



Machine Learning



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Lecture: Dimensionality Reduction

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Dimensionality Reduction

- Dimensionality reduction is the process of converting a dataset that typically has a large number of features into a dataset with a <u>reduced number of features</u> (in some cases a significantly reduced number of features).
- This is <u>not</u> the same thing as feature selection as the reduced set of features may not be a subset of the original features.
- Dimensionality reduction algorithms such as PCA performs a linear
 transformation of d dimensional input to d dimensional feature vectors such that k < d, while attempting to that explain a maximum amount of the variance.
- Can be used very effectively for reducing the dimensionality when you have a very large number of features. A typical application would be image analysis.

PCA

- PCA is by far the most popular method used for dimensionality reduction.
- The objective of the PCA algorithm is to find a surface on to which to project the data so as to minimize the projection error
- More specify we want to identify k vectors that minimize the projection error where the projection error is defined as:

$$\rightarrow \frac{1}{m} \sum_{i=1}^{m} |x^i - x_{mapped}^i|^2$$

Note the PCA expects that you data has been standardized in advance.

PCA

- ▶ PCA is a parameterized algorithm. We specify in advance the a parameter value k, which specifies the number of features vectors the algorithm will return.
- Typically the selection of a value for k is made depending on the <u>retained</u> <u>variance from running PCA with the value k</u>. You can think of the retained variance as a measure of how effective the projection process employed by PCA.
- You will hear phrases such as selecting a value of k resulted in "99% of the variance retained". Another common measure would be "95% of the variance retained".

```
from sklearn import decomposition
from sklearn import datasets
from sklearn import cross_validation
from sklearn import tree
from sklearn.preprocessing import StandardScaler
import numpy as np
iris = datasets.load iris()
                                                     Original Accuracy 0.95
X = iris.data
                                                     Explained Variance: 0.958
y = iris.target
                                                     Accuracy Using Two New Features 0.91
scaler = StandardScaler()
X = scaler.fit transform(X)
clf = tree.DecisionTreeClassifier()
scores = cross validation.cross val score(clf, X, iris.target, cv=10)
print ("Original Accuracy", np.mean(scores))
pca = decomposition.PCA(n components=2)
pca.fit(X)
X = pca.transform(X)
print ("Explained Variance: ",np.sum(pca.explained_variance ratio ))
clf = tree.DecisionTreeClassifier()
scores = cross validation.cross_val_score(clf, X, iris.target, cv=10)
```

print (np.mean(scores))