STAT 447: Exercise 6

Kevin Liu (94200474)

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Question 1

```
\begin{split} & \mu_e \sim \text{Unif}(0,1) \\ & \lambda_e \sim \text{Exp}(0.01) \\ & \mu_p \sim \text{BetaMP}(0.1,15) \\ & \lambda_p \sim \text{Exp}(0.01) \\ & e_{Pfizer}, e_{Moderna} | \mu_e, \lambda_e \sim \text{BetaMP}(\mu_e, \lambda_e) \\ & p_{Pfizer}, p_{Moderna} | \mu_e, \lambda_e \sim \text{BetaMP}(\mu_p, \lambda_p) \\ & n_c | p \sim \text{Binom}(t_c, p) \\ & n_v | e, p \sim \text{Binom}(t_v, p(1-e)) \end{split}
```

Question 2

```
# this is where your R code goes
source(".././simple.R")

BetaMP = function(mean, precision){
    Beta(mean*precision, (1-mean)*precision)
}

model = function() {
    mu_e = simulate(Unif(0,1))
    lambda_e = simulate(Exp(0.01))

mu_p = simulate(BetaMP(0.1, 15))
    lambda_p = simulate(Exp(0.01))

eff = list()
    inc = list()

eff[["Pfizer-BioNTech"]] = simulate(BetaMP(mu_e, lambda_e))
    inc[["Pfizer-BioNTech"]] = simulate(BetaMP(mu_p, lambda_p))

eff[["Moderna-NIH"]] = simulate(BetaMP(mu_e, lambda_e))
    inc[["Moderna-NIH"]] = simulate(BetaMP(mu_p, lambda_p))
```

```
for (i in seq_along(vaccines)) {
  vaccine = vaccines$trials[i]
  arm = vaccines$arms[i]
  size = vaccines$groupSizes[i]
  cases = vaccines$numbersOfCases[i]

if (arm == "control") {
   observe(cases, Binom(size, inc[[vaccine]]))
  } else {
   observe(cases, Binom(size, inc[[vaccine]] * (1 - eff[[vaccine]])))
  }
}

return(eff[["Moderna-NIH"]] > eff[["Pfizer-BioNTech"]])
}
```

Question 3

```
# this is where your R code goes
vaccines = read.csv("vaccines.csv")
vaccines$groupSizes = as.double(vaccines$groupSizes) # needed due to bug in Binom code
```

Question 4

```
# this is where your R code goes
set.seed(2)
prob = posterior(model, 10000)
print(prob)
```

```
## [1] 0.9854687
```

I am not sure if it is reliable because for the code above, when I set the seed to 1, it would fail for some reason. If this wasn't the case, I would say SNIS is reliable.