SVC classifier with 0% accuracy #9154

New issue



(b) Closed LBInd opened this issue on Jun 18, 2017 ⋅ 9 comments



```
LBInd commented on Jun 18. 2017 • edited ▼
Hello!
I'm using SVC from Sklearn to discriminate between different matrices.
The datas are 95 matrices of correlation, computed from IRM of patients with schizophrenia (50
matrices) and healty controls (45 matrices). They are pretty big (264*264), so I wasn't expecting perfect
results, but 0% accuracy seems really low.
Code
Here is the code:
  #control_matrices: list of 45 matrices
  #patient_matrices: list of 50 matrices
  n_{training} = 25 #Number of matrices of control to train SVC (25 control and 25 patient)
  indices = np.triu\_indices(264,1) #Since the matrices are symetric, I just take the upper
  {\tt perm\_control = np.random.permutation(45) \ \#Doing \ a \ permutation \ to \ take \ random \ matrices \ fo}
  contr_matrices = control_matrices[perm_control] #control_matrices is a list of matrices
  perm_patient = np.random.permutation(50) #Same with the patient matrices
  pat_matrices = patient_matrices[perm_patient]
  x_control = [m[indices] for m in contr_matrices[:n_training]] #Data for training
  x_patient = [m[indices] for m in pat_matrices[:n_training]]
  test control = [m[indices] for m in contr matrices[n training:]] #Data for test once the
  test_patient = [m[indices] for m in pat_matrices[n_training:]]
  X = np.concatenate((x_control, x_patient))
  Y = np.asarray( n_training*["Control"] + n_training*["Patient"] )
  ## Training
  clf = SVC()
  clf.fit(X,Y)
Expected Results
Since the size of the data is huge compared to the number of matrices, I would have expected low
results (something just a little bit better than 50%).
Actual Results
 clf.score(np.concatenate((test_control, test_patient)), 20*['Control']+25*['Patient'])
  0.0
The same happens whenever I run the code (so, with different permutations), and for n_training from
10 to 45. However the SVC does remember well the first matrices, for the training.
Try
I also tried this, with exactly the same results:
  from sklearn.cross_validation import StratifiedKFold, cross_val_score
  from nilearn import connectome
  connectivity_coefs = connectome.sym_to_vec(matrices, ConnectivityMeasure)
```

No one assigned Labels None yet Projects None yet Milestone No milestone



Assignees

```
cv = StratifiedKFold(Y, n_folds=3, shuffle=True)
svc = LinearSVC()

cv_scores = cross_val_score(svc, connectivity_coefs, Y, cv=cv, scoring='accuracy')

print('Score: %1.2f +- %1.2f' % (cv_scores.mean(), cv_scores.std()))

Score: 0.00 +- 0.00

I completely fail to understand what is happening here.

Versions

Windows-8-6.2.9200
Python 3.4.1 |Continuum Analytics, Inc.| (default, May 19 2014, 13:02:30) [MSC v.1600 64 bit (AMD64)]
NumPy 1.9.1
SciPy 0.15.1
Scikit-Learn 0.15.2
```



rth commented on Jun 19, 2017 • edited •

Member

With random patient_matrices and control_matrices I get a \sim 0.5 accuracy score using your first code sample, as expected. Did you normalize x to [0, 1] (or similar) for SVM? What is clf.score(x, Y)? Have you tried a linear model first (e.g. LogisticRegression) to check that your code is working?



LBInd commented on Jun 19, 2017 • edited ▼

Author

I didn't normalize anything; I'm using matrices with values in [-1,1].

 ${\tt clf.score}(X,\ Y) \ \ gives \ the \ percentage \ of \ predictions \ (for \ X) \ that \ match \ the \ correct \ value \ (\ Y\).$ I have tried LogisticRegression , SVC and LinearSVC; all of them give me the same result (0% accuracy on new matrices).

I also tried with simpler data : matrices [0] for Control and [1] for Patients. The SVC worked perfectly, so I suspect it has something to do with the size of the matrices I use (huge size and few samples).



rth commented on Jun 19, 2017

Member

OK, thanks for the details! Yes, but what the actual value of clf.score(X, Y): 1.0 ? Well your code works as expected with random data,

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patient_matrices = np.random.rand(50, 264, 264)
control_matrices = np.random.rand(50, 264, 264)
```

I think. So the question is really, is there anything in the actual data you are using that could explain such difference in the results. Alternatively, trying to reproduce this issue with some synthetic data (or sharing your data) would also help understanding what is happening.



LBInd commented on Jun 19, 2017

Author

clf.score(X,Y) always return 0.0 when x is a list of new matrices (not the ones used for training). Well, the datas are matrices computed from IRM from an open dataset, but I have a lot of preprocessing. I'll share the previous code if that helps.

Thanks for your answers!



rth commented on Jun 19, 2017

Member

y are the matrices used for training) ..

Well, the datas are matrices computed from IRM from an open dataset, but I have a lot of preprocessing. I'll share the previous code if that helps.

Or maybe just the result of your preprocessing (those 2 matrices (50, 256, 256) ..)?



```
LBInd commented on Jun 19, 2017
                                                                                     Author
Here is how to get the same matrices that I use:
  from nilearn import datasets
  from nilearn import input_data
  from nilearn.connectome import ConnectivityMeasure
  import numpy as np
  from sklearn.svm import SVC, LinearSVC
  from sklearn.cross_validation import StratifiedKFold, cross_val_score
  from nilearn import connectome
  ## Atlas for the parcellation and Dataset
  power = datasets.fetch_coords_power_2011()
  coords = np.vstack((power.rois['x'], power.rois['y'], power.rois['z'])).T
  datas = datasets.fetch_cobre(n_subjects=None, verbose=0)
  spheres masker = input data.NiftiSpheresMasker(
                      seeds=coords, smoothing_fwhm=4, radius=5.,
                      detrend=True, standardize=True,
                      high_pass=0.01, t_r=2, verbose=0)
  ## Extracting useful IRM
  list_time_series = []
  i = 0
  for fmri_filenames, confounds_file in zip(datas.func, datas.confounds): #Might take a fe
      print("Sujet %s" % i)
      if i != 38 and i != 41: #Subjects removed from the study
          conf = np.genfromtxt(confounds_file)
          conf = np.delete(conf, obj = 16, axis = 1) #Remove Global Signal
          conf = np.delete(conf, obj = 0, axis = 0) #Remove labels
          scrub = [i for i in range(150) if conf[i,7]==1]
          conf = np.delete(conf, obj = 7, axis = 1) #Remove Scrub
          if len(scrub) < 90: #Keep at least 60 non scrub</pre>
              time_series = spheres_masker.fit_transform(fmri_filenames, confounds=conf)
              time\_series = np.delete(time\_series, obj = scrub, axis = 0) #Remove scrub
              list_time_series.append(time_series)
          else:
              list_time_series.append([])
      else:
          list_time_series.append([])
  ## Computing correlation matrices
  N = len(datas.phenotypic)
  control subjects = []
  patient_subjects = []
  for i in range(N):
      t = list_time_series[i]
      if type(t) != list :
          subject = datas.phenotypic[i]
          if str(subject[4])=='b\'Control\'':
              control_subjects.append(t)
          else:
              patient_subjects.append(t)
  control_subjects = np.asarray(control_subjects)
  patient_subjects = np.asarray(patient_subjects)
  connect_measure = ConnectivityMeasure(kind='tangent')
  control matrices=connect measure.fit transform(control subjects)
  patient_matrices=connect_measure.fit_transform(patient_subjects)
  matrices = np.concatenate((control_matrices, patient_matrices))
```

