DAT300_CA1_16

October 17, 2019

1 Compulsory assignment 1 - Using Dask-ml on large data

1.1 Group #16

1.1.1 Thomas Moen and Jørgen Kongsro

1.2 Import libraries

```
[1]: %matplotlib notebook
   import numpy as np
   import pandas as pd
   import os
   import dask
   import dask.dataframe as dd
   import scipy
   import matplotlib.pyplot as plt
   import skimage.io
   import dask.array as da
   import joblib
   from dask.diagnostics import ProgressBar
   from dask_ml.linear_model import LogisticRegression
   from dask_ml.model_selection import train_test_split
   from dask_ml.datasets import make_classification
   from dask_ml.model_selection import train_test_split
   from dask_ml.linear_model import LogisticRegression
   from dask_ml.metrics import accuracy_score
   from dask_ml.model_selection import IncrementalSearchCV
   from dask.distributed import Client
   from sklearn.linear_model import SGDClassifier
   from sklearn.metrics import f1_score
   from sklearn.metrics import confusion_matrix
   from sklearn import metrics
   print (dask.__version__)
```

1.3 Install Kaggle API and download Kaggle data

```
[]: # Install Kaggle API

# How to setup: https://github.com/Kaggle/kaggle-api

# or visit: https://adityashrm21.github.io/Setting-Up-Kaggle/

#!pip install kaggle

[]: # Download Kaggle data using Kaggle API

#!kaggle competitions download -c dat300-ca1-autumn-2019

[]: # Unzip Kaggle data

#!unzip "dat300-ca1-autumn-2019.zip" -d "/tmp/whatever"
```

1.4 List files in directory (adjust for different operating systems)

```
if os.name == 'nt':
    workdir = 'C://Users//thomoe//Documents//myDAT300//dat300-ca1-autumn-2019//'
elif os.name == 'posix':
    workdir = '/Users/jorgenkongsro/Downloads/dat300-ca1-autumn-2019/'
    os.listdir(workdir)
```

[2]: ['y_test_sampleSubmission.csv', 'X_train.csv', 'y_train.csv', 'X_test.csv']

1.5 Import data

```
[131]: def import_data(csv_file):
    """ Import data from csv file

    :param data: a .csv separated dataset
    :return: a pandas data array, df

    """

    df = dd.read_csv(csv_file)
    return df

x_train_df = import_data(workdir + 'X_train.csv')
x_test_df = import_data(workdir + 'X_test.csv')
y_train_df = import_data(workdir + 'y_train.csv')
```

x_train_df
y_train_df

[131]: Dask DataFrame Structure:

Target

npartitions=1

float64

. .

Dask Name: from-delayed, 3 tasks

1.5.1 View data and study the structure

[10]: x_train_df

[10]:	Dask	DataFrame	Structure:							
			f1	f2	f3	f4	f5	f6	f7	
	f8	f9	f10	f11	f12	f13	f14	f15	f16	
	f17	f18	f19	f20	f21	f22	f23	f24	f25	
	f26	f27	f28	f29	f30	f31	f32	f33	f34	
	f35	f36	f37	f38	f39	f40	f41	f42	f43	
	f44	f45	f46	f47	f48	f49	f50	f51	f52	
	f53	f54	f 55	f56	f57	f58	f59	f60	f61	
	f62	f63	f64	f65	f66	f67	f68	f69	f70	
	f71	f72	f73	f74	f75	f76	f77	f78	f79	
	f80	f81	f82	f83	f84	f85	f86	f87	f88	
	f89	f90	f91	f92	f93	f94	f95	f96	f97	
	f98	f99	f100	f101	f102	f103	f104	f105	f106	
	f107	f108	f109	f110	f111	f112	f113	f114	f115	
	f116	f117	f118	f119	f120	f121	f122	f123	f124	
	f125	f126	f127	f128	f129	f130	f131	f132	f133	
	f134	f135	f136	f137	f138	f139	f140	f141	f142	
	f143	f144	f145	f146	f147	f148	f149	f150	f151	
	f152	f153	f154	f155	f156	f157	f158	f159	f160	
	f161	f162								
	npart	titions=83								

float64 float64

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. .

Dask Name: from-delayed, 249 tasks

[11]: y_train_df

[11]: Dask DataFrame Structure:

Target

npartitions=1

float64

Dask Name: from-delayed, 3 tasks

1.6 Check for missing data

```
[132]: def percent_missing(dataframe):
         """ Check for percent missing values in dataframe
         :param data: dataframe
         :return: dataframe
         missing_values = dataframe.isnull().sum()
         with ProgressBar():
            percent_missing = ((missing_values / dataframe.index.size) * 100).
      →compute()
         return percent_missing
     print(percent_missing(x_train_df))
     note: the results indicate that the features come in "tripets", e.g. f1 to f3_{\sqcup}
     →have quite similar missing%.
     →values within the triplet,
     if only one or two values are missing
     11 11 11
```

```
[############################### | 100% Completed | 35.0s
f1
        3.628002
f2
       3.754723
f3
       3.836795
f4
       4.133142
f5
       4.299061
f6
       4.377457
f7
       9.171993
f8
       9.276405
f9
       9.410288
f10
       3.628002
       3.754723
f11
f12
       3.836795
f13
       4.133142
f14
       4.299061
       4.377457
f15
f16
       9.171993
f17
       9.276405
       9.410288
f18
f19
       3.628002
```

```
f20
        3.754723
f21
        3.836795
f22
        4.133142
f23
        4.299061
f24
        4.377457
f25
        9.171993
f26
        9.276405
f27
        9.410288
f28
        3.628002
f29
        3.754723
        3.836795
f30
f133
        9.171993
f134
        9.276405
f135
        9.410288
f136
        3.628002
f137
        3.754723
        3.836795
f138
f139
        4.133142
f140
        4.299061
f141
        4.377457
f142
        9.171993
f143
        9.276405
f144
        9.410288
f145
        3.628002
f146
        3.754723
f147
        3.836795
f148
        4.133142
f149
        4.299061
f150
        4.377457
f151
        9.171993
f152
        9.276405
        9.410288
f153
f154
        3.628002
f155
        3.754723
f156
        3.836795
f157
        4.133142
f158
        4.299061
f159
        4.377457
f160
        9.171993
f161
        9.276405
f162
        9.410288
Length: 162, dtype: float64
```

[132]: '\nnote: the results indicate that the features come in "tripets", e.g. f1 to f3 have quite similar missing%. \nWe could impute some values very precisely by insert the mean of the other values within the triplet, \nif only one or two

1.7 Drop rows due to missing values in target

```
[14]: # Delete rows in x_train_df and y_train_df where
     # 1) y is NaN and/or
     # 2) x has more than N missing features
     # Delete the same rows in x_train_df and y_train_df
     \#First inspect the distribution of missing values in rows of x\_train\_df, in
     →order to decide on a threshold for culling rows:
     if dir().count('nacounts rows') == 0:
         nacounts_rows = x_train_df.isna().sum(axis=1).compute()
     nacounts_rows.plot.hist(bins=20, grid=True)
     #Based on the histogram we chose to remove rows of x_train_df which had more_
     →than 150 missing features:
     nacount_thresh = 150
     #Make boolean list where True means the number of NaN's exceeds nacount thresh
     \hookrightarrow for that row (of x_train_df):
     rows_to_drop_x = list(nacounts_rows > nacount_thresh)
     #Make boolean list where True means thaty train df is NaN for that row:
     rows_to_drop_y = list(y_train_df['Target'].isna().compute())
     \#Combine\ rows\_to\_drop\_x\ and\ rows\_to\_drop\_y\ and\ negate\ the\ list\ so\ that\ it_{\sqcup}
      →becomes a boolean list of rows to keep:
     rows_to_keep = [not(rows_to_drop_x[a] or rows_to_drop_y[a]) for a in_
      →range(len(rows_to_drop_y))]
     #Delete rows from x_train_df and y_train_df (we did not manage to do thisu
      →without pandas dataframes from the dask dataframes):
     x_train_droprows = x_train_df.compute()[rows_to_keep]
     y_train_droprows = y_train_df.compute()[rows_to_keep]
     #...therefore we had to revert back to dask dataframe afterwards:
     x_train_df_droprows=dd.from_pandas(x_train_droprows, npartitions=249) #_J
      →npartitions based on viewing data structure in previous section
     y_train_df droprows=dd.from_pandas(y_train_droprows, npartitions=3) #_
      →npartitions based on viewing data structure in previous section
     # Delete obsolete variables
     del x_train_df, y_train_df
[21]: type(x_train_droprows)
```

[21]: pandas.core.frame.DataFrame

1.8 Impute missing data

```
[15]: #choose here which imputation methods to use:
     imputation_methods = ['correlated_columns', 'col_means']
     ,, ,, ,,
     We decided to use col means as imputation method since we managed to get this.
      \rightarrowto work. There is a lot of possible solutions here and hopefully better_{\sqcup}
      \hookrightarrow outcomes,
     but we chose to focus on what was actually working.
     11 II II
[16]: #impute by inserting the mean of the column in question, for all columns:
     if 'col_means' in imputation_methods:
         #calculate mean (note that axis needs to be 0 to get columns, which is \Box
      \rightarrow weird)
         miin = x_train_df_droprows.mean(axis = 0).compute() # Fill with mean value
         x_train_df_impute = x_train_df_droprows.fillna(dict(miin))
     type(x_train_df_impute)
[16]: dask.dataframe.core.DataFrame
 []: #impute by inserting value from most closely correlated column:
     We have tried to impute by inserting value from closely correlated column.
     We can't understand why this one doesn't work. The idea was to fill in NaN's_{\sqcup}
      \rightarrow from correlated columns.
     The problem might be in the last line
     11 11 11
     #will only correct if the correlation between columns is above this threshold:
     lowest_allowed_corr = 0.995
     if 'correlated_columns' in imputation_methods:
```

#if correlation matrix does not exist, first try to read it from file, if $_{ extsf{L}}$

→that does not work calculate it:
if dir().count('corrs') == 0:

```
try:
           corrs = pd.read_csv(workdir + 'features_correlation_matrix.txt')
       except:
           corrs = x_train_df.corr('pearson')
   #impute for each feature feat:
  for feat in x_train_df_colnames:
       #order the feature names according to (absolute value of) correlations,
\rightarrow to feat:
       abscorr = [abs(a) for a in list(corrs[feat])]
       order = np.argsort(abscorr)[::-1]
       topfeatures = [x_train_df_colnames[a] for a in order]
       #remove features which are not sufficiently correlated to feat:
      mapper = dict(zip(x_train_df_colnames, abscorr))
       topfeatures = [a for a in topfeatures if mapper[a] >=__
→lowest_allowed_corr]
       #correct using each feature in topfeatures, starting with the most \sqcup
→strongly correlated otherfeature:
       for otherfeat in [a for a in topfeatures if a != feat]:
           print(feat, otherfeat)
           x_train_df_trim[feat] = x_train_df_trim[feat].
→fillna(x train df trim[otherfeat]).compute()
```

1.9 Build model and train

```
[82]: # Create dask array with chunks. Delete obsolete variables
X = x_train_df_impute
X = X.to_dask_array(lengths=True)
x_test = x_test_df
x_test = x_test_to_dask_array(lengths=True)
y = y_train_df_droprows
y = y.to_dask_array(lengths=True)
X
#del x_train_df_imean, x_train_df, y_train_df_imean, y_train_df
[82]: dask.array<values, shape=(2858793, 162), dtype=float64, chunksize=(11550, 162)>
```

```
[91]: # Convert blocks in dask array x for new chunks
X = X.rechunk((100000, 162))
y = y.rechunk((100000, 1))
y = y.flatten()
y
```

[91]: dask.array<reshape, shape=(2858793,), dtype=float64, chunksize=(100000,)>

```
[93]: X_train, X_val, y_train, y_val = train_test_split(X, y, test_size=0.1)
#del X, y
X_train
```

[93]: dask.array<concatenate, shape=(2572913, 162), dtype=float64, chunksize=(90000, 162)>

1.9.1 Parallelize Scikit-Learn

```
[]: # Model 1: Random forest skikit-learn parallelized

client = Client() # Connect to a Dask Cluster

from sklearn.ensemble import RandomForestClassifier
model = RandomForestClassifier(n_estimators=100, max_depth=2,random_state=0)
with joblib.parallel_backend('dask'):
    model.fit(X_train, y_train)
    y_true = y_test
    y_pred = model.predict(X_test)

"""

We ran into memory error for the model.predict part of the code
"""
```

1.9.2 Reimplement Scalable Algorithms with Dask Array

```
[122]: # Model 2: Logistic regression Dask style
from dask.diagnostics import ProgressBar

lr = LogisticRegression()
with ProgressBar():
    lr.fit(X_train, y_train)
```

```
| 100% Completed | 54.4s
100% Completed | 53.9s
100% Completed | 55.5s
100% Completed | 54.0s
100% Completed | 53.8s
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100% Completed | 52.1s
```

```
[############################### | 100% Completed | 50.7s
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[################################ | 100% Completed | 45.6s
```

[122]: '\n\nTakes a lot of time to compute\n\n'

[108]: # Model 3: Logistic regression Using grid search to tune hyperparameters

```
from dask_ml.model_selection import GridSearchCV

parameters = {'penalty': ['11', '12'], 'C': [0.01, 0.1, 1, 10, 100]}

lr = LogisticRegression()
tuned_lr = GridSearchCV(lr, parameters)

with ProgressBar():
    tuned_lr.fit(X_train, y_train)
```

```
[################################## | 100% Completed | 30min 52.6s
[################################ | 100% Completed | 1min 53.9s
```

1.9.3 Run diagnostics

```
[128]: y_true = y_val
y_pred = tuned_lr.predict(X_val)
accuracy_score(y_true, y_pred)
fpr, tpr, thresholds = metrics.roc_curve(y_true, y_pred)
AUC = metrics.auc(fpr, tpr)
F1 = f1_score(y_true, y_pred)

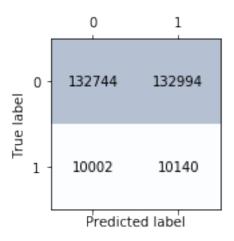
print (AUC)
print (F1)
```

- 0.5001943914080547
- 0.1257484708371596

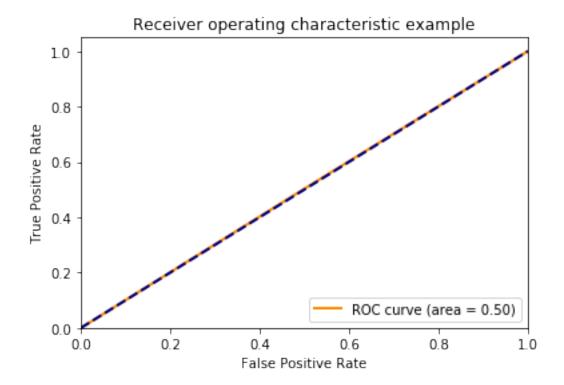
```
[124]: # Confusion matrix of test data
%matplotlib inline
confmat_test = confusion_matrix(y_true, y_pred)
fig, ax = plt.subplots(figsize=(2.5, 2.5))
ax.matshow(confmat_test, cmap=plt.cm.Blues, alpha=0.3)
for i in range(confmat_test.shape[0]):
    for j in range(confmat_test.shape[1]):
        ax.text(x=j, y=i, s=confmat_test[i, j], va='center', ha='center')

plt.xlabel('Predicted label')
plt.ylabel('True label')

plt.tight_layout()
plt.show()
```



```
[125]: # Plot the ROC for your best model on the training data
      # Compute ROC curve and ROC area for each class
      %matplotlib inline
      roc_auc=metrics.auc(fpr, tpr)
      plt.figure()
      lw = 2
      plt.plot(fpr, tpr, color='darkorange',
               lw=lw, label='ROC curve (area = %0.2f)' % roc_auc)
      plt.plot([0, 1], [0, 1], color='navy', lw=lw, linestyle='--')
      plt.xlim([0.0, 1.0])
      plt.ylim([0.0, 1.05])
      plt.xlabel('False Positive Rate')
      plt.ylabel('True Positive Rate')
      plt.title('Receiver operating characteristic example')
      plt.legend(loc="lower right")
     plt.show()
```



1.9.4 tuned_lr using GridSearch gave the best model, and we have submitted that to Kaggle

1.10 Make ready file to be submitted to Kaggle

```
[127]: # Predict using model and X_test standardized
y_pred = lr.predict(x_test)
YPRED = y_pred.compute()

np.mean(YPRED)
# Make list of IDs from 0 to 9999
ID = list(range(len(YPRED)))

# Make dataframes from predicted values, IDs and concatenate in result
pred1 = pd.DataFrame(YPRED, index=None, columns=['Target'])
pred2 = pd.DataFrame(ID, index=None, columns=['id'])
result = pd.concat([pred2, pred1], axis=1)
result = result.astype(int) # change type to integers

# Read results to csv for publication on Kaggle
result.to_csv('CA1_16.csv',index=False)
```