practical_exercise_10, Methods 3, 2021, autumn semester

[FILL IN YOUR NAME]

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Exercises and objectives

- 1) Use principal component analysis to improve the classification of subjective experience
- 2) Use logistic regression with cross-validation to find the optimal number of principal components

REMEMBER: In your report, make sure to include code that can reproduce the answers requested in the exercises below (MAKE A KNITTED VERSION)

REMEMBER: This is Assignment 4 and will be part of your final portfolio

EXERCISE 1 - Use principal component analysis to improve the classification of subjective experience

We will use the same files as we did in Assignment 3 The files megmag_data.npy and pas_vector.npy can be downloaded here (http://laumollerandersen.org/data_methods_3/megmag_data.npy) and here (http://laumollerandersen.org/data_methods_3/pas_vector.npy)

The function equalize_targets is supplied - this time, we will only work with an equalized data set. One motivation for this is that we have a well-defined chance level that we can compare against. Furthermore, we will look at a single time point to decrease the dimensionality of the problem

- 1) Create a covariance matrix, find the eigenvectors and the eigenvalues
 - i. Load megmag_data.npy and call it data using np.load. You can use join, which can be imported from os.path, to create paths from different string segments
 - ii. Equalize the number of targets in y and data using equalize_targets
 - iii. Construct times=np.arange(-200, 804, 4) and find the index corresponding to 248 ms then reduce the dimensionality of data from three to two dimensions by only choosing the time index corresponding to 248 ms (248 ms was where we found the maximal average response in Assignment 3)
 - iv. Scale the data using StandardScaler
 - v. Calculate the sample covariance matrix for the sensors (you can use np.cov) and plot it (either using plt.imshow or sns.heatmap (import seaborn as sns))
 - vi. What does the off-diagonal activation imply about the independence of the signals measured by the 102 sensors?
 - vii. Run np.linalg.matrix_rank on the covariance matrix what integer value do you get? (we'll use this later)
 - viii. Find the eigenvalues and eigenvectors of the covariance matrix using np.linalg.eig note that some of the numbers returned are complex numbers, consisting of a real and an imaginary part

(they have a j next to them). We are going to ignore this by only looking at the real parts of the eigenvectors and -values. Use np.real to retrieve only the real parts

- 2) Create the weighting matrix W and the projected data, Z
 - i. We need to sort the eigenvectors and eigenvalues according to the absolute values of the eigenvalues (use np.abs on the eigenvalues).
 - ii. Then, we will find the correct ordering of the indices and create an array, e.g. sorted_indices that contains these indices. We want to sort the values from highest to lowest. For that, use np.argsort, which will find the indices that correspond to sorting the values from lowest to highest. Subsequently, use np.flip, which will reverse the order of the indices.
 - iii. Finally, create arrays of sorted eigenvalues and eigenvectors using the sorted_indices array just created. For the eigenvalues, it should like this eigenvalues = eigenvalues[sorted_indices] and for the eigenvectors: eigenvectors = eigenvectors[:, sorted_indices]
 - iv. Plot the log, np.log, of the eigenvalues, plt.plot(np.log(eigenvalues), 'o') are there some values that stand out from the rest? In fact, 5 (noise) dimensions have already been projected out of the data how does that relate to the matrix rank (Exercise 1.1.vii)
 - v. Create the weighting matrix, W (it is the sorted eigenvectors)
 - vi. Create the projected data, Z, Z = XW (you can check you did everything right by checking whether the X you get from $X = ZW^T$ is equal to your original X, np.isclose may be of help)
 - vii. Create a new covariance matrix of the principal components (n=102) plot it! What has happened off-diagonal and why?

```
def equalize_targets(data, y):
np.random.seed(7)
 targets = np.unique(y)
 counts = list()
 indices = list()
 for target in targets:
     counts.append(np.sum(y == target))
     indices.append(np.where(y == target)[0])
min_count = np.min(counts)
first_choice = np.random.choice(indices[0], size=min_count, replace=False)
 second_choice = np.random.choice(indices[1], size=min_count, replace=False)
third_choice = np.random.choice(indices[2], size=min_count, replace=False)
 fourth_choice = np.random.choice(indices[3], size=min_count, replace=False)
 new_indices = np.concatenate((first_choice, second_choice,
                              third_choice, fourth_choice))
new_y = y[new_indices]
new_data = data[new_indices, :, :]
return new_data, new_y
```

EXERCISE 2 - Use logistic regression with cross-validation to find the optimal number of principal components

- 1) We are going to run logistic regression with in-sample validation
 - i. First, run standard logistic regression (no regularization) based on $Z_{d\times k}$ and y (the target vector). Fit (.fit) 102 models based on: k = [1, 2, ..., 101, 102] and d = 102. For each fit get the

- classification accuracy, (.score), when applied to $Z_{d\times k}$ and y. This is an in-sample validation. Use the solver newton-cg if the default solver doesn't converge
- ii. Make a plot with the number of principal components on the x-axis and classification accuracy on the y-axis what is the general trend and why is this so?
- iii. In terms of classification accuracy, what is the effect of adding the five last components? Why do you think this is so?
- 2) Now, we are going to use cross-validation we are using cross_val_score and StratifiedKFold from sklearn.model_selection
 - i. Define the variable: cv = StratifiedKFold() and run $cross_val_score$ (remember to set the cv argument to your created cv variable). Use the same estimator in $cross_val_score$ as in Exercise 2.1.i. Find the mean score over the 5 folds (the default of StratifiedKFold) for each k, k = [1, 2, ..., 101, 102]
 - ii. Make a plot with the number of principal components on the x-axis and classification accuracy on the y-axis how is this plot different from the one in Exercise 2.1.ii?
 - iii. What is the number of principal components, $k_{max_accuracy}$, that results in the greatest classification accuracy when cross-validated?
 - iv. How many percentage points is the classification accuracy increased with relative to the to the full-dimensional, d, dataset
 - v. How do the analyses in Exercises 2.1 and 2.2 differ from one another? Make sure to comment on the differences in optimization criteria.
- 3) We now make the assumption that $k_{max_accuracy}$ is representative for each time sample (we only tested for 248 ms). We will use the PCA implementation from *scikit-learn*, i.e. import PCA from sklearn.decomposition.
 - i. For each of the 251 time samples, use the same estimator and cross-validation as in Exercises 2.1.i and 2.2.i. Run two analyses one where you reduce the dimensionality to $k_{max_accuracy}$ dimensions using PCA and one where you use the full data. Remember to scale the data (for now, ignore if you get some convergence warnings you can try to increase the number of iterations, but this is not obligatory)
 - ii. Plot the classification accuracies for each time sample for the analysis with PCA and for the one without in the same plot. Have time (ms) on the x-axis and classification accuracy on the y-axis
 - iii. Describe the differences between the two analyses focus on the time interval between 0 ms and 400 ms describe in your own words why the logistic regression performs better on the PCA-reduced dataset around the peak magnetic activity