HW6 Solution

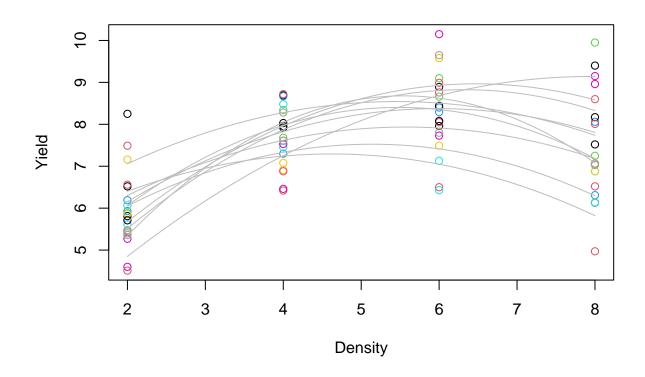
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2023-04-25

Question 1

```
a)
```

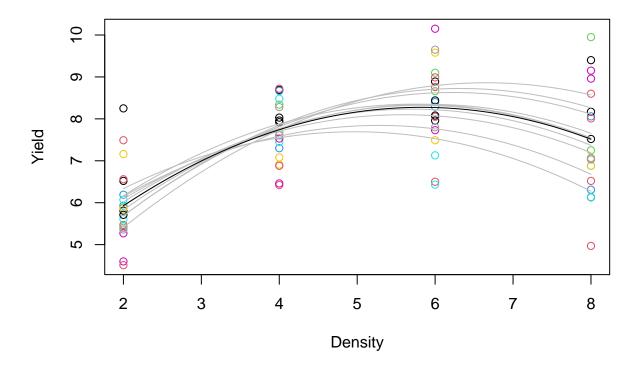
```
data_p1 = read.table("http://www2.stat.duke.edu/~pdh10/FCBS/Exercises/pdensity.dat", header = TRUE)
ids<-sort(unique(data_p1$plot))</pre>
m<-length(ids)</pre>
Y<-list(); X<-list(); N<-NULL
for(j in 1:m)
  Y[[j]]<-data_p1[data_p1$plot==ids[j], 3]
  N[j]<- sum(data_p1$plot==ids[j])</pre>
  xj<-data_p1[data_p1$plot==ids[j], 2]</pre>
 X[[j]] \leftarrow cbind(rep(1,N[j]), xj, xj^2)
GRID = data.frame(cbind(1, seq(2,8,0.1), seq(2,8,0.1)^2))
x_{grid} = seq(2,8,0.1)
#### OLS fits
FIT.LINE = NULL
S2.LS<-BETA.LS<-NULL
for(j in 1:m) {
 fit<-lm(Y[[j]]~-1+X[[j]])
  BETA.LS<-rbind(BETA.LS,c(fit$coef))</pre>
  S2.LS<-c(S2.LS, summary(fit)$sigma^2)
}
PRED = BETA.LS %*% t(GRID)
plot(data_p1[,2], data_p1[,3], col = data_p1[,1], xlab = 'Density', ylab = 'Yield')
for(j in 1:m){
  lines(x_grid, PRED[j,], type ='l', col = 'gray')
}
```



```
apply(BETA.LS,2,mean)
     X[[j]] X[[j]]xj X[[j]]
    2.86875 1.85485 -0.15925
mean(S2.LS)
## [1] 0.7878099
cov(BETA.LS)
                 X[[j]]
                            X[[j]]xj
                                            X[[j]]
##
             2.00120764 -0.69321312 0.044309549
## X[[j]]
## X[[j]]xj -0.69321312  0.27555421 -0.020742679
## X[[j]]
             0.04430955 -0.02074268 0.001968451
b, c)
## mvnormal simulation
rmvnorm<-function(n,mu,Sigma)</pre>
  E<-matrix(rnorm(n*length(mu)),n,length(mu))</pre>
  t( t(E%*%chol(Sigma)) +c(mu))
## Wishart simulation
rwish<-function(n,nu0,S0)</pre>
{
  sSO <- chol(SO)
```

```
S<-array( dim=c( dim(S0),n ) )</pre>
  for(i in 1:n)
    Z <- matrix(rnorm(nu0 * dim(S0)[1]), nu0, dim(S0)[1]) %*% sS0
    S[,,i] \leftarrow t(Z) %*%Z
  S[,,1:n]
## Setup
p<-dim(X[[1]])[2]</pre>
theta<-mu0<-apply(BETA.LS,2,mean)
nu0 < -2; s2 < -s20 < -mean(S2.LS)
etaO<-4; Sigma<-SO<-LO<-cov(BETA.LS); BETA<-BETA.LS
THETA.b<-S2.b<-NULL
iL0<-solve(L0) ; iSigma<-solve(Sigma)</pre>
Sigma.ps<-matrix(0,p,p)</pre>
SIGMA.PS<-NULL
BETA.ps<-BETA*0
BETA.pp<-NULL
set.seed(1)
## MCMC
for(s in 1:10000) {
  ##update beta_j
  for(j in 1:m)
    Vj<-solve( iSigma + t(X[[j]])%*%X[[j]]/s2 )</pre>
    Ej < -Vj\%*\%( iSigma\%*\%theta + t(X[[j]])\%*\%Y[[j]]/s2 )
    BETA[j,]<-rmvnorm(1,Ej,Vj)</pre>
  }
  ##
  ##update theta
  Lm<- solve( iL0 + m*iSigma )</pre>
  mum<- Lm%*%( iL0%*%mu0 + iSigma%*%apply(BETA,2,sum))</pre>
  theta<-t(rmvnorm(1,mum,Lm))</pre>
  ##
  ##update Sigma
  mtheta<-matrix(theta,m,p,byrow=TRUE)</pre>
  iSigma<-rwish(1, eta0+m, solve(S0+t(BETA-mtheta)%*%(BETA-mtheta)))
  ##update s2
  RSS<-0
  for(j in 1:m) { RSS<-RSS+sum( (Y[[j]]-X[[j]]%*%BETA[j,] )^2 ) }</pre>
  s2<-1/rgamma(1,(nu0+sum(N))/2,(nu0*s20+RSS)/2)
  ##
  ##store results
  if(s\%10==0)
    # cat(s, s2, "\n")
    S2.b<-c(S2.b,s2); THETA.b<-rbind(THETA.b,t(theta))
    Sigma.ps<-Sigma.ps+solve(iSigma); BETA.ps<-BETA.ps+BETA
    SIGMA.PS<-rbind(SIGMA.PS,c(solve(iSigma)))</pre>
    BETA.pp<-rbind(BETA.pp,rmvnorm(1,theta,solve(iSigma)) )</pre>
  }
```

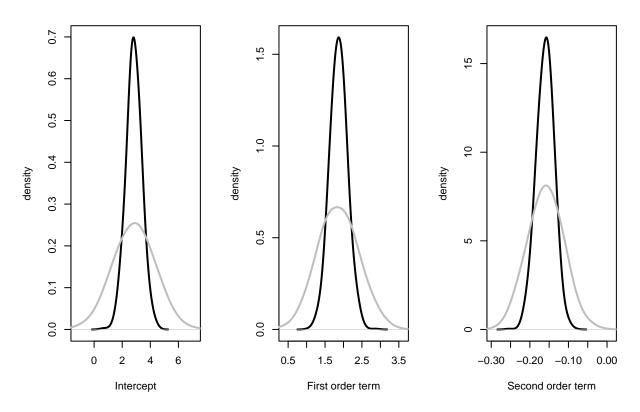
```
BETA.PM <- BETA.ps/1000
BETA.PM.mean <- apply(BETA.PM, 2, mean)
PRED.bayesian = BETA.PM %*% t(GRID)
pred.bayesian.mean = BETA.PM.mean %*% t(GRID)
plot(data_p1[,2], data_p1[,3], col = data_p1[,1],xlab = 'Density', ylab = 'Yield')
for(j in 1:m){
   lines(x_grid, PRED.bayesian[j,], type ='l', col = 'gray')
}
lines(x_grid, pred.bayesian.mean, type ='l', col = 'black')</pre>
```



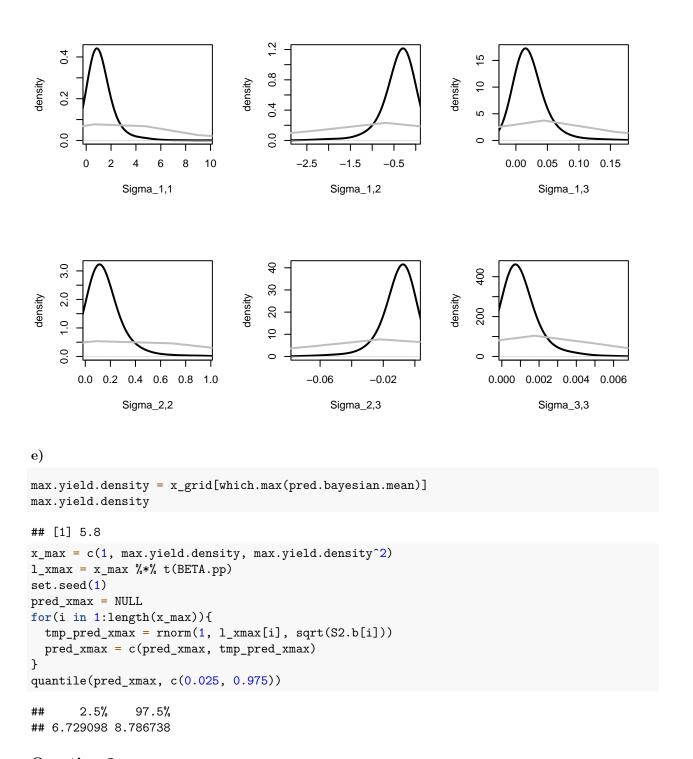
The fitted functions for different groups are closer to the overall fitted function.

d)

```
prior_theta = rmvnorm(1000, mu0, Sigma)
par(mfrow=c(1,3))
plot(density(THETA.b[,1],adj=2),xlim=range(BETA.pp[,1]),
main="",xlab="Intercept",ylab="density",lwd=2)
lines(density(prior_theta[,1],adj=2),col="gray",lwd=2)
plot(density(THETA.b[,2],adj=2),xlim=range(BETA.pp[,2]),
main="",xlab="First order term",ylab="density",lwd=2)
lines(density(prior_theta[,2],adj=2),col="gray",lwd=2)
plot(density(THETA.b[,3],adj=2),xlim=range(BETA.pp[,3]),
main="",xlab="Second order term",ylab="density",lwd=2)
```



```
SIGMA.prior.Sample = NULL
for(i in 1:1000){
  tmp = rwish(1, 4, solve(Sigma))
  SIGMA.prior.Sample<-rbind(SIGMA.prior.Sample,c(solve(tmp)))</pre>
par(mfrow=c(2,3))
plot(density(SIGMA.PS[,1],adj=5),xlim=range(SIGMA.PS[,1]),
main="",xlab="Sigma_1,1",ylab="density",lwd=2, col = 'black')
lines(density(SIGMA.prior.Sample[,1],adj=5),lwd=2, col = 'gray')
plot(density(SIGMA.PS[,2],adj=5),xlim=range(SIGMA.PS[,2]),
main="",xlab="Sigma_1,2",ylab="density",lwd=2, col = 'black')
lines(density(SIGMA.prior.Sample[,2],adj=5),lwd=2, col = 'gray')
plot(density(SIGMA.PS[,3],adj=5),xlim=range(SIGMA.PS[,3]),
main="",xlab="Sigma 1,3",ylab="density",lwd=2, col = 'black')
lines(density(SIGMA.prior.Sample[,3],adj=5),lwd=2, col = 'gray')
plot(density(SIGMA.PS[,5],adj=5),xlim=range(SIGMA.PS[,5]),
main="",xlab="Sigma_2,2",ylab="density",lwd=2, col = 'black')
lines(density(SIGMA.prior.Sample[,5],adj=5),lwd=2, col = 'gray')
plot(density(SIGMA.PS[,6],adj=5),xlim=range(SIGMA.PS[,6]),
main="",xlab="Sigma_2,3",ylab="density",lwd=2, col = 'black')
lines(density(SIGMA.prior.Sample[,6],adj=5),lwd=2, col = 'gray')
plot(density(SIGMA.PS[,9],adj=5),xlim=range(SIGMA.PS[,9]),
main="",xlab="Sigma_3,3",ylab="density",lwd=2, col = 'black')
```



Question 2

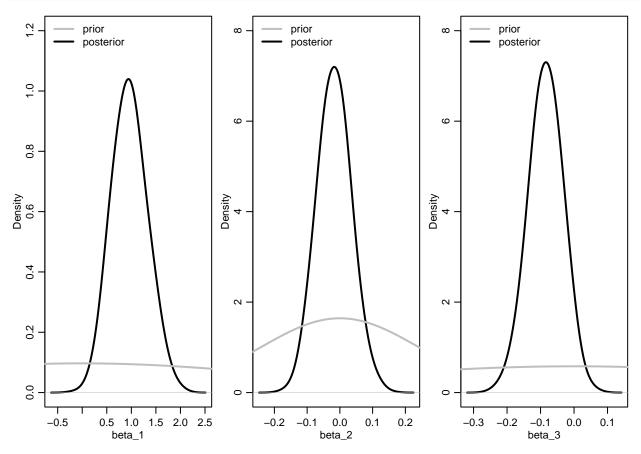
```
data_p2 = read.table("http://www2.stat.duke.edu/~pdh10/FCBS/Exercises/prayer.dat", header = TRUE)
y = data_p2[,4]
```

```
X = as.matrix(data_p2[,1:3])
# Rank likelihood regression
treg<-function(y,X,NSCAN=25000,seed=1) {</pre>
  set.seed(seed)
  n < -dim(X)[1]; p < -dim(X)[2]
  iXX < -solve(t(X)% * %X); V < -iXX * (n/(n+1)); cholV < -chol(V)
  ranks<-match(y,sort(unique(y))) ; uranks<-sort(unique(ranks))</pre>
  z<-qnorm(rank(y,ties.method="random")/(n+1))</pre>
  b<-matrix(0,p,1); BETA<-matrix(NA,1000,p); Z<-matrix(NA,1000,n)
  for(nscan in 1:NSCAN) {
    ###update b
    E \leftarrow V%*%(t(X)%*%z)
    b<- cholV%*%rnorm(p) + E
    ###update z
    mu < -X%*%b
    for(r in sample(uranks)) {
      ir<-(1:n)[ranks==r]
      lb<-suppressWarnings(max(z[ranks<r]))</pre>
      ub<-suppressWarnings(min(z[ranks>r]))
      z[ir] <-qnorm(</pre>
        runif( length(ir), pnorm(lb,mu[ir],1), pnorm(ub,mu[ir],1) ),
      )
    }
    ###help mixing
    zp < -z + rnorm(1, 0, n^(-1/3))
    lhr<- sum(dnorm(zp,mu,1,log=T) - dnorm(z,mu,1,log=T) )</pre>
    if(\log(\text{runif}(1)) < \text{lhr}) \{ z < -zp ; ac < -ac + 1 \}
    ###output
    if(nscan%%(NSCAN/1000)==0) {
      BETA[nscan/(NSCAN/1000),] <- t(b)
      Z[nscan/(NSCAN/1000),] < -z
    }
  }
  list( BETA=BETA, Z=Z )
treg.adhoc<-function(y,X,z=qnorm(rank(y,ties.method="random")/(length(z)+1)))
  ranks<-match(y,sort(unique(y))); uranks<-sort(unique(ranks))</pre>
  for(s in 1:10)
    fit < -lm(z \sim -1 + X); b<-fit$coef; e<-fit$res
    for(r in uranks)
      ir<-(1:n)[ranks==r]
      lb<-suppressWarnings(max(z[ranks<r]))</pre>
      ub<-suppressWarnings(min(z[ranks>r]))
      z[ir] <-sort(z[ir] )[rank(-e[ir]) ]</pre>
    }
  }
  z
}
```

```
rfit <- treg(y,X)
n<-dim(X)[1]
sd<-sqrt(solve(t(X)%*%X/n))</pre>
```

Warning in sqrt(solve(t(X) %*% X/n)): NaNs produced

```
x<-seq(-5,5,length=1000)
par(mfrow=c(1,3),mar=c(2.75,2.75,.5,.5),mgp=c(1.7,.7,0))
plot(density(rfit$BETA[,1],adj=2),lwd=2,main="",
xlab='beta_1',col="black",ylim=c(0,1.2),ylab='Density')
lines(x, dnorm(x, 0, sd[1,1]), lwd = 2, col = 'gray')
legend('topleft',legend=c("prior","posterior"),lwd=c(2,2),col=c("gray","black"),bty="n")
plot(density(rfit$BETA[,2],adj=2),lwd=2,main="",
xlab='beta_2',col="black",ylim=c(0,8),ylab='Density')
lines(x, dnorm(x, 0, sd[2,2]), lwd = 2, col = 'gray')
legend('topleft',legend=c("prior","posterior"),lwd=c(2,2),col=c("gray","black"),bty="n")
plot(density(rfit$BETA[,3],adj=2),lwd=2,main="",
xlab='beta_3',col="black",ylim=c(0,8),ylab='Density')
lines(x, dnorm(x, 0, sd[3,3]), lwd = 2, col = 'gray')
legend('topleft',legend=c("prior","posterior"),lwd=c(2,2),col=c("gray","black"),bty="n")</pre>
```



d)
quantile(rfit\$BETA[,1], c(0.025, 0.975))

2.5% 97.5%

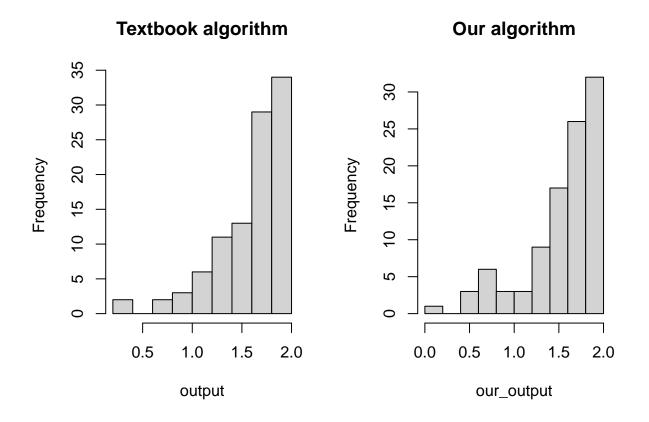
```
## 0.3154128 1.6833093
quantile(rfit$BETA[,2], c(0.025, 0.975))
##
          2.5%
                      97.5%
## -0.11285892 0.08219409
quantile(rfit$BETA[,3], c(0.025, 0.975))
           2.5%
                        97.5%
## -0.184092059 0.008249275
The coefficient
                                              \beta_1
is significant, which means female pray more than male.
Question 3
n = 100
mu = 8
sigma2 = 3
a = 0
b = 2
# Textbook algorithm
u = runif(100, pnorm((a-mu)/sqrt(sigma2)), pnorm((b-mu)/sqrt(sigma2)))
output = mu + sqrt(sigma2)*qnorm(u)
set.seed(1)
# Our naive algorithm
i = 0
iter = 0
our_output = NULL
while(i < n){</pre>
  iter = iter + 1
 x = rnorm(1, mu, sqrt(sigma2))
 if(x > a && x < b){
    our_output = c(our_output, x)
    i = i+1
  }
```

```
## [1] "Our algorithm iterates 343578 times."
```

print(paste('Our algorithm iterates', iter, 'times.'))

}

```
par(mfrow = c(1, 2))
hist(output, main = 'Textbook algorithm')
hist(our_output, main = 'Our algorithm')
```



The textbook algorithm significantly outperform the algorithm implemented here.

Question 4

```
library(sbgcop)
##
## Attaching package: 'sbgcop'
## The following object is masked _by_ '.GlobalEnv':
##
##
       rwish
load("socmob.RData")
data_p4 = socmob
n = dim(data_p4)[1]
p = dim(data_p4)[2]
nu0 = p + 2
S0 = (p+2)*diag(p)
n_{iter} = 1000
res_copula <- sbgcop.mcmc(</pre>
  Y = data_p4,
  SO = SO,
  n0 = nu0,
  nsamp = 10000,
  odens = 10,
```

```
seed = 1,
  verb = FALSE
cov_all= res_copula$C.psamp
apply(cov_all, c(1,2), mean)
##
                INCOME
                            DEGREE
                                     CHILDREN
                                                PINCOME
                                                          PDEGREE
                                                                    PCHILDREN
## INCOME
             1.00000000 \quad 0.47070246 \quad 0.28694099 \quad 0.1260760 \quad 0.1697058 \quad -0.04924544
## DEGREE
             0.47070246 1.00000000 -0.04059447 0.2014234
                                                        0.4553500 -0.19899551
## CHILDREN
             0.28694099 \ -0.04059447 \ 1.00000000 \ -0.1474823 \ -0.2470654 \ 0.21871297
## PINCOME
             0.4255160 -0.20717398
## PDEGREE
             0.16970583 \quad 0.45534996 \quad -0.24706536 \quad 0.4255160 \quad 1.0000000 \quad -0.27950214
## PCHILDREN -0.04924544 -0.19899551 0.21871297 -0.2071740 -0.2795021 1.00000000
## AGE
             ##
                   AGE
             0.32376685
## INCOME
## DEGREE
             0.04794096
## CHILDREN
             0.57404790
```

PINCOME

PDEGREE

AGE

-0.12405462

-0.21943953

1.00000000

PCHILDREN 0.11530131