

Principal component methods

Readings for today

- Chapter 6: Linear model selection and regularization. James, G., Witten, D., Hastie, T., & Tibshirani, R. (2013). An introduction to statistical learning: with applications in R (Vol. 6). New York: Springer

Topics

1. Principal components analysis
2. Principal component regression
3. Partial least squares

Principal component analysis

Dimensionality

Dimensionality of a model: $n \times p$

As $n \rightarrow p$, dimensionality increases & model variance increases

$$\begin{pmatrix} x_{1,1} \\ \dots \\ x_{n,1} \end{pmatrix} \rightarrow \begin{pmatrix} x_{1,1} & x_{1,2} & \dots & x_{1,15} \\ \dots & & & \\ x_{n,1} & x_{n,2} & \dots & x_{n,15} \end{pmatrix} \rightarrow \uparrow \text{model flexibility}$$

How do you reduce the dimensionality of your model?

How to deal with high model dimensionality

So far we have covered:

1. Feature selection by comparing lower dimensional variants of your model.
 - Best subset selection
 - Forward/Backward stepwise selection
2. Apply a sparsity constraint to your model during fitting.
 - Ridge regression
 - LASSO
 - Elastic Net

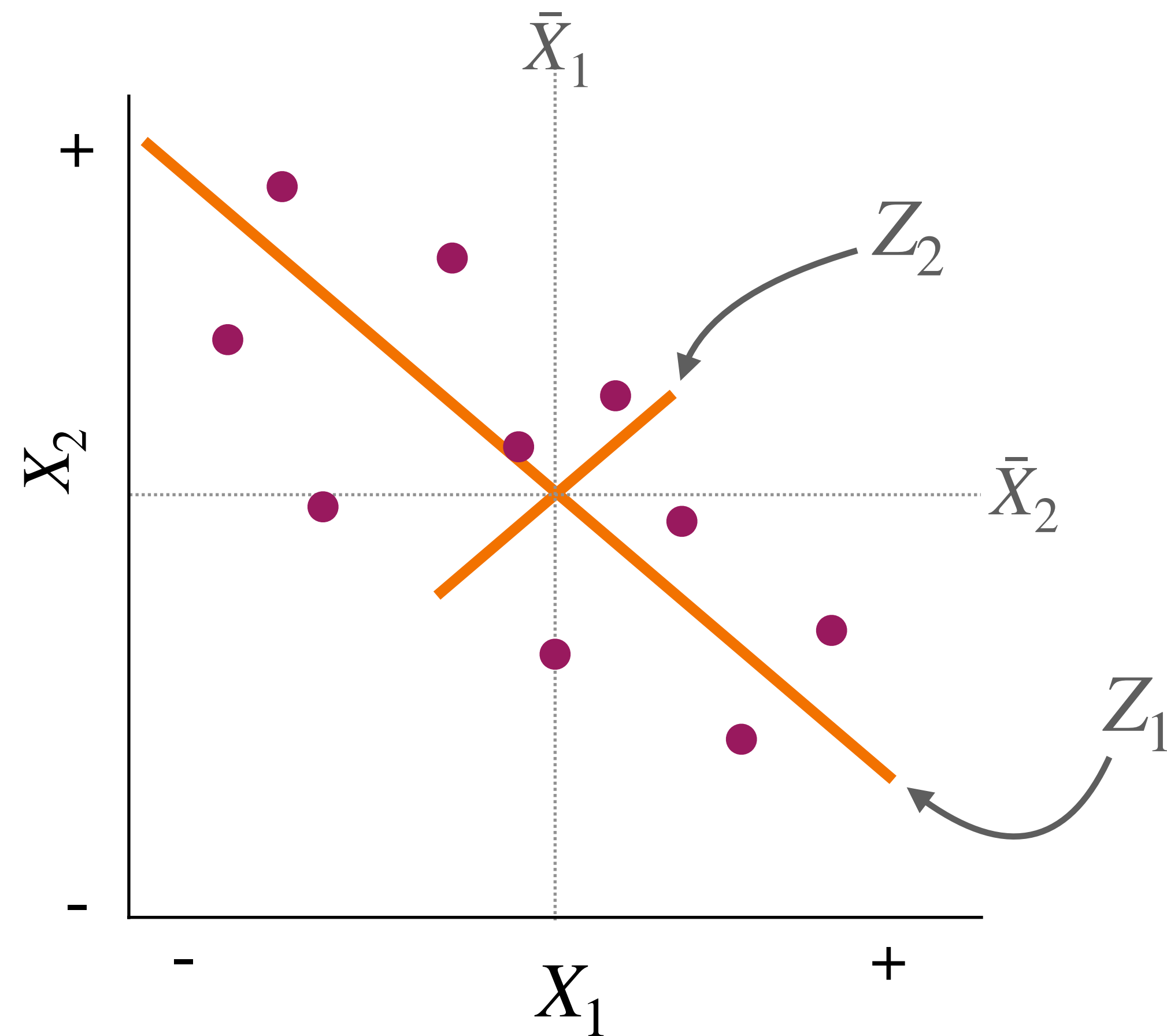
Principal component analysis (PCA)

What if you reduce the dimension of X itself?

PCA:

$$\underbrace{Z_m}_{\text{lower dimensional projection (component)}} = \sum_{j=1}^p \underbrace{\phi_{j,m}}_{\text{component loading}} \underbrace{X_j}_{\text{original data variable}}$$

Low dimensional components



The first principal component (Z_1) explains the most variance about the relationship between X_1 and X_2 .

PCA algorithm

Step 1: Find the first component (Z_1) loading.

$$\phi_1 = \arg \max \left(\frac{\phi_1' X' X \phi_1}{\phi_1' \phi_1} \right)$$

Step 2: Take the residuals after accounting for Z_1 .

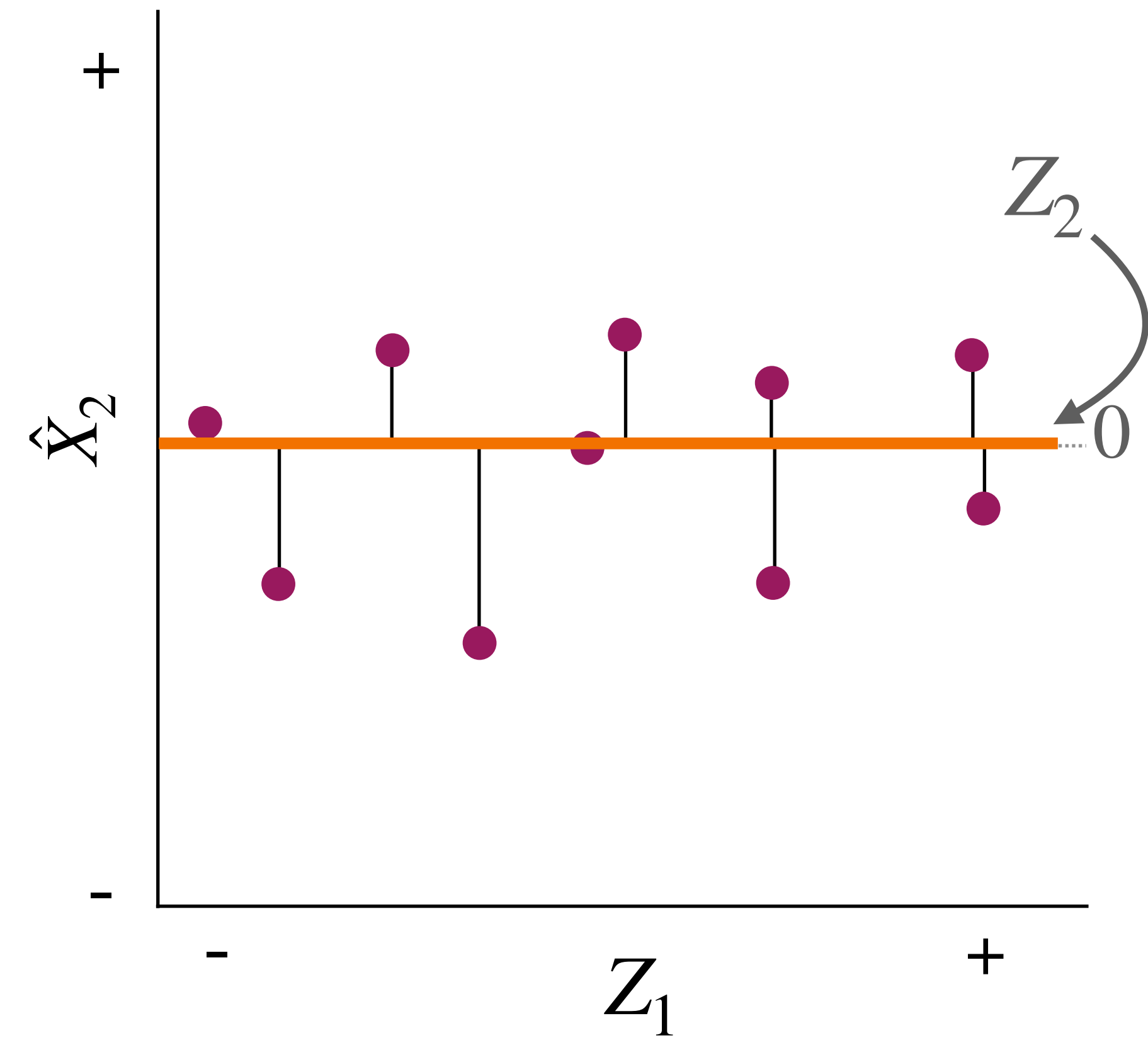
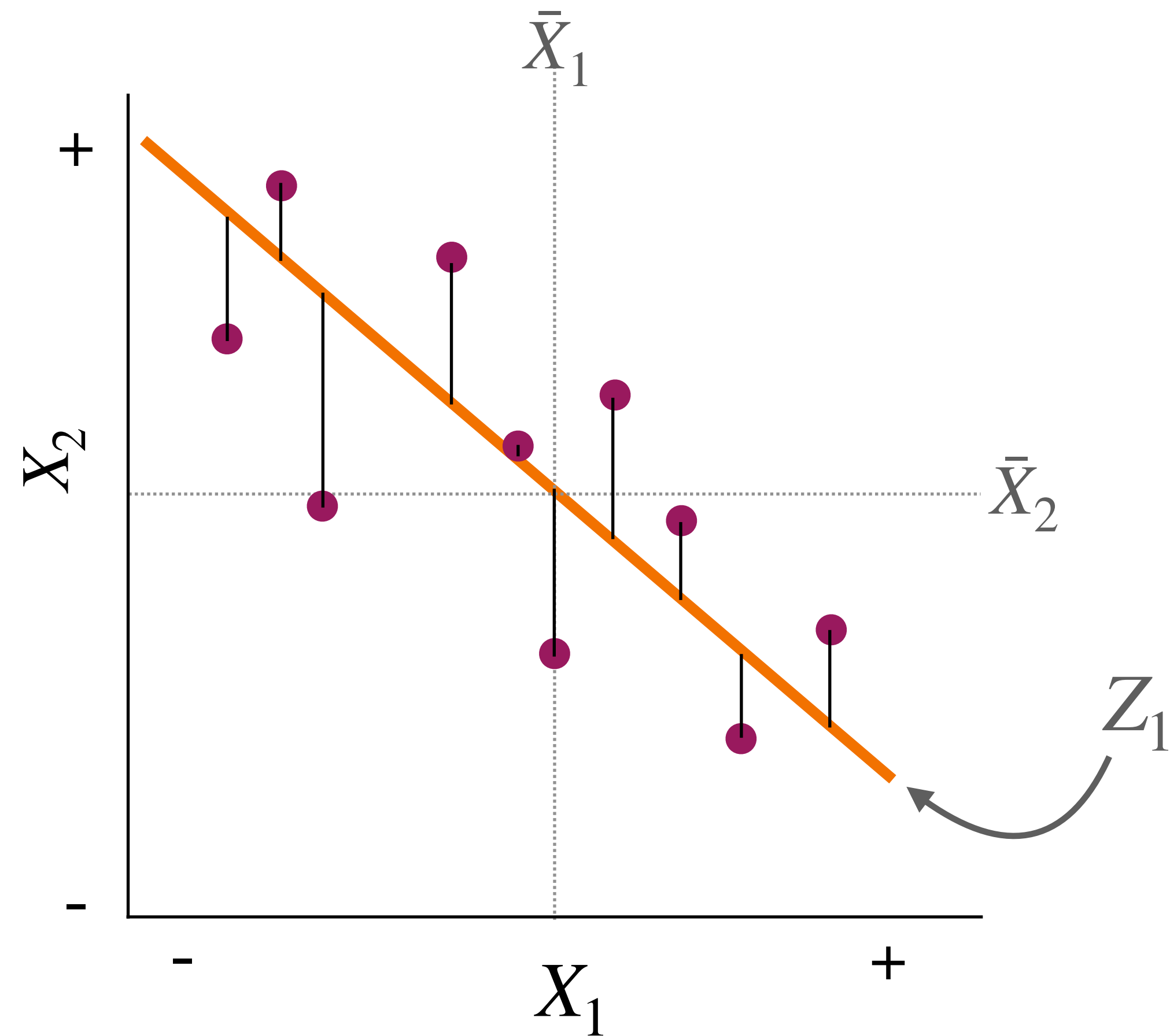
$$\hat{X}_m = X - \sum_{s=1}^{m-1} X \phi_s \phi_s'$$

Step 3: Calculate the next component (Z_m).

$$\phi_m = \arg \max \left(\frac{\phi_m' \hat{X}_m' \hat{X}_m \phi_m}{\phi_m' \phi_m} \right)$$

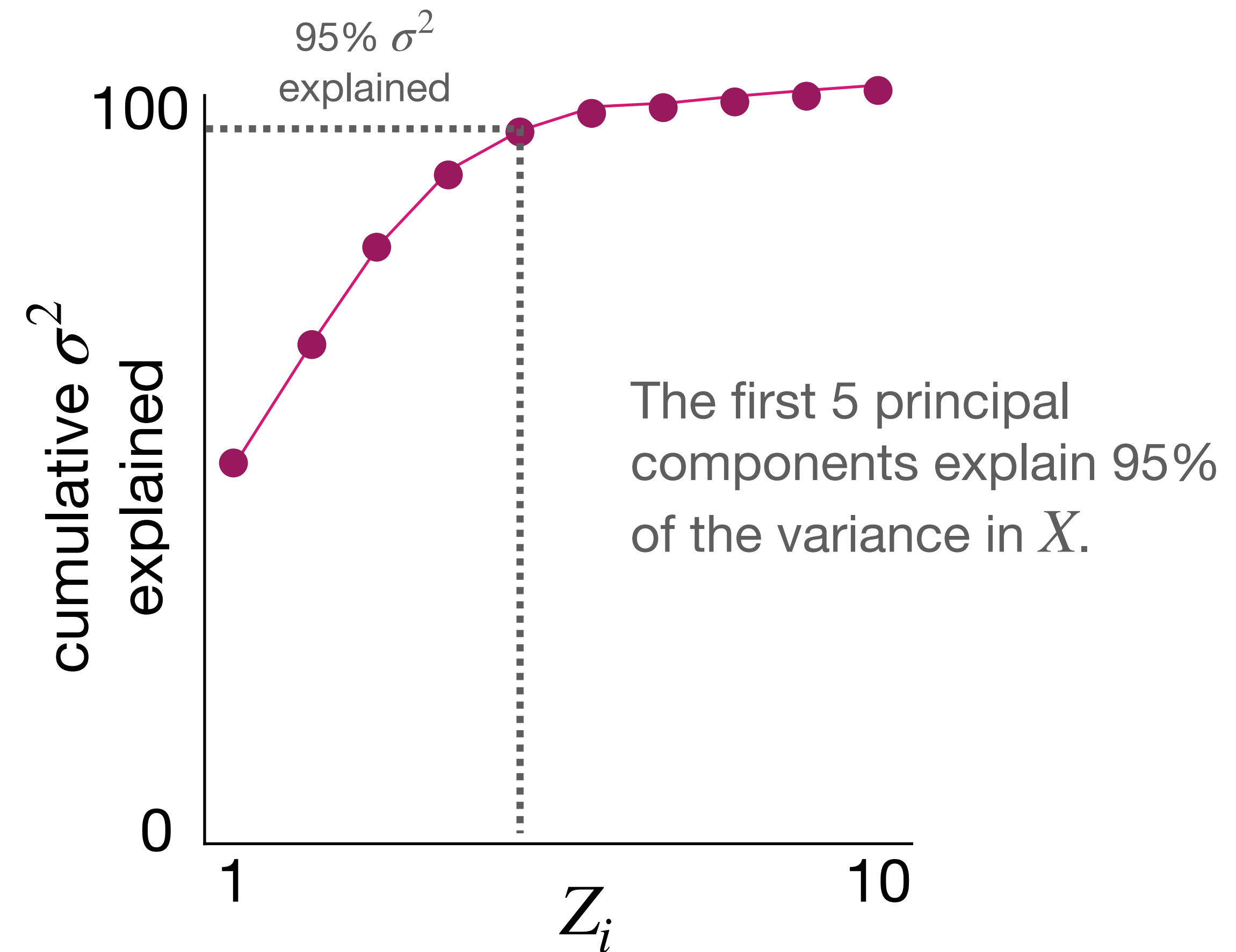
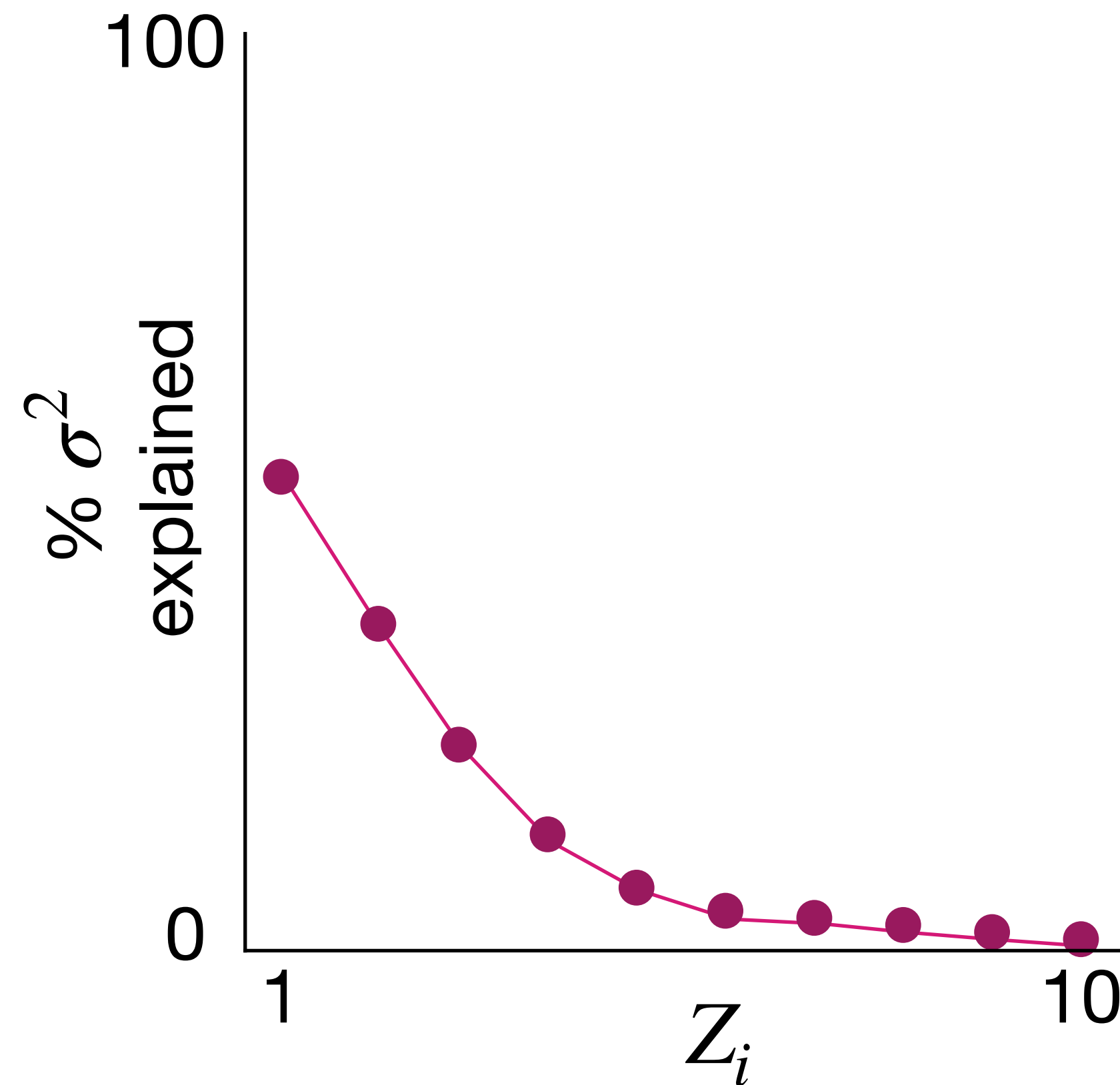
Step 4: Repeat Steps 2-3 until $m = p$

PCA algorithm



Example: 10 dimensional X

$$\begin{pmatrix} x_{1,1} & \dots & x_{1,p} \\ \vdots & & \\ x_{n,1} & \dots & x_{n,p} \end{pmatrix} \rightarrow n = 15, p = 10$$



PCA vs. Factor Analysis

PCA: $\underbrace{X}_{n \times p} = \underbrace{Z}_{n \times m} \underbrace{\phi^{-1}}_{m \times p}$

Factor Analysis (FA): $\underbrace{X^T}_{p \times n} = \underbrace{L}_{p \times m} \underbrace{F}_{m \times n}$

Factor loading
matrix with
 m specified up front.

- PCA is better for exploratory analysis.
- FA is better for hypothesis testing.
- PCA explains all variance in X .
- FA assumes lower dimensionality in X .

Principal component regression

Principal component regression (PCR)

Goal: Reduce the dimensions of X using PCA and use the strongest components in Z as your predictor variables.

PCA + linear regression:

$k < p$ # of components that explain $1 - \alpha$ percent variance in X

regression coefficients learned on Z_1, \dots, Z_k

principal component m

$$\hat{y}_i = \sum_{m=1}^k \hat{\theta}_m Z_{i,m}$$

PCR algorithm

Step 1: Calculate $Z = \phi X$.

Step 2: Identify the k components that explain $1 - \alpha$ (e.g., $(1 - 0.05) = 95\%$) of the variance in X .

Step 3: Fit your regression model with the k components identified in Step 2 (i.e., learn $\hat{Y} = \hat{f}(Z_{1\dots k})$)

Projecting back to X

Can determine the regression weights in X (i.e., $\hat{\beta}_j$) that best resolve the bias-variance tradeoff via the coefficients learned in Z (i.e., $\hat{\theta}_j$).

PCR to linear regression:

$$\begin{aligned}\hat{y}_i &= \sum_{m=1}^k \hat{\theta}_m Z_{i,m} \longrightarrow Z_m = \sum_{j=1}^p \phi_{j,m} X_j \\ &= \sum_{m=1}^k \hat{\theta}_m \sum_{j=1}^p \phi_{j,m} X_j \\ &= \sum_{j=1}^p \underbrace{\sum_{m=1}^k \hat{\theta}_m \phi_{j,m}}_{\hat{\beta}_j} X_j\end{aligned}$$

Best regression model:

$$\hat{\beta}_j = \sum_{m=1}^k \hat{\theta}_m \phi_{j,m}$$

Even when n is close to p .

Partial least squares

Partial least squares (PLS)

Goal: Find the lower dimensions in X that *maximize* $Cov[X, Y]$

of variables in X $\rightarrow p$

of variables in Z ($k = p$) $\rightarrow k$

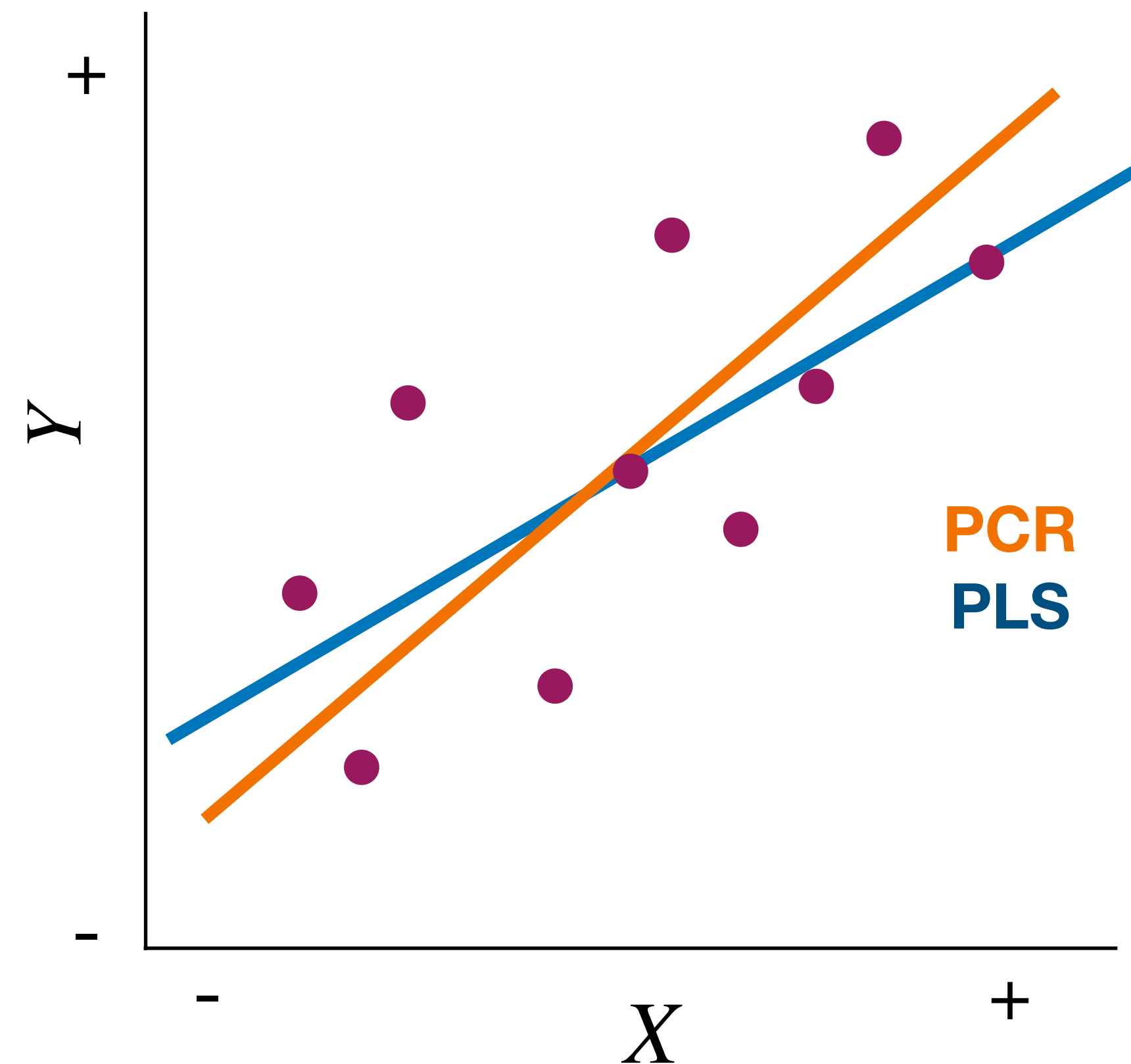
regression coefficients learned on Z_1, \dots, Z_k $\rightarrow \hat{\theta}_m$

principal component loading learned by minimizing $\sum_{m=1}^k (y - \sum_{m=1}^k \hat{\theta}_m Z_{i,m})^2$ $\rightarrow \hat{\phi}_{j,m}$

$$\hat{y}_i = \sum_{j=1}^p \sum_{m=1}^k \hat{\theta}_m \hat{\phi}_{j,m} x_{i,j}$$

$\hat{\phi}$ and $\hat{\theta}$ estimated in one step.

PLS vs. PCR



- PLS and PCR produce qualitatively different results depending on how the low dimensional components in X associate with Y .

Take home message

- Principal component methods offer an easy way of resolving the bias-variance tradeoff in high dimensionality (i.e., high variance) contexts by leveraging any correlational structure in your predictor variables.