# TDDE07 - Lab 1

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# 1. Bernoulli ... again

### 1a)

En massa text här.

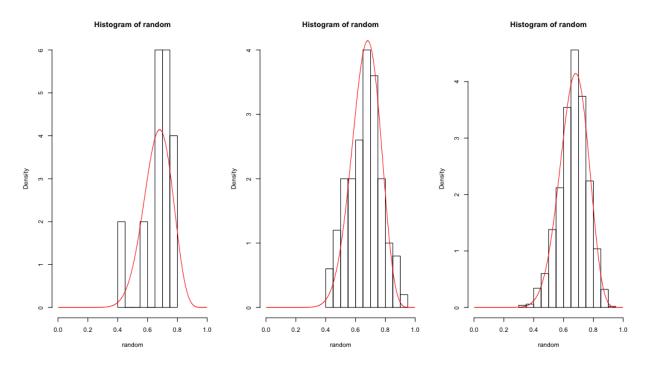


Figure 1: Convergence

### 1b)

The posterior was simulated with 10000 draws, resulting in  $P(\theta < 0.4|y) = 0.0043$ .

### 1c)

Visualization of the log odds  $\phi = log(\frac{\phi}{1-\phi})$  by simulation, using 10000 draws.

# density.default(x = random)

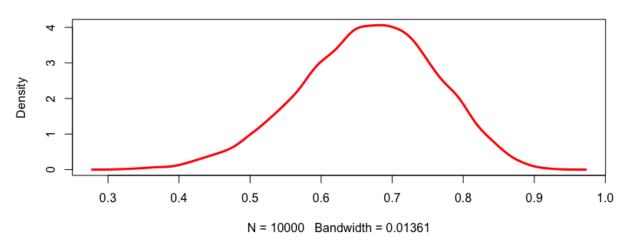


Figure 2: Posterior distribution of the log-odds

# 2. Log-normal distribution and the Gini coefficient

**2**a)

# Histogram of sigma\_sq 4 0.0 0.5 1.0 1.5 2.0 sigma\_sq

Figure 3: Posterior of  $\sigma^2$  by draws and the theoretical posterior

# Histogram of G

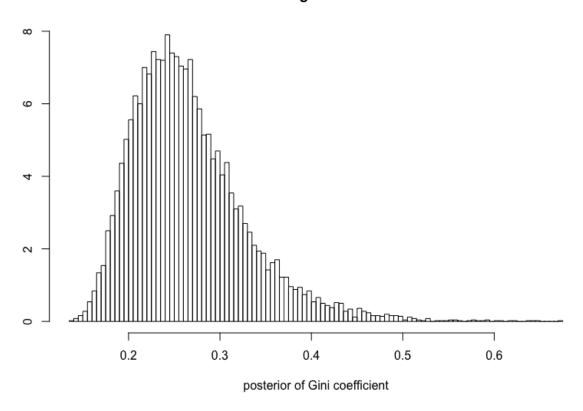


Figure 4: Gini Coefficient

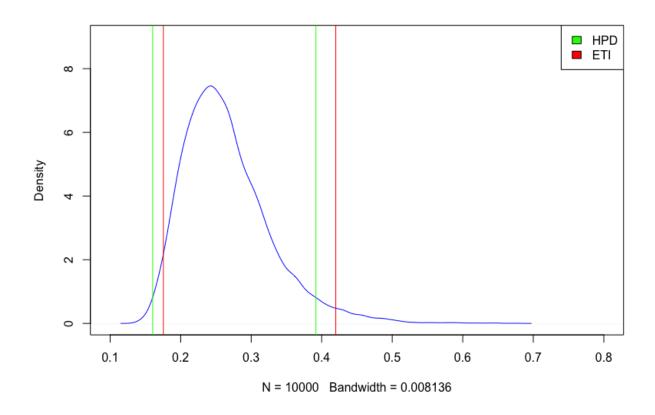


Figure 5: 95% Credibility interval

# 3. Bayesian inference for the concentration parameter in the von Mises distribution

The distribution of  $\kappa$  was calculated as the product of the likelihood and the prior, where the likelihood was assumed to be the product of the individual probabilities due to independent observations.

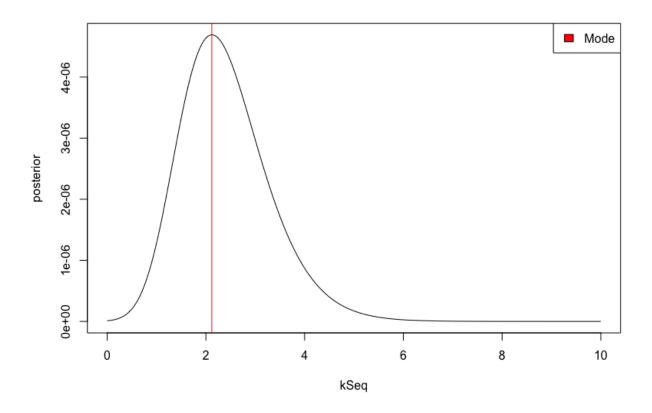


Figure 6: Posterior distribution and mode

### 1 - Code

```
alpha <- 2
beta <- 2
s <- 14
n <- 20
nDraws <- c(10, 100, 1000)
# 1a
xGrid \leftarrow seq(0.001, 0.999, by=0.001)
posterior = dbeta(xGrid, alpha + s, beta + (n-s))
par(mfrow=c(1,3))
for (draws in nDraws) {
 random = rbeta(draws, alpha+s, beta+(n-s))
 hist(random, xlim = c(0,1), freq = FALSE, breaks=10)
 lines(xGrid, posterior, type='l', col ='red')
}
dev.off()
# 1b
nDraws <- 10000
random = rbeta(nDraws, alpha+s, beta+(n-s))
val <- random * (random < 0.4)</pre>
print(sum(val>0)/length(val))
# 1c
nDraws <- 10000
random = rbeta(nDraws, alpha+s, beta+(n-s))
phi <- log(random/(1 - random))</pre>
plot(density(random), lwd=3, type='l', col='red')
```

### 2 - Code

```
# -----A-----
income \leftarrow c(14,25,45,25,30,33,19,50,34,67)
\#income \leftarrow c(1,1,1,1,1,1,1,1,1,1)
m < -3.5
n <- length(income)</pre>
n draws = 10000
t <- 0
for (v in income) {
 t <- t + (log(v) - m)^2
tSquared <- t / n
# simulate postarior draws
X_draw <- rchisq(n_draws, n)</pre>
sigma_sq = n*tSquared / X_draw
#theoretical
theoretical <- function(theta, v, s) {</pre>
 return (((v/2)^{(v/2)})/gamma(v/2)*(s^v)*(theta^{(-(v/2+1))})*exp((-v*s^2)/(2*theta)))
}
range <- seq(0,10,by=0.01)
mx <- 0 * range
for (i in 1:length(range)) {
 mx[i] <- theoretical(range[i], length(income), sqrt(tSquared))</pre>
}
# plot
hist(sigma_sq, 100, freq=FALSE)
lines(range, mx, lwd=3, type='l', col='red', xlab = 'sigma')
# -----B-----
z <- sqrt(sigma_sq/2)</pre>
G \leftarrow 2*pnorm(z)-1
\# Plot where y = values and x = index of the value in the vector
hist(G, 100, freq = FALSE, xlab="posterior of Gini coefficient", ylab="")
# -----C-----
cred_int \leftarrow quantile(G, probs = c(0.025, 0.975))
G_dens = density(G)
y_ordered = G_dens$y[order(-G_dens$y)]
x_ordered = G_dens$x[order(-G_dens$y)]
dens_mass = sum(G_dens$y)
```

```
sum <- 0
current_mass <- 0</pre>
for (i in 1:length(y_ordered)) {
 current_mass <- y_ordered[i] + sum</pre>
  if ((current_mass/dens_mass) > 0.95) {
    break
 } else {
    sum <- current_mass</pre>
}
a <- min(x_ordered[1:i])</pre>
b <- max(x_ordered[1:i])</pre>
plot(density(G),
     col='blue',
     xlim=c(0.1,0.8),
     ylim=c(0,9),
     main='')
legend('topright',
       legend = c('Highest Posterior Density', 'Equal Tail Interval'),
       fill = c('green', 'red'))
abline(v=cred_int[1], col='red')
abline(v=cred_int[2], col='red')
abline(v=a, col='green')
abline(v=b, col='green')
```

#### 3 - Code

```
\# A: Plot the posterior distribution of k
# posterior(k | y, mu) = likelihood * prior(k) = prod_prob * prior
y \leftarrow c(-2.44, 2.14, 2.54, 1.83, 2.01, 2.33, -2.79, 2.23, 2.07, 2.02)
kSeq \leftarrow seq(0,10, by=0.01)
mu <- 2.39
lambda <- 1
# kappa <- dexp(kSeq, lambda)</pre>
mises <- function(k, y, mu) {</pre>
 I <- besselI(k,0)</pre>
 return ((exp(k * cos(y - mu))) / (2 * pi * I))
kPos <- function(k, mu,y) {</pre>
  #prod since independent
 return ( prod( mises(k, y, mu) ) * dexp(k))
}
posterior = c()
for (k in kSeq){
  posterior = c(posterior, c(kPos(k, mu, y)))
plot(kSeq, posterior,type='l')
legend('topright', legend='Mode', fill='red')
\# B: Compute the posterior mode of k
kPosMode <- kSeq[which.max(posterior)]</pre>
abline(v=kPosMode, col='red', lwd=1)
```