# Compression Normal

March 28, 2017

Compute the performance of MAB methods

```
In [4]: import numpy as np
    import time
    import sys
    import matplotlib.pyplot as plt
    from sklearn import metrics
    %matplotlib inline
    #plt.rcParams['figure.figsize'] = (15, 6)
```

### 0.1 Load BOKEH libariry

## 1 Compare the accuracy of the models

#### 1.1 Load the pruned algorithm from normal prune

In [7]: ucb1\_Multiple = np.load('/Users/salemameen/Desktop/banditsbook\_Prune\_many\_one\_play/pytho

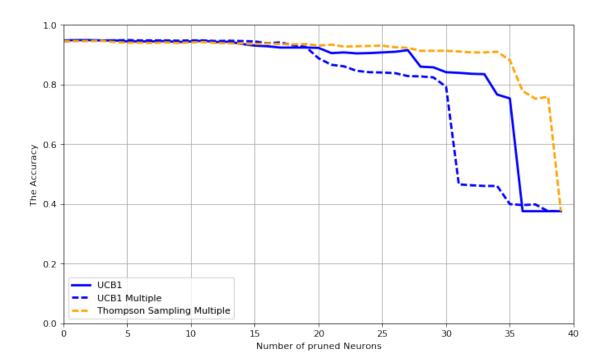
#### 1.2 Load the pruned algorithm from Multiple prune

```
ThompsonSampling_Multiple = np.load('/Users/salemameen/Desktop/banditsbook_Prune_many_on

In [8]: fig = plt.figure(figsize=(10, 6), dpi=80)
    ax = fig.add_subplot(111)
    N = len(ucb1)
    ## necessary variables
    ind = np.arange(N)  # the x locations for the groups
    ### Normal algoritms
    plt.plot(ind , ucb1 , color="blue", linewidth=2.5, linestyle="-", label="UCB1")
    ### Absuluate once algoritms
```

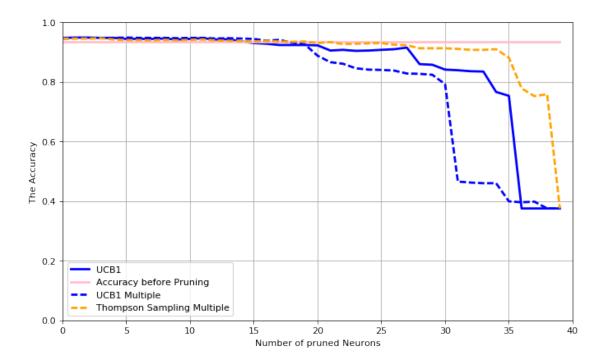
```
plt.plot(ind , ThompsonSampling_Multiple, color="orange", linewidth=2.5, linestyle="--",
##############################
plt.legend(loc = 3)
plt.axis([0, 40, 0, 1])
plt.xlabel('Number of pruned Neurons')
plt.ylabel('The Accuracy')
plt.grid(True)
plt.show()
```

plt.plot(ind , ucb1\_Multiple , color="blue", linewidth=2.5, linestyle="--", label="UCB1



#### 1.3 Comparing All algorithms with the model before pruning

```
N = len(ucb1)
Acc = [Accuracy for col in range(N)]
## necessary variables
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 , Acc, color="pink", linewidth=2.5, linestyle="-", label="Accuracy before
### Absuluate once algoritms
plt.plot(ind , ucb1_Multiple , color="blue", linewidth=2.5, linestyle="--", label="UCB1
plt.plot(ind , ThompsonSampling_Multiple, color="orange", linewidth=2.5, linestyle="--"
plt.legend(loc = 3)
plt.axis([0, 40, 0, 1])
plt.xlabel('Number of pruned Neurons')
plt.ylabel('The Accuracy')
plt.grid(True)
plt.show()
```



p1.circle(ind, ucb1\_Multiple, legend="ucb1 Multiple", line\_color="red", line\_width=2)

```
p1.line(ind, ucb1_Multiple, legend="ucb1 Multiple", line_color="red", line_width=2)

p1.circle(ind, ThompsonSampling_Multiple, legend="Thompson Sampling Multiple", line_color

p1.line(ind, ThompsonSampling_Multiple, legend="Thompson Sampling Multiple", line_color

p1.line(ind, Acc, legend="Accuracy", line_dash=(4, 4), line_color="orange", line_width=

p1.title.align = "center"

show(p1)
```

In []: