Gr?ficas fourier nystroem

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```
In [1]: %matplotlib inline
        import matplotlib.pyplot as plt
        import numpy as np
        from time import time
        # Import datasets, classifiers and performance metrics
        from sklearn import datasets, svm, pipeline
        from sklearn.kernel_approximation import (RBFSampler,
                                                  Nystroem)
        from sklearn.decomposition import PCA
        from sklearn.tree import DecisionTreeClassifier
In [2]: # The digits dataset
       digits = datasets.load_digits(n_class=9)
In [3]: # To apply an classifier on this data, we need to flatten the image, to
        # turn the data in a (samples, feature) matrix:
       n_samples = len(digits.data)
        data = digits.data / 16.
        data -= data.mean(axis=0)
In [4]: # We learn the digits on the first half of the digits
        data_train, targets_train = (data[:n_samples // 2],
                                     digits.target[:n_samples // 2])
        # Now predict the value of the digit on the second half:
        data_test, targets_test = (data[n_samples // 2:],
                                   digits.target[n_samples // 2:])
        # data_test = scaler.transform(data_test)
In [5]: # Create a classifier: a support vector classifier
       kernel_svm = svm.SVC(gamma=.2)
        linear_svm = svm.LinearSVC()
In [6]: # create pipeline from kernel approximation
        # and linear sum
```

Código propio: hago el pipeline con un DecisionTree Nota: de momento uso el mismo RBF-Sampler que usan ellos, la misma instancia. Si afecta puedo usar uno nuevo. No lo creo

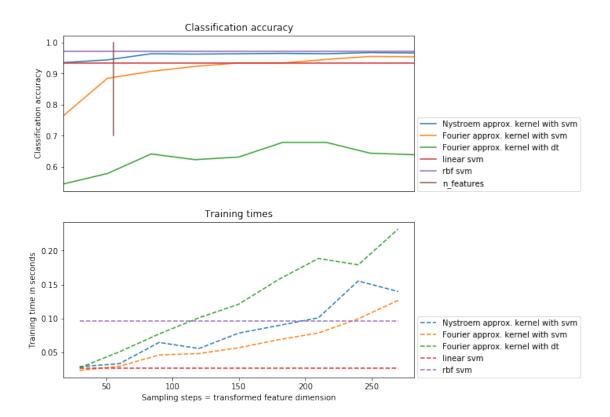
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In [8]: fourier_approx_dt = pipeline.Pipeline([("feature_map", feature_map fourier),
                                               ("dt", DecisionTreeClassifier())])
In [9]: # fit and predict using linear and kernel sum:
       kernel_svm_time = time()
       kernel_svm.fit(data_train, targets_train)
       kernel_svm_score = kernel_svm.score(data_test, targets_test)
       kernel_svm_time = time() - kernel_svm_time
        linear_svm_time = time()
        linear_svm.fit(data_train, targets_train)
        linear_svm_score = linear_svm.score(data_test, targets_test)
        linear_svm_time = time() - linear_svm_time
In [10]: sample_sizes = 30 * np.arange(1, 10)
         fourier_scores = []
         nystroem_scores = []
         fourier_times = []
         nystroem_times = []
```

Código propio: hago las listas para mi nuevo modelo. Nota: ellos usan un nombre ambiguo. Les llaman fourier_scores y fourier_times porque solamente la svm usa fourier. Nosotros usamos fourier con dt.

```
nystroem_times.append(time() - start)
             # Fourier sum
             start = time()
             fourier_approx_svm.fit(data_train, targets_train)
             fourier times.append(time() - start)
             # Código propio: el tiempo del fourier dt
             start = time()
             fourier_approx_dt.fit(data_train, targets_train)
             fourier_dt_times.append(time() - start)
             # Cálculo de scores
             fourier_score = fourier_approx_svm.score(data_test, targets_test)
             nystroem_score = nystroem_approx_svm.score(data_test, targets_test)
             # Código propio: calculo sl score con mi pipeline
             fourier_dt_score = fourier_approx_dt.score(data_test, targets_test)
             # Poner los scores en las listas de score
             nystroem_scores.append(nystroem_score)
             fourier_scores.append(fourier_score)
             # Código propio: pongo mis propios scores y times
             fourier_dt_scores.append(fourier_dt_score)
In [13]: # plot the results:
         plt.figure(figsize=(8, 8))
         accuracy = plt.subplot(211)
         # second y axis for timeings
         timescale = plt.subplot(212)
         # Accuracy y tiempo de nystroem
         accuracy.plot(sample_sizes, nystroem_scores,
                       label="Nystroem approx. kernel with svm")
         timescale.plot(sample_sizes, nystroem_times, '--',
                        label='Nystroem approx. kernel with svm')
         # Accuracy y tiempo de fourier (este es ambiguo, se refiere al sum)
         accuracy.plot(sample_sizes, fourier_scores,
                       label="Fourier approx. kernel with svm")
         timescale.plot(sample_sizes, fourier_times, '--',
                        label='Fourier approx. kernel with svm')
         # Accuracy y tiempo de fourier dt
         accuracy.plot(sample_sizes, fourier_dt_scores,
```

nystroem_approx_svm.fit(data_train, targets_train)

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label="Fourier approx. kernel with dt")
         timescale.plot(sample_sizes, fourier_dt_times, '--',
                        label='Fourier approx. kernel with dt')
         # horizontal lines for exact rbf and linear kernels:
         accuracy.plot([sample_sizes[0], sample_sizes[-1]],
                       [linear_svm_score, linear_svm_score], label="linear_svm")
         timescale.plot([sample_sizes[0], sample_sizes[-1]],
                        [linear svm time, linear svm time], '--', label='linear svm')
         accuracy.plot([sample_sizes[0], sample_sizes[-1]],
                       [kernel_svm_score, kernel_svm_score], label="rbf svm")
         timescale.plot([sample_sizes[0], sample_sizes[-1]],
                        [kernel_svm_time, kernel_svm_time], '--', label='rbf svm')
         # vertical line for dataset dimensionality = 64
         accuracy.plot([64, 64], [0.7, 1], label="n_features")
         # legends and labels
         accuracy.set_title("Classification accuracy")
         timescale.set_title("Training times")
         accuracy.set_xlim(sample_sizes[0], sample_sizes[-1])
         accuracy.set_xticks(())
         #accuracy.set_ylim(np.min(fourier_scores), 1)
         timescale.set_xlabel("Sampling steps = transformed feature dimension")
         accuracy.set_ylabel("Classification accuracy")
         timescale.set_ylabel("Training time in seconds")
         accuracy.legend(loc=(1.01,0)) # Modifico la posición de la leyenda, que no se ve bien
         timescale.legend(loc=(1.01,0))
Out[13]: <matplotlib.legend.Legend at 0x7ff30aec33c8>
```



```
In [14]: ## visualize the decision surface, projected down to the first
         ## two principal components of the dataset
         #pca = PCA(n_components=8).fit(data_train)
         \#X = pca.transform(data_train)
         ## Generate grid along first two principal components
         \#multiples = np.arange(-2, 2, 0.1)
         ## steps along first component
         #first = multiples[:, np.newaxis] * pca.components [0, :]
         ## steps along second component
         #second = multiples[:, np.newaxis] * pca.components_[1, :]
         ## combine
         #qrid = first[np.newaxis, :, :] + second[:, np.newaxis, :]
         #flat_grid = grid.reshape(-1, data.shape[1])
         ## title for the plots
         #titles = ['SVC with rbf kernel',
                    'SVC (linear kernel) \n with Fourier rbf feature map \n'
                    'n_components=100',
         #
                    'SVC (linear kernel)\n with Nystroem rbf feature map\n'
         #
                    'n_components=100']
```

```
#plt.tight_layout()
#plt.figure(figsize=(12, 5))
## predict and plot
#for i, clf in enumerate((kernel_sum, nystroem_approx_sum,
                          fourier_approx_sum)):
     # Plot the decision boundary. For that, we will assign a color to each
#
    # point in the mesh [x_min, x_max]x[y_min, y_max].
#
    plt.subplot(1, 3, i + 1)
    Z = clf.predict(flat_grid)
#
#
   # Put the result into a color plot
    Z = Z.reshape(grid.shape[:-1])
# plt.contourf(multiples, multiples, Z, cmap=plt.cm.Paired)
#
    plt.axis('off')
#
#
    # Plot also the training points
    plt.scatter(X[:, 0], X[:, 1], c=targets_train, cmap=plt.cm.Paired,
#
#
                 edgecolors=(0, 0, 0))
    plt.title(titles[i])
#plt.tight_layout()
#plt.show()
```