

Stat 570 Final

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Q1

a

During each cycle, the likelihood can be calculated by multiplying together the probability $p(1-p)^{t-1}$ of pregnancy for Y_t conceptions (to the power of Y_t), and for the last case when we calculate the likelihood for the number of women that have not conceived by cycle N , the probability that they do not conceive for N cycles is $(1-p)^N$, and there are Y_{N+1} of them.

Therefore, we obtain the likelihood function as follows:

$$L(p) = f(t=1)f(t=2)\dots f(t=N)f(t>N) = (p(1-p)^{1-1})^{Y_1}(p(1-p)^{2-1})^{Y_2}\dots(p(1-p)^{N-1})^{Y_N}((1-p)^N)^{Y_{N+1}} = [\prod_{t=1}^N (p(1-p)^{t-1})^{Y_t}]((1-p)^N)^{Y_{N+1}}$$

b

$$\text{Log likelihood } l(p) = \log p \sum_{t=1}^N Y_t + \log(1-p) \sum_{t=1}^N Y_t(t-1) + N Y_{N+1} \log(1-p)$$

Taking derivative wrt p ,

$$\frac{1}{p} \sum_{t=1}^N Y_t - \frac{1}{1-p} \sum_{t=1}^N Y_t(t-1) - \frac{1}{1-p} N Y_{N+1} = 0$$

$$\text{Solving, we have } \hat{p} = \frac{\sum_{t=1}^N Y_t}{\sum_{t=1}^N (Y_t(t-1) + N Y_{N+1})}$$

c

$$\text{Taking derivative of the score function, we have } -\frac{1}{p^2} \sum_{t=1}^N Y_t - \frac{1}{(1-p)^2} \sum_{t=1}^N Y_t(t-1) - \frac{1}{(1-p)^2} N Y_{N+1}$$

$$\text{Therefore, the observed FIN is } \frac{1}{p^2} \sum_{t=1}^N Y_t + \frac{1}{(1-p)^2} \sum_{t=1}^N Y_t(t-1) + \frac{1}{(1-p)^2} N Y_{N+1} \text{ And theoretically it would be } \frac{N}{p^3} + \frac{N(t-1)}{(1-p)^2 p} + \frac{N}{(1-p)}$$

$$\text{Also, the asymptotic variance is } \frac{1}{\text{FIN}} = \frac{1}{\frac{1}{p^2} \sum_{t=1}^N Y_t + \frac{1}{(1-p)^2} \sum_{t=1}^N Y_t(t-1) + \frac{1}{(1-p)^2} N Y_{N+1}}$$

d

```
##               MLE      MLE sd   CI lower   CI upper
## Smoker result    0.2203791 0.01957709 0.1820088 0.2587495
## Nonsmoker result 0.3289382 0.01148539 0.3064273 0.3514492
```

e

$$P(Y|p) \propto [\prod_{t=1}^N (p(1-p)^{t-1})^{Y_t}]((1-p)^N)^{Y_{N+1}} p^a (1-p)^b = p^{\sum_{t=1}^N Y_t + a} (1-p)^{\sum_{t=1}^N (t-1)Y_t + N Y_{N+1} + b} \sim \text{Beta}(\sum_{t=1}^N Y_t + a, \sum_{t=1}^N (t-1)Y_t + N Y_{N+1} + b)$$

$$\text{Prior mean pm is } \frac{a}{a+b}$$

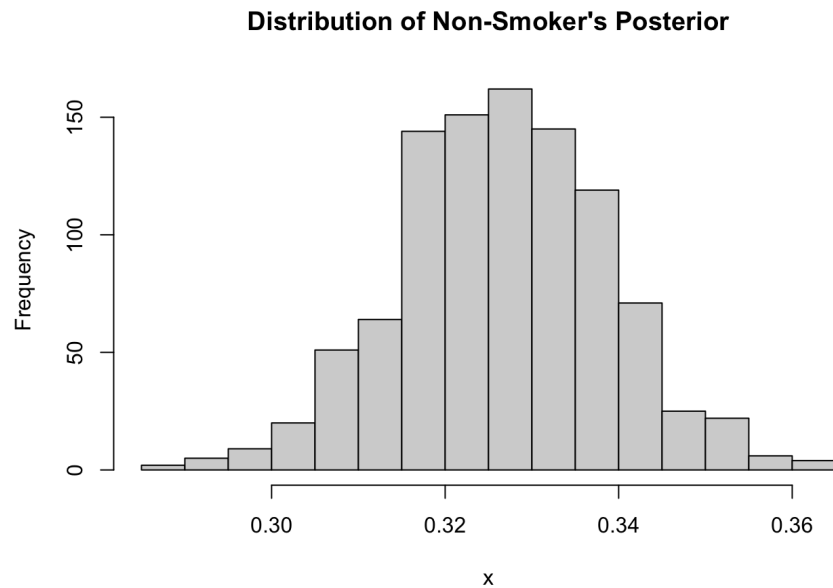
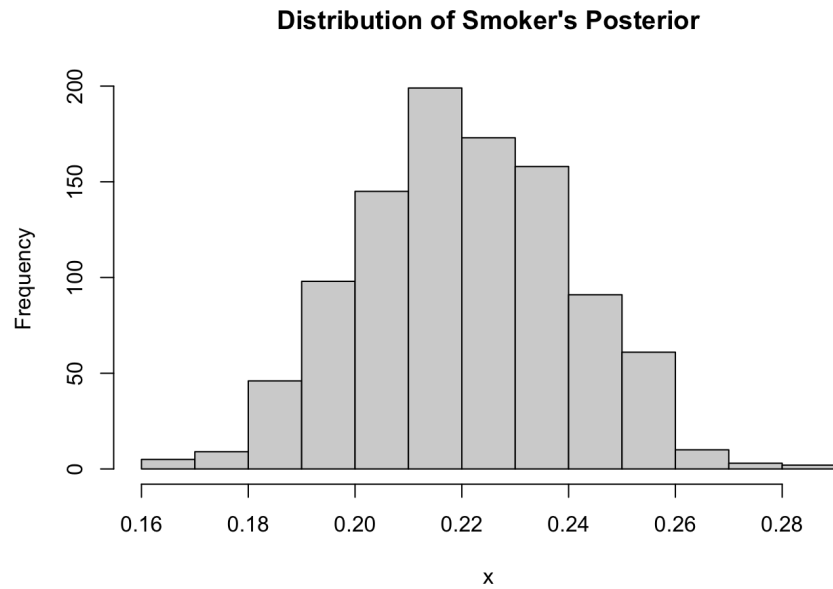
$$\text{The posterior mean is } \frac{\sum_{t=1}^N Y_t + a}{a + \sum_{t=1}^N (t-1)Y_t + N Y_{N+1} + b} = \frac{\sum_{t=1}^N (t-1)Y_t + N Y_{N+1}}{a + \sum_{t=1}^N (t-1)Y_t + N Y_{N+1} + b} \hat{p} + \frac{a+b}{a + \sum_{t=1}^N (t-1)Y_t + N Y_{N+1} + b} * \text{pm}$$

f

$$\mu = \frac{a}{a+b} \text{ and } \sigma^2 = \frac{ab}{(a+b)^2(a+b+1)}.$$

$$\text{Writing } a = \frac{\mu}{1-\mu} b \text{ we obtain } b = \frac{\mu - \frac{\sigma^2}{1-\mu}}{\frac{\sigma^2}{(1-\mu)^2}} \text{ so we have } a = \frac{\mu}{1-\mu} b = \frac{\mu}{1-\mu} \frac{\mu - \frac{\sigma^2}{1-\mu}}{\frac{\sigma^2}{(1-\mu)^2}}$$

g



```
##               Posterior a Posterior b Posterior mean Posterior sd  CI lower
## Smoker result      97.8      348.2      0.2192825    0.01957020 0.1809256
## Nonsmoker result   478.8      986.2      0.3268259    0.01225053 0.3028153
##               CI upper
## Smoker result      0.2576394
## Nonsmoker result   0.3508365
```

Q2

a

By definition,

$$P(T = t) = E(p(1-p)^{t-1}) = E((1-p)^{t-1}) - E((1-p)^{t-1}) + E(p(1-p)^{t-1}) = E((1-p)^{t-1}) - E((1-p)^{t-1} - p(1-p)^{t-1}) = E((1-p)^{t-1}) - E((1-p)^t)$$

Also, from the equation above, $P(T \leq t) = P(T = 1) + P(T = 2) + \dots + P(T = t) = \sum_{i=1}^t E((1-p)^{i-1}) - E((1-p)^i) = 1 - E((1-p)^t)$ and therefore $P(T > t) = E((1-p)^t)$

b

$$P(T = t | \alpha, \beta) = E(p(1-p)^{t-1}) = \int_0^1 p(1-p)^{t-1} \frac{p^{\alpha-1}(1-p)^{\beta-1}}{B(\alpha, \beta)} dp = \frac{1}{B(\alpha, \beta)} \int_0^1 p^\alpha (1-p)^{t+\beta-2} dp = \frac{B(\alpha+1, \beta+t-1)}{B(\alpha, \beta)} = \frac{\alpha \Gamma(\alpha+\beta) \Gamma(\beta+t-1)}{\Gamma(\beta) \Gamma(\beta+\alpha+t)}$$

$$P(T > t | \alpha, \beta) = E((1-p)^t) = \int_0^1 (1-p)^t \frac{p^{\alpha-1}(1-p)^{\beta-1}}{B(\alpha, \beta)} dp = \frac{B(\alpha, \beta+t)}{B(\alpha, \beta)} = \frac{\Gamma(\alpha+\beta)\Gamma(\beta+t)}{\Gamma(\beta)\Gamma(\beta+\alpha+t)}$$

C

Substituting the original parts of the probability in 1(a) for results in 2(a), we switch $L(p) = \{\prod_{i=1}^N [p(1-p)^{t-1}]^{Y_i}\}[(1-p)^N]^{Y_{N+1}}$ to

$$L(\alpha, \beta) = \{\prod_{i=1}^N [\frac{\alpha\Gamma(\alpha+\beta)\Gamma(\beta+t-1)}{\Gamma(\beta)\Gamma(\beta+\alpha+t)}]^{Y_i}\}[\frac{\Gamma(\alpha+\beta)\Gamma(\beta+N)}{\Gamma(\beta)\Gamma(\beta+\alpha+N)}]^{Y_{N+1}}$$

d

Calculating the MLE, we obtain $l(\alpha, \beta) = -Y_{N+1} \log B(\alpha, \beta) + Y_{N+1} \log B(\alpha, \beta + N) + \sum_{t=1}^N Y_t \log B(\alpha + 1, \beta + t - 1) - \sum_{t=1}^N Y_t \log B(\alpha, \beta)$ which is equivalent to $l(\alpha, \beta) = -N \log B(\alpha, \beta) + Y_{N+1} \log B(\alpha, \beta + N) + \sum_{t=1}^N Y_t \log B(\alpha + 1, \beta + t - 1)$

To write in gamma form, we have

$$l(\alpha, \beta) = -n(\log \Gamma(\alpha) + \log \Gamma(\beta) - \log \Gamma(\alpha + \beta)) + Y_{N+1}(\log \Gamma(\alpha) + \log \Gamma(\beta + N) - \log \Gamma(\alpha + \beta + N)) + \sum_{t=1}^N Y_t(\log \Gamma(\alpha + 1) + \log \Gamma(\beta + t - 1) - \log \Gamma(\alpha + \beta + t))$$

where $n = \sum_{t=1}^{N+1} Y_t$

Then we use the optim function to solve for answers.

```
## [1] 2.022436e+00 -5.551115e-17
```

```
## [1] 1.664322e+00 -5.551115e-17
```

```
##          alpha MLE      beta MLE
## Smoker      2.022436 -5.551115e-17
## Nonsmoker    1.664322 -5.551115e-17
```

Q3

a

$L(p) = \{\prod_{i=1}^N [p(1-p)^{t-1}]^{Y_i}\}[(1-p)^N]^{Y_{N+1}} = \prod_{i=1}^N p^{Y_i} (1-p)^{(t-1)Y_i + Y_{N+1}} \propto \prod_{i=1}^N \binom{tY_i + Y_{N+1}}{Y_i} p^{Y_i} (1-p)^{(t-1)Y_i + Y_{N+1}}$ which follows a product of the binomial distribution.

b

The log likelihood is almost the same as calculated in Q1. $l(p) = c + \log p \sum_{t=1}^N Y_t + \log(1-p) \sum_{t=1}^N Y_t(t-1) + N Y_{N+1} \log(1-p)$ where $c = \log \prod_{i=1}^N \binom{tY_i + Y_{N+1}}{Y_i}$. Therefore, after taking derivative wrt p , the result of MLE of p will remain the same. Same logic applies for the fisher information as well as asymptotic variance. Therefore, as discussed in Q1, the stats are as follows.

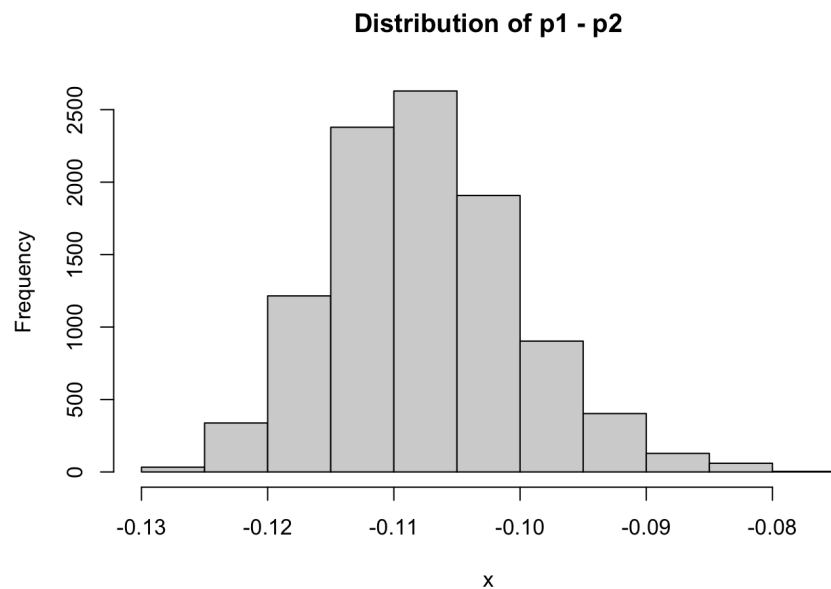
```
##          MLE      MLE sd  CI lower  CI upper
## Smoker result  0.2203791 0.01957709 0.1820088 0.2587495
## Nonsmoker result 0.3289382 0.01148539 0.3064273 0.3514492
```

C

Since the likelihood function of this problem only differs from the likelihood function from Q1, we know that posterior distribution has form

$$P(Y|p) \propto [\prod_{i=1}^N (p(1-p)^{t-1})^{Y_i}]((1-p)^N)^{Y_{N+1}} p^a (1-p)^b = p^{\sum_{i=1}^N Y_i + a} (1-p)^{\sum_{i=1}^N (t-1)Y_i + N Y_{N+1} + b} \sim \text{Beta}(\sum_{i=1}^N Y_i + a, \sum_{i=1}^N (t-1)Y_i + N Y_{N+1} + b).$$

In this case, we have prior $\text{Beta}(1, 1)$ so we plug in $a = 1, b = 1$.



d

```
## [1] 10000
```

The probability is 100%.

e

The predicted probability is as follows:

```
##           [,1]      [,2]      [,3]      [,4]      [,5]      [,6]
## Smokers      0.2900000 0.1600000 0.1700000 0.0400000 0.03000000 0.09000000
## Nonsmokers    0.4074074 0.2201646 0.1131687 0.0781893 0.03703704 0.04526749
##           [,7]      [,8]      [,9]      [,10]     [,11]     [,12]
## Smokers      0.04000000 0.05000000 0.01000000 0.01000000 0.01000000 0.03000000
## Nonsmokers    0.01440329 0.01851852 0.01028807 0.00617284 0.01234568 0.01234568
##           [,13]
## Smokers      0.07000000
## Nonsmokers    0.02469136
```

Appendix

Q1

```
time = seq(1,13,1)
smoker = c(29, 16,17,4,3,9,4,5,1,1,1,3,7)
nonsmoker = c(198,107,55,38,18,22,7,9,5,3,6,6,12)

part1_1 = smoker[c(1:12)]
part2_1 = smoker[c(1:12)]*time[c(1:12)]
part3_1 = length(smoker) * smoker[13]

mle1 = sum(part1_1)/(sum(part2_1) + part3_1)

fin1 = 1/mle1^2 * sum(part1_1) + 1/(1-mle1)^2 * sum(part2_1) + 1/(1-mle1)^2 * part3_1
var1 = 1/fin1
sd1 = sqrt(var1)
ci1 = c(mle1 - sqrt(var1) * qnorm(0.975), mle1 + sqrt(var1) * qnorm(0.975))

part1_2 = nonsmoker[c(1:12)]
part2_2 = nonsmoker[c(1:12)]*time[c(1:12)]
part3_2 = length(nonsmoker) * nonsmoker[13]

mle2 = sum(part1_2)/(sum(part2_2) + part3_2)

fin2 = 1/mle2^2 * sum(part1_2) + 1/(1-mle2)^2 * sum(part2_2) + 1/(1-mle2)^2 * part3_2
var2 = 1/fin2
sd2 = sqrt(var2)
ci2 = c(mle2 - sqrt(var2) * qnorm(0.975), mle2 + sqrt(var2) * qnorm(0.975))

smoker_res = c(mle1, sd1,ci1)
nonsmoker_res = c(mle2, sd2,ci2)

res_ld = rbind(smoker_res, nonsmoker_res)
rownames(res_ld) = c("Smoker result", "Nonsmoker result")
colnames(res_ld) = c("MLE", "MLE sd", "CI lower", "CI upper")
res_ld

mu = 0.2
sigma = 0.08

b = (mu - sigma^2/(1-mu))/(sigma^2/(1-mu)^2)
a = mu/(1-mu) * b

#sanity check
#a/(a+b)
#sqrt(a*b/((a+b)^2*(a+b+1)))

#post dist
post1_a = sum(part1_1) + a
post1_b = sum(part2_1) + part3_1 + b -sum(part1_1)
post2_a = sum(part1_2) + a
post2_b = sum(part2_2) + part3_2 + b -sum(part1_2)

#post mean
post1_mean = (sum(part1_1) + a)/(sum(part2_1) + part3_1 + a + b)
post2_mean = (sum(part1_2) + a)/(sum(part2_2) + part3_2 + a + b)

#post sd
post1_sd = sqrt(post1_a * post1_b/((post1_a+post1_b)^2*(post1_a+post1_b+1)))
post2_sd = sqrt(post2_a * post2_b/((post2_a+post2_b)^2*(post2_a+post2_b+1)))

#ci
ci1 = c(post1_mean - post1_sd * qnorm(0.975), post1_mean + post1_sd * qnorm(0.975))
ci2 = c(post2_mean - post2_sd * qnorm(0.975), post2_mean + post2_sd * qnorm(0.975))

#hist
hist(rbeta(1000, post1_a, post1_b), xlab='x', main = "Distribution of Smoker's Posterior")
plot.new()
hist(rbeta(1000, post2_a, post2_b), xlab='x', main = "Distribution of Non-Smoker's Posterior")

#make table
par(mfrow = c(2, 1))
post_1 = c(post1_a, post1_b, post1_mean, post1_sd, ci1[1], ci1[2])
post_2 = c(post2_a, post2_b, post2_mean, post2_sd, ci2[1], ci2[2])
```

```

res_1 = rbind(post_1, post_2)
colnames(res_1) = c("Posterior a", "Posterior b", "Posterior mean", "Posterior sd", "CI lower", "CI upper")
rownames(res_1) = c("Smoker result", "Nonsmoker result")
res_1

```

Q2

```

getLogL <- function(parameters){
  a = parameters[1]
  b = parameters[2]
  part1 = 0
  for (i in 1:12){
    part1 = part1 + smoker[i] *(lgamma(a+1) + lgamma(b+i -1) - lgamma(a+b+i))
  }
  result = -sum(smoker) * ( lgamma(a) + lgamma(b) - lgamma(a+b) )+
    smoker[13] * (lgamma(a) + lgamma(b+12) - lgamma(a+b+12)) + part1
  return(result)
}

ab1 <- optim(par=c(1,1),fn=getLogL)$par
ab1

getLogL2 <- function(parameters){
  a = parameters[1]
  b = parameters[2]
  part1 = 0
  for (i in 1:12){
    part1 = part1 + nonsmoker[i] *(lgamma(a+1) + lgamma(b+i -1) - lgamma(a+b+i))
  }
  result = -sum(nonsmoker) * ( lgamma(a) + lgamma(b) - lgamma(a+b) )+
    nonsmoker[13] * (lgamma(a) + lgamma(b+12) - lgamma(a+b+12)) + part1
  return(result)
}

ab2 <- optim(par=c(1,1),fn=getLogL2)$par
ab2

res_2d = rbind(ab1, ab2)
colnames(res_2d) = c("alpha MLE", "beta MLE")
rownames(res_2d) = c("Smoker", "Nonsmoker")
res_2d

```

Q3

```

post_a1 = sum(part1_1) + 1
post_b1 = sum(part2_1) + part3_1 + 1 -sum(part1_1)
post_a2 = sum(part1_2) + 1
post_b2 = sum(part2_2) + part3_2 + 1 -sum(part1_2)

post_1_sorted = sort(rbeta(10000, post_a1, post_b1))
post_2_sorted = sort(rbeta(10000, post_a2, post_b2))

hist(post_1_sorted - post_2_sorted, xlab ='x', main = "Distribution of p1 - p2")

set.seed(42)
res_3d1 = rbeta(10000, post_a1, post_b1)
res_3d2 = rbeta(10000, post_a2, post_b2)
sum(res_3d1<res_3d2)

```