Principal Component Analysis (PCA) Application to images

Václav Hlaváč

Czech Technical University in Prague
Center for Machine Perception *(bridging groups of the)*
Czech Institute of Informatics, Robotics and Cybernetics and
Faculty of Electrical Engineering, Department of Cybernetics
`http://people.ciirc.cvut.cz/hlavac`, `hlavac@ciirc.cvut.cz`

Outline of the lecture:

- Principal components, informal idea.

- Needed linear algebra.

- Least-squares approximation.

- PCA derivation, PCA for images.

- Drawbacks. Interesting behaviors live in manifolds.
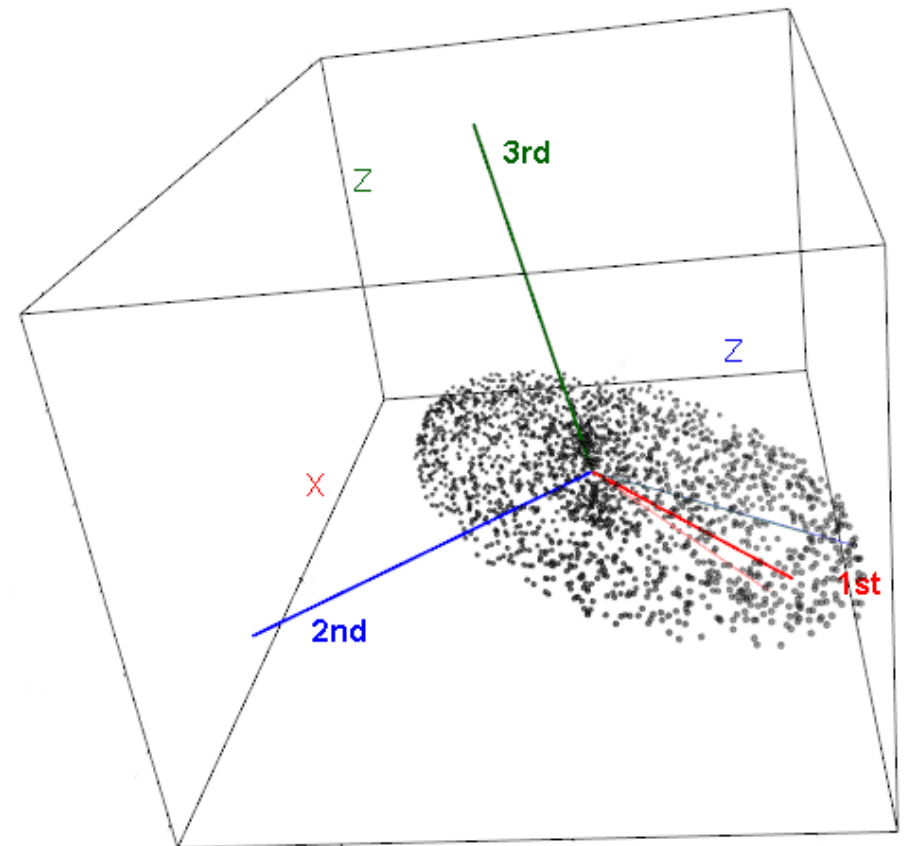
- Subspace methods, LDA, CCA, . . .

◆ PCA seeks to represent observations (or signals, images, and general data) in a form that enhances the mutual independence of contributory components.

◆ One observation is assumed to be a point in a p-dimensional linear space.

◆ This linear space has some 'natural' orthogonal basis vectors. It is of advantage to express observation as a linear combination with regards to this 'natural' base (given by eigen-vectors as we will see later).

◆ PCA is mathematically defined as an orthogonal linear transformation that transforms the data to a new coordinate system such that the greatest variance by some projection of the data comes to lie on the first coordinate (called the first principal component), the second greatest variance on the second coordinate, and so on.

PCA objective is to rotate rigidly the coordinate axes of the p-dimensional linear space to new 'natural' positions (principal axes) such that:

◆ Coordinate axes are ordered such that principal axis 1 corresponds to the highest variance in data, axis 2 has the next highest variance, ..., and axis p has the lowest variance.

◆ The covariance among each pair of principal axes is zero, i.e. they are uncorrelated.

◆ Two-dimensional vector space of observations, (x_1, x_2).

◆ Each observation corresponds to a single point in the vector space.

◆ The goal:
Find another basis of the vector space, which treats variations of data better.

_____

◆ We will see later:

Data points (observations) are represented in a rotated orthogonal coordinate system. The origin is the mean of the data points and the axes are provided by the eigenvectors.
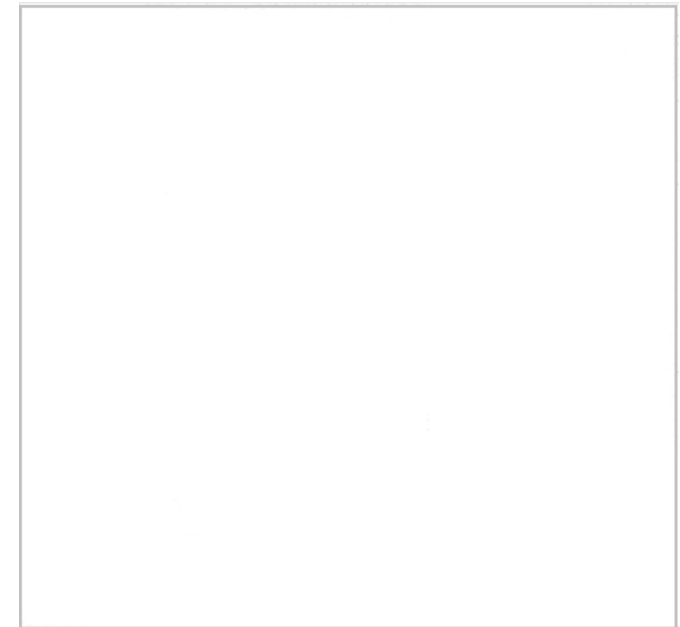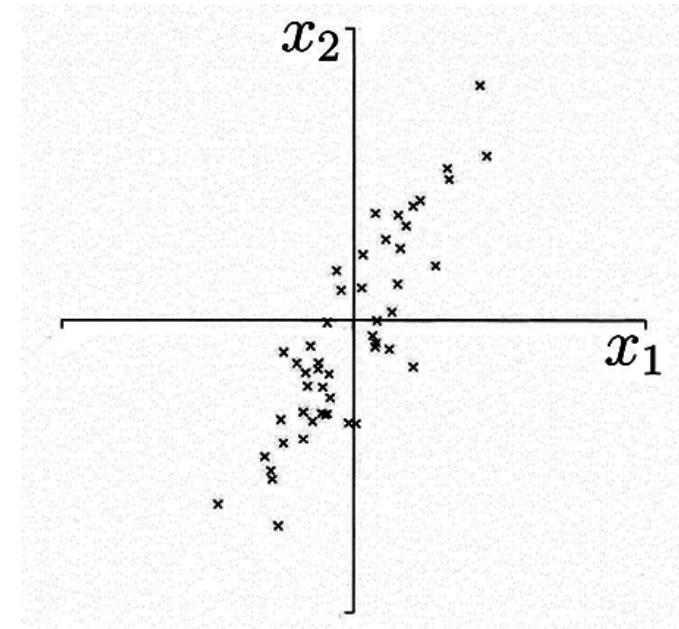
Geometric motivation, principal components (2)

◆ Assume a single straight line approximating best the observation in the least-square sense, i.e. by minimizing the sum of distances between data points and the line.

◆ The first principal direction (component) is the direction of this line. Let it be a new basis vector z_1.

◆ The second principal direction (component, basis vector) z_2 is a direction perpendicular to z_1 and minimizing the distances to data points to a corresponding straight line.

◆ For higher dimensional observation spaces, this construction is repeated.
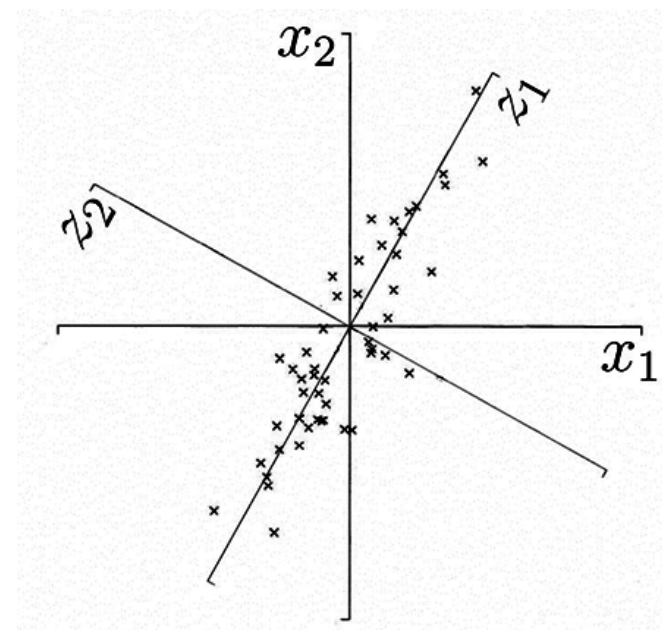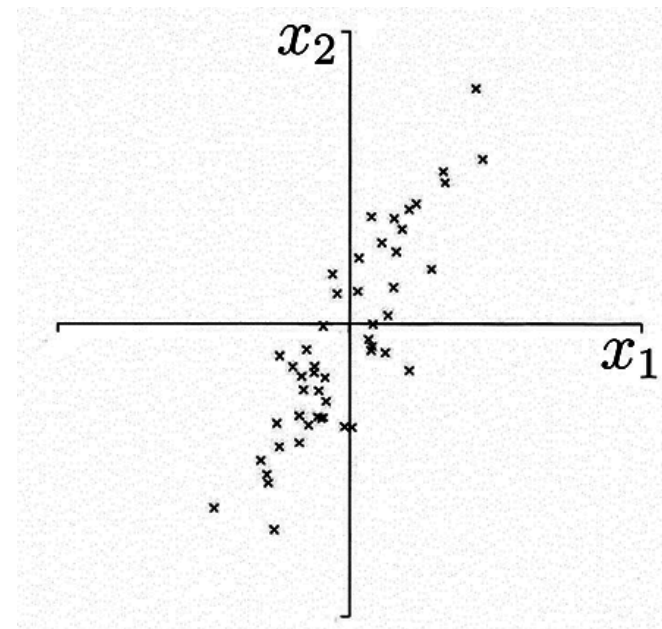
Eigen-values, eigen-vectors

◆ Assume a square $n \times n$ regular matrix A.

◆ Eigen-vectors are solutions of the eigen-equation $A\mathbf{x} = \lambda\mathbf{x}$, where the vector λ is contains eigen-values λ_i, $i = 1, \ldots, n$, (which may be complex).

◆ Let us derive: $A\mathbf{x} = \lambda\mathbf{x} \Rightarrow A\mathbf{x} - \lambda\mathbf{x} = 0 \Rightarrow (A - \lambda I)\mathbf{x} = 0$. Matrix I is the identity matrix. The equation $(A - \lambda I)\mathbf{x} = 0$ has the non-zero solution $\mathbf{x}$ if and only if $\det(A - \lambda I) = 0$. The polynomial $\det(A - \lambda I)$ is called the characteristic polynomial of the matrix A. The fundamental theorem of algebra implies that the characteristic polynomial can be factored, i.e. $\det(A - \lambda I) = 0 = (\lambda_1 - \lambda)(\lambda_2 - \lambda) \ldots (\lambda_n - \lambda)$. Eigen-values λ_i are not necessarily distinct. Multiple eigen-values arise from multiple roots of the characteristic polynomial.

◆ We start reviewing eigen-analysis from a deterministic, linear algebra standpoint.

◆ Later, we will develop a statistical view based on covariance matrices and principal component analysis.

♦ A system of linear equations can be expressed in a matrix form as $A\mathbf{x} = \mathbf{b}$, where A is the matrix of the system.

Example:

$$\left.\begin{array}{rrrrrrr} x & + & 3y & - & 2z & = & 5 \\ 3x & + & 5y & + & 6z & = & 7 \\ 2x & + & 4y & + & 3z & = & 8 \end{array}\right\} \implies A = \begin{bmatrix} 1 & 3 & -2 \\ 3 & 5 & 6 \\ 2 & 4 & 3 \end{bmatrix}, \quad \mathbf{b} = \begin{bmatrix} 5 \\ 7 \\ 8 \end{bmatrix}.$$

♦ The augmented matrix of the system is created by concatenating a column vector $\mathbf{b}$ to the matrix A, i.e., $[A|\mathbf{b}]$.

$$\text{Example:} \quad [A|\mathbf{b}] = \left[\begin{array}{rrr|r} 1 & 3 & -2 & 5 \\ 3 & 5 & 6 & 7 \\ 2 & 4 & 3 & 8 \end{array}\right].$$

♦ This system has a unique solution if and only if the rank of the matrix A is equal to the rank of the extended matrix $[A|\mathbf{b}]$.

◆ Let A be a regular matrix.

◆ Matrices A and B with real or complex entries are called similar if there exists an invertible square matrix P such that $P^{-1}A\,P = B$.

◆ Matrix P is called the change of basis matrix.

◆ The similarity transformation refers to a matrix transformation that results in similar matrices.

◆ Similar matrices have useful properties: they have the same rank, determinant, trace, characteristic polynomial, minimal polynomial and eigen-values (but not necessarily the same eigen-vectors).

◆ Similarity transformations allow us to express regular matrices in several useful forms, e.g. Jordan canonical form, Frobenius normal form (called also rational canonical form).

◆ Any complex square matrix is similar to a matrix in the Jordan canonical form

$$
\begin{bmatrix} J_1 & & 0 \\ & \ddots & \\ 0 & & J_p \end{bmatrix}
, \text{ where } J_i \text{ are Jordan blocks }
\begin{bmatrix} \lambda_i & 1 & & 0 \\ 0 & \lambda_i & \ddots & 0 \\ 0 & 0 & \ddots & 1 \\ 0 & 0 & & \lambda_i \end{bmatrix},
$$

in which λ_i are the multiple eigen-values.

◆ The multiplicity of the eigen-value gives the size of the Jordan block.

◆ If the eigen-value is not multiple then the Jordan block degenerates to the eigen-value itself.

Least-square approximation

◆ Assume that abundant data comes from many observations or measurements. This case is very common in practice.

◆ We intent to approximate the data by a linear model - a system of linear equations, e.g. a straight line in particular.

◆ Strictly speaking, the observations are likely to be in a contradiction with respect to the system of linear equations.

◆ In the deterministic world, the conclusion would be that the system of linear equations has no solution.

◆ There is an interest in finding the solution to the system, which is in some sense 'closest' to the observations, perhaps compensating for noise in observations.

◆ We will usually adopt a statistical approach by minimizing the least square error.

◆ PCA is a powerful and widely used linear technique in statistics, signal processing, image processing, and elsewhere.

◆ Several names: the (discrete) Karhunen-Loève transform (KLT, after Kari Karhunen, 1915-1992 and Michael Loève, 1907-1979) or the Hotelling transform (after Harold Hotelling, 1895-1973). Invented by Pearson (1901) and H. Hotelling (1933).

◆ In statistics, PCA is a method for simplifying a multidimensional dataset to lower dimensions for analysis, visualization or data compression.

◆ PCA represents the data in a new coordinate system in which basis vectors follow modes of greatest variance in the data.

◆ Thus, new basis vectors are calculated for the particular data set.

◆ The price to be paid for PCA's flexibility is in higher computational requirements as compared to, e.g., the fast Fourier transform.

◆ Suppose a data set comprising N observations, each of M variables (dimensions). Usually $N \gg M$.

◆ The aim: to reduce the dimensionality of the data so that each observation can be usefully represented with only L variables, $1 \leq L < M$.

◆ Data are arranged as a set of N column data vectors, each representing a single observation of M variables: the n-th observations is a column vector $\mathbf{x}_n = (x_1, \ldots, x_M)^\top$, $n = 1, \ldots, N$.

◆ We thus have an $M \times N$ data matrix X. Such matrices are often huge because N may be very large: this is in fact good, since many observations imply better statistics.

◆ This procedure is not applied to the raw data R but to normalized data X as follows.

◆ The raw observed data is arranged in a matrix R and the empirical mean is calculated along each row of R. The result is stored in a vector $\mathbf{u}$ the elements of which are scalars

$$u(m) = \frac{1}{N} \sum_{n=1}^{N} R(m, n) , \quad \text{where } m = 1, \ldots, M .$$

◆ The empirical mean is subtracted from each column of R: if $\mathbf{e}$ is a unitary vector of size N (consisting of ones only), we will write

$$X = R - \mathbf{u}\,\mathbf{e} .$$

If we approximate higher dimensional space X (of dimension M) by the lower dimensional matrix Y (of dimension L) then the mean square error ε^2 of this approximation is given by

$$\varepsilon^2 = \frac{1}{N} \sum_{n=1}^{N} |\mathbf{x}_n|^2 - \sum_{i=1}^{L} \mathbf{b}_i^\top \left(\frac{1}{N} \sum_{n=1}^{N} \mathbf{x}_n \, \mathbf{x}_n^\top \right) \mathbf{b}_i \, ,$$

where $\mathbf{b}_i$, $i = 1, \ldots, L$ are basis vector of the linear space of dimension L.

If ε^2 has to be minimal then the following term has to be maximal

$$\sum_{i=1}^{L} \mathbf{b}_i^\top \operatorname{cov}(\mathbf{x}) \, \mathbf{b}_i \, , \quad \text{where} \ \operatorname{cov}(\mathbf{x}) = \sum_{n=1}^{N} \mathbf{x}_n \, \mathbf{x}_n^\top \, ,$$

is the covariance matrix.

◆ The covariance matrix $\mathrm{cov}(\mathbf{x})$ has special properties: it is real, symmetric and positive semi-definite.

◆ So the covariance matrix can be guaranteed to have real eigen-values.

◆ Matrix theory tells us that these eigen-values may be sorted (largest to smallest) and the associated eigen-vectors taken as the basis vectors that provide the maximum we seek.

◆ In the data approximation, dimensions corresponding to the smallest eigen-values are omitted. The mean square error ε^2 is given by

$$\varepsilon^2 = \mathrm{trace}\left(\,\mathrm{cov}(\mathbf{x})\right) - \sum_{i=1}^{L} \lambda_i = \sum_{i=L+1}^{M} \lambda_i \,,$$

where $\mathrm{trace}(A)$ is the *trace*—sum of the diagonal elements—of the matrix A. The trace equals the sum of all eigenvalues.

Can we use PCA for images?

◆ It took a while to realize (Turk, Pentland, 1991), but yes.

◆ Let us consider a 321×261 image.

◆ The image is considered as a very long 1D vector by concatenating image pixels column by column (or alternatively row by row), i.e. $321 \times 261 = 83781$.

◆ The huge number 83781 is the dimensionality of our vector space.

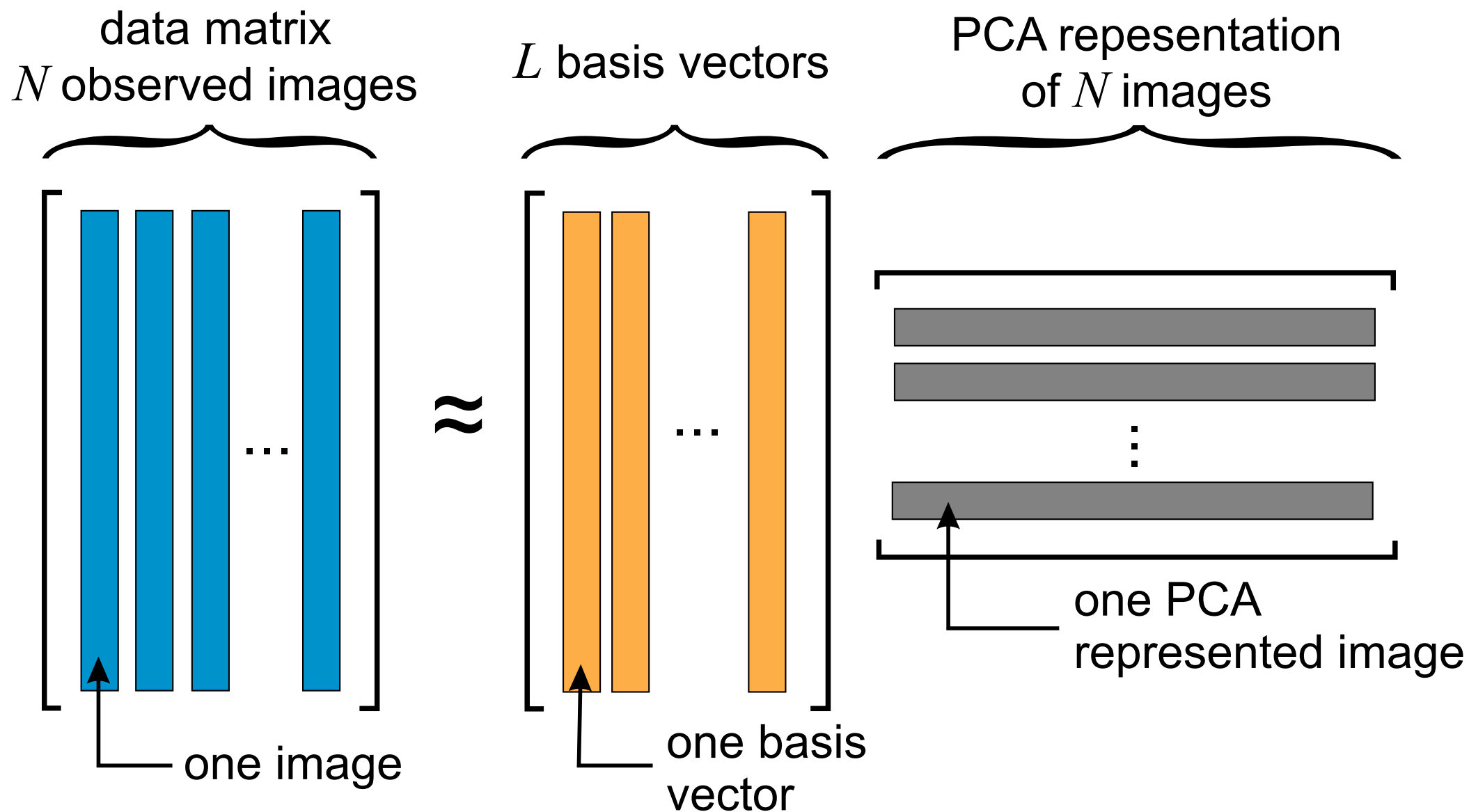◆ The intensity variation is assumed in each pixel of the image.

What if we have 32 instances of images?

◆ We have only 32 observations and 83781 unknowns in our example!

◆ The induced system of linear equations is not over-constrained but under-constrained.

◆ PCA is still applicable.

◆ The number of principle components is less than or equal to the number of observations available (32 in our particular case). This is because the (square) covariance matrix has a size corresponding to the number of observations.

◆ The eigen-vectors we derive are called eigen-images, after rearranging back from the 1D vector to a rectangular image.

◆ Let us perform the dimensionality reduction from 32 to 4 in our example.

PCA, graphical illustration

data matrix
N observed images

L basis vectors

PCA repesentation
of N images

$\approx$

...

...

one image

one basis
vector

one PCA
represented image

◆ Reconstruction of the image from four basis vectors $\mathbf{b}_i$, $i = 1, \ldots, 4$ which can be displayed as images.

◆ The linear combination was computed as $q_1\mathbf{b}_1 + q_2\mathbf{b}_2 + q_3\mathbf{b}_3 + q_4\mathbf{b}_4 = 0.078\,\mathbf{b}_1 + 0.062\,\mathbf{b}_2 - 0.182\,\mathbf{b}_3 + 0.179\,\mathbf{b}_4$.

$= q_1$ $+ q_2$ $+ q_3$ $+ q_4$

◆ By rearranging pixels column by column to a 1D vector, relations of a given pixel to pixels in neighboring rows are not taken into account.

◆ Another disadvantage is in the global nature of the representation; small change or error in the input images influences the whole eigen-representation. However, this property is inherent in all linear integral transforms.

Reconstructive (also generative) representation

♦ Enables (partial) reconstruction of input images (hallucinations).

♦ It is general. It is not tuned for a specific task.

♦ Enables closing the feedback loop, i.e. bidirectional processing.

Discriminative representation

♦ Does not allow partial reconstruction.

♦ Less general. A particular task specific.

♦ Stores only information needed for the decision task.

◆ Images, as we saw, lead to enormous dimensionality.

◆ The data of interest often live in a much lower-dimensional subspace called the manifold.

◆ Example (courtesy Thomas Brox):
The 100×100 image of the number 3 shifted and rotated, i.e. there are only 3 degrees of variations.
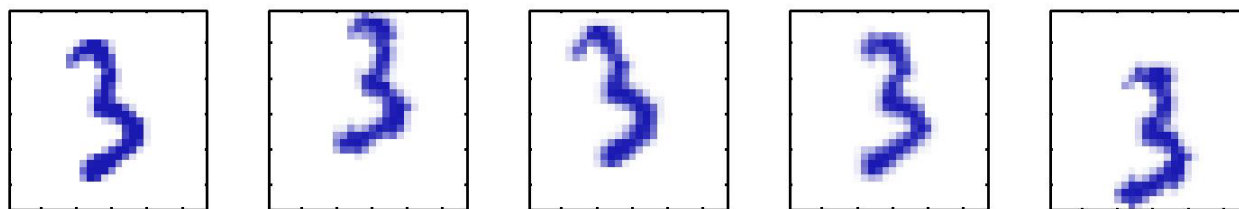
All data points live in a 3-dimensional manifold of the 10,000-dimensional observation space.

◆ The difficulty of the task is to find out empirically from the data in which manifold the data vary.

Subspace methods explore the fact that data (images) can be represented in a subspace of the original vector space in which data live.

Different methods examples:

Method (abbreviation)	Key property
Principal Component Analysis (PCA)	reconstructive, unsupervised, optimal reconstruction, minimizes squared reconstruction error, maximizes variance of projected input vectors
Linear Discriminative Analysis (LDA)	discriminative, supervised, optimal separation, maximizes distance between projection vectors
Canonical Correlation Analysis (CCA)	supervised, optimal correlation, motivated by regression task, e.g. robot localization
Independent Component Analysis (ICA)	independent factors
Non-negative matrix factorization (NMF)	non-negative factors
Kernel methods for nonlinear extension	local straightening by kernel functions