

syn_classification_notebook

January 22, 2018

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In [123]: # Synapse Classification Challenge
          # Introduction to Connectomics 2017
          # W. Gray Roncal (your name here)

          your_name = 'ramsdén_devin'

In [124]: # Load data

import numpy as np

data = np.load('./synchallenge2017_training.npz')

imtrain = data['imtrain']
annotrain = data['annotrain']
ytrain = data['ytrain']

data = np.load('./synchallenge2017_validation.npz')

imvalid = data['imvalid']
annovalid = data['annovalid']
yvalid = data['yvalid']

In [125]: # Define feature extraction code

import skimage.feature as skif

def extract_features(imdata):
    xtrain = []
    for im in imdata:
        fvector = []
        # 50th percentile based on intensity
        fvector.append(np.percentile(im,50))

        #corner_kitchen_rosenfeld
        ckr = skif.corner_kitchen_rosenfeld(im)
        ckr = np.ravel(ckr)
        for i in ckr:
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        fvector.append(i)

    #corner_shi_tomasi
    csh = skif.corner_shi_tomasi(im)
    csh = np.ravel(csh)
    for i in csh:
        fvector.append(i)

    #structure_tensor
    st = skif.structure_tensor(im)
    st = np.ravel(st)
    for i in st:
        fvector.append(i)

    # add a contrast feature
    g = skif.greycomatrix(im, [1, 2], [0, np.pi/2], normed=True, symmetric=True)
    homogeneity = skif.greycoprops(g, 'homogeneity')

    # explicit way to add feature elements one at a time
    homogeneity = np.ravel(homogeneity)
    for i in homogeneity:
        fvector.append(i)

    fvector = np.asarray(fvector)
    xtrain.append(fvector)

    return np.asarray(xtrain)

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In [126]: *# Extract Features from training*

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xtrain = extract_features(imtrain)
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In [127]: *# Train Classifier*

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from sklearn.ensemble import RandomForestClassifier
clf = RandomForestClassifier(n_estimators=200)
clf = clf.fit(xtrain, ytrain)

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In [128]: *# Extract features from validation set*

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xvalid = extract_features(imvalid)
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In [129]: *# Run Classifier on validation set*

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scoresvalid = clf.predict_proba(xvalid)
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In [130]: *# Best f1 score report on validation set*

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from sklearn.metrics import f1_score
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# Can add post-processing here if desired

prob_syn = scoresvalid[:,1]

# default threshold
print('default f1 score: {}'.format(np.round(f1_score(yvalid, prob_syn >=0.5),2)))

f1_out = 0
thresh = 0
for i in np.arange(0.0, 1, 0.05):
    f1_test = f1_score(yvalid, prob_syn > i)
    if f1_test > f1_out:
        f1_out = f1_test
        thresh = i

print('My best validation f1-score is: {} at {} threshold.'.format(np.round(f1_out,2),
                                                                    thresh))

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default f1 score: 0.84
My best validation f1-score is: 0.85 at 0.45 threshold.

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/opt/conda/lib/python3.6/site-packages/sklearn/metrics/classification.py:1113: UndefinedMetricWarning: Precision is undefined because there is no predicted label in the dataset
  'precision', 'predicted', average, warn_for)

```

In [131]: *# Changing a parameter*
Train Classifier

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from sklearn.ensemble import RandomForestClassifier
clf = RandomForestClassifier(n_estimators=200)
clf = clf.fit(xtrain, ytrain)

# Run Classifier on validation set
scoresvalid = clf.predict_proba(xvalid)

# Best f1 score report on validation set

from sklearn.metrics import f1_score

# Can add post-processing here if desired

prob_syn = scoresvalid[:,1]

# default threshold
print('default f1 score: {}'.format(np.round(f1_score(yvalid, prob_syn >=0.5),2)))

f1_out = 0
thresh = 0

```

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for i in np.arange(0.0, 1, 0.05):
    f1_test = f1_score(yvalid, prob_syn > i)
    if f1_test > f1_out:
        f1_out = f1_test
        thresh = i

print('My best validation f1-score is: {} at {} threshold.'.format(np.round(f1_out,2),

```

default f1 score: 0.84

My best validation f1-score is: 0.84 at 0.5 threshold.

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/opt/conda/lib/python3.6/site-packages/sklearn/metrics/classification.py:1113: UndefinedMetricWarning:
'precision', 'predicted', average, warn_for)

```

In [132]: *# here we can inspect results*

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valid_labels = np.asarray(prob_syn > thresh, dtype='int')
# find images we did well on
idx_correct_syn = np.where((valid_labels == yvalid) & (yvalid == 1))[0]
idx_correct_nosyn = np.where((valid_labels == yvalid) & (yvalid == 0))[0]
# find images we did poorly on

idx_wrong_syn = np.where((valid_labels != yvalid) & (yvalid == 1))[0]
idx_wrong_nosyn = np.where((valid_labels != yvalid) & (yvalid == 0))[0]

import ndparse as ndp

print('synapse present - true positive')
ndp.plot(imvalid[idx_correct_syn[3]])

print('no synapse present - true negative')
ndp.plot(imvalid[idx_correct_nosyn[3]])

print('synapse present - false negative')
ndp.plot(imvalid[idx_wrong_syn[3]])

print('no synapse present - false positive')
ndp.plot(imvalid[idx_wrong_nosyn[3]])

```

ModuleNotFoundError Traceback (most recent call last)

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<ipython-input-132-3ab3cf437d5e> in <module>()
   10 idx_wrong_nosyn = np.where((valid_labels != yvalid) & (yvalid == 0))[0]
   11

```

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---> 12 import ndparse as ndp
      13
      14 print('synapse present - true positive')

```

ModuleNotFoundError: No module named 'ndparse'

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In [ ]: # Validate performance on test set (should only run/score once!)

data = np.load('./synchallenge2017_test_notruth.npz')

imtest = data['imtest']
annotest = data['annotest']

# Extract features from test set
xtest = extract_features(imtest)

# Run classifier on test set
scoretest = clf.predict_proba(xvalid)

# Post-processing
prob_syntest = scoretest[:,1]
syntest_predict = prob_syntest > thresh
syntest_predict = np.asarray(syntest_predict, dtype = 'uint8')

# save file and upload to google docs with label vector
np.save(your_name+'_synchallenge_testdata.npy',syntest_predict)

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