

```

{r, message=FALSE, warning=FALSE}
model_unif_cov <- template.jags(
  formula = pa ~ (1 | unit) + (1 | hosp) + #INTERCEPTS
    expe + full + #level 1
    unitsur + we + #level 2
    tech + teach + beds, #level 3

  data = RN4CAST_complete,
  file = "JAGSmodel_unif_cov.txt",
  precision.prior = "dunif(0.001, 100)",
  family = "poisson",
  write.data = FALSE,
  n.chains = 5
)

```

$$\begin{aligned}
 y_i &\sim \text{Poisson}(\lambda) \\
 \mu &\sim \text{Normal}(0, \tau) \\
 \tau &\sim \text{Uniform}(0.001, 100)
 \end{aligned}$$

