STAT 3690 Lecture 32

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Misclassification/error rate

- Population: $Pr(Y \neq h(\mathbf{X}))$
 - $-h(\cdot)$: the classifier to be evaluated
- Apparent estimation
 - Implementation
 - 1. Fit a classifier according to training data
 - 2. Apply the fitted classifier to training data as well
 - 3. Estimate the error rate by the misclassification proportion
 - Comments
 - * Training and testing with identical data points
 - * Severe underestimation likely
- Parametric estimation
 - Implementation
 - 1. Express $Pr(Y \neq h(\mathbf{X}))$ in terms of unknown parameters
 - 2. Plug in estimates of unknown parameters
 - Comment
 - * Able to derive the analytical form of $Pr(Y \neq h(\mathbf{X}))$ in rare cases
 - * Underestimation likely

```
For K=2 and X/Y=k~MVN (Ak. I).
                       the error roote of LDA classifier hIX)
                   = \mathcal{P}_r \left( Y \neq h(X) \right)
                     = Pr (Y=1, h(x)=2) + Pr (Y=2, h(x)=1)
                   = P_r(Y=1, \delta_1(x) < \delta_2(x)) + P_r(Y=1, \delta_1(x) > \delta_2(x))
                   = T.Pr (3,(x) < 3,(x) | Y=1) + T. Pr (3,(x) > 3,(x) | Y=1)
let U = \hat{\beta}_1(N) - \hat{\beta}_2(X) = X^T \Sigma^{-1}(R_1 - R_2) - \frac{1}{2} \mathcal{A}_1^T \Sigma^{-1} \mathcal{U}_1 + \frac{1}{2} \mathcal{A}_1^T \Sigma^{-1} \mathcal{U}_1 + \ln(\pi_1/\pi_1), then
      E\left(U\mid Y=1\right)=\mathcal{M}_{i}^{T}\Sigma^{T}\left(\mathcal{M}_{i}-\mathcal{M}_{i}\right)-\frac{1}{2}\mathcal{M}_{i}^{T}\Sigma^{T}\mathcal{M}_{i}+\frac{1}{2}\mathcal{M}_{i}^{T}\Sigma^{T}\mathcal{M}_{i}+\ln 1\pi ./\pi _{z})
                                   =\frac{1}{2}\mathcal{M}_1^T\Sigma^{-1}\mathcal{M}_1-\mathcal{M}_1^T\Sigma^{-1}\mathcal{M}_1+\frac{1}{2}\mathcal{M}_2^T\Sigma^{-1}\mathcal{M}_2+\ln(\pi_1/\pi_2)
                                    = 1 (M1-M2) T 5-1 (M1-M2) + h (71/72)
       E(U|Y=2) = - = (M,-M) = - (M,-M2) + (m(T/T))
      vor (U | Y=1) = (M, -M,) TI-1 = > (M,-M)
       vw(U|Y=2) = (M, -M_L)^T \Sigma^{-1} (M, -M_L)
               \frac{\bigcup_{i} - \frac{1}{2} \lfloor \mathcal{M}_{i} \rfloor^{n} \sum_{i} \frac{1}{2} \lfloor \mathcal{M}_{i} - \mathcal{M}_{i} \rfloor - \ln \left( \pi_{i} / \pi_{i} \right)}{\sqrt{\left[ \mathcal{M}_{i} - \mathcal{M}_{i} \right]^{T} \sum_{i} \frac{1}{2} \left[ \mathcal{M}_{i} - \mathcal{M}_{i} \right]}} \mid Y = I \quad \sim \mathcal{N}(0, I)
               \frac{1}{\sqrt{\left[\left(\mu_{1}-\mu_{k}\right)^{T}\Sigma^{-1}\left(\mu_{1}-\mu_{k}\right)-\left[\mu_{1}+\pi_{k}/\pi_{k}\right)}\right]}} \Big| Y_{z} \chi \sim N(0,1)
    So, Pr(Y = h(x))
          \begin{split} &=\pi,\;\mathcal{P}_{r}\left(\bigcup \circ (Y=1)+\pi_{k}\mathcal{P}_{r}\left(\bigcup \circ (Y=2)\right)\right.\\ &=\pi_{1}\;\;\overline{\Phi}\left(\frac{-\frac{1}{4}\left[\mu_{1}-\mu_{k}\right]^{T}\boldsymbol{\Sigma}^{-1}\left[\mu_{1}-\mu_{k}\right]-\left[\mu_{k}\mathcal{T}/\pi_{k}\right]}{\sqrt{\left[\mu_{1}-\mu_{k}\right]^{T}\boldsymbol{\Sigma}^{-1}\left[\mu_{1}-\mu_{k}\right]}}\right) \end{split}
             + \pi_{3} \overline{\Phi} \left( \frac{-\frac{1}{2} (M-M_{3})^{T} \Sigma^{-1} (M-M_{3}) + (n(\pi_{1}/\pi_{1}))}{\sqrt{(M-M_{3})^{T} \Sigma^{-1} (M-M_{3})}} \right)
where \overline{\Phi}(\cdot) is the studentd normal colf
```

- Estimation via M-fold cross validation (CV)
 - Implementation
 - 1. The dataset is randomly partitioned into M chunks.
 - 2. Train one classifier upon each combination of M-1 chunks.
 - Apply each classifier to the corresponding remaining chunk and compute the empirical error rate.
 - 4. Estimate the population error rate by averaging these M empirical error rates.
 - Comment
 - * Leave-one-out $CV \Leftrightarrow n$ -fold CV
- Estimation via $M \times L$ -fold CV
 - Implementation
 - 1. Repeat the four steps of M-fold CV L times.
 - 2. Average all the ML resulting empirical error rates.
 - Comment
 - * $M \times 1$ -fold CV $\Leftrightarrow M$ -fold CV

```
options(digits = 4)
set.seed(3690)
L = 1; M = nrow(iris) # Leave-one-out CV
\# L = 1; M = 10 \# 10 \text{ fold } CV
\# L = 10; M = 10 \# 10by10 fold CV
# initiation
errLda1 = matrix(0, nrow = L, ncol = M)
errLda2 = matrix(0, nrow = L, ncol = M)
errQda1 = matrix(0, nrow = L, ncol = M)
errQda2 = matrix(0, nrow = L, ncol = M)
for (1 in 1:L){
  idx_new = sample(1:nrow(iris), size = nrow(iris))
  folds = cut(1:nrow(iris), breaks = M, labels=FALSE)
  # follow formulas
  for (m in 1:M){
    # Segement your data by fold using the which() function
    picked = idx_new[which(folds == m, arr.ind=TRUE)]
    train = iris[-picked,]
    Xtrain = train[, !(names(train) %in% c("Species"))]
    Ytrain = train$Species
    test = iris[picked,]
    Xtest = test[, !(names(test) %in% c("Species"))]
    Ytest = test[, names(test) %in% c("Species")]
    labels = unique(iris$Species)
    K = length(labels)
    p = ncol(Xtrain)
    n = nrow(Xtrain)
    nks = numeric(K)
    piks = numeric(K)
    Muks = matrix(0, nrow = K, ncol = p)
    Sigmaks = list()
    for (k in 1:K){
```

```
Xtrain_k = Xtrain[Ytrain == labels[k],]
    nks[k] = nrow(Xtrain_k)
    piks[k] = nks[k]/n
    Muks[k,] = colMeans(Xtrain_k)
    Sigmaks[[k]] = cov(Xtrain_k)
    if (k==1){
     SigmaPool = Sigmaks[[k]] * (nks[k]-1)
     SigmaPool = SigmaPool + Sigmaks[[k]] * (nks[k]-1)
 }
 SigmaPool = SigmaPool/(n-1)
 SigmaPoolInv = solve(SigmaPool)
 deltaksLda = matrix(0, nrow = nrow(Xtest), ncol = K)
 deltaksQda = matrix(0, nrow = nrow(Xtest), ncol = K)
 for (k in 1:K){
    # LDA
    deltaksLda[,k] = as.matrix(Xtest) %*% SigmaPoolInv %*% Muks[k,] -
      .5* as.vector(t(Muks[k,]) %*% SigmaPoolInv %*% Muks[k,]) +
     log(piks[k])
    # QDA
   SigmakInv = solve(Sigmaks[[k]])
    deltaksQda[,k] = -diag(as.matrix(Xtest) %*% SigmakInv %*% t(as.matrix(Xtest))) +
     2* as.matrix(Xtest) %*% SigmakInv %*% Muks[k,] -
     as.vector(t(Muks[k,]) %*% SigmakInv %*% Muks[k,]) +
      2* log(piks[k]) -
     log(det(Sigmaks[[k]]))
 }
 resLda = apply(deltaksLda, 1, FUN = function(x){labels[which.max(x)]})
 resQda = apply(deltaksQda, 1, FUN = function(x){labels[which.max(x)]})
 errLda1[1, m] = mean(Ytest != resLda)
  errQda1[1, m] = mean(Ytest != resQda)
}
# use MASS
for (m in 1:M){
  # Segment your data using the which() function
 picked = idx_new[which(folds == m, arr.ind=TRUE)]
 train = iris[-picked,]
 Xtrain = train[, !(names(train) %in% c("Species"))]
 Ytrain = train$Species
 test = iris[picked,]
 Xtest = test[, !(names(test) %in% c("Species"))]
 Ytest = test[, names(test) %in% c("Species")]
 for (k in 1:K){
    # LDA
    objLda = MASS::lda(Xtrain, Ytrain, method = "moment")
   objQda = MASS::qda(Xtrain, Ytrain, method = "moment")
```

```
errLda2[1, m] = mean(Ytest != predict(objLda, Xtest)$class)
  errQda2[1, m] = mean(Ytest != predict(objQda, Xtest)$class)
}
mean(errLda1)
mean(errLda2)
mean(errQda1)
mean(errQda2)
```