

Stat 8003: Homework 5

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Solution to Problem 1

We have that $x_i \sim f(x_i) = \pi_0 f_0(x_i) + \pi_1 f_1(x_i)$ where $f_0(x_i) = \mathbb{1}(0 \leq x_i \leq 1)$ with a uniform density and $f_1(x_i) = \beta(1 - x_i)^{\beta-1} = \text{Beta}(1, \beta)$. The complete data is represented by $y_i = (x_i, z_i)$ and $\theta = (\pi_0, \beta)$, the vector of parameters.

- (a) Derive the completely likelihood function $L_n(\theta|x_i, z_i)$

$$L_n(\theta|x_i, z_i) = \prod_{i=1}^n \sum_{j=0}^1 \pi_j f_j(x_i) \mathbb{1}(z_i = j)$$

the completely log-likelihood function can be represented by,

$$l_n(\theta|x_i, z_i) = \sum_{i=1}^n \sum_{j=0}^1 \mathbb{1}(z_i = j) \log(\pi_j f_j(x_i))$$

- (b) Using the EM Algorithm to derive the estimators for π_0 and β

Expectation Step:

$$Q(\theta, \theta^t) = E[l_n(\theta|x_i, z_i)] = E\left[\sum_{i=1}^n \sum_{j=0}^1 \mathbb{1}(z_i = j) \log(\pi_j f_j(x_i))\right] = \sum_{i=1}^n \sum_{j=0}^1 P(z_i = j|x_i, \theta) \log(\pi_j f_j(x_i))$$

$$\text{where: } P(z_i = j|x_i, \theta) = \frac{P(x_i|z_i = j)P(z_i = j)}{\sum_{j=0}^1 P(x_i|z_i = j)P(z_i = j)} = \frac{\pi_j f_j(x_i)}{\pi_0 f_0(x_i) + \pi_1 f_1(x_i)} = T_{ji}^t$$

therefore:

$$Q(\theta, \theta^t) = \sum_{i=1}^n \sum_{j=0}^1 T_{ji}^t \log(\pi_j f_j(x_i))$$

Maximization Step:

(Maximize π_0):

if $j = 0$: $\sum_{i=1}^n T_{0i}^t \log(\pi_0 f_0(x_i))$

if $j = 1$: $\sum_{i=1}^n T_{1i}^t \log((1 - \pi_0) f_1(x_i))$

therefore:

$$\Delta = \sum_{i=1}^n T_{0i}^t \log(\pi_0 f_0(x_i)) + \sum_{i=1}^n T_{1i}^t \log((1 - \pi_0) f_1(x_i))$$

$$\frac{\partial \Delta}{\partial \pi_0} \left(\sum_{i=1}^n T_{0i}^t \log(\pi_0 f_0(x_i)) + \sum_{i=1}^n T_{1i}^t \log((1 - \pi_0) f_1(x_i)) \right) = 0$$

$$= \left(\sum_{i=1}^n T_{0i}^t \right) \frac{f_0(x_i)}{\pi_0 f_0(x_i)} + \left(\sum_{i=1}^n T_{1i}^t \right) \frac{-f_1(x_i)}{f_1(x_i) \pi_0 f_1(x_i)} = 0$$

$$\left(\sum_{i=1}^n T_{0i}^t \right) \frac{f_0(x_i)}{\pi_0 f_0(x_i)} = \left(\sum_{i=1}^n T_{1i}^t \right) \frac{f_1(x_i)}{(1 - \pi_0) f_1(x_i)}$$

$$\left(\sum_{i=1}^n T_{0i}^t \right) \frac{1}{\pi_0} = \left(\sum_{i=1}^n T_{1i}^t \right) \frac{1}{1 - \pi_0}$$

Thus,

$$\pi_0^{t+1} = \frac{1}{n} \sum_{i=1}^n T_{0i}^t$$

(Maximize β):

Given:

$$Q(\theta, \theta^t) = \sum_{i=1}^n T_{0i}^t \log(\pi_0 f_0(x_i)) + \sum_{i=1}^n T_{1i}^t \log(\pi_1 \beta (1 - x_i)^{\beta-1})$$

Let

$$\Delta = \sum_{i=1}^n T_{1i}^t (\log(\pi_1) + \log(\beta) + (\beta - 1) \log(1 - x_i))$$

$$\frac{\partial \Delta}{\partial \beta} = \sum_{i=1}^n T_{1i}^t \left(\frac{1}{\beta} + \log(1 - x_i) \right) = 0$$

$$\sum_{i=1}^n T_{1i}^t \frac{1}{\beta} = - \sum_{i=1}^n T_{1i}^t \log(1 - x_i)$$

$$\boxed{\beta^{t+1} = - \frac{\sum_{i=1}^n T_{1i}^t}{\sum_{i=1}^n T_{1i}^t \log(1 - x_i)}}$$

(c) Our R-program to estimate π_0^{t+1} and β^{t+1} using EM algorithm has been attached to this homework assignment. Our computed estimates are:

$$\pi_0^{t+1} = 0.696794$$

$$\beta^{t+1} = 11.093275$$

Using these estimates, we fit our beta-uniform mixture model to the pvalues dataset:

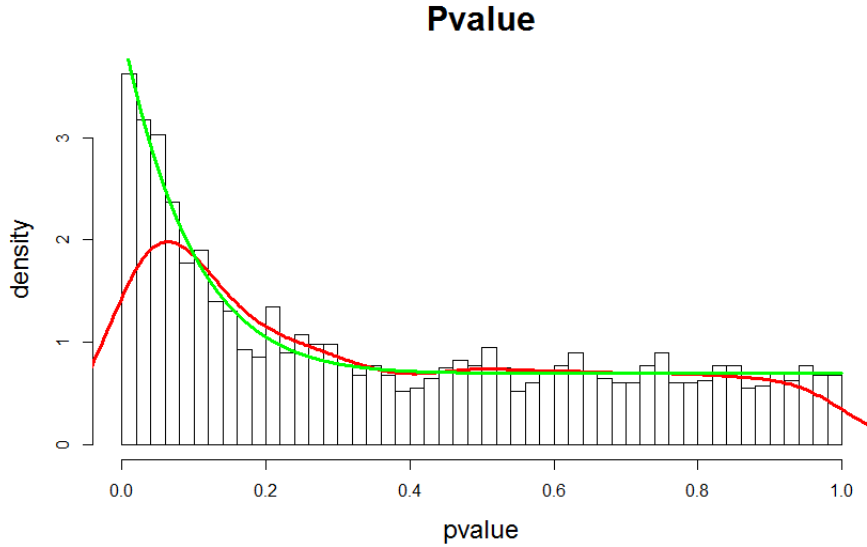


Figure 1: Beta-uniform mixture fit to pvalues; generated in R. Note: the green curve represents the fitted density

Given:

$$fdr_i(x_i) = P(Z_i = 0|x_i) = \frac{\pi_0 f_0(x_i)}{\pi_0 f_0(x_i) + \pi_1 f_1(x_i)} = \frac{\pi_0 f_0(x_i)}{\pi_0 f_0(x_i) + (1 - \pi_0) \beta (1 - x_i)^{\beta-1}}$$

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we can substitute our computed estimates, π_0^{t+1} and β^{t+1} for π_0 and β and compute the local fdr score for each x_i using the following R-script:

```
####X is a vector of pvalues
fdrlocal<-(pi0*dunif(X,0,1)) / (pi0*dunif(X,0,1)+pi1*dbeta(X,1,beta))
```

- (d) Our R-script attached to this homework yields **321** (117 + 204) falsely classified data points when $fdr_i(x_i) > 0.5$

	<i>group_{EM}</i>	
	0	1
<i>group</i>		
0	1182	204
1	117	497

Solution to Problem 2

Given the local fdr score as:

$$fdr_i = \frac{\pi_0 f_0(x_i)}{f(x_i)}$$

where $f(x_i)$ is the marginal density of x_i and assuming $\pi_0 = 0.7$

- (a) Given: $\hat{f}_h(X) = \frac{1}{nh} \sum_{i=1}^n k\left(\frac{x-x_i}{h}\right)$ and $k(u) = \frac{1}{\sqrt{2\pi}} e^{-\frac{1}{2}u^2}$:

results in:

$$\hat{f}_h(X) = \frac{1}{n} \sum_{i=1}^n \frac{1}{h} \phi\left(\frac{x-x_i}{h}\right)$$

using Silverman's h:

$$h = 1.06\hat{\sigma}n^{-\frac{1}{5}}$$

Below is a density plot generated in R using the Gaussian-Kernel method with Silverman's h to estimate $f(x_i)$ from the pvalue dataset:

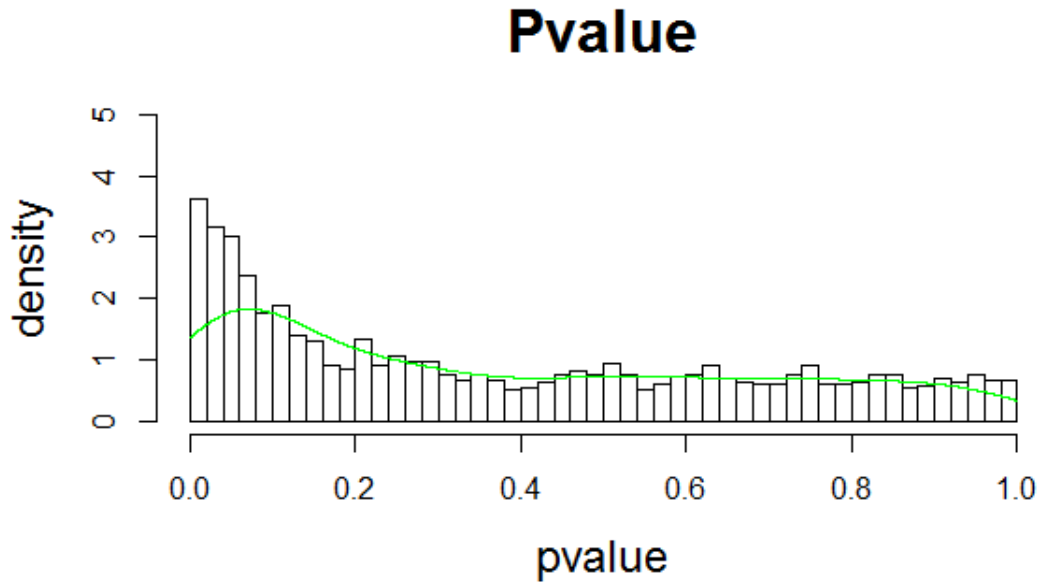


Figure 2: Gaussian-Kernel with Silverman's h , fit to pvalues generated in R. Note: the green curve represents the fitted density

- (b) We can estimate the local fdr score using the Gaussian-Kernel with Silverman's h method with the following R-script:

```
#estimate local fdr score using Gaussian-Kernel method
#X is a vector of pvalues
n <- length(X)
h <- 1.06 * sqrt( var(X) ) / (n^(1/5)) # Silverman's h
fnorm.hat.h <- X
for( i in 1:length( X ) )
{
  fnorm.hat.h[i] <- mean(dnorm( X[i]-X, 0, h ))
}
```

```
fdr_local_kernel <- (0.7*dunif(pvalue$X,0,1)) / fnorm.hat.h #yields a
2000x1 vector of local fdr scores
```

```
plot(X,fnorm.hat.h) #plots pvalue against Gaussian-Kernel density estimate
```

- (c) Our R-script attached to this homework yields **335** ($117 + 218$) falsely classified data points when $fdr_i(x_i) > 0.5$

	<i>group</i> _{Kernel}	
	0	1
<i>group</i>		
0	1168	218
1	117	497

(d) Using maximum likelihood cross-validation method:

$$\hat{f}_h(X) = \frac{1}{nh} \sum_{i=1}^n k\left(\frac{x - x_i}{h}\right)$$

$$\hat{f}_h(x_j) = \frac{1}{nh} \sum_{i=1}^n k\left(\frac{x_j - x_i}{h}\right)$$

Likelihood function:

$$\prod_{j=1}^n \hat{f}_h(x_j) = \frac{1}{nh} \prod_{j=1}^n \sum_{i=1}^n k\left(\frac{x_j - x_i}{h}\right)$$

using the "leave-one-out" method:

$$\hat{f}_{h,i}(x_j) = \frac{1}{(n-1)h} \sum_{i \neq j} k\left(\frac{x_j - x_i}{h}\right)$$

therefore:

$$MLCV = \frac{1}{n} \sum_i \log\left(\sum_{i \neq j} k\left(\frac{x_j - x_i}{h}\right) \frac{1}{(n-1)h}\right)$$

The following R-script using the "kedd" package will generate a density curve using the MLCV method:

#X is a vector of pvalues

library(kedd)

h.cv <- **h.mlcv**(X)\$h

xaxis_new <- **seq**(**min**(X), **max**(X), 0.00001)

fnorm.cv.hat <- **xaxis_new**

for(i in 1:**length**(**xaxis_new**))

{

fnorm.cv.hat[i] <- **mean**(**dnorm**(**xaxis_new**[i]-X, 0, **h.cv**))

}

#plot density

hist(X, freq=F, br=40, main="Pvalue", xlab="pvalue", ylab="density", cex.n

points(**xaxis_new**, **fnorm.cv.hat**, main=**paste**("h=",**h.cv**, sep=""), xlab="pvalue

Figure 3 depicts densities of 3 methods fit to pvalues: 1) EM-algorithm 2) Gaussian-Kernel with Silverman's h 3) Maximum likelihood cross-validation

Our R-script attached to this homework yields **501** (334 + 167) falsely classified data points when compared to the original group classification, and when $fdr_i(x_i) > 0.5$

	<i>group</i> _{MLCV}	
	0	1
<i>group</i>		
0	1219	167
1	334	280

(e) The EM-algorithm worked the best in terms of having the lowest classification error.

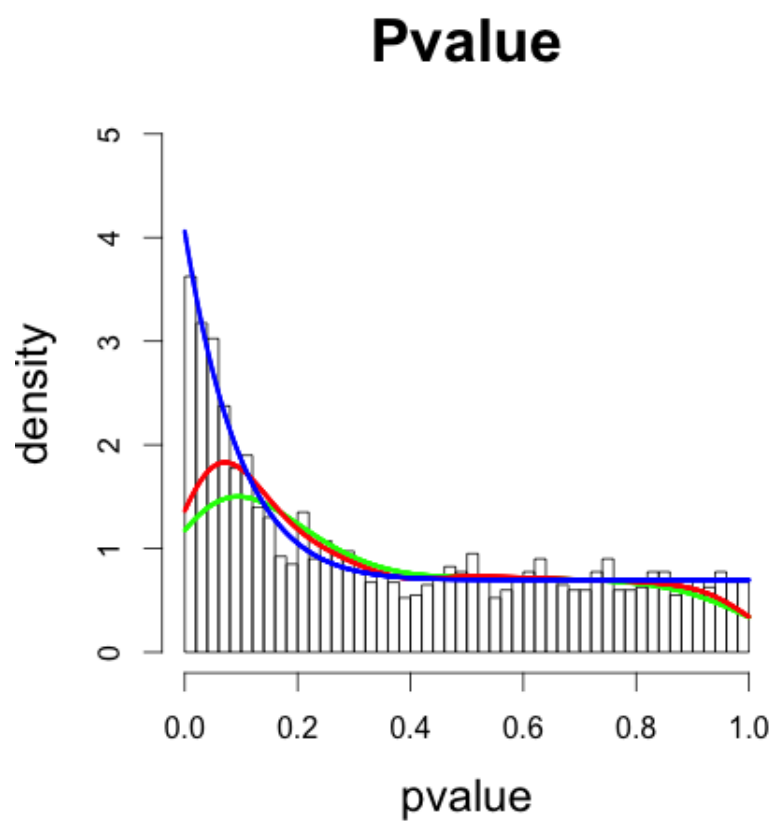


Figure 3: Note: Blue = EM-algorithm, Red= Gaussian-Kernel w Silverman's h , Green=MLCV