

ETC3250

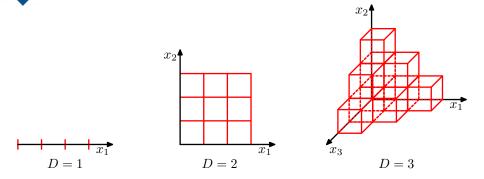
Business Analytics

Week 7
Principal Components Analysis

17 April 2018

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If we divide a region of a space into regular cells, then the number of such cells grows exponentially with the dimensionality of the space \implies We would need an exponentially large quantity of training data points in order to ensure that the cells are not empty.

Consider the **hyper-cube** $[-a, a]^d$ and the inscribed **hyper-sphere**. What does your intuition tell you about the relative sizes of these two objects?

- volume of sphere \approx volume of cube
- volume of sphere >> volume of cube
- volume of sphere << volume of cube</p>

$$s_d = rac{ ext{Volume(hyper-sphere)}}{ ext{Volume(hyper-cube)}} = rac{rac{a^d \pi^{d/2}}{\Gamma(rac{d}{2}+1)}}{(2a)^d} = \left(rac{\sqrt{\pi}}{2}
ight)^d rac{1}{\Gamma(rac{d}{2}+1)}$$

where $\Gamma(\cdot)$ is the Gamma function.

d	1	2	3	4	5	6
Sd	1	.785	.524	.308	.164	.080

- \blacksquare s_d does not depend on a_i just on d!
- As the dimension increases the volume of the area between the cube and the sphere becomes larger
- High dimensional spaces are strange → never trust your intuition in high dimensions!

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- Prediction is much more difficult near the edges of the training sample: extrapolation vs interpolation
 - \rightarrow Can we learn in high dimension?

- Most data points are closer to the boundary of the sample space than to any other data point
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Yes!

- First, real data will often be confined to a region of the space having *lower intrinsic dimensionality*. The data *lives* in a low dimensional subspace.
- Second, real data will typically exhibit some smoothness properties (at least locally)
 - → Dimensionality reduction

Wy dimensionality reduction?

- Curse of dimensionality
- Intrinsic dimensionality
- Visualization
- Reduce computation and storage

We avoid unnecessary dimensions which can be measured in two ways:

- features are not useful
- features are not independent

Dimension reduction methods

- Feature selection vs feature extraction
- Unsupervised vs supervised
- Linear vs nonlinear

Principal components analysis (PCA) is an unsupervised linear feature extraction method.

Principal components analysis

PCA produces a low-dimensional representation of a dataset. It finds a sequence of linear combinations of the variables that have maximal variance, and are mutually uncorrelated.

Why?

- We may have too many predictors for a regression. Instead, we can use the first few principal components.
- Understanding relationships between variables.
- We can plot a small number of variables

Principal components analysis

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Principal components analysis

The first <u>principal component</u> of a set of features x_1, x_2, \dots, x_p is the linear combination

$$z_1 = \phi_{11}x_1 + \phi_{21}x_2 + \cdots + \phi_{p1}x_p$$

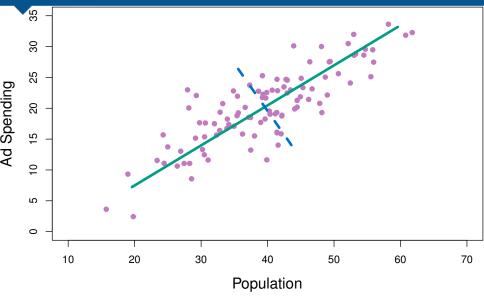
that has the <u>largest variance</u> such that $\sum_{j=1}^{r} \phi_{j1}^2 = 1$.

 $\overline{\phi_{11},\ldots,\phi_{p1}}$ are the <u>loadings</u> of the first principal component.

Geometry of PCA

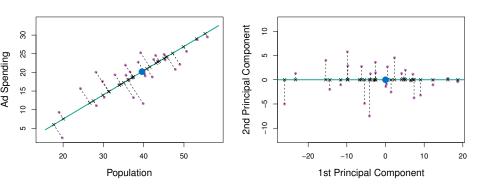
- The loading vector $\phi_1 = [\phi_{11}, \dots, \phi_{p1}]'$ defines a direction in feature space along which the data vary the most.
- If we project the n data points $\mathbf{x}_1, \dots, \mathbf{x}_n$ onto this direction, the projected values are the principal component scores z_{11}, \dots, z_{n1} .
- The second principal component is the linear combination $z_{i2} = \phi_{12}x_{i1} + \phi_{22}x_{i2} + \cdots + \phi_{p2}x_{ip}$ that has maximal variance among all linear combinations that are *uncorrelated* with z_1 .
- Equivalent to constraining ϕ_2 to be orthogonal (perpendicular) to ϕ_1 . And so on.
- There are at most min(n-1,p) PCs.

PCA Example



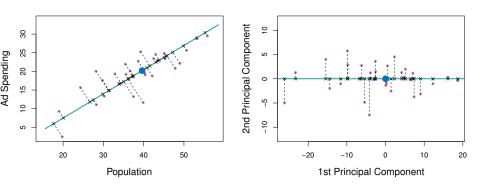
Green: first PC. Blue: second PC

Further principal components



PCA can be thought of as fitting an *n*-dimensional ellipsoid to the data, where each axis of the ellipsoid represents a principal component.

Further principal components



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Suppose we have a $n \times p$ data set $\mathbf{X} = [x_{ij}]$.

- Centre each of the variables to have mean zero (i.e., the column means of X are zero).
- $z_{i1} = \phi_{11}x_{i1} + \phi_{21}x_{i2} + \cdots + \phi_{p1}x_{ip}$
- Sample variance of z_{i1} is $\frac{1}{n} \sum_{i=1}^{n} z_{i1}^2$.

Compute the covariance matrix (after scaling the columns of X)

$$C = X'X$$

Find eigenvalues and eigenvectors:

$$C = VDV'$$

where columns of \mathbf{V} are orthonormal (i.e., $\mathbf{V}'\mathbf{V} = \mathbf{I}$)

3 Compute PCs: $\Phi = V$. $Z = X\Phi$.

Singular Value Decomposition

$$X = U\Lambda V'$$

- **X** is $n \times p$ matrix
- **U** is $n \times r$ matrix with orthonormal columns $(\mathbf{U}'\mathbf{U} = \mathbf{I})$
- Λ is $r \times r$ diagonal matrix with non-negative elements.
- V is $p \times r$ matrix with orthonormal columns (V'V = I).

It is always possible to uniquely decompose a matrix in this way.

- **1** Compute SVD: $\mathbf{X} = \mathbf{U}\Lambda\mathbf{V}'$.
- **2** Compute PCs: $\Phi = V$. $Z = X\Phi$.

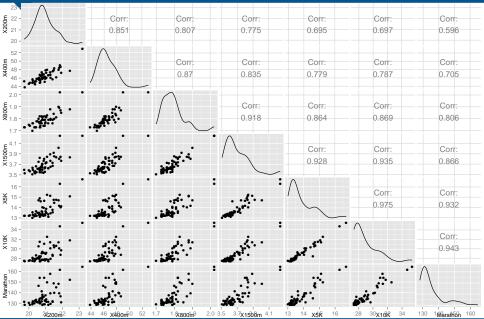
Relationship with covariance:

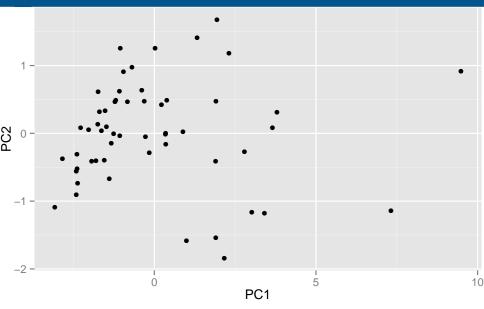
$$\mathbf{C} = \mathbf{X}'\mathbf{X} = \mathbf{V}\Lambda\mathbf{U}'\mathbf{U}\Lambda\mathbf{V}' = \mathbf{V}\Lambda^2\mathbf{V}' = \mathbf{V}\mathbf{D}\mathbf{V}'$$

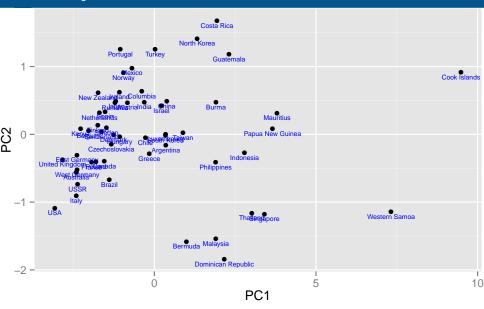
- Eigenvalues of C are squares of singular values of X.
- Eigenvectors of **C** are right singular vectors of **X**.
- The PC directions $\phi_1, \phi_2, \phi_3, \ldots$ are the right singular vectors of the matrix \mathbf{X} .

The data on national track records for men are listed in the following table (as at 1984):

Country	100m	200m	400m	800m	1500m	5000m	10000m	Marathon
	<i>(s)</i>	<i>(s)</i>	<i>(s)</i>	(min)	(min)	(min)	(min)	(min)
Argentina	10.39	20.81	46.84	1.81	3.70	14.04	29.36	137.72
Australia	10.31	20.06	44.84	1.74	3.57	13.28	27.66	128.30
Austria	10.44	20.81	46.82	1.79	3.60	13.26	27.72	135.90
Belgium	10.34	20.68	45.04	1.73	3.60	13.22	27.45	129.95
Bermuda	10.28	20.58	45.91	1.80	3.75	14.68	30.55	146.62
Brazil	10.22	20.43	45.21	1.73	3.66	13.62	28.62	133.13
:								
Turkey	10.71	21.43	47.60	1.79	3.67	13.56	28.58	131.50
USA	9.93	19.75	43.86	1.73	3.53	13.20	27.43	128.22
USSR	10.07	20.00	44.60	1.75	3.59	13.20	27.53	130.55
W.Samoa	10.82	21.86	49.00	2.02	4.24	16.28	34.71	161.83







> pca

Standard deviations:

[1] 2.573 0.937 0.399 0.352 0.283 0.261 0.215 0.150

Rotation:

	PC1	PC2	PC3	PC4	PC5	PC6	PC7	PC8
X100m	0.318	0.5669	0.332	-0.1276	0.263	-0.5937	0.13624	-0.105542
X200m	0.337	0.4616	0.361	0.2591	-0.154	0.6561	-0.11264	0.096054
X400m	0.356	0.2483	-0.560	-0.6523	-0.218	0.1566	-0.00285	0.000127
X800m	0.369	0.0124	-0.532	0.4800	0.540	-0.0147	-0.23802	0.038165
X1500m	0.373	-0.1398	-0.153	0.4045	-0.488	-0.1578	0.61001	-0.139291
X5K	0.364	-0.3120	0.190	-0.0296	-0.254	-0.1413	-0.59130	-0.546697
X10K	0.367	-0.3069	0.182	-0.0801	-0.133	-0.2190	-0.17687	0.796795
Marathon	0.342	-0.4390	0.263	-0.2995	0.498	0.3153	0.39882	-0.158164

Proportion of variance explained

Total variance in data (assuming variables centered at 0):

$$TV = \sum_{j=1}^{p} Var(x_j) = \sum_{j=1}^{p} \frac{1}{n} \sum_{i=1}^{n} x_{ij}^2$$

Variance explained by mth PC:

$$V_m = \text{Var}(z_m) = \frac{1}{n} \sum_{i=1}^n z_{im}^2$$

$$\mathsf{TV} = \sum_{m=1}^M V_m$$
 where $M = \min(n-1, p)$.

Proportion of variance explained:

$$PVE_m = V_m/TV$$

Scree plots and biplots

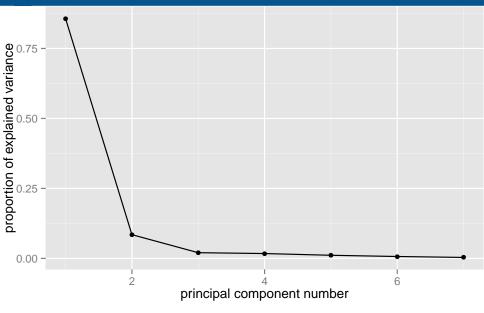
Scree plot

Plot of variance explained by each component vs number of component.

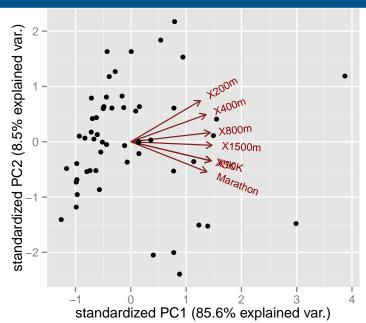
Biplot

Plot of PC2 vs PC1, overlaid with directions of the loading vectors (ϕ_{i1}, ϕ_{i2}) .

Scree plot



Biplot



Scaling

- If the variables are in different units, scaling each to have standard deviation equal to one is recommended.
- If they are in the same units, you might or might not scale the variables.