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title: STAT 457 Homework 04
author: Martha Eichlersmith
date: 2019-11-05
output:
 pdf document:
   fig caption: yes
header-includes:
 - \usepackage{color}
 - \usepackage{mathtools}
 - \usepackage{amsbsy} #bold in mathmode
 - \usepackage{nicefrac} # for nice fracs
```{r, echo=FALSE, results="hide", warning=FALSE, message=FALSE}
library(ggplot2) #ggplot
library(readr) #import CSV
library(gridExtra) #organize plots
library(grid) #organize plots
library(latex2exp) #latex in ggplot titles
library(knitr) #to help make tables
library(matlib) \#A = matrix, inv(A) = A^{-1}
library(numDeriv) #calculate numerical first and second order
derivatives
decimal <- function(x, k) trimws(format(round(x, k), nsmall=k))</pre>
dec <- 5
## Problem 1
Let y_1, cdots, y_n be an iid sample from the Poisson
distribution with parameter $\lambda$. Derive Jeffery's
(noninformative) prior. This prior corresponds to the gamma
distribution with which parameters?
$$
\begin{aligned}
L(\lambda y) &= \frac{y!}{u}
\propto \lambda^y e^{-\lambda}
\\[.5ex]
\left( \right) & = y \log \alpha - \beta
\frac{\partial \ell(\lambda \mid Y)}{\partial \lambda }
& = \frac{y}{\lambda} - 1
\[.5ex]
\frac{\partial^2 \ell(\lambda \mid Y)}{\partial \lambda^2}
& = - \frac{y}{\lambda^2}
```

```
\right]
 $$
 and |X| is the determinant of the matrix X.
Let y 1, cdots, y n be an iid sample from the mathcal{N}
  (\mu, \sigma^2)$ distribution, where $\mu$ and $\sigma$ are both
unknown. Derive the invariant prior. How does it compare with
the prior $p(\theta, \sigma^2) \propto \nicefrac{1}{\sigma^2}$?
 $$
 \begin{aligned}
L(\mu, \sigma^2 \mb{Y}) &= \prod_{i=1}^n \frac{1}{\sqrt{}}
2 \pi \sigma^2}} \exp \left\{ \frac{ (y i - \mu)^2}{2 \sigma^2}}
 \right\}
 \[.5ex]
 \left( \sum_{i=1}^n \right) 
 \frac{1}{\sqrt{2 \cdot j}} \exp \left( \frac{1}{j} \right)
 \[.5ex]
 & = - \frac{n}{2} \log (2 \pi^2) - \frac{n}{2} \log (\frac{n}{2} - \frac{n}{2} \log (\frac{n}{2} - \frac{n}{2} \log n)
 \frac{(y i - \mu)^2}{2 \sigma^2}
 \[ .5ex]
 \frac{2 \left( \frac{n}{sigma^2} \right) }{ \left( \frac{n}{sigma^2} \right) }
 \[.5ex]
 \frac{2 \cdot \frac{1}{2} 
 \frac{ \int (y i - \mu)^2}{\sin^4} = \frac{\rho}{\mu}^2 \left(\frac{1}{\mu}\right)^2}
  {\partial \sigma^2 \partial \mu}
 \[.5ex]
 \frac{\alpha^2 \cdot \beta^2}{\beta^2} 
  { \langle n \rangle } = \frac{n}{2 \sin^4} - \frac{\sin^4}{\sin^4} 
  {\sigma^6}
 \\[1ex]
p(\mu, \sigma^2) \in p(\mu, \sigma^2) \in p(\mu, \sigma^2) = p(\mu,
 \[.5ex] & = \left[ .5et \right] 
- \frac{1}{n} \text{det} \begin{bmatrix}
-\nicefrac{n}{\sigma^2} & 0 \
 0 & - \nicefrac{n}{2 \sigma^4}
 \end{bmatrix}
 \right)^{\frac{1}{2}}
 \[.5ex]
 & = \frac{1}{\sqrt{3}}
 \end{aligned}
 This is different than commonly used prior of $\nicefrac{1}
  {\sigma^2}$. This shows that the Jeffery's prior does not work
```

```
for (i in 1:3) tg$grobs[[font.vec[i]]] <-</pre>
editGrob(tg$grobs[[font.vec[i]]], gp=gpar(col=colors[i]))
for (i in 1:3) tg$grobs[[bg.vec[i]]] <-</pre>
editGrob(tg$grobs[[bg.vec[i]]], gp=gpar(fill=colors[i]))
x < - seq(0,1, len=100)
qplot(x, geom="blank")+
  annotation custom(tg, xmin=0.4, ymin=1)+
  Uniform+Beta1+Beta2+
  ggtitle("Graph of Priors")+
 theme(axis.title.x = element blank())+
  scale colour manual("", values = c(colors[3], colors[2],
colors[1])) +
  scale linetype manual("", values=c(line[3], line[2], line[1]))
  theme(legend.position = "bottom")
The Uniform has a mean of 0.27 and both beta distributions have a
mean of 0.30. The uniform distribution has the lowest variance
and a block shape. The Beta(10.2,23.8) has the highest variance
and a lower peak than the Beta(20.4,47.6).
\newpage
### Problem 3b
Suppose a player gets 5 hits in 40 at-bats. For each of the
above priors: plot the likelihood, posterior and prior; compute
the probability that he player is better than a .200 hitter;
compute your best guess as to the batting average of the player;
compute a 95% credible interval for $p$.
```{r, echo=FALSE}
dist <- c("Likelihood", "Posterior", "Prior")</pre>
colors <- c("darkgrey", "navy", "maroon")</pre>
p < -5/40
n < -40
it <- 10000
Binomial.Likelihood <- data.frame("L"=rbinom(it, 1, p))</pre>
func distplot <- function (Likelihood, prior, posterior, a, b,
name) {
plot1 <- ggplot(Likelihood, aes(L))</pre>
+geom density(aes(y=..density.., col=dist[1]), lwd=1.5)+
 xlim(0, 1) + ylim(0, 8) +
  prior+posterior+
```

```
n < -40
y <- 5
a0 < - 0.19
b0 < -0.35
a < -y + a0
b < - n - y + b0
func postdist <- function(x) \{dunif(x, .19, .35) * dbeta(x, .19, .35) \}
shape1=a, shape2=b) }
            stat function(fun = dunif, args =
prior <-
list (min=0.19, max=0.35), lwd = 1.5, linetype="dashed",
aes(col=dist[3]))
posterior <-stat function(fun = func postdist, lwd = 1.5,</pre>
linetype="dotted", aes(col=dist[2]))
set.seed(040302)
x.sim < - seq(0, 1, length=1000)
y.sim <- func postdist(x.sim)</pre>
df <- data.frame("Xsim" = x.sim, "Ysim"=y.sim)</pre>
plot1 <- ggplot(Binomial.Likelihood, aes(L))</pre>
+geom density(aes(y=..density.., col=dist[1]), lwd=1.5)+
  xlim(0, 1) + ylim(0, 20) +
  prior+posterior+
  ggtitle(paste("Unifrom(.19, .35) Prior, Binomial Likelihood,
Beta Posterior"))+
  theme(axis.title.x = element blank())+
  scale colour manual("", values = c(colors[1], colors[2],
colors[3])) +
  theme(legend.position = "bottom")
#For the probability of > 0.2, we are essentially finding P(X)=
8), because 0.2 is **8**/40. We do this analytically with
integral, for each value of y and then add those probabilites up
lower <- 0.19
upper <- 0.35
f <- function(theta) {dbinom(y, n, theta) / (upper - lower) }</pre>
num integral <- integrate(f, .20, .35)</pre>
den integral <- integrate(f, .19, .35)</pre>
pval <- num integral$value/den integral$value</pre>
```{r, echo=FALSE, eval=FALSE }
```

```
```{r, echo=FALSE, warning=FALSE, fig.width=10, fig.height=4}
CI.up <- 0.3057802 #run beforehand (above)
CI.low <- 0.1910938 #run beforehand (above)
plot2 <- ggplot(df, aes(Xsim, Ysim)) + geom line(lwd = 1.5,</pre>
linetype="dotted", aes(col=dist[2]))+xlim(0, 1)+
  geom ribbon(data=subset(df, Xsim>0.2), aes(ymax=Ysim), ymin=0,
fill=colors[2], color=NA, alpha=0.4)+
  annotate("text", x=0.3, y=5, label=paste("P(X > 0.200 | Y):",
decimal(pval, dec)), hjust=0, size=5)+
  annotate ("text", x=0.3, y=8, label=paste ("CI: (",
decimal(CI.low, dec), ",", decimal(CI.up, dec), ")"), size=5,
hjust=0) +
  ggtitle("Probability X > 0.200, Credible Interval")+
  theme(axis.title.x = element blank())+
  scale colour manual("", values = c(colors[2])) +
  theme(legend.position = "bottom")
grid.arrange(plot1, plot2, nrow=1)
```{r, echo=FALSE, warning=FALSE, fig.width=10, fig.height=4}
#BETA PRIOR Beta(10.2,23.8)
n < -40
y <- 5
a0 < -10.2
b0 < -23.8
a < -y + a0
b < - n - y + b0
prior <-
         stat function(fun = dbeta, args = list(shape1=a0,
shape2=b0), lwd = 1.5, linetype="dashed", aes(col=dist[3]))
posterior <-stat function(fun = dbeta, args = list(shape1=a,</pre>
shape2=b), lwd = 1.5, linetype="dotted", aes(col=dist[2]))
func distplot (Binomial.Likelihood, prior, posterior, a, b,
"Beta(10.2, 23.8) Prior, Binomial Likelihood, Beta Posterior")
```

```
g0 <- 89 #number of games played
n0 <- 47 #number of games won
p0 < - n0/g0
g1 <- 73 #remaining games
it <- 1000
Binomial.Likelihood <- data.frame("L"=rbinom(it, 162, p0))/162
#random gen for est games won in next 73 games
func distplot3c <- function(Likelihood, a0, b0, name) {</pre>
a < - n0 + a0
b < -q0 - n0 + b0
prior <- stat function(fun = dbeta, args = list(shape1=a0,</pre>
shape2=b0), lwd = 1.5, linetype="dashed", aes(col=dist[3]))
posterior <-stat function(fun = dbeta, args = list(shape1=a,</pre>
shape2=b), lwd = 1.5, linetype="dotted", aes(col=dist[2]))
mean.post <-a/(a + b)
mode.post <- (a - 1)/(a + b - 2)
med.post <- (a - (1/3))/(a + b - 2/3)
stat <- c("Mean", "Mode", "Est. Median")</pre>
stat.val <- c(mean.post, mode.post, med.post)</pre>
est.games <- stat.val*162 #total games played is 162
actual.games <- rep(84, 3) #actual games won = 89
difference <- actual.games - est.games</pre>
labels <- c("Posterior Value", "Est Games Won", "Actual Games
Won", "Difference")
vec.blank <- rep(NA, length(labels))</pre>
compare <- data.frame("Mean"=vec.blank, "Mode"=vec.blank, "Est.</pre>
Median" = vec.blank)
for (i in 1:3) {
compare[,i] <- c(</pre>
                  decimal(stat.val[i], dec),
                  decimal(est.games[i], 3),
                  actual.games[i],
                  decimal(difference[i], 3)
}
rownames(compare) <- labels</pre>
plot <- ggplot(Likelihood, aes(L))</pre>
+geom density(aes(y=..density.., col=dist[1]), lwd=1.5)+
  xlim(0, 1) +
```

```
```{r, echo=FALSE, warning=FALSE, fig.width=10, fig.height=4}
a0 <- 0
b0 <- 0
func distplot3c (Binomial.Likelihood, a0, b0, "Haldane's Prior:
Beta(0, 0)")
\newpage
## Problem 4
The following data represents the number of arrivals for 45 time
intervals of length 2 minutes at a cashier's desk at a
supermarket and are taken from Andersen (1980):
```{r}
Arrival \leftarrow c(rep(0,6), rep(1,18), rep(2,9), rep(3,7), rep(4, 4),
rep(5, 1))
### Problem 4a
For a Gamma(2,1) prior, obtain the posterior distribution under a
Poisson ($\lambda$) model for the data. Draw the prior and the
posterior. Note on your plot the mean, variance and mode of the
posterior.
```{r, echo=FALSE, fig.width=10, fig.height=4}
dist <- c("Likelihood", "Posterior", "Prior")</pre>
colors <- c("darkgrey", "navy", "maroon")
n <- length(Arrival)</pre>
a0 < -2
b0 <- 1
a <- sum(Arrival) + a0</pre>
b < - n + b0
lam <- mean(Arrival)</pre>
prior <- stat function(fun = dgamma, args =</pre>
list(shape=a0,rate=b0), lwd = 1.5, aes(col=dist[3],
linetype=dist[3]))
posterior <- stat function(fun = dgamma, args = list(shape=a,</pre>
rate=b), lwd = 1.5, aes(col=dist[2], linetype=dist[2]))
mean.post <- a/b</pre>
var.post <- a/b^2</pre>
mode.post <- (a-1)/b
```

it <- 10000

```
a <- sum(Arrival) + a0
b < - n + b0
lam <- mean(Arrival)</pre>
prior <- stat function(fun = dgamma, args =</pre>
list(shape=a0, rate=b0), lwd = 1.5, aes(col=dist[3],
linetype=dist[3]))
posterior <- stat function(fun = dgamma, args = list(shape=a,
rate=b), lwd = 1.5, aes(col=dist[2], linetype=dist[2]))
mean.post <- a/b</pre>
var.post <- a/b^2
mode.post <- (a-1)/b
 ggplot(Poisson.Likelihood, aes(L))+
 prior+posterior+
  ggtitle("Gamma(.5, .00001) Prior, Poisson Likelihood, Gamma
Posterior")+
  geom point(aes(x=mode.post, y=dgamma(mode.post, a, b)), size=2,
shape=21, color="grey45", stroke=1.5)+
   annotate("text", x=mode.post, y=dgamma(mode.post, a, b),
label=paste("Mode:", decimal(mode.post, dec)), hjust=1.1, size=5,
color="grey45", fontface="bold")+
  geom point(aes(x=mean.post, y=dgamma(mean.post, a, b)), size=2,
shape=22, color="grey35", stroke=1.5)+
   annotate ("text", x=mean.post, y=dgamma (mean.post, a, b),
label=paste("Mean:", decimal(mean.post, dec)), hjust=-.1,
size=5.5, color="grey35", fontface="bold")+
   annotate ("text", x=2, y=dgamma(2, a, b),
label=paste("Variance:", decimal(var.post, dec)), hjust=-.1,
size=5, fontface="bold")+
  theme(axis.title.x = element blank())+
  scale colour manual("", values = c(colors[2], colors[3])) +
scale linetype manual("", values = c("dotted", "dashed"))+
 theme(legend.position = "bottom")
\newpage
## Problem 5
197 animals are distributed into four categories: $Y = (y 1, y 2,
y 3, y 4)$ according ot the genetic linkage model $\left( \frac{2}
+ \hat{4}, \frac{1 - \hat{4}}{4}, \frac{1 - \hat{4}}{4},
\frac{4}{4} \cdot \frac{4}{4}
```

```
root.approx <- tail(k, n=1)</pre>
    it.completed <- length(k)</pre>
    # Once the difference between x0 and x1 becomes sufficiently
small, output the results.
    if (abs(x1 - x0) < tol & !is.na(abs(x1-x0)))
      {print(paste("Start at", start, ": Root approximation is",
root.approx, "with", it.completed, "iterations"))
      break}
    else if( it.completed == it) {print(paste("Start at", start,
": diverges"))}
    else{x0 <- x1}
 }
tol <- 1e-5
it <- 1000
### Problem 5a
$$
L(\theta \neq 0 ) = (125, 18, 20, 34)) \propto (2+
\hat{34} \cdot (125) \cdot (1 - \theta)^{38} \cdot (\theta)^{34}
\[.5ex]
$$
### Problem 5b
$$
L(\theta \neq \theta) = (14, 0, 1, 5) \propto (2+ \theta)^{14}
\cdot (1 - \theta)^{1} \cdot (\theta)^{5}
$$
### Problem 5c
Use th Newton-Raphson to obtain the MLE (\hat t) for Y = (\hat t)
(125, 18, 20, 34)$. Start the algorithm at $\theta = .1, .2, .3,
.4, .6, .8$.
```{r, echo=FALSE, eval=FALSE}
#How do you assess convergence of the algorithm.
```{r, echo=FALSE}
y1 <- 125
y2 <- 18
```

```
```{r, echo=FALSE}
y1 <- 14
y2 <- 0
y3 <- 1
y4 <- 5
func b<-function(x){</pre>
 y1/(2+x) - (y2+y3)/(1-x)+y4/x
 ) / (
 -y1/(2+x)^2-(y2+y3)/(1-x)^2-y4/x^2
  )
 \#-(14/(2+x)-1/(1-x)+5/x)/(14/(2+x)^2+1/(1-x)^2+5/x^2)
func newton.raphson(func b, 0.1, it, tol)
func_newton.raphson(func_b, 0.2, it, tol)
func newton.raphson(func b, 0.3, it, tol)
func newton.raphson(func b, 0.4, it, tol)
func newton.raphson(func b, 0.6, it, tol)
func newton.raphson(func b, 0.8, it, tol)
```