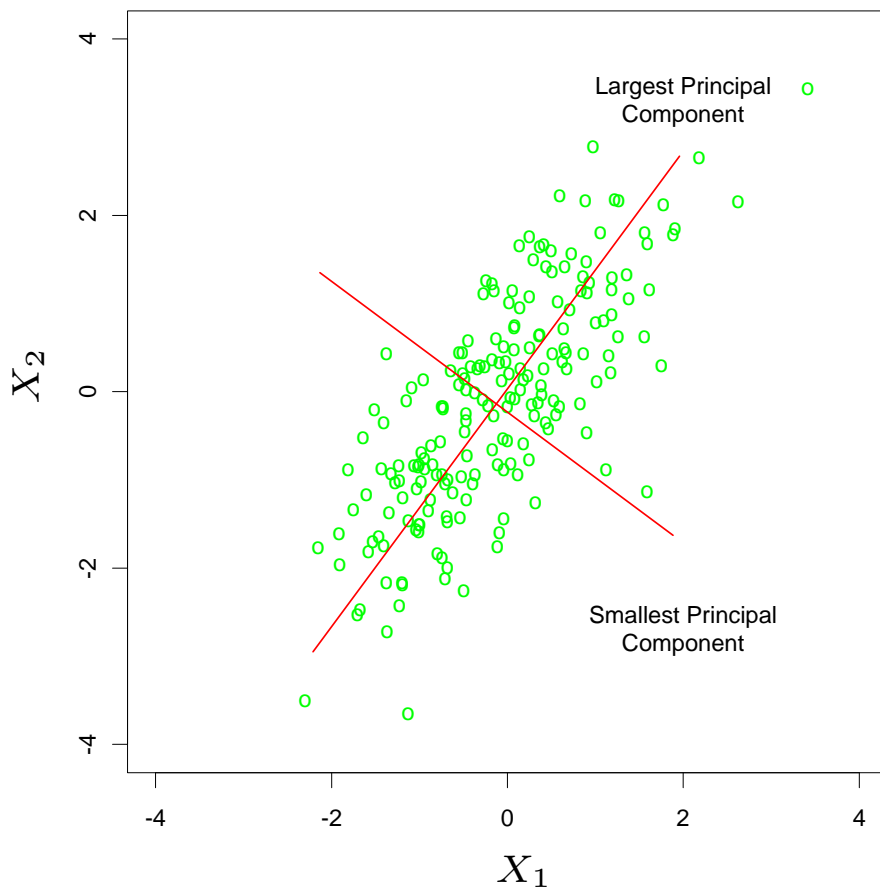


Principal Components

Suppose we have N measurements on each of p variables X_j , $j = 1, \dots, p$. There are several equivalent approaches to principal components:

- Given $X = (X_1, \dots, X_p)$, produce a derived (and small) set of uncorrelated variables $Z_k = X\alpha_k$, $k = 1, \dots, q < p$ that are linear combinations of the original variables, and that explain most of the variation in the original set.
- Approximate the original set of N points in \mathbb{R}^p by a least-squares optimal linear manifold of co-dimension $q < p$.
- Approximate the $N \times p$ data matrix \mathbf{X} by the best rank- q matrix $\hat{\mathbf{X}}_{(q)}$. This is the usual motivation for the SVD.

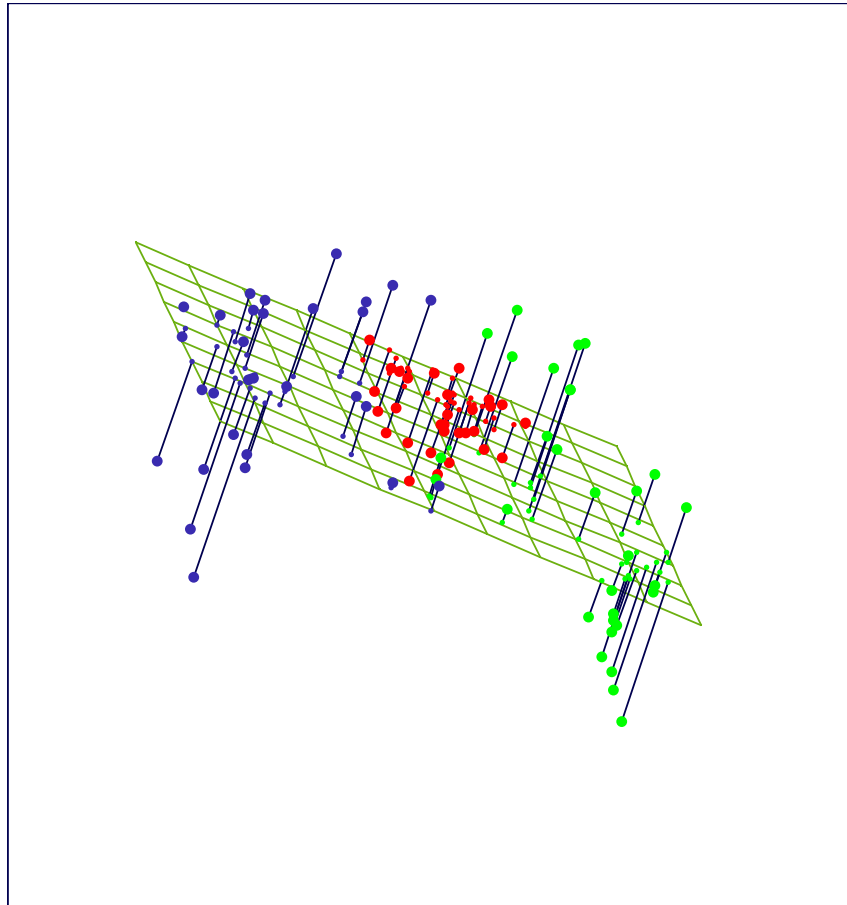
PC: Derived Variables



$Z_1 = X\alpha_1$ is the projection of the data onto the longest **direction**, and has the largest variance amongst all such normalized projections.

α_1 is the eigenvector corresponding to the largest eigenvalue of $\hat{\Sigma}$, the sample covariance matrix of X . Z_2 and α_2 correspond to the second-largest eigenvector.

PC: Least Squares Approximation



Find the linear manifold $f(\lambda) = \mu + \mathbf{V}_q \lambda$ that best approximates the data in a least-squares sense:

$$\min_{\mu, \{\lambda_i\}, \mathbf{V}_q} \sum_{i=1}^N \|x_i - \mu - \mathbf{V}_q \lambda_i\|^2.$$

Solution: $\mu = \bar{x}$, $v_k = \alpha_k$, $\lambda_k = \mathbf{V}_q^T (x_i - \bar{x})$.

PC: Singular Value Decomposition

Let $\tilde{\mathbf{X}}$ be the $N \times p$ data matrix with centered columns (assume $N > p$).

$$\tilde{\mathbf{X}} = \mathbf{U}\mathbf{D}\mathbf{V}^T$$

is the **SVD** of $\tilde{\mathbf{X}}$, where

- \mathbf{U} is $N \times p$ orthogonal, the left singular vectors.
- \mathbf{V} is $p \times p$ orthogonal, the right singular vectors.
- \mathbf{D} is diagonal, with $d_1 \geq d_2 \geq \dots \geq d_p \geq 0$, the singular values.

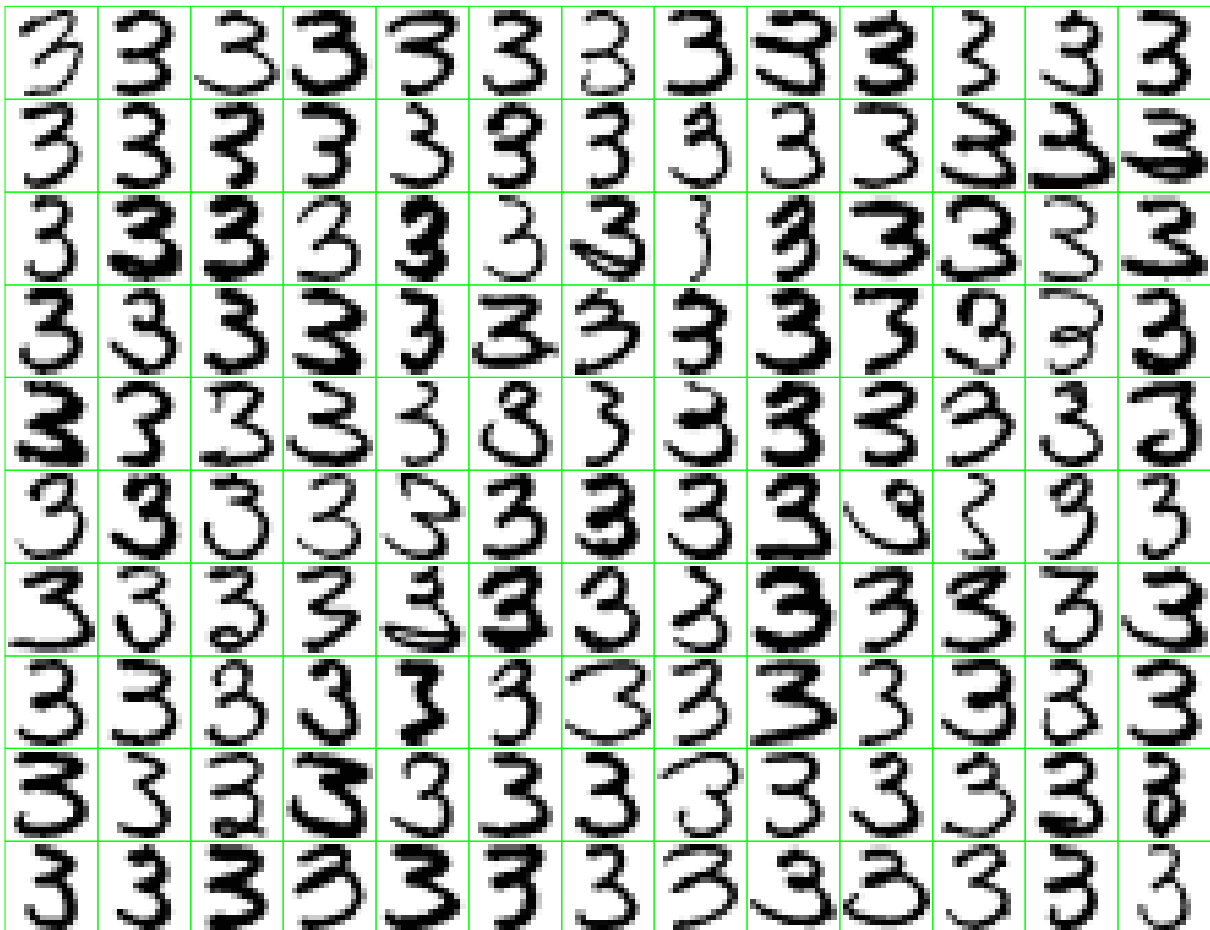
The SVD always exists, and is unique up to signs.

The columns of \mathbf{V} are the principal components, and $Z_j = U_j d_j$.

Let \mathbf{D}_q be \mathbf{D} , with all but the first q diagonal elements set to zero. Then $\hat{\mathbf{X}}_q = \mathbf{U}\mathbf{D}_q\mathbf{V}^T$ solves

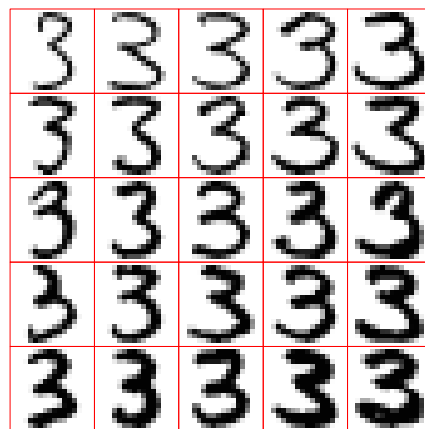
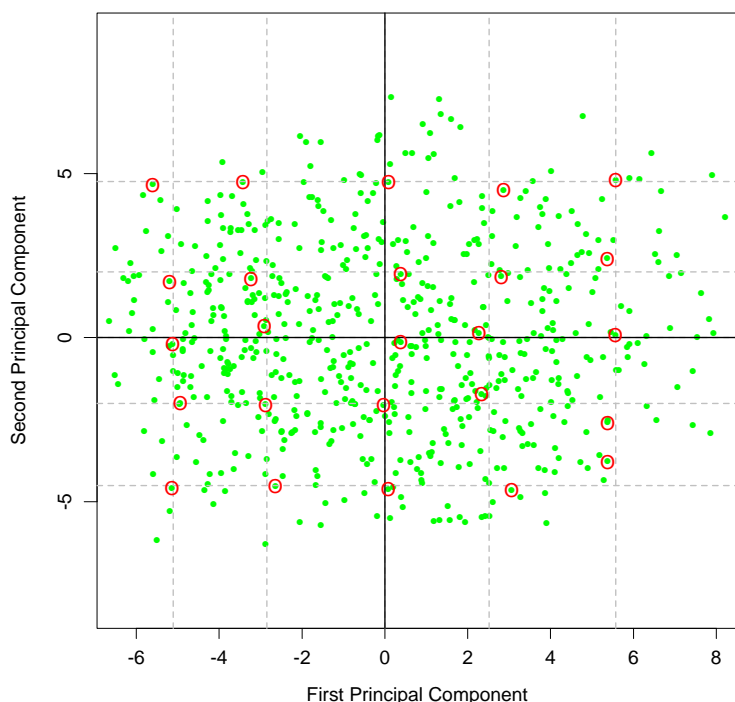
$$\min_{\text{rank}(\hat{\mathbf{X}}_q)=q} \|\tilde{\mathbf{X}} - \hat{\mathbf{X}}_q\|$$

PC: Example — Digit Data



130 threes, a subset of 638 such threes and part of the handwritten digit dataset. Each three is a 16×16 greyscale image, and the variables X_j , $j = 1, \dots, 256$ are the greyscale values for each pixel.

Rank-2 Model for Threes



Two-component model has the form

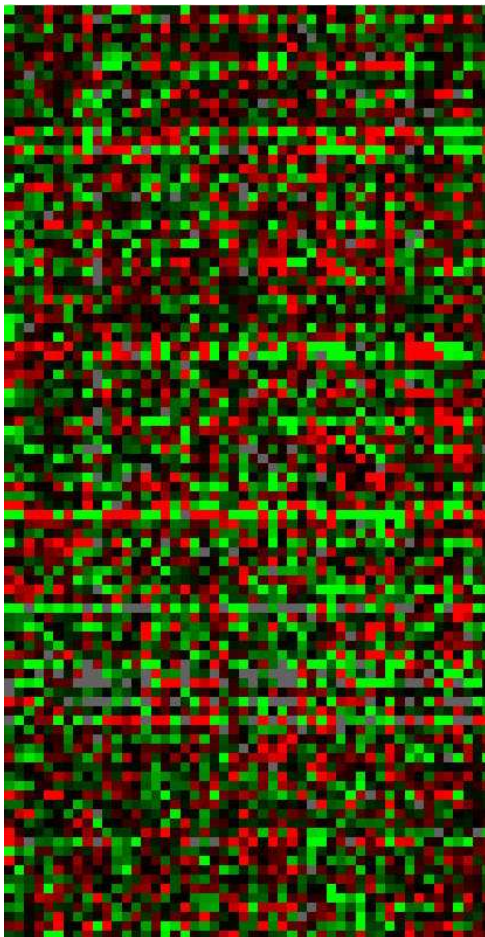
$$\begin{aligned}\hat{f}(\lambda) &= \bar{x} + \lambda_1 v_1 + \lambda_2 v_2 \\ &= \text{[Image of } \bar{x}] + \lambda_1 \cdot \text{[Image of } v_1] + \lambda_2 \cdot \text{[Image of } v_2].\end{aligned}$$

Here we have displayed the first two principal component directions, v_1 and v_2 , as images.

SVD: Expression Arrays

The rows are genes (variables) and the columns are observations (samples, DNA arrays).

Typically 6-10K genes, 50 samples.



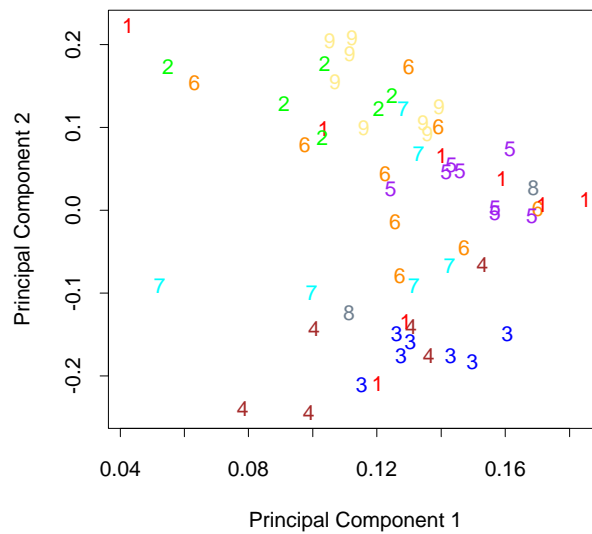
Eigengenes

- The first principal component or **eigengene** is the linear combination of the genes showing the most variation over the samples.
- The individual gene loadings for each eigengene or **eigenarrays** can have biological meaning.
- The sample values for the eigengenes show useful low-dimensional projections.

Example: NCI Cancer Data

First two eigengenes

Points are colored according to NCI cancer classes



First two eigenarrays

