Compression Normal

March 28, 2017

Compute the performance of MAB methods

```
In [1]: 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

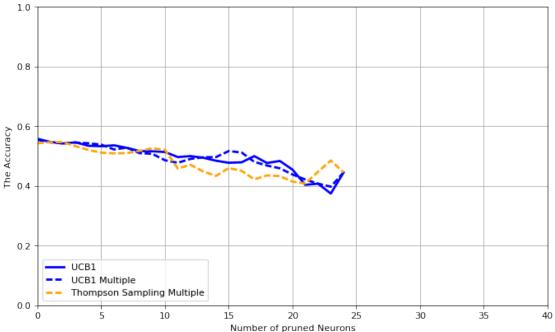
1.1 Load the pruned algorithm from normal prune

1.2 Load the pruned algorithm from Multiple prune

Absuluate once algoritms

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

```
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()
```



1.3 Comparing All algorithms with the model before pruning

```
In [7]: fig = plt.figure(figsize=(10, 6), dpi=80)
    ax = fig.add_subplot(111)
```

```
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 F
 ### 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()
1.0
0.8
0.6
```

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

Number of pruned Neurons

Accuracy

UCB1

UCB1 Multiple

Accuracy before Pruning

Thompson Sampling Multiple

10

```
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=2

p1.title.align = "center"

show(p1)
```

In []: