

clinical

May 3, 2023

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[1]: import pandas as pd
import os
import numpy as np
import ut
import gc
from sklearn.linear_model import LogisticRegression
from sklearn.model_selection import GridSearchCV
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[2]: def get_metadata(data_path):
    # Clean metadata dataframe
    md = pd.read_csv(os.path.join(data_path, 'ADNI1_Complete_2Yr_3T_4_18_2023.
↳csv'))

    md = md.rename(columns={'Image Data ID': 'Img_ID'})

    md = md.drop(columns=['Modality', 'Description', 'Type', 'Format', '
↳Downloaded', 'Acq Date'])

    md['Group'] = md['Group'].map({'CN':0, 'MCI':1, 'AD':2})
    md['Sex'] = md['Sex'].map({'F':0, 'M':1})
    md = md.rename(columns = {'Visit':'VISCODE'})

    return md
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[3]: def clean_data(md):
    c1 = pd.read_csv("/scratch/users/neuroimage/conda/data/clinical/ADNIMERGE.
↳csv", low_memory=False, na_values=[-4])
    c1 = c1[c1.COLPROT=='ADNI1']

    # could add EXAMDATE for time series
    filter = ['PTID', 'PTEDUCAT', 'PTETHCAT', 'PTRACCAT', 'PTMARRY', 'MMSE', '
↳AGE', 'VISCODE']
    c1 = c1[filter]
    c1 = c1.rename(columns={'PTID':'Subject', 'PTEDUCAT':'Educate', 'PTRACCAT':
↳'Race', 'PTETHCAT':'Ethn', 'PTMARRY':'Marry'})

    comb = md.merge(c1, how='left', on=['Subject', 'VISCODE'])
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comb['MMSE'] = comb['MMSE'].astype(np.int32)
comb = comb.drop(columns=['VISCODE'])

# one-hot encoding
one_hot = pd.get_dummies(comb, columns=['Ethn', 'Race', 'Marry'])

return one_hot

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[4]: def full_pipeline(X_matrix):

    df = ut.matrix_to_df(X_matrix)

    X, y = df.pipe(ut.clean_data, md=clean)

    X_train_pca, y_train, X_test_pca, y_test = ut.perform_pca(X,y)

    return X_train_pca, y_train, X_test_pca, y_test

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[5]: data_path = '/scratch/users/neuroimage/conda/data'
img_path = os.path.join(data_path, 'preprocessed/imgsss')

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[6]: smt_files_gm, smt_files_wm = ut.get_ordered_files(img_path, "smt")
md = get_metadata(data_path)
clean = clean_data(md)

X_wm, X_gm = ut.imgs_to_matrix(img_path, smt_files_gm, smt_files_wm,
    ↪combine=False)

X_comb = ut.imgs_to_matrix(img_path, smt_files_gm, smt_files_wm, combine=True)

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[7]: X_train_wm, y_train_wm, X_test_wm, y_test_wm = full_pipeline(X_wm)
gc.collect()
X_train_gm, y_train_gm, X_test_gm, y_test_gm = full_pipeline(X_gm)
gc.collect()
X_train_cb, y_train_cb, X_test_cb, y_test_cb = full_pipeline(X_comb)
gc.collect()

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n components:147
n components:160
n components:175

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[7]: 0

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[8]: base_pca_dir = os.path.join(data_path, 'full_pca_data')
os.makedirs(base_pca_dir, exist_ok=True)

arrs = {'X_train_wm': X_train_wm, 'y_train_wm': y_train_wm,

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        'X_test_wm': X_test_wm, 'y_test_wm': y_test_wm,
        'X_train_gm': X_train_gm, 'y_train_gm': y_train_gm,
        'X_test_gm': X_test_gm, 'y_test_gm': y_test_gm,
        'X_train_cb': X_train_cb, 'y_train_cb': y_train_cb,
        'X_test_cb': X_test_cb, 'y_test_cb': y_test_cb,
    }

    # loop through the dictionary and save each dataframe to CSV file
    for name, arr in arrs.items():
        np.savetxt(os.path.join(base_pca_dir, f'{name}.csv'), arr, delimiter=',')

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[9]: logreg_wm_acc = ut.perform_logreg(X_train_wm, y_train_wm, X_test_wm, y_test_wm)
    logreg_gm_acc = ut.perform_logreg(X_train_gm, y_train_gm, X_test_gm, y_test_gm)
    logreg_cb_acc = ut.perform_logreg(X_train_cb, y_train_cb, X_test_cb, y_test_cb)

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[10]: print(f"Classification accuracy under WM data: {logreg_wm_acc}")
    print(f"Classification accuracy under GM data: {logreg_gm_acc}")
    print(f"Classification accuracy under Combined data: {logreg_cb_acc}")

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Classification accuracy under WM data: 0.8620689655172413
 Classification accuracy under GM data: 0.7241379310344828
 Classification accuracy under Combined data: 0.7126436781609196

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[11]: import matplotlib.pyplot as plt

    # Create a bar plot
    accuracies = [logreg_wm_acc, logreg_gm_acc, logreg_cb_acc]
    labels = ['White Matter', 'Gray Matter', 'White + Gray Matter']
    plt.bar(labels, accuracies)

    # Add labels and title
    plt.xlabel('Matter')
    plt.ylabel('Accuracy')
    # Show the plot
    plt.show()

```

