4주차 복습 스터디

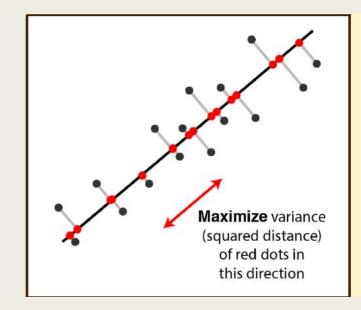
전형준

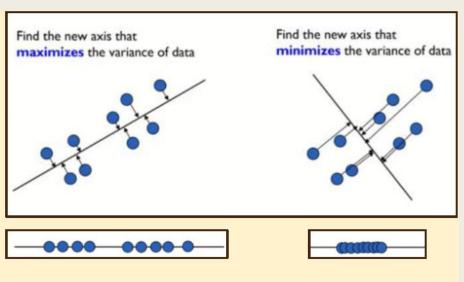
Dimension Reduction

- Feature selection : 기존 설명 변수들 중 소수의 예측 변수만을 <mark>선택</mark> ex) Lasso, ···
- Feature extraction : 기존 설명 변수들의 변환을 통해 새로운 예측 변수를 추출
- ex) PCA (Principal Component Analysis): unsupervised feature extraction
 PLS (Partial Least Squares): supervised feature extraction

PCA

- D개의 feature를 가진 data를 orthogonal한 q개의 변수로 구성된 데이터(PC)로 요약
- 가정 : 기존 변수들을 선형결합하여 새로운 변수 추출, 데이터들은 centered and scaled
- 원래 데이터의 분산을 최대한 보존하도록, Reconstruction error가 최소가 되도록





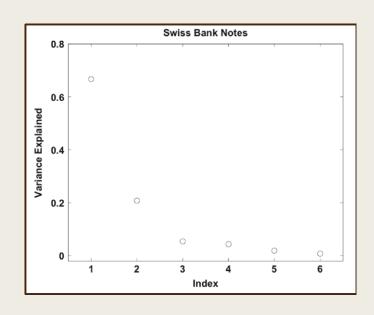
PCA

- How? \sum (Covariance martix)의 eigenvector가 가중치, eigenvalue가 분산값
- In sample : $\mathcal{Y} = (\mathcal{X} 1_n \bar{x}^T)\mathcal{G}$
- Proof : HW!

PCA

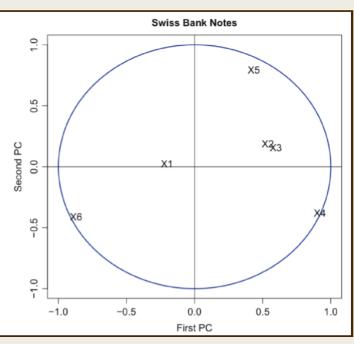
- How to select q?
- Method 1: 고유값 감소율이 유의미하게 낮아지는 Elbow point 선택
- Method 2: 일정 수준 이상의 분산 비를 보존하는 최소의 주성분 선택 (보통 70%)

$$\psi_q = \frac{\sum\limits_{j=1}^q \lambda_j}{\sum\limits_{j=1}^q \text{Var}(Y_j)} = \frac{\sum\limits_{j=1}^q \text{Var}(Y_j)}{\sum\limits_{j=1}^p \text{Var}(Y_j)}.$$



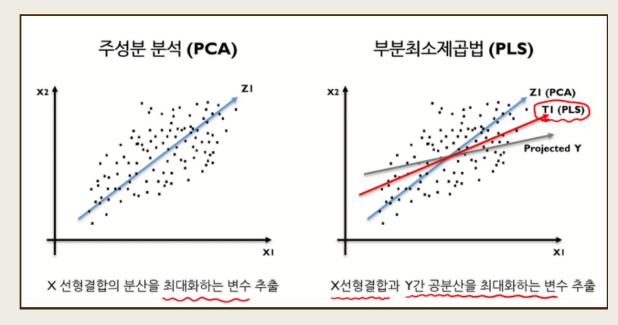
Interpretation of PC

■ In sample, $r_{X_iY_j} = g_{ij} \left(\frac{l_j}{s_{X_iX_i}}\right)^{1/2}$



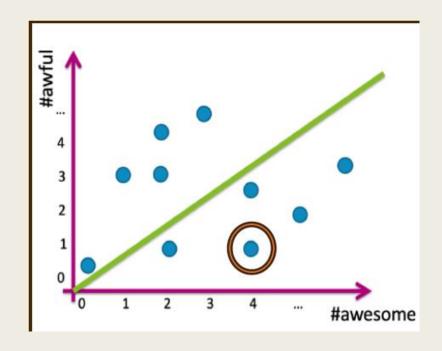
PLS

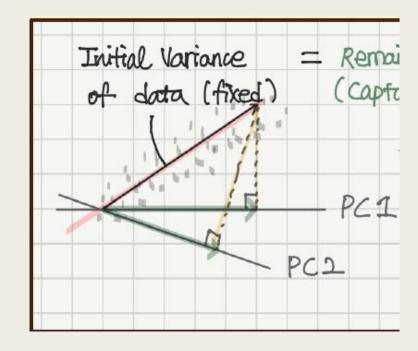
- PCA에서는 반영하지 못했던 Y와의 상관관계를 반영
- $t = Xw \Rightarrow Cov(t,Y) = Cov(Xw,Y) = E[(Xw E[Xw])(Y E[Y])] = E(Xw \cdot Y)$
- $E(Xw \cdot Y) = \frac{1}{n} \sum_{i=1}^{n} (Xw)_{i} \cdot Y_{i} = \frac{1}{n} (Xw)^{T} Y = \frac{1}{n} w^{T} (X^{T} Y)$



PCA in terms of Reconstruction error

- 결국 Remaining Variance를 최대화하는 앞의 방법과 근본적으로 같음



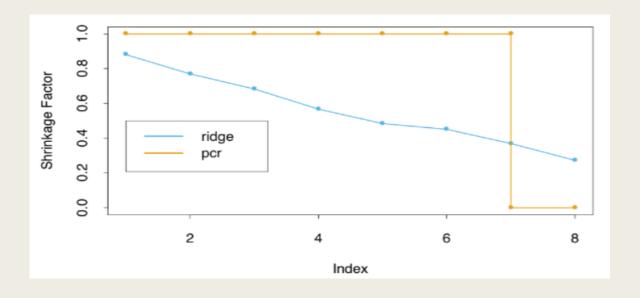


PCA in terms of Reconstruction error

- $f(\lambda) = \mu + V_q \lambda \Rightarrow \min_{\mu, \{\lambda_i\}, V_q} \sum_{i=1}^n ||x_i \mu V_q \lambda||^2$
- $\hat{\mu} = \bar{x}, \hat{\lambda} = V_q^T(x_i \bar{x}) \Rightarrow \min_{V_q} \sum_{i=1}^n \left\| (x_i \bar{x}) V_q V_q^T(x_i \bar{x}) \right\|^2$
- V_q : first q columns of V where $X = UDV^T$
- Columns of UD: principal components of X
- In terms of Unsupervised Learning : $L(v) = \frac{1}{n} \sum_{i=1}^{n} ||x_i decode(encode(x_i; V); V)||_2^2$

PCR

- $\hat{y}_{PCR} = \bar{y} + X\hat{\beta}_{PCR} \text{ where } \hat{\beta}_{PCR} = \sum_{j=1}^{q} \hat{\theta}_{j} v_{j}$
- $\widehat{Y}_{ridge} = \sum_{j=1}^{d} u_j \frac{d_j^2}{d_j^2 + \lambda} u_j^T Y$
- $\widehat{Y}_{PCR} = \sum_{j=1}^{q} u_j 1 u_j^T Y + \sum_{j=q+1}^{d} u_j 0 u_j^T Y$



When should / should not I use PCA?

- ✓ 변수를 줄이고는 싶은데, predictor들의 구조도 모르고 뭘 drop해야 할지도 잘 모르겠을 때..
- ✓ You want to ensure your variables are independent of one another.
- ➤ You are not comfortable making your independent variables less interpretable.

 (PCA 쓰려면 설명력은 포기해야...)
- ➤ PCA assumes there is a lower dimensional linear subspace that represents the data well.
 (Doesn't work well with non-linear manifold) → solutions include t-SNE, Isomap, etc.
 (PC들은 기존 변수들의 선형결합이기 때문에 비선형 data에서는 잘 작동하지 않을수도 있음.)

