Al and Machine Learning Zhiyun Lin



Principal Components Analysis (PCA)

- Principal Components Analysis
- Autoencoders

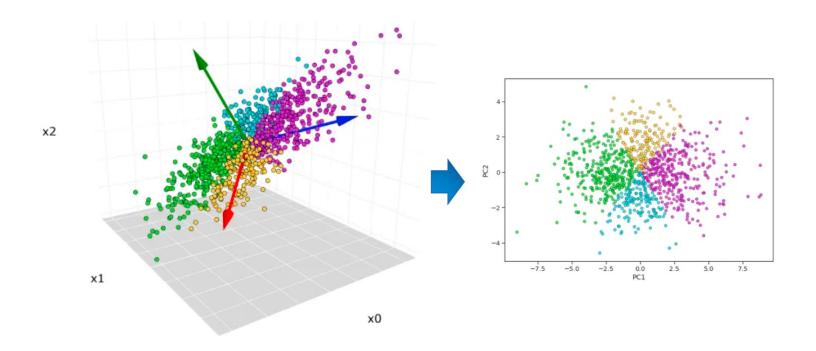
Principal Components Analysis

- PCA: most popular instance of second main class of unsupervised learning methods, **projection methods**, aka **dimensionality-reduction** methods
- Aim: find a **small number** of "directions" in input space that explain variation in input data; re-represent data by projecting along those directions
- Important assumption: variation contains information.
- Data is assumed to be continuous:
 - linear relationship between data and the learned representation

PCA: Common Tool

- Handles high-dimensional data
 - If data has thousands of dimensions, can be difficult for a classifier to deal with
- Often can be described by much lower dimensional representation
- Useful for:
 - Visualization
 - Preprocessing
 - Modeling prior for new data
 - Compression

PCA: Intuition



PCA: Intuition

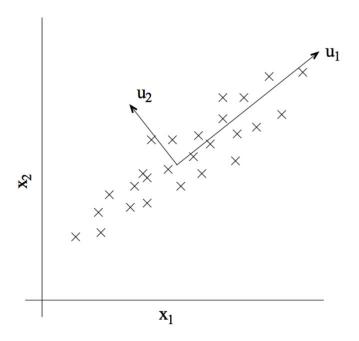
- As in the previous lecture, training data has N vectors, $\{x^{(n)}\}_{n=1}^N$, of dimensionality D, so $x^{(n)} \in \mathbb{R}^D$
- Aim to reduce dimensionality
 - linearly project to a much lower dimensional space, M << D::

$$xpprox U_{pca}z+a$$

66 where U_{pca} is a D imes M matrix and z is an M-dimensional vector

PCA: Intuition

- Search for orthogonal directions in space with the **highest variance**
 - o project data onto this subspace
- Structure of data vectors is encoded in sample covariance



Finding Principal Components

- To find the principal component directions, we center the data (subtract the sample mean from each variable)
- Calculate the **empirical covariance matrix**:

$$C = rac{1}{N} \sum_{n=1}^{N} (x^{(n)} - ar{x}) (x^{(n)} - ar{x})^T$$

- 66 with \bar{x} the mean
- What's the dimensionality of *C*?

Finding Principal Components

- Find the M eigenvectors with largest eigenvalues of C: these are the **principal components**
- Assemble these eigenvectors into a D imes M matrix U_{pca}
- We can now express D-dimensional vectors x by projecting them to M-dimensional

$$z = U_{pca}^T x$$

Standard PCA

- Algorithm: to find M components underlying D-dimensional data
- 1. Select the top M eigenvectors of C (data covariance matrix):

$$C = \; rac{1}{N} \sum_{n=1}^{N} (x^{(n)} - ar{x}) (x^{(n)} - ar{x})^T = U \Sigma U^T \; pprox \; U_{1:M} \Sigma_{1:M} U_{1:M}^T$$

66 where U is orthogonal, columns are unit-length eigenvectors

66

$$U^TU = UU^T = I$$

66 and Σ is a matrix with eigenvalues on the diagonal, representing the variance in the direction of each eigenvector

Standard PCA

• Matrix form of C:

$$C = \; rac{1}{N} (\mathbf{X} - ar{x}) (\mathbf{X} - ar{x})^T$$

66 where \mathbf{X} is a D-by-N matrix of data vectors, with each column being a data sample

Standard PCA

2. Project each input vector x into this subspace, e.g.,

$$z_j = u_j^T x; \qquad z = U_{1:M}^T x$$

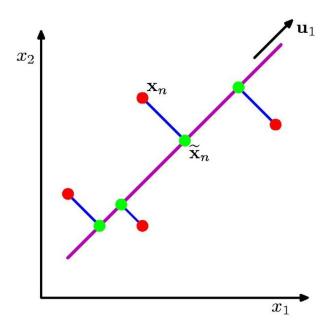
66 Let $U_{pca}=U_{1:M}.$ Then we have the principal components

66

$$z = U_{pca}^T x$$

Two Derivations of PCA

- Two views/derivations:
 - Maximize variance (scatter of green points)
 - Minimize error (red-green distance per datapoint)



PCA: Minimizing Reconstruction Error

- We can think of PCA as projecting the data onto a lower-dimensional subspace
- One derivation is that we want to find the projection such that the best linear reconstruction of the data is as close as possible to the original data

$$J(u,z,b) = \sum_n ||x^{(n)} - ilde{x}^{(n)}||^2$$

66 where

66

$$ilde{x}^{(n)} = \sum_{j=1}^{M} z_j^{(n)} u_j + \sum_{j=M+1}^{D} b_j u_j$$

PCA: Minimizing Reconstruction Error

• Objective minimized when first M components are the eigenvectors with the maximal eigenvalues

$$egin{align} z_j^{(n)} &= u_j^T x^{(n)}, \; orall j = 1, \ldots, M; \ b_j &= ar{x}^T u_j, \; orall j = M+1, \ldots, D. \end{cases}$$

66 In the maxtirx form:

66

$$x^{(n)} pprox \tilde{x}^{(n)} = U_{1:M} z^{(n)} + a$$

66 where
$$a=U_{M+1:D}b, \qquad b=U_{M+1:D}^Tar{x}$$

• If the mean \bar{x} is zero, then b=0 and a=0.

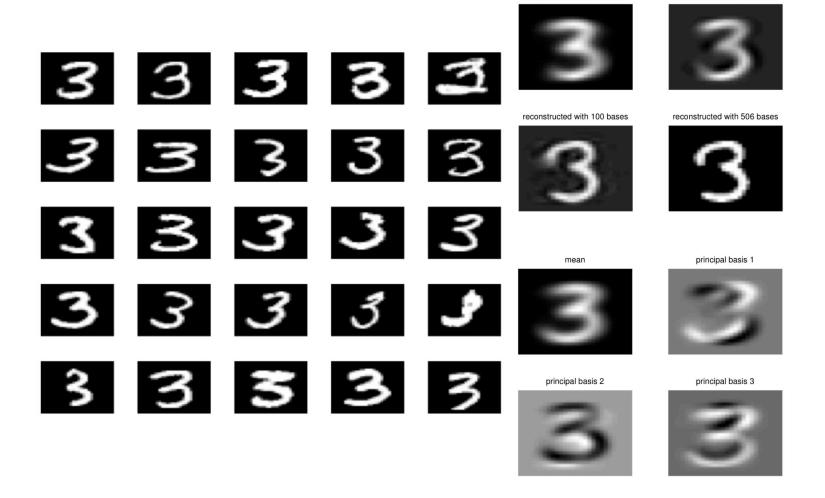
Applying PCA to Faces

- Run PCA on 2429 19×19 grayscale images
- Compresses the data: can get good reconstructions with only 3 components



- PCA for pre-processing: can apply classifier to latent representation
 - PCA with 3 components obtains **79% accuracy** on face/non-face discrimination on test data vs.76.8% for GMM with 84 states
- Can also be good for visualization

Applying PCA to Digits

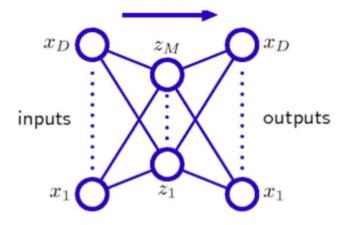


reconstructed with 2 bases

reconstructed with 10 bases

Relation to Neural Networks

- PCA is closely related to a particular form of neural network
- An autoencoder is a neural network whose outputs are its own inputs



• The goal is to minimize **reconstruction error**

Autoencoders

Define

$$z = f(Wx); \quad \hat{x} = g(Vz)$$

• Goal:

$$\min_{W,V} rac{1}{2N} \sum_{n=1}^N ||x^{(n)} - \hat{x}^{(n)}||^2$$

• If g and f are linear

$$\min_{W,V} \; rac{1}{2N} \sum_{n=1}^N ||x^{(n)} - VWx^{(n)}||^2$$

- In other words, the **optimal solution is PCA** for the case when the mean of the data is 0.
- What if the mean of the data is not 0?

Autoencoders: Nonlinear PCA

- What if g() is not linear?
- Then we are basically doing **nonlinear PCA**
- Some subtleties but in general this is an accurate description

Comparing Reconstructions



Real data

30-d deep autoencoder

30-d logistic PCA

30-d PCA

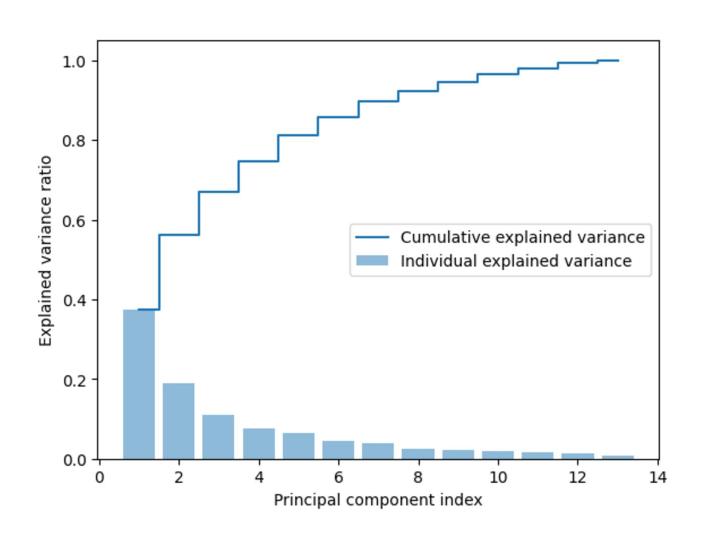
	0	1	2	3	4	5	6	7	8	9	10	11	12	13
0	1	14.23	1.71	2.43	15.6	127	2.80	3.06	0.28	2.29	5.64	1.04	3.92	1065
1	1	13.20	1.78	2.14	11.2	100	2.65	2.76	0.26	1.28	4.38	1.05	3.40	1050
2	1	13.16	2.36	2.67	18.6	101	2.80	3.24	0.30	2.81	5.68	1.03	3.17	1185
3	1	14.37	1.95	2.50	16.8	113	3.85	3.49	0.24	2.18	7.80	0.86	3.45	1480
4	1	13.24	2.59	2.87	21.0	118	2.80	2.69	0.39	1.82	4.32	1.04	2.93	735

```
. .
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
X, y = df_wine.iloc[:, 1:].values, df_wine.iloc[:, 0].values
X_train, X_test, y_train, y_test = train_test_split(
    X, y, test_size=0.3,
    stratify=y, random_state=0
# standardize the features
sc = StandardScaler()
X_train_std = sc.fit_transform(X_train)
X_test_std = sc.transform(X_test)
```

```
import numpy as np

cov_mat = np.cov(X_train_std.T)
eigen_vals, eigen_vecs = np.linalg.eig(cov_mat)
```

```
. .
import matplotlib.pyplot as plt
# 累加解释方差(特征值)之和
tot = sum(eigen_vals)
var_exp = [(i / tot) for i in sorted(eigen_vals, reverse=True)]
cum_var_exp = np.cumsum(var_exp)
plt.bar(range(1,14), var_exp, alpha=0.5,
        align='center', label='individual explained variance')
plt.step(range(1,14), cum_var_exp, where='mid',
         label='cumulative explained variance')
plt.ylabel('Explained variance ratio')
plt.xlabel('Principal component index')
plt.legend(loc='best')
plt.show()
```



```
●●●

# 生成(特征值,特征向量)的特征对, tuple类型

eigen_pairs = [(np.abs(eigen_vals[i]), eigen_vecs[:, i]) for i in range(len(eigen_vals))]

# 特征对按特征值降序排序

eigen_pairs.sort(key=lambda k: k[0], reverse=True)
```

```
Eigenvalues in descending order:
[(4.892308303273746, array([ 0.14669811, -0.24224554, -0.02993442, -0.25519002, 0.12079772,
       0.38934455, 0.42326486, -0.30634956, 0.30572219, -0.09869191, 0.30032535, 0.36821154, 0.29259713])),
(2.4663503157592244, array([ 0.50417079, 0.24216889, 0.28698484, -0.06468718, 0.22995385,
       0.09363991, 0.01088622, 0.01870216, 0.03040352, 0.54527081, -0.27924322, -0.174365, 0.36315461])),
(1.4280997275048464, array([-0.11723515, 0.14994658, 0.65639439, 0.58428234, 0.08226275,
       0.18080442, 0.14295933, 0.17223475, 0.1583621, -0.14242171, 0.09323872, 0.19607741, -0.09731711])),
(1.0123346209044963, array([ 0.20625461, 0.1304893 , 0.01515363, -0.09042209, -0.83912835,
       0.19317948, 0.14045955, 0.33733262, -0.1147529 , 0.07878571, 0.02417403, 0.18402864, 0.05676778])),
(0.8490645933450235, array([-0.18781595, 0.56863978, -0.29920943, -0.04124995, -0.02719713,
       0.14064543, 0.09268665, -0.08584168, 0.56510524, 0.01323461, -0.37261081, 0.08937967, -0.21752948])),
(0.6018151434229905, array([-0.14885132, -0.26905276, -0.09333861, -0.10134239, 0.11256735,
       0.01222488, -0.05503452, 0.69534088, 0.49835441, 0.15945216, 0.21651535, -0.23517236, 0.10562138])),
(0.5225154620639972, array([-0.17926366, -0.59263673, 0.06073346, 0.25032387, -0.28524056,
       0.05314553, 0.07989941, -0.29737172, 0.20251913, 0.39736411, -0.38465475, -0.08629033, -0.13029829])),
(0.33051429173094055, array([-0.40305492, -0.10183371, 0.35184142, -0.50045728, 0.08373917,
       0.13511146, 0.00336\overline{0}17, 0.19012076, -0.17602994, -0.21493067, -0.51725944, 0.13645604, 0.16775843])),
(0.29595018365934656, array([-0.41719758, 0.21710149, 0.12854985, 0.04733441, -0.27891878,
      -0.28098565, -0.0391443 , -0.27862219, 0.14853946, -0.00410241, 0.19781412, -0.23813815, 0.63735021])),
(0.23995530477949092, array([ 4.13320786e-04, -0.0878560762, -0.452518598, 0.486169765, 0.114764951,
       0.0945645138, -0.100444099, 0.200128778, -0.139942067, -0.115349466, -0.302254353, 0.318414303, 0.503247839])),
-0.61860015, -0.13968028, 0.00163324, 0.38856849, -0.3083459, -0.20045639, 0.28410033, 0.03755468])),
(0.16831253504096216, array([ 0.27566086, -0.0813845 , -0.01297513, 0.0989088 , -0.09592977,
       0.28389764, 0.11672921, -0.03965663, 0.08606027, -0.57165189, -0.19884453, -0.65086971, 0.07123771])),
(0.08414845672679461, array([-0.05546872, 0.03327316, -0.10061857, 0.05616586, 0.09584239,
      -0.42126512, 0.8472247, 0.1662568, -0.16619747, 0.03961736, -0.10538369, -0.09950556, -0.01606618]))]
```

```
w = np.hstack((eigen_pairs[0][1][:, np.newaxis], eigen_pairs[1][1][:, np.newaxis]))
print('Matrix W:\n', w)
```

```
Matrix U:
  [[ 0.14669811    0.50417079]
  [-0.24224554    0.24216889]
  [-0.02993442    0.28698484]
  [-0.25519002    -0.06468718]
  [ 0.12079772    0.22995385]
  [ 0.38934455    0.09363991]
  [ 0.42326486    0.01088622]
  [-0.30634956    0.01870216]
  [ 0.30572219    0.03040352]
  [-0.09869191    0.54527081]
  [ 0.36821154    -0.174365   ]
  [ 0.29259713    0.36315461]]
```

• In math

$$z = U^T x$$

• In python codes

$$z = x^T U$$

X_train_0_pca = X_train_std[0].dot(U) print(X_train_0_pca)

[2.59891628 0.00484089]

 $X_{train_pca} = X_{train_std.dot(U)}$

```
. . .
colors = ['r', 'b', 'g']
markers = ['s', 'x', 'o']
for l, c, m in zip(np.unique(y_train), colors, markers):
    plt.scatter(X_train_pca[y_train==1, 0],
                X_train_pca[y_train==1, 1],
                 c=c, label=l, marker=m)
plt.xlabel('PC 1')
plt.ylabel('PC 2')
plt.legend(loc='lower left')
plt.show()
```

