Ridge Regression

OLS finds regression coefficients that minimize the SSE:

$$SSE = \sum_{i=1}^{n} (y_i - \beta_0 - \beta_1 x_{1i} - \beta_2 x_{2i} - etc.)^2$$

· Ridge regression finds coefficients that minimize:

$$SSE(R) = SSE + \text{shrinkage penalty} = SSE + \lambda (\beta_1^2 + \beta_2^2 + \beta_3^2 + etc.)$$

- This seems like a complicated idea but the concept is simple:
 - Ridge regression fits a line that minimizes SSE(R)
 - That is, Ridge minimizes SSE plus a penalty
 - ➤ We can vary the penalty A thus controlling the shrinkage
 - If we set λ = 0, Ridge minimizes SSE → same as OLS
 - If we set λ very large, then the resulting β's have to be very small → i.e., we shrink the coefficients
 - ➤ So if $\lambda = \infty$ Ridge yields the null model $y = β_0$
 - \succ The goal is to select the λ that minimizes the Test MSE

LASSO Regression



(Least Absolute Shrinkage and Selection Operator)

Again, OLS finds regression coefficients that minimize the SSE:

$$SSE = \sum_{i=1}^{n} (y_i - \beta_0 - \beta_1 x_{1i} - \beta_2 x_{2i} - etc.)^2$$

LASSO regression finds coefficients that minimize:

$$SSE(L) = SSE + \text{shrinkage penalty} = SSE + \lambda (|\beta_1| + |\beta_2| + |\beta_3| + etc.)$$

- That is, the penalty λ is applied over the sum of the absolute values of the coefficients, rather than over the sum of their squared values
- The effect is similar to Ridge regression:
 - If we set λ = 0, LASSO minimizes SSE → same as OLS
 - ➤ If we set $\lambda = \infty$, LASSO yields the null model $y = β_0$
 - Again, the goal is to select the λ that minimizes the Test MSE
- One important and interesting difference: mathematically, the Ridge **coefficients** can never be shrunk to **0** (except when $\lambda = \infty$), but some **LASSO coefficients** do become exactly $\mathbf{0}$ eventually as λ increases
 - → LASSO falls in between Subset Selection and Ridge.



Dimension Reduction Methods

- The basic idea is if we have P somewhat correlated predictors it is
 possible to transform these into M linear combinations, such that P >
 M, thus reducing the number of variables in a model.
- Dimension reduction = reduce the estimation of P+1 coefficients (β₀, β₁, β₂,... β_P) to estimating M+1 coefficients (α₀, α₁, α₂,... α_M)
- Example: if we suspect that a vehicle's volume, horsepower, and weight affect the vehicle's gas mileage, but these 3 variables are highly correlated, we could combine them into a new variable called something like "size" composed of some percentage of volume, plus some of horsepower, plus some of weight, reducing the model variables from 3 to 1.
- Naturally, we also lose some interpretability, so it is a tradeoff
- Two popular dimension reduction methods are Principal
 Components Analysis (PCA) and Partial Least Squares (PLS), both
 of which use the correlation matrix of P predictors to find M (<P)
 linear combinations of the P predictors
- These methods are may increase bias but substantially reduce variance of the coefficients, particularly when P is large relative to N

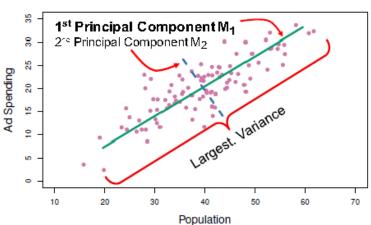


Principal Components: Illustration

This can be **illustrated** with **2 variables** like population size and advertising expenditures (as predictors of sales), resulting in **2 Principal Components**



Think of Principal component as rotating the axes into the highest variance direction and then moving the origin to the mean (i.e., centering) of the variables involved.



It is clear from this plot that Ad Spending and Population are highly correlated. But Principal Components M_1 and M_2 are not



Variables, Components & Scores

- For a set of **P** variables $\rightarrow X_b X_b \dots X_p$
- There are P "orthogonal" PC's:

$$M_1 = l_{11}X_1 + l_{12}X_2 + \dots + l_{1P}X_P$$

 $M_2 = l_{21}X_1 + l_{22}X_2 + \dots + l_{2P}X_P$
.....
 $M_P = l_{P1}X_1 + l_{P2}X_2 + \dots + l_{PP}X_P$

• Each with a sum of squared factor loadings = 1:

$$l_{11}^{2} + l_{12}^{2} + \dots + l_{1p}^{2} = 1$$

$$l_{21}^{2} + l_{22}^{2} + \dots + l_{2p}^{2} = 1$$

$$\dots$$

$$l_{P1}^{2} + l_{P2}^{2} + \dots + l_{pp}^{2} = 1$$

For a data point X_{i,D} X_{i,D} X_{ip} the corresponding value m_{i,I} is called the PC "score" for that data point



Principal Components Regression (PCR)

- The main purpose of PCR is to reduce the dimensionality (i.e., number of variables) of a model without removing variables.
- Since the PC's are sorted from highest to lowest variance, it is very likely that the first few PC's are sufficient to represent the variance in the data.
- So, for a given OLS regression:

$$Y = \beta_{\theta} + \beta_{1}(X_{1}) + \beta_{2}(X_{2}) + \dots + \beta_{p}(X_{p}) + \varepsilon$$

We can construct another OLS model on M PC's:

$$Y = \alpha_{\theta} + \alpha_{1}(M_{1}) + \alpha_{2}(M_{2}) + \dots + \alpha_{M}(M_{M}) + \varepsilon$$

- M can be anywhere between 1 and P (all PC's → same as OLS)
- Since each PC is a linear transformation of all variables using factor loadings, all variables are represented
- The key is to find the optimal number of M PC's (i.e., a "tuning" parameter) to include in the model
- As with regular OLS, as M ↑ → Bias ↓ and Variance ↑







Partial Least Squares (PLS): Intuition

- With PCR, the X_p X₂, X_p variables are transformed into M_p M₂
 M_p components in an "unsupervised" way
- That is, the independent variable dimensions are rotated to find directions in which the data exhibits highest variance.
- This is an "unsupervised" method because the outcome variable
 Y is not taken into account when doing PCA
- While PCR does well in general, there is no guarantee the first few M components will be the best directions to predict Y
- In contrast, PLS is a "supervised" method
- Like PCR, PLS is a dimension reduction method, but unlike PCR, PLS does further rotation of the dimensions to maximize the correlation with Y
- In sum, PLS attempts to find directions that not only explain the predictors, but also the outcome variable
- It does this by placing stronger weight on variables that are more strongly correlated with Y