

**Problem 1:**
 $X \sim \text{Gamma}(a1, b1)$ 
 $Y \sim \text{Gamma}(a2, b2)$ 
 $a1, b1, a2, b2 \sim \text{Gamma}(0.001, 0.001)$ 
 $\text{mean1} \leftarrow (a1/b1)$ 
 $\text{mean2} \leftarrow (a2/b2)$ 
 $\text{diff} \leftarrow (\text{mean2} - \text{mean1})$ 

Posterior distribution for difference in mean blood percentages for the 2 procedures are  $N(2.45, 0.7679)$

95% credible set for the difference is  $[0.9526, 3.978]$ , thus it does not contain 0.

**Blood Volume in Infants**

```
model{
  for (i in 1:n){
    x[i] ~ dgamma(a1, b1)
    y[i] ~ dgamma(a2, b2)
  }

  a1 ~ dgamma(0.001, 0.001)
  a2 ~ dgamma(0.001, 0.001)
  b1 ~ dgamma(0.001, 0.001)
  b2 ~ dgamma(0.001, 0.001)
  mean1 <- (a1/b1)
  mean2 <- (a2/b2)
  diff <- (mean1-mean2)
}
```

**DATA**

```
list(n=16, x=c(13.8, 8.0, 8.4, 8.8, 9.6, 9.8, 8.2, 8.0, 10.3, 8.5, 11.5, 8.2, 8.9, 9.4, 10.3, 12.6),
      y=c(10.4, 13.1, 11.4, 9.0, 11.9, 16.2, 14.0, 8.2, 13.0, 8.8, 14.9, 12.2, 11.2, 13.9, 13.4, 11.9))
```

**INITS**

```
list(a1 = 5, a2 = 5, b1 = 1, b2 = 1 ) # inits here
```

node	mean	sd	MC error	2.5%	median	97.5%	start	sample
a1	32.66	12.26	1.014	13.63	31.18	60.09	1001	10000
a2	27.93	9.164	0.6951	12.85	26.93	48.04	1001	10000
b1	3.384	1.28	0.1058	1.411	3.223	6.216	1001	10000
b2	2.31	0.7663	0.05821	1.038	2.225	4.004	1001	10000
diff	-2.45	0.7679	0.007766	-3.978	-2.447	-0.9526	1001	10000
mean1	9.671	0.454	0.004821	8.814	9.653	10.62	1001	10000
mean2	12.12	0.6187	0.006967	10.97	12.11	13.4	1001	10000

## Problem 2:

Arctic  $\sim \text{Bern}(p[i])$

$\text{Logit}(p) = \text{beta}[1] + \text{beta}[2] * \text{gender} + \text{beta}[3] * x3 + \text{beta}[4] * x7$

$\text{Beta}[j] \sim N(0, 0.01)$

Probability that a female wolf with measures  $x3 = 5.28$  and  $x7 = 1.78$  comes from Arctic habitat is 49.93%

node	mean	sd	MC error	2.5%	median	97.5%	start	sample
beta[1]	0.1047	10.01	0.07112	-19.65	0.09353	19.66	1001	20000
beta[2]	0.04629	10.01	0.06725	-19.56	0.09329	19.62	1001	20000
beta[3]	-0.03984	9.93	0.06369	-19.36	0.005805	19.19	1001	20000
beta[4]	-0.109	9.991	0.07341	-19.81	-0.04389	19.71	1001	20000
pp	0.4993	0.4926	0.004775	0.0	0.4847	1.0	11001	10000
ympred	0.4973	0.5	0.003403	0.0	0.0	1.0	1001	20000

```

model{
  eps <- 0.0000001
  for(i in 1:n){
    arctic[i] ~ dbern(p[i])
    logit(p[i]) <- beta[1] + beta[2] * gender[i] +
      beta[3] * x3[i] + beta[4] * x7[i]
    devres[i] <- 2*arctic[i]*log(arctic[i]/p[i] + eps) +
      2*(1-arctic[i])*log((1-arctic[i])/(1-p[i]) + eps)
  }
  for(j in 1:4){
    beta[j] ~ dnorm(0,0.01)
  }
  dev <- sum(devres[])
  #Prediction
  ympred ~ dbern(pp)
  logit(pp) <- -beta[1] + beta[2] * 1 +
    beta[3] * 5.28 + beta[4] * 1.78
}

```

DATA

list(n=25 )

```

arctic[] gender[] x3[] x7[]
0 0 5.55 2
0 0 5.94 2.07
0 0 5.98 1.94
0 0 5.55 1.9

```

### Problem 3:

$y \sim \text{Poiss}(\text{lambda})$

$\text{lambda} \leftarrow \exp(\text{beta}[1] + \text{beta}[2] * x)$

$\text{beta} \sim N(0, 0.001)$

After 1000 burn-out and 4000 simulations

Beta = [ -2.816, 0.6715]

Average number of nuclei for 3.5G dose is mean = 0.6232

95% credible set =[ 0, 2]

```
model{
  for(i in 1:n){
    y[i] ~dpois(lambda[i])
    lambda[i] <- exp(beta[1] + beta[2]*x[i])
  }
  for(j in 1:2){
    beta[j] ~dnorm(0,0.001)
  }

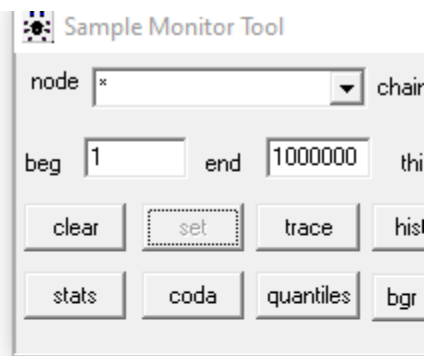
  lambdastar <- exp(beta[1] + beta[2] * 3.5)
  ystar ~dpois(lambdastar)
}
```

**DATA**

**list(n=6000)**

**INITS**

**list(beta =c(1,1))**



Node statistics								
node	mean	sd	MC error	2.5%	median	97.5%	start	sample
beta[1]	-2.816	0.06139	0.002911	-2.942	-2.816	-2.696	1001	5000
beta[2]	0.6715	0.01883	8.842E-4	0.6342	0.672	0.7095	1001	5000
lambdastar	0.6279	0.01579	2.026E-4	0.5974	0.6275	0.6591	1001	5000
ystar	0.6232	0.778	0.01049	0.0	0.0	2.0	1001	5000