**S2 APPENDIX.** R/JAGS code for temporary emigration model

model {

# Priors

 for (k in 1:6){

alpha.om ~ dnorm(2.2,6) #conditional detection probability

 for (i in 1:R){

 alpha.lam[i,k] ~ dnorm(0, 0.01) #for model TE[k], alpha.lam[k]

 alpha.nu[i,k] ~ dunif(-4.6, 4.6) #same as (0.01, 0.99) on prob scale

 }

 }

for (i in 1:R){

 lam.site[i] ~ dnorm(0,tau.site)T(-20,20) #random site effect

}

 tau.nu <- 1 / (sd.nu \* sd.nu) #random availability effect

 sd.nu ~ dunif(0,5)

 tau.site<-1/(sd.site\*sd.site)

 sd.site~dunif(0,5)

 beta.aspect ~ dunif(-3,3)

 beta.slope ~ dunif(-3,3)

beta.aw ~ dunif(-3,3)

beta.TSI ~ dunif(-3,3)

beta.LFI ~ dunif(-3,3)

 beta.nu.r ~ dunif(-3,3) #rain

 beta.nu.ti ~ dunif(-3,3) #time of day

 beta.nu.te ~ dunif(-3,3) #soil temperature

beta.nu.t2 ~ dunif(-3,3) #(time of day)^2

 beta.om.rk ~ dunif(-3,3) #rocks

 beta.om.w ~ dunif(-3,3) #woody cover

 beta.om.ll ~ dunif(-3,3) #leaf litter

# State process model for abundance

 for (i in 1:R){ # Loop over R sites (40)

 for (k in 1:6){ # Loop over k seasons (6)

log(lambda[i,k]) <- alpha.lam[i,k] + beta.aspect\*aspect[i] + beta.slope\*slope[i] + beta.aw\*aw[i] + beta.TSI\*TSI[i] + beta.LFI\*LFI[i] + beta.lfm\*lfm[i] + lam.site[i]

 N[i,k] ~ dpois(lambda[i,k])

# Observation model

 for (j in 1:Tr[k]){ # Loop over surveys (3-5)

 y[i,j,k] ~ dbin(p[i,j,k], N[i,k])

 p[i,j,k] <- (nu[i,j,k])\*(om[i,j,k])

#availability

 nu[i,j,k] <- exp(logit.nu[i,j,k])/(1+exp(logit.nu[i,j,k]))

 logit.nu[i,j,k] ~ dnorm(mu.ln[i,j,k], tau.nu)

 mu.ln[i,j,k] <- alpha.nu[i,k] + beta.nu.r\*rain[i,j,k] + beta.nu.ti\*time[i,j,k] + beta.nu.te\*temp[i,j,k] + beta.nu.t2\*time2[i,j,k]

#conditional det prob

 om[i,j,k] <- exp(logit.om[i,j,k])/(1+exp(logit.om[i,j,k]))

 logit.om[i,j,k] <- alpha.om[k] + beta.om.rk\*rocks[i,j,k] + beta.om.ll\*litter[i,j,k] + beta.om.w\*wco[i,j,k]

# Assess model fit adequacy using Chi-squared discrepancy (for Bayesian P)

# Compute fit statistic FIT for observed data

 expect[i,j,k] <- p[i,j,k]\*N[i,k] # Expected values

 FIT[i,j,k] <- pow((y[i,j,k] - expect[i,j,k]),2)/(expect[i,j,k] + 0.5)

# Generate replicate data and compute fit stats for them

 y.new[i,j,k] ~ dbin(p[i,j,k], N[i,k])

 FIT.new[i,j,k] <- pow((y.new[i,j,k] - expect[i,j,k]),2)/(expect[i,j,k] + 0.5)

 } #j

 } #i

 } #k

# Derived and other quantities

 for (k in 1:6){

 totalN[k] <- sum(N[,k]) # Total pop. size across all sites

 mean.abundance[k] <- mean(lambda[,k])

 }

 fit <- (sum(FIT[ ,1:5,1:4])) + (sum(FIT[ ,1:3,5])) + (sum(FIT[ ,1:4,6]))

 fit.new <- (sum(FIT.new[ ,1:5,1:4])) + (sum(FIT.new[ ,1:3,5])) + (sum(FIT.new[ ,1:4,6])

 }