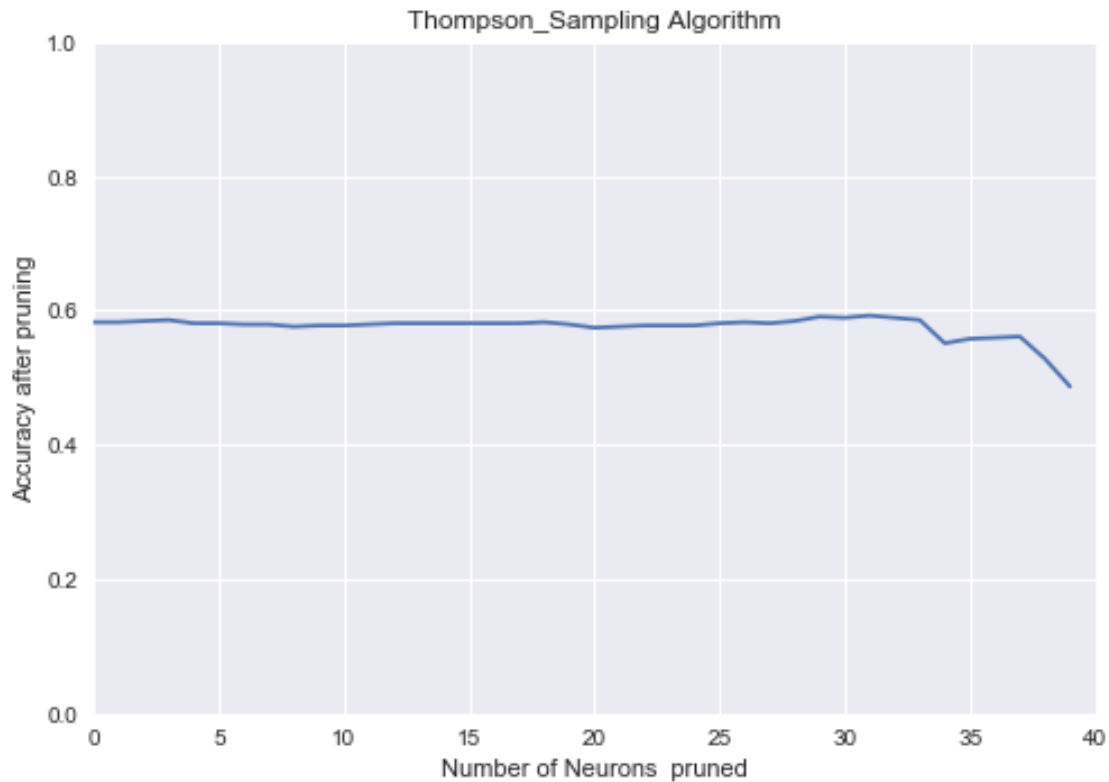
Number of training examples 484  
Number of validation examples 122  
Number of testing examples 606

```
In [4]: exec(open("core.py").read()) # python 3x
```

```
In [5]: algo = Thompson_Sampling([], [])
        Alg_name = 'Thompson_Sampling Algorithm'
        path = './Thompson_Sampling/'
        sys.path.append("./Thompson_Sampling")
        exec(open("mnist_cnnFORTESTING.py").read())
```

[illegible]

Test after pruning= 0.59  
 Test after pruning= 0.59  
 Test after pruning= 0.55  
 Test after pruning= 0.56  
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 Test after pruning= 0.53  
 Test after pruning= 0.49



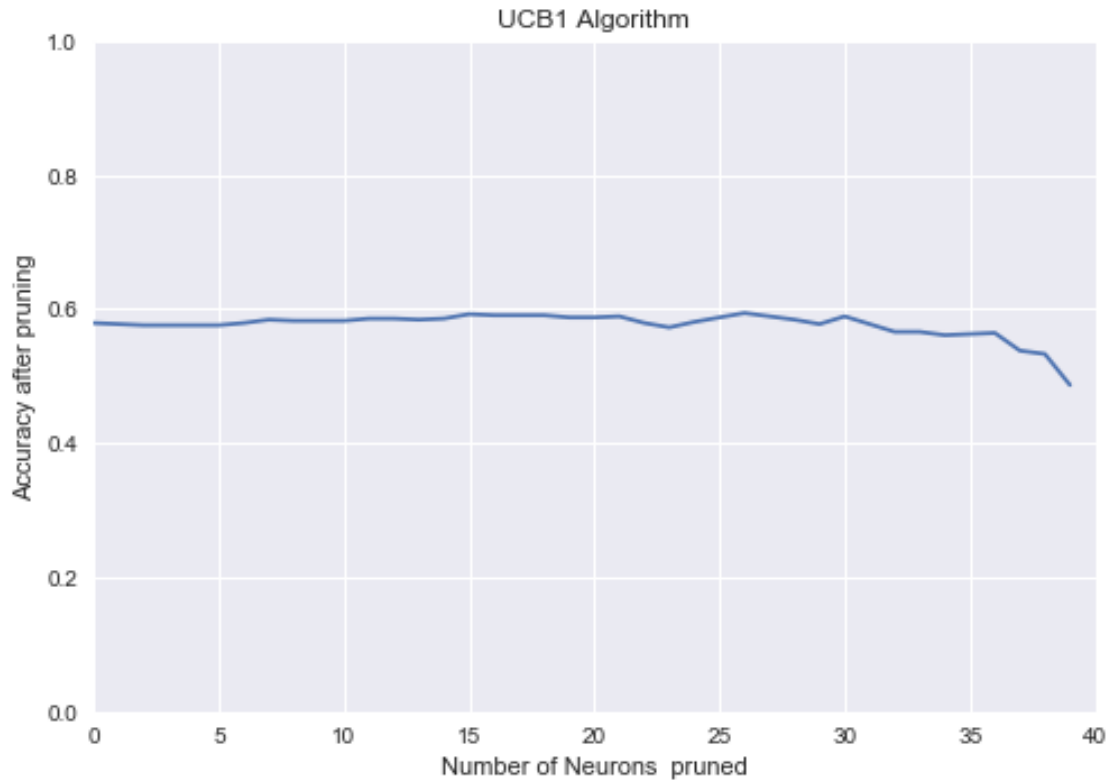
## 2.2 Run UCB1 pruning Algorithm

```

In [6]: algo = UCB1([], [])
        Alg_name = 'UCB1 Algorithm'
        path = './UCB1/'
        sys.path.append("./UCB1")
        exec(open("mnist_cnnFORTESTING.py").read())
  
```

Test fraction correct (NN-Score) = 0.70  
 Test fraction correct (NN-Accuracy) = 0.58  
 The time for running this method is 8.914093971252441 seconds  
 Finsh playing start pruning:

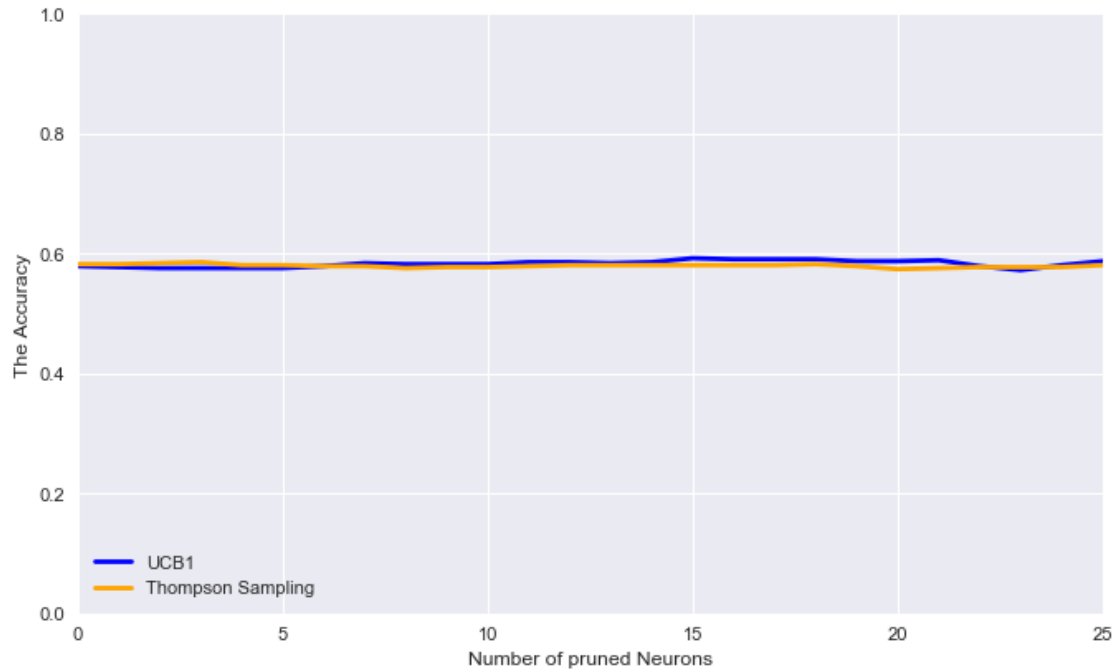
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Test after pruning= 0.49



### 3 Compare the accuracy

```
In [7]: ucb1 = np.load('./UCB1/AccuracyAfterPrune.npy')
        ThompsonSampling = np.load('./Thompson_Sampling/AccuracyAfterPrune.npy')
        Accuracy = np.load('AccuracyBeforePruning.npy')
```

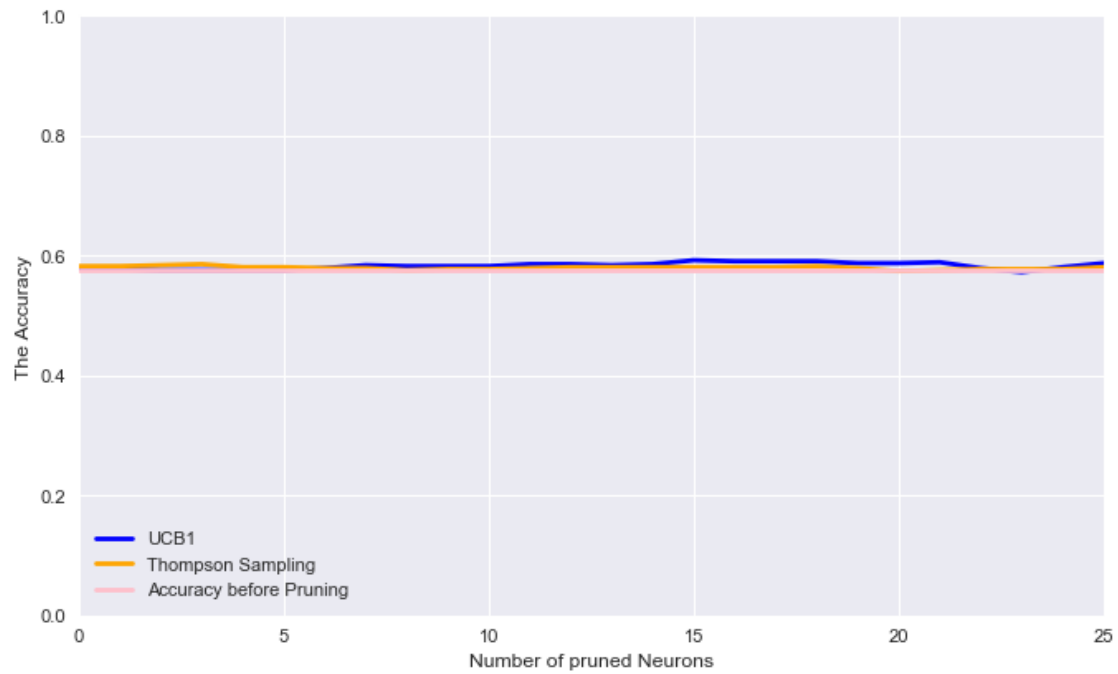
```
In [8]: fig = plt.figure(figsize=(10, 6), dpi=80)
        ax = fig.add_subplot(111)
        N = len(ucb1)
        ind = np.arange(N)                                # the x locations for the groups
        plt.plot(ind, ucb1, color="blue", linewidth=2.5, linestyle="-", label="UCB1")
        plt.plot(ind, ThompsonSampling, color="orange", linewidth=2.5, linestyle="-", label="ThompsonSampling")
        plt.legend(loc = 3)
        plt.axis([0, 25, 0, 1])
        plt.xlabel('Number of pruned Neurons')
        plt.ylabel('The Accuracy')
        plt.grid(True)
        plt.show()
```



```
In [9]: p1 = figure(title="The Performance over the number of neurons' pruned", tools=TOOLS)
p1.line(ind, ucb1, legend="ucb1", line_color="blue", line_width=2)
p1.line(ind, ThompsonSampling, legend="Thompson Sampling", line_color="red", line_width=2)
p1.title.align = "center"
show(p1)
```

## 4 Comparing All algorithms with the model before pruning

```
In [10]: fig = plt.figure(figsize=(10, 6), dpi=80)
ax = fig.add_subplot(111)
N = len(ucb1)
Acc = [Accuracy for col in range(N)]
ind = np.arange(N) # the x locations for the groups
plt.plot(ind, ucb1, color="blue", linewidth=2.5, linestyle="-", label="UCB1")
plt.plot(ind, ThompsonSampling, color="orange", linewidth=2.5, linestyle="-", label="Thompson Sampling")
plt.plot(ind, Acc, color="pink", linewidth=2.5, linestyle="-", label="Accuracy before pruning")
plt.legend(loc = 3)
plt.axis([0, 25, 0, 1])
plt.xlabel('Number of pruned Neurons')
plt.ylabel('The Accuracy')
plt.grid(True)
plt.show()
```



```
In [11]: p1 = figure(title="The Performance over the number of neurons' pruned", tools=TOOLS)
p1.line(ind, ucb1, legend="ucb1", line_color="green", line_width=2)
p1.line(ind, ThompsonSampling, legend="Thompson Sampling", line_color="red", line_width=2)
p1.line(ind, Acc, legend="Accuracy", line_dash=(4, 4), line_color="blue", line_width=2)
p1.title.align = "center"
show(p1)
```