### Final Exam

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#### 1

Selecting a beta distribution because  $0 \le \theta \le 1$ . The  $E[\theta] = \alpha/\beta = .03$  and  $2\sqrt{\text{var}(\theta)} = 2\sqrt{\alpha/\beta^2} = .07$ .

$$\beta = \frac{E[\theta]}{var(\theta)}$$

$$= \sqrt{\frac{.07}{2}}$$

$$= .187 * .03$$

$$= .00561.$$

I have  $Y \sim bin(25, \theta)$ . This sample Y = 0. Then

$$p(\theta|y) \propto p(y = 0|\theta)p(\theta)$$

$$\propto \theta^{y}(1-\theta)^{25-y}\theta^{\alpha-1}(1-\theta)^{\beta-1}$$

$$\propto \theta^{\alpha-1}(1-\theta)^{25+\beta-1}$$

$$\{\theta|y=0\} \sim \text{beta}(\alpha, 25+\beta)$$

Point estimate of  $\theta$  is  $\hat{\theta} = E[\theta|y=0] = \alpha/(25+\beta) = 0.014\,845\,36$ . 95% upper bound for  $\theta$  is global global global estimate of  $\theta$  is global glob  $0.048\,569\,43$ , giving interval for  $\theta$  is  $(0, 0.048\,569\,43)$ .

#### 2

Took  $S = 1\,000\,000$  Monte Carlo samples of posterior distribution of  $\beta$ . Marginal posterior means  $\hat{\beta}_1 = 0.016\,699\,77$ ,  $\hat{\beta}_2 = 0.016\,699\,77$  $0.020\,701\,44$ . Marginal 95% HPD intervals showed  $\beta_1 \in (0.007\,630\,409, 0.025\,895\,26), \beta_2 \in (0.011\,814\,688, 0.029\,731\,22)$ . 95% HPD interval for  $E[Y^*|X_1^*=70,X_2^*=80] \in (2.613\,156,3.033\,227)$ . 95% HPD for  $\{Y^*|X_1^*=70,X_2^*=80\} \in (1.576\,744,4.090\,318)$ . Used the Monte Carlo sample as the posterior distri-

bution of  $\{\beta_1, \beta_2, \sigma^2 | Y\}$  and found 95% HPD using HPDinterval. Predicted Mary's GPA from each Monte Carlo

# 9.0 0.5 4. 0.3 0.2 0.1 0.0 2

GPA

Histogram of Fred's predicted GPA

sample and subtracted Fred's predicted GPA from the same sample. The proportion of these that were greater than 0 are the probability of Mary having a higher GPA and is equal to 0.555 257.

Comparing the data in menchild30nobach.dat to a new mixture model. The sampling and prior distributions for the mixture model are

$$\begin{split} \{Y_i|X_i,\theta\} \sim \begin{cases} 0 & \text{individual } i \text{ is unmarried}(X_i=0) \\ \text{Poisson}(\theta) & \text{individual } i \text{ is married}(X_i=1) \end{cases} \\ \{X_i|p\} \sim \text{Bernoulli}(p) \\ p \sim \text{Beta}(1,1) \\ \theta \sim \text{Gamma}(1,\frac{1}{3}). \end{split}$$

The prior on p was selected as a minimal prior, while the prior on  $\theta$  loosely represented my idea that families have 2-3 children on average. The goal is to see if this model fits the data better than a Poisson model. In order to sample from the new model, I made a Gibbs sampler. The full conditional distributions derived for p and  $\theta$  are

$$p(p|y, x, \theta) \propto p(x|p)p(p)$$

$$\propto \prod_{i=1}^{n} p^{x_i} (1-p)^{1-x_i}$$

$$\propto p^{\sum x_i} (1-p)^{n-\sum x_i}$$

$$\{p|Y, X, \theta\} \sim \text{Beta}\left(\sum_{i=1}^{n} x_i + 1, n - \sum_{i=1}^{n} x_i + 1\right)$$

$$p(\theta|y, x, p) \propto p(y|x, \theta)p(\theta)$$

$$\propto \prod_{\{i|x_i=1\}} (\theta^{y_i} e^{-\theta})e^{-\frac{\theta}{2}}$$

$$\propto \theta^{n_1\bar{y}_1} e^{-(n_1 + \frac{1}{2})\theta}$$

$$\{\theta|y, x, p\} \sim \text{Gamma}\left(n_1\bar{y}_1 + 1, n_1 + \frac{1}{2}\right).$$

For x, I used basic probability theory,

$$p(x_{i} = 0, y_{i} = 0 | \theta, p) = p$$

$$p(x_{i} = 1, y_{i} = 0 | \theta, p) = (1 - p)e^{-\theta}$$

$$p(x_{i} = 1, y_{i} > 0 | \theta, p) = (1 - p)(1 - e^{-\theta})$$

$$p(x_{i} = 1, y_{i} > 0 | \theta, p) = (1 - p)(1 - e^{-\theta})$$

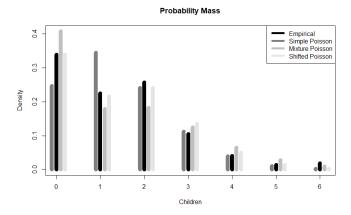
$$p(x_{i} = 0 | y_{i} = 0, \theta, p) = \frac{p}{p + (1 - p)\exp(-\theta)}$$

$$p(x_{i} = 0 | y_{i} > 0, \theta, p) = 0$$

$$p(x_{i} = 1 | y_{i} > 0, \theta, p) = 0$$

$$p(x_{i} = 1 | y_{i} > 0, \theta, p) = 1$$

I get a Bayes factor of 56.13282 (log 4.027721) to show a preference for the new mixture model over the Poisson model, leading me to prefer the new model, although graphically it is not ideal. The mixture model mean is 1.40308, close to the sample mean of 1.399083, with a variance of 2.35191, not as far from the sample variance 1.91834 as the simple Poisson model (also not as far in standard deviation). I also decided to check a similar (and simpler) mixture model, where Y = 0 with probability p, otherwise  $Y - 1 \sim \text{Poisson}(\theta)$ . This shifted Poisson model gave much better results visually, and had a Bayes factor of 81.83272 (log 4.404677) against the mixture model.



## 4 Appendix

```
#### Math 457 ####
#### Final Exam ####
#### Miles Keppler ####
setwd("U:/Winter_2018/M457/")
library (coda)
#### 1 ####
mean.theta = .03
sd.theta = .07/2
beta = mean.theta / sd.theta^2
alpha = beta * mean.theta
alpha/(25+beta)
qbeta(.95, alpha, beta+25)
curve(dbeta(x,alpha,beta),lty=2,col="gray60")
curve(dbeta(x,alpha,beta+25),add=T)
legend("topright", legend=c("Prior", "Posterior"), lty=c(2,1), col=c("gray60", "black"))
#### 2 ####
# gprior function from
# http://wwwlegacy.stat.washington.edu/people/pdhoff/Book/ComputerCode/regression_gprior.r
lm.gprior<-function(y, X, g=dim(X)[1], nu0=1,</pre>
                     s20=try(summary(lm(y^-1+X))$sigma^2,silent=TRUE),S=1000) {
  n < -dim(X)[1]; p < -dim(X)[2]
  Hg < - (g/(g+1)) *X% * % solve(t(X) % * % X) % * % t(X)
  SSRg<- t(y) %*% (diag(1, nrow=n)-Hg) %*%y
  s2 < -1/rgamma(S, (nu0+n)/2, (nu0*s20+SSRg)/2)
  Vb < - g * solve(t(X) % * % X) / (g+1)
  Eb<- Vb%*%t(X)%*%y
  E<-matrix(rnorm(S*p, 0, sqrt(s2)), S, p)</pre>
  beta<-t(t(E%*%chol(Vb))+c(Eb))
```

```
list (beta=beta, s2=s2)
}
data = read.table("GPAdata.dat", header = T)
Y = data$y
X = cbind(data$x1, data$x2)
n = length(Y)
S = 1000000
myglm = lm.gprior(y = Y, X = X, g = n, nu0 = 2, s20 = .4^2, S = S)
colMeans (myglm$beta)
HPDinterval(as.mcmc(myglm$beta))
fredmean = 70*myglm$beta[,1]+80*myglm$beta[,2]
HPDinterval(as.mcmc(fredmean))
fredsample = fredmean+rnorm(S,0,sqrt(myglm$s2))
HPDinterval(as.mcmc(fredsample))
hist(fredsample,probability=T,main="Histogram_of_Fred's_predicted_GPA",xlab="GPA")
marysample = 65*myglm\$beta[,1]+90*myglm\$beta[,2]+rnorm(S,0,sqrt(myglm$s2))
sum (marysample>fredsample) /S
#### 3 ####
set.seed(111)
Y = scan("menchild30nobach.dat")
findMass = function(x, t)  {
  n = as.numeric(names(t))
  for (i in 1:length(n)) {
   if (n[i] == x) {
      return(as.numeric(t)[i])
  return(0)
}
# Gibbs sampler (mixture model)
# should look like scaled poisson plus mass at 0
n = length(Y)
p = .5
theta = 2
M = 100000
PHI = matrix(NA, ncol = 2, nrow = M)
for (i in 1:M) {
  X = NULL
  for (j in 1:n) {
    if (Y[j] == 0) {
     X[j] = rbinom(1, 1, (1-p) *exp(-theta) / (p+(1-p) *exp(-theta)))
    } else {
      X[j] = 1
    }
```

```
}
  sx = sum(X)
  p = rbeta(1, sx+1, n-sx+1)
  theta = rgamma(1, sum(Y[X==1])+1, sx + 1/3)
  PHI[i,] = c(theta, p)
NY = rep(NA, M)
for (i in 1:M) {
  theta = PHI[i,1]
  p = PHI[i,2]
  x = rbinom(1, 1, p)
  ny = 0
  if (x==1) {
   ny = rpois(1, theta)
 NY[i] = ny
# Plotting
# Empirical
taby = table(Y)/n
tabny = table(NY)/M
yind = as.numeric(names(taby))
plot(as.numeric(yind),c(taby),type="h",ylim = c(0,max(c(taby,tabny))),lwd=10,
     main = "Probability_Mass", xlab = "Children", ylab = "Density", xlim = c(0-.1, max(yind))
# Mixture Poisson
nyind = as.numeric(names(tabny))
points(as.numeric(nyind)+.1,c(tabny),type="h",lwd=10,col="gray75")
# Poisson model
points (0:max(yind)-.1, dpois(min(yind):max(yind), mean(Y)), type="h", lwd=10, col="gray50")
p = sum(Y==0)/n
mytab = as.table(c(p,(1-p)*dpois(as.numeric(names(taby))[-1]-1,mean(Y[Y!=0]-1))))
names(mytab) = mytab.names = 0:(length(mytab)-1)
points(mytab.names+.2, mytab, type="h", lwd=10, col="gray90")
legend("topright", legend = c("Empirical", "Simple_Poisson", "Mixture_Poisson", "Shifted_Poisson
       lwd = c(5,5,5,5), col = c("black", "gray50", "gray75", "gray90"))
LL = c(0,0,0)
for (i in 1:n) {
  LL[1] = LL[1] + log(findMass(Y[i], tabny))
  LL[2] = LL[2] + log(dpois(Y[i], mean(Y)))
 LL[3] = LL[3] + log(findMass(Y[i], mytab))
bf = c(exp(LL[1]-LL[2]), exp(LL[3]-LL[1]))
bf ; bf/(1+bf)
nLL = 0
for (i in 1:n) {
  nLL = nLL + log(findMass(Y[i], taby))
}
```