analysis

December 5, 2024

1 Analysis of BTSP-enabled local sensitive hashing performance

```
[1]: # import dependencies
import os
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn.decomposition import IncrementalPCA
import xgboost as xgb
```

1.1 Load dataset and check data

```
[2]: # load results.csv
results = pd.read_csv('merged_results.csv')

# print dataset length
print(len(results))

# print schema
print(results.dtypes)
```

Unnamed: 0 int64object dataset_name training_data_num int64hash_length int64space_ratio int64 binary_mode bool sampling_ratio float64 random_seed int64 experiment_index int64 input_dim int64 embedding_size int64 btsp_fq float64 btsp_mAP float64 fly_mAP float64

float64

1680

 wta_mAP

```
lsh_mAP float64 dtype: object
```

1.2 Relative importance analysis of each parameter

To provide a high-level comprehension of the dataset, we use XGBoost library to analyze the impact of each parameter on the performance of the model.

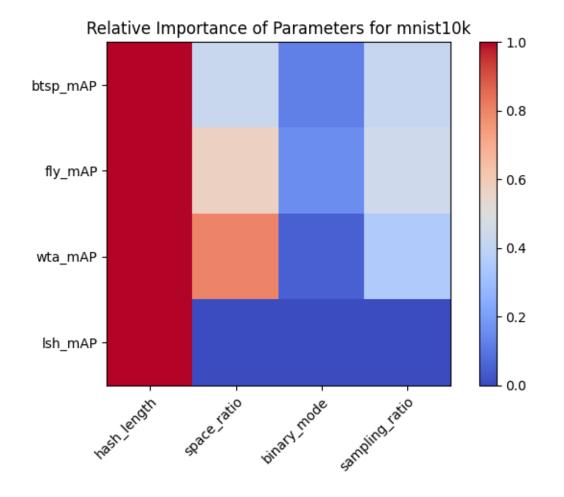
This section is for empirical analysis only, as we use a simple logistic regression model that may not precisely reflect the causal relationship between the parameters and the performance of the model.

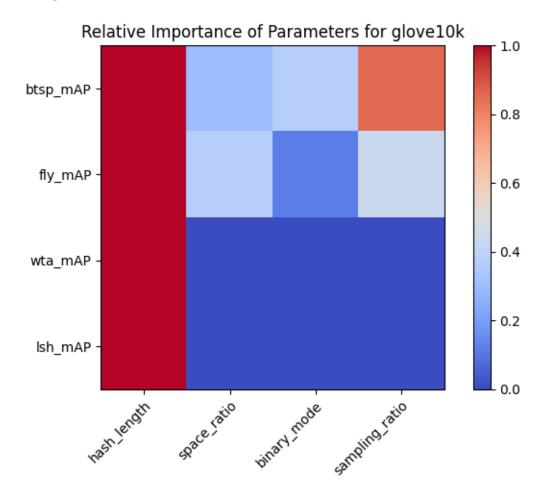
```
[]: # data cleaning
     # keep the columns we need
     # dataset_name, hash_length, space_ratio, binary_mode, sampling_ratio, and all_u
      →mAP columns
     meta_results = results.drop(columns=["training_data_num", "random_seed",_

¬"experiment_index", "input_dim", "embedding_size", "btsp_fq"])

     # remove index column
     meta_results = meta_results.drop(columns=["Unnamed: 0"])
     # separate data for each dataset
     datasets = meta_results["dataset_name"].unique()
     for dataset in datasets:
         # XGB analysis
         curr_dataset_data = meta_results[meta_results["dataset_name"] == dataset]
         # remove dataset name column
         curr_dataset_data = curr_dataset_data.drop(columns=["dataset_name"])
         # separate parameters and mAP columns
         mAP_types = ["btsp_mAP", "fly_mAP", "wta_mAP", "lsh_mAP"]
         curr_dataset_params = curr_dataset_data.drop(mAP_types, axis=1)
         # normalize the parameters
         curr dataset params = (curr dataset params - curr dataset params.mean()) / ____
      ⇔curr_dataset_params.std()
         # prepare XGBoost data
         X = curr_dataset_params
         # for each mAP type, use XGBoost to find the importance of each parameter
         mAP types importance = []
         for mAP_type in mAP_types:
             y = curr_dataset_data[mAP_type]
             dtrain = xgb.DMatrix(X, label=y)
             # set parameters
             # choose a parameter suitable for non-linear regression
             params = {
                 'objective': 'reg:logistic',
                 'booster': 'gbtree',
                 'eta': 0.05,
```

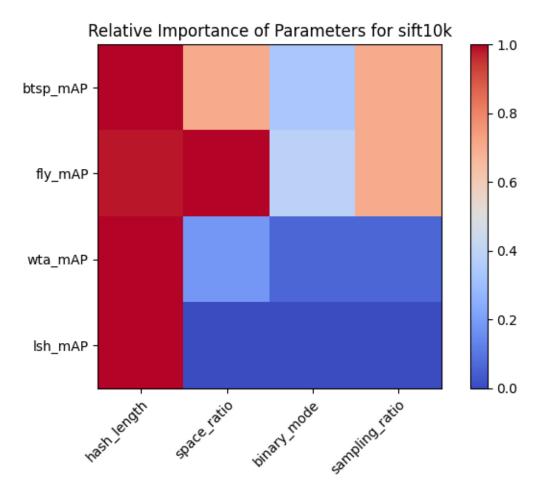
```
# train the model
        bst = xgb.train(params, dtrain)
        # print error
        print(bst.eval(dtrain))
        # get feature importance and convert to numpy array
        importance = bst.get_score(importance_type='weight')
        # check if each parameter is in the importance
        # if not, add it with O importance
        for param in curr_dataset_params.columns:
            if param not in importance:
                importance[param] = 0
        # for debugging
        # print(importance)
        # normalize the importance
        importance = np.array(list(importance.values()))
        importance = importance / importance.max()
        # print(importance)
        mAP_types_importance.append(importance)
    # plot importance heatmap
    norm = plt.Normalize(0, 1)
    fig, ax = plt.subplots()
    im = ax.imshow(mAP_types_importance, norm=norm, cmap='coolwarm')
    ax.set_xticks(np.arange(len(curr_dataset_params.columns)))
    ax.set_yticks(np.arange(len(mAP_types)))
    ax.set_xticklabels(curr_dataset_params.columns)
    ax.set_yticklabels(mAP_types)
    plt.setp(ax.get_xticklabels(), rotation=45, ha="right", __
 →rotation_mode="anchor")
    fig.tight_layout()
    # add title
    plt.title("Relative Importance of Parameters for " + dataset)
    # add colorbar
    cbar = ax.figure.colorbar(im, ax=ax)
    plt.show()
    # close the plot
    plt.close()
[0]
        eval-rmse:0.10103211259350435
{'hash_length': 81.0, 'space_ratio': 35.0, 'binary_mode': 10.0,
'sampling_ratio': 34.0}
           0.43209877 0.12345679 0.41975309]
[1.
```



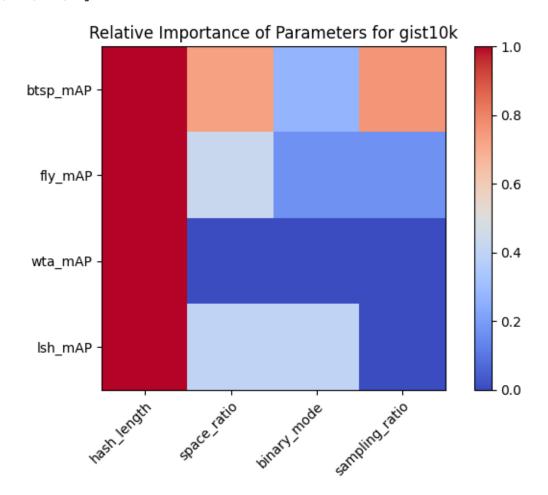


```
[0]
        eval-rmse:0.11770251840221084
{'hash_length': 30.0, 'space_ratio': 21.0, 'binary_mode': 10.0,
'sampling_ratio': 21.0}
[1.
            0.7
                       0.33333333 0.7
                                            ]
[0]
        eval-rmse:0.08375268210964371
{'hash_length': 56.0, 'space_ratio': 57.0, 'binary_mode': 22.0,
'sampling_ratio': 40.0}
[0.98245614 1.
                      0.38596491 0.70175439]
ΓοΊ
        eval-rmse:0.06654027230914542
```

```
{'hash_length': 50.0, 'space_ratio': 9.0, 'binary_mode': 3.0, 'sampling_ratio':
3.0}
[1.     0.18     0.06     0.06]
[0]     eval-rmse:0.07999836120665565
{'hash_length': 50.0, 'space_ratio': 0, 'binary_mode': 0, 'sampling_ratio': 0}
[1.     0.     0.]
```



```
[0] eval-rmse:0.03474972803155780
{'hash_length': 50.0, 'space_ratio': 20.0, 'sampling_ratio': 20.0, 'binary_mode': 0}
[1. 0.4 0.4 0.]
```



We verify the results of the analysis by examining the importance <code>space_ratio</code>, <code>binary_mode</code> and <code>sampling_ratio</code> to <code>wta_mAP</code> and <code>lsh_mAP</code>. These two algorithms are independent to these parameters, so we expect the importance of these parameters to be 0. Despite some exceptions, the results are overall consistent with our expectations.

One key observation is that binary_mode seems to have a neglegible impact on the performance of all algorithms and hash_length is the most important parameter for all algorithms. Other parameters have a relatively small impact on the performance of the algorithms and their impact may be dataset-dependent.

1.3 Numerical performance analysis

1.3.1 The impact of binary_mode on the performance of the algorithms

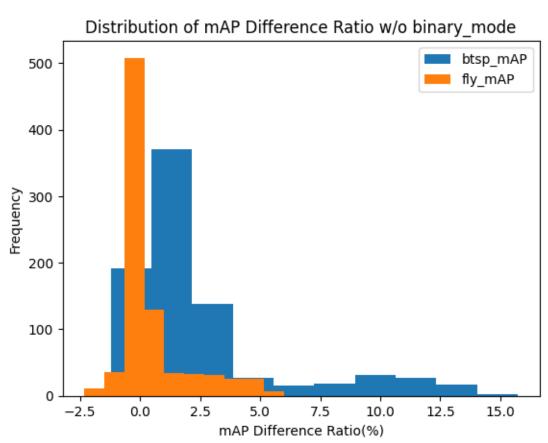
As binary_mode has no effect on wta_mAP and lsh_mAP, we only examine its impact on btsp_mAP and fly_mAP, respectively. We compare the performance of the algorithms with and without binary_mode and calculate the difference ratio of performance with:

```
\label{eq:definition} Difference\ ratio = \frac{mAP\ with\ binary\_mode - mAP\ without\ binary\_mode}{min(mAP\ with\ binary\_mode, mAP\ without\ binary\_mode)}
```

```
[23]: # load results.csv
      results = pd.read_csv('merged_results.csv')
      # keep the columns we need
      # dataset name, hash length, space ratio, binary mode, sampling ratio, btsp mAP_
       \hookrightarrow and fly mAP
      meta_results = results.drop(columns=["training_data_num", "random_seed",__
       ⇔"experiment_index", "input_dim", "embedding_size", "btsp_fq", "wta_mAP", □

¬"lsh mAP"])
      # remove index column
      meta_results = meta_results.drop(columns=["Unnamed: 0"])
      # compare the mAP of each algorithm w/o binary_mode on all datasets
      # separate data for binary mode enabled and disabled
      binary_mode_enabled = meta_results[meta_results["binary_mode"] == True]
      binary_mode_disabled = meta_results[meta_results["binary_mode"] == False]
      # sort two datasets by same order, ensure other parameters are the same for
       ⇔same index
      binary_mode_enabled = binary_mode_enabled.sort_values(by=["dataset_name",_
       →"hash_length", "space_ratio", "sampling_ratio"])
      binary_mode_disabled = binary_mode_disabled.sort_values(by=["dataset_name",__
       →"hash_length", "space_ratio", "sampling_ratio"])
      # calculate mAP difference ratio
      btsp_mAP_diff = binary_mode_enabled["btsp_mAP"].to_numpy() -__
       ⇔binary_mode_disabled["btsp_mAP"].to_numpy()
      fly mAP diff = binary mode enabled["fly mAP"].to numpy() - ____
       ⇔binary_mode_disabled["fly_mAP"].to_numpy()
      btsp_mAP_min = np.minimum(binary_mode_enabled["btsp_mAP"].to_numpy(),__
       ⇔binary_mode_disabled["btsp_mAP"].to_numpy())
      fly_mAP_min = np.minimum(binary_mode_enabled["fly_mAP"].to_numpy(),__
       dbinary_mode_disabled["fly_mAP"].to_numpy())
      # for debugging
      # print(btsp_mAP_diff)
      btsp_mAP_diff_ratio = btsp_mAP_diff / btsp_mAP_min
      fly_mAP_diff_ratio = fly_mAP_diff / fly_mAP_min
      # plot distribution of mAP difference ratio
```

```
fig, ax = plt.subplots()
ax.hist(btsp_mAP_diff_ratio, label='btsp_mAP')
ax.hist(fly_mAP_diff_ratio, label='fly_mAP')
ax.legend()
ax.set_xlabel("mAP Difference Ratio(%)")
ax.set_ylabel("Frequency")
plt.title("Distribution of mAP Difference Ratio w/o binary_mode")
plt.show()
# close the plot
plt.close()
```



The results show that binary_mode has a negligible impact on the performance of the algorithms. We also show that adding binary_mode will slightly increase the performance of the algorithms, given the majority of the difference ratios are positive.

1.3.2 Parameter-specific performance analysis between BTSP and FLY algorithms

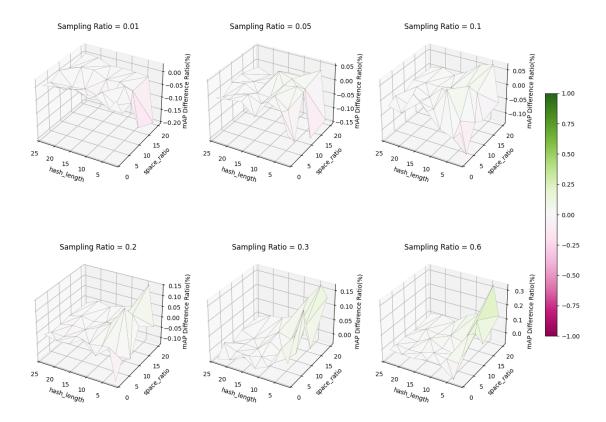
From empirical analysis, we find that hash_length, space_ratio and sampling_ratio all have considerable impact on the performance of the algorithms, and their impact may be dataset-dependent. Therefore, we plot the difference ratio of performance between the BTSP and FLY

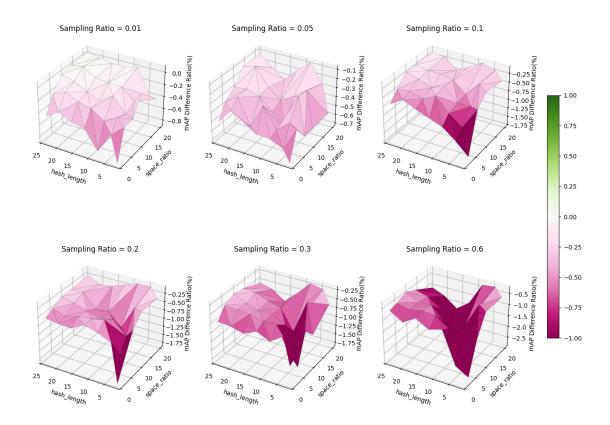
algorithms to find out in which range of these parameters the BTSP algorithm outperforms the FLY algorithm.

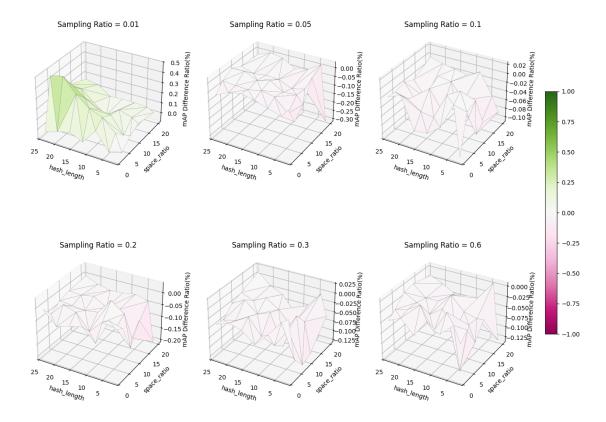
```
[13]: # load results.csv
     results = pd.read_csv('merged_results.csv')
     # keep the columns we need
     # dataset name, hash length, space ratio, binary mode, sampling ratio, btsp mAP_
      ⇔and fly mAP
     meta_results = results.drop(columns=["training_data_num", "random_seed",__

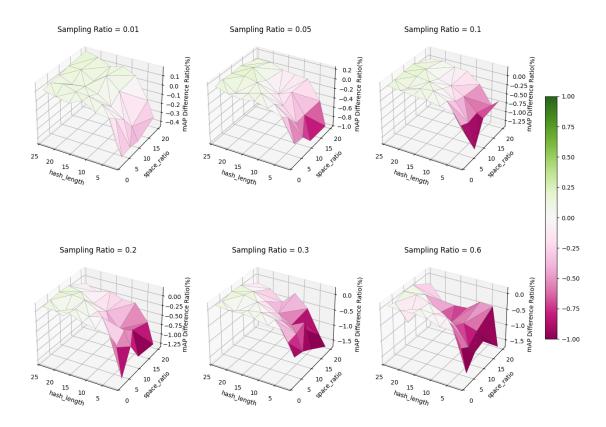
¬"lsh mAP"])
     # remove index column
     meta_results = meta_results.drop(columns=["Unnamed: 0"])
     # discard all rows with binary mode disabled
     meta_results = meta_results[meta_results["binary_mode"] == True]
     # calculate difference ratio
     mAP_diff = meta_results["btsp_mAP"].to_numpy() - meta_results["fly_mAP"].
      →to_numpy()
     mAP_min = np.minimum(meta_results["btsp_mAP"].to_numpy(),__
      →meta_results["fly_mAP"].to_numpy())
     mAP_diff_ratio = mAP_diff / mAP_min
     meta_results["mAP_diff_ratio"] = mAP_diff_ratio
     # separate data for each dataset
     datasets = meta_results["dataset_name"].unique()
     for dataset in datasets:
         curr_dataset_data = meta_results[meta_results["dataset_name"] == dataset]
         # start plotting
         # get the number of unique sampling_ratio
         sampling_ratios = curr_dataset_data["sampling_ratio"].unique()
         # create a figure with multiple subplots
         subplots_rows = int(np.sqrt(len(sampling_ratios)))
         subplots_cols = int(np.ceil(len(sampling_ratios) / subplots_rows))
         subplot_width = 6
         subplot_height = 6
         fig, axs = plt.subplots(subplots_rows, subplots_cols,_
       →subplot_kw={'projection': '3d'}, figsize=(subplots_cols * subplot_width, __
      subplots_rows * subplot_height))
         # use uniform norm
         norm = plt.Normalize(-1, 1)
         # plot each subplot
         for index, sampling_ratio in enumerate(sampling_ratios):
             curr sampling data = ___
       curr_dataset_data[curr_dataset_data["sampling_ratio"] == sampling_ratio]
             # get the subplot
```

```
if subplots_rows > 1:
          ax = axs[index // subplots_cols, index % subplots_cols]
      else:
          ax = axs[index]
      # plot trisurface
      x = curr_sampling_data["hash_length"].to_numpy()
      y = curr_sampling_data["space_ratio"].to_numpy()
      z = curr_sampling_data["mAP_diff_ratio"].to_numpy()
      ax.plot_trisurf(x, y, z, cmap='PiYG', norm=norm, edgecolor='black',
→linewidth=0.1)
      # invert hash_length axis
      ax.invert_xaxis()
      # add titles
      ax.set_title("Sampling Ratio = " + str(sampling_ratio))
      ax.set_xlabel("hash_length")
      ax.set_ylabel("space_ratio")
      ax.set_zlabel("mAP Difference Ratio(%)")
  # add a colorbar
  fig.colorbar(plt.cm.ScalarMappable(norm=norm, cmap='PiYG'), ax=axs,
⇔orientation='vertical', shrink=0.6)
  # add figure title
  fig.suptitle("mAP Difference Ratio between BTSP and Fly on " + dataset)
  # show and close
  plt.show()
  plt.close()
```









From the results, we confirm that the impact of these parameters on the performance of the algorithms is dataset-dependent. For example, on gist10k a larger space ratio generally makes Fly algorithm outperform BTSP algorithm, while the opposite is true for mnist10k. On glove10k Fly outperforms BTSP in almost all cases, while on minst10k the overall scale of the difference ratio is significantly smaller.