```
ISYE 6420 Bayesian Statistics
Final Exam
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1.
a)
To handle the missing data for the chol variable, we will use the suggested distribution, dnorm(200,\ 0.01)
```

Provided is the WinBugs formulation for the problem:

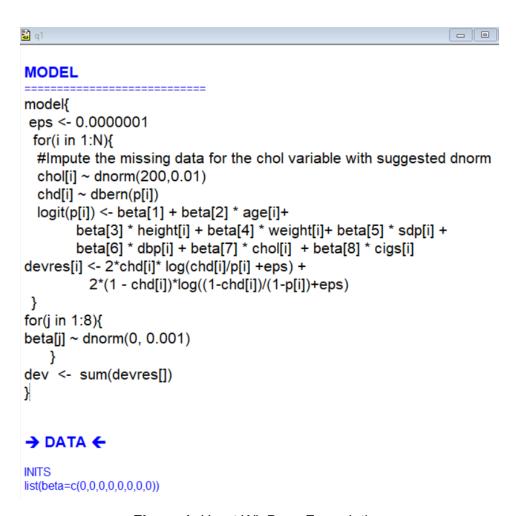


Figure 1: Heart WinBugs Formulation

b)
Provided are the posterior densities for the seven predictors start from the top right downwards, since the top left plot is in relation to b[1] or the intercept:

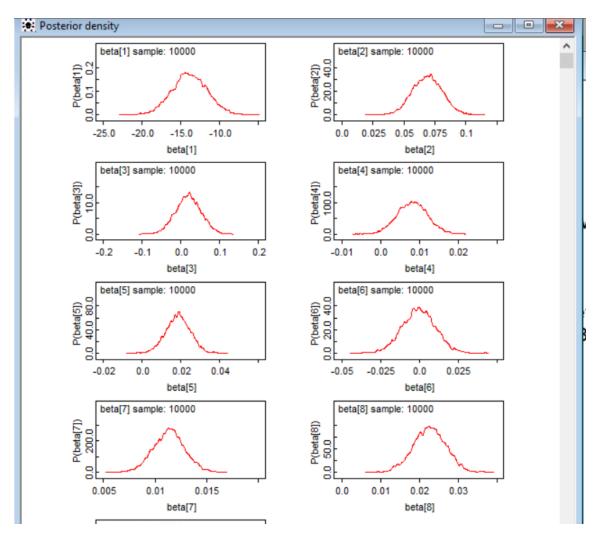


Figure 2: Density Plots Predictor Variables

Provided are the boxplots for these variables:

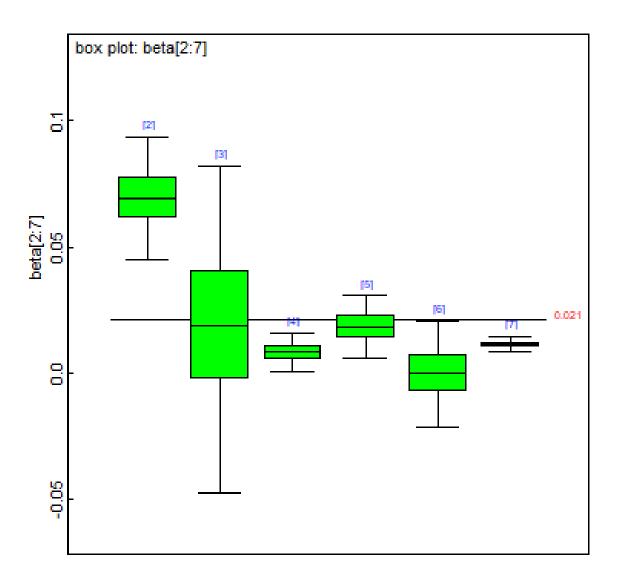


Figure 3: Boxplot Predictor Variables

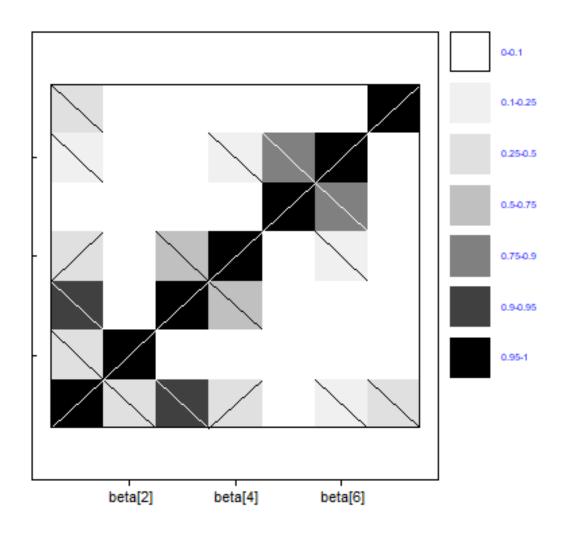


Figure 4: Correlation Plot Predictor Variables

We can also see that there's a bit of correlation exhibited between some of our variables when looking at the matrix plot.

c) Provided is the summary output statistics for the the model:

	mean	sd	MC_error	val2.5pc	median	val97.5pc	start	sample
beta[1]	-13.86	2.29	0.03253	-18.28	-13.88	-9.367	1001	10000
beta[2]	0.0689	0.01197	1.639E-4	0.04553	0.06903	0.0924	1001	10000
beta[3]	0.01914	0.03294	4.747E-4	-0.0448	0.01941	0.08328	1001	10000
beta[4]	0.008208	0.003877	6.049E-5	6.583E-4	0.008201	0.01579	1001	10000
beta[5]	0.01853	0.006324	9.609E-5	0.00602	0.01865	0.0307	1001	10000
beta[6]	-3.034E-4	0.01078	1.723E-4	-0.02146	-2.87E-4	0.02054	1001	10000
beta[7]	0.01136	0.001531	2.256E-5	0.008346	0.01139	0.01438	1001	10000
beta[8]	0.02266	0.004204	5.995E-5	0.01421	0.02267	0.0308	1001	10000
dev	1612.0	4.068	0.06047	1606.0	1611.0	1622.0	1001	10000

Figure 5: Summary Output Heart Model

We can see that our 95% credible sets for each variable are as follows:

```
 age = [0.04553, \, 0.0924] \\ height = [-0.0448, \, 0.08328] \\ weight = [0.0006583 \, (Converted \, from \, scientific \, notation \, 6.583E - 4), \, 0.01579] \\ sdp = [0.00602, \, 0.0307] \\ dbp = [-0.02146, \, 0.02054] \\ chol = [0.008346 \, , \, 0.01438] \\ cigs = [0.01421 \, , \, 0.0308] \\ d) \\ When taking the median values to form a prediction, we're left with the following equation: <math display="block"> chd = -13.88 \, + \, 0.06903*\beta_2 + \, 0.01941*\beta_3 + \, 0.008201*\beta_4 + \, 0.01865*\beta_5 - \, 0.000287*\beta_6 \\ + \, 0.01139*\beta_7 \, + \, 0.02267*\beta_8
```

Therefore, we will make the necessary code changes to get our p value for this prediction, based on the median values for each of the 7 predictors (calculated in excel):

```
OpenBUGS
 File Edit Attributes Tools Info Model Inference Doodle Map Text Window Examples Manuals Help
          for(i in 1:N){
            #Impute the missing data for the chol variable with suggested dnorm
            chol[i] ~ dnorm(200,0.01)
            chd[i] ~ dbern(p[i])
            logit(p[i]) \leftarrow beta[1] + beta[2] * age[i] +
                             beta[3] * height[i] + beta[4] * weight[i]+ beta[5] * sdp[i] +
                               beta[6] * dbp[i] + beta[7] * chol[i] + beta[8] * cigs[i]
      devres[i] \leftarrow 2*chd[i]*log(chd[i]/p[i] + eps) + 2*(1 - chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log((1-chd[i])*log
      chd[i])/(1-p[i])+eps)
       }
      for(j in 1:8){
      beta[j] \sim dnorm(0, 0.001)
      dev <- sum(devres[])
      #Prediction
      median age <- 45
      median_height <- 70
      median_weight <- 170
      median_sdp <- 126
      median_dbp <- 80
      median chol <- 223
      median cigs <- 0
      logit(chd_pred) <- beta[1] + beta[2] * median_age+ beta[3] *
      median_height + beta[4] * median_weight + beta[5] * median_sdp +
      beta[6] * median_dbp + beta[7] * median_chol + beta[8] * median_cigs
      → DATA ←
      INITS
      list(beta=c(0,0,0,0,0,0,0,0))
```

Figure 6: WinBugs Code Prediction Median Values

After running this we get the following result:

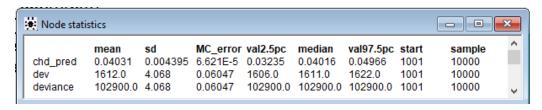


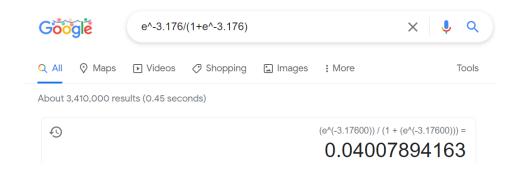
Figure 7: Prediction Result

We can confirm this is correct because when we remove the logit() from the result we get the -3.176. Then when plugging this into the p-value calculation where I is the result from the

prediction:

$$p = \frac{e^l}{\left(1 + e^l\right)}$$

Which results in the following:



Therefore we can be confident that WinBugs is storing the result of the p-value.

Provided is our density plot for the posterior distribution of the prediction:

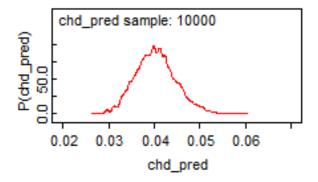


Figure 8: Density Plot chd_pred

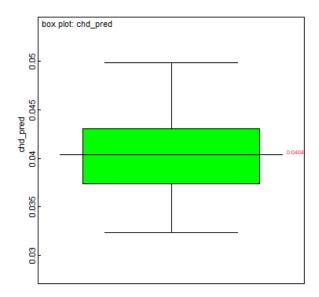


Figure 9: Boxplot chd pred

From a mean value of 0.04031 we can conclude that this person is NOT predicted to have heart disease, assuming the default threshold of 0.5

2.

Pulp brightness. Consider the following experiment, which was performed at a pulp mill. Plant performance is based on pulp brightness as measured by a reflectance meter. Each of the four shift operators (denoted by A, B, C, and D) made five pulp handsheets from unbleached pulp. Reflectance was read for each of the handsheets using a brightness tester

Solve the problem as a Bayesian one-way ANOVA. Use STZ constraints on treatment effects.

1. Do the operators differ in making the pulp handsheets and reading their brightness? Look at the 95% credible sets for the differences between treatment effects.

We'll use the following formulation in OpenBugs to construct this problem: We will map each operator to a given number with the following mapping:

- $A \rightarrow 1$
- $B \rightarrow 2$
- $C \rightarrow 3$
- $D \rightarrow 4$

```
MODEL
model{
for (i in 1:ntotal) {
reflectance[i] ~ dnorm( mu[i], tau )
mu[i] <- mu0 + alpha[operator[i]]</pre>
#alpha[1] <- 0.0;
                          #CR constraints
alpha[1] <- -sum( alpha[2:a] ); #STZ Constraint
mu0 \sim dnorm(0, 0.0001)
alpha[2] \sim dnorm(0, 0.0001)
alpha[3] \sim dnorm(0, 0.0001)
alpha[4] \sim dnorm(0, 0.0001)
tau ~ dgamma(0.001, 0.001)
sigma <- sqrt(1/tau)</pre>
#pairwise
  for(i in 1:3) {
    for(j in i+1:4){
    adiff[i,j] <- alpha[i]-alpha[j]
}
DATA
list(ntotal = 20, a=4,
reflectance =
c(59.8,60,60.8,60.8,59.8,59.8,60.2,60.4,59.9,60,60.7,60.7,60.5,60
.9,60.3,61,60.8,60.6,60.5,60.5),
operator = c(1,1,1,1,1,2,2,2,2,2,3,3,3,3,3,4,4,4,4,4,4)
INITS
list(mu0=0, alpha = c(NA,0,0,0), tau=1)
```

Figure 10: OpenBugs Code Problem 2

When we run our code while burning the first 1000 samples on 100,000 iterations we get the following result:

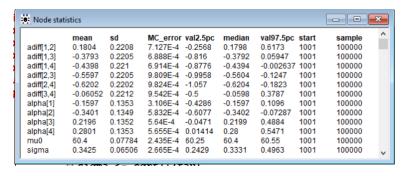


Figure 11: Reflectance Summary Output

When analyzing the output, we can confirm that some of the operators differ in making pulp handsheets as some of the credible sets for our pairwise differences do not contain 0. When we look at the combinations:

A and D seem to differ since the credible set ranges from -0.8776 to -0.002637 Meaning $\alpha_a < \alpha_d$

B and C seem to differ since the credible set ranges from -0.9958 to -0.1247 Meaning $\alpha_b < \alpha_c$

B and D seem to differ since the credible set ranges from -1.057 to -0.1823 Meaning $\alpha_b < \alpha_d$

Since the credible sets are all negative, H_0 being the assumption that the treatment means are equal is rejected.

b) Find the 95% credible set for the contrast $\mu 1 - \mu 2 - \mu 3 + \mu 4$, where $\mu 1$, $\mu 2$, $\mu 3$, and $\mu 4$ are the mean pulp brightness for the operators A, B, C, and D, respectively.

When we reformulate our WinBugs code, we initialize this case to a fixed index of adif f[7,7]. This results in the following code change:

tester. **MODEL** model{ for (i in 1:ntotal) { reflectance[i] ~ dnorm(mu[i], tau) mu[i] <- mu0 + alpha[operator[i]]</pre> #alpha[1] <- 0.0; #CR constraints alpha[1] <- -sum(alpha[2:a]); #STZ Constraint $mu0 \sim dnorm(0, 0.0001)$ alpha[2] ~ dnorm(0, 0.0001) alpha[3] ~ dnorm(0, 0.0001) alpha[4] ~ dnorm(0, 0.0001) tau ~ dgamma(0.001, 0.001) sigma <- sqrt(1/tau) #pairwise for(i in 1:3){ for(j in i+1:4){ adiff[i,j] <- alpha[i]-alpha[j]</pre> adiff[7,7] <- (alpha[1]-alpha[2])-(alpha[3]-alpha[4])

When we run we get the same output but with the additional adiff entry added:

}

	mean	sd	MC_error val2.5pc	median	val97.5pc	start	sample	
adiff[1,2]	0.1804	0.2208	7.127E-4 -0.2568	0.1798	0.6173	1001	100000	
adiff[1,3]	-0.3793	0.2205	6.888E-4 -0.816	-0.3792	0.05947	1001	100000	
adiff[1,4]	-0.4398	0.221	6.914E-4 -0.8776	-0.4394	-0.002637	1001	100000	
adiff[2,3]	-0.5597	0.2205	9.809E-4 -0.9958	-0.5604	-0.1247	1001	100000	
adiff[2,4]	-0.6202	0.2202	9.824E-4 -1.057	-0.6204	-0.1823	1001	100000	
adiff[3,4]	-0.06052	0.2212	9.542E-4 -0.5	-0.0598	0.3787	1001	100000	
adiff[7,7]	0.2409	0.3119	0.001191 -0.3752	0.2413	0.8573	1001	100000	
alpha[1]	-0.1597	0.1353	3.106E-4 -0.4286	-0.1597	0.1096	1001	100000	
alpha[2]	-0.3401	0.1349	5.832E-4 -0.6077	-0.3402	-0.07287	1001	100000	
alpha[3]	0.2196	0.1352	5.64E-4 -0.0471	0.2199	0.4884	1001	100000	
alpha[4]	0.2801	0.1353	5.655E-4 0.01414	0.28	0.5471	1001	100000	
mu0	60.4	0.07784	2.435E-4 60.25	60.4	60.55	1001	100000	
sigma	0.3425	0.06506	2.665E-4 0.2429	0.3331	0.4963	1001	100000	

Figure 12: Summary Output Ctd

Therefore our output for the credible set is [-0.3752, 0.8573]

3.

a)

Provided is the winbugs formulation for the problem:

```
🔁 q3
                                                                            The data in the file enzyme.csv gives the initial rate of reaction of an
enzyme (y) and the substrate concentration (x).
MODEL
model{
for (i in 1:N) {
y[i] ~ dnorm(mu[i],tau)
mu[i] \leftarrow (theta1*x[i]/(theta2+x[i]))+epsilon
theta1 \sim dgamma(0.01, 0.01)
theta2 \sim dgamma(0.01, 0.01)
epsilon ~ dnorm(0,sigma2)
tau ~ dgamma(0.01, 0.01)
sigma2 <- 1/tau
y_pred \leftarrow (theta1*1/(theta2+1))+epsilon
list(N=12,
x=c(0.02,0.02,0.06,0.06,0.11,0.11,0.22,0.22,0.56,0.56,1.1,1.1),
y=c(76,47,97,107,123,139,159,152,191,201,207,200)
INITS
```

Figure 13: WinBugs Formulation Enzymes

This produced the following plots for our densities:

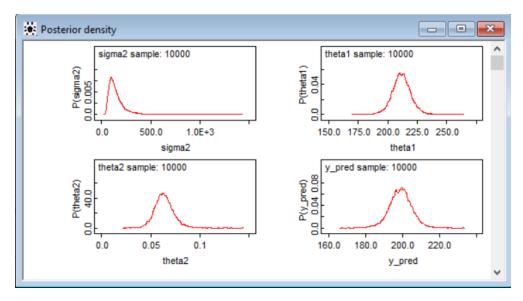


Figure 14: Marginal Densities for each variable

b)
Running this model gives us the following summary output:

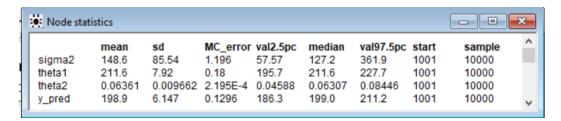


Figure 15: Enzymes Summary Output

Therefore our credible sets for each 3 variables are:

$$sigma2$$
 ($sigma$ $squared$) = [55.57, 361.9]
 θ_1 = [195.7, 227.7]
 θ_2 = [0.04588, 0.08446]

c) When we plot the density for where x = 1, we get the following:

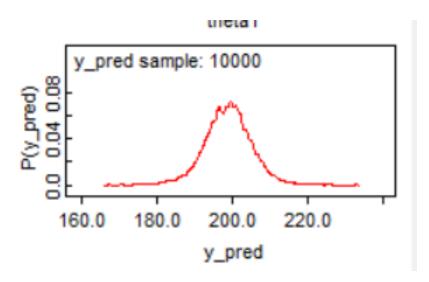


Figure 16: Prediction Density Plot

Which also produces the following credible set:

 $y_{pred} = [186.3, 211.2]$