

Chapter 12

WAIC balances how well a model predicts, with flexibility of the model

- How well does a model predict the data?

$$\begin{aligned}\sum_{j=1}^n \log(p(y_j|y_1, \dots, y_n)) &= \sum_{j=1}^n \log(\mathbb{E}[p(y_j|\alpha, \beta, \sigma, y_1, \dots, y_n)]) \\ &= \sum_{j=1}^n \log \left(\int p(y_j|\tilde{\alpha}, \tilde{\beta}, \tilde{\sigma}) \cdot \right. \\ &\quad \left. \cdot p(\tilde{\alpha}, \tilde{\beta}, \tilde{\sigma}|y_1, \dots, y_n) d\tilde{\alpha} d\tilde{\beta} d\tilde{\sigma} \right)\end{aligned}$$

- How flexible is the model?

$$p_{WAIC} = \sum_{j=1}^n \mathbb{V}[\log(p(y_j|\alpha, \beta, \sigma, y_1, \dots, y_n))]$$

How to get WAIF in STAN, use generate quantities to get log likelihood

```
data{
  ...
}
parameters{
  ...
}
model{
  ...
}
generated quantities {
  vector[N_obs] log_lik;
  for (i in 1 : N_obs)
    log_lik[i] = binomial_logit_lpmf(y[i] | n[i],
      alpha + beta_P * P[i] + beta_V * V[i] + beta_A * A[i] + beta_PA * PA[i]);
}
```

Then in R:

```
colVars <- function(a) {  
  #helpful function exists in resample I believe  
  n <- dim(a)[[1]]  
  c <- dim(a)[[2]]  
  return(.colMeans(((a - matrix(.colMeans(a, n, c), nrow = n, ncol = c, byrow = TRUE))^2),  
    n, c) * n/(n - 1))  
}  
list_param1 <- extract(stan_model1)  
log_lik      <- list_param1.1$log_lik #collect the log lik created in generated quantities  
lpd          <- log_sum_exp(log_lik) # prediction part  
p_waic       <- sum(colVars(log_lik)) # flexiablity part  
WAIC         <- -2 * lpd + 2 * p_waic.
```

Generate quantities can be used to much more:

```
data{
  ...
}
parameters{
  ...
}
model{
  ...
}
generated quantities {
  vector[N_obs] predict_y;
  for (i in 1 : N_obs)
    predict_y[i] = exp( alpha + beta * x[i]);
}
```

Several equivalent names:

- Multilevel models
- Hierarchical models
- Random effect models

Data survival of hatching of reed frogs.



© L. Mahler and B. Zinkus

Figure: African reed frog, *Hyperolius spinigularis*

Ecology, 86(6), 2005, pp. 1580–1591
© 2005 by the Ecological Society of America

COMPENSATORY LARVAL RESPONSES SHIFT TRADE-OFFS ASSOCIATED WITH PREDATOR-INDUCED HATCHING PLASTICITY

JAMES R. VONESH¹ AND BENJAMIN M. BOLKER

Department of Zoology, University of Florida, 223 Bartram Hall, Gainesville, Florida 32611 USA

Abstract. Many species with complex life histories can respond to risk by adaptively altering the timing of key life history switch points, including hatching. It is generally thought that such hatching plasticity involves a trade-off between embryonic and hatchling predation risk, e.g., hatching early to escape egg predation comes at the cost of increased vulnerability to hatchling predators. However, most empirical work has focused on simply detecting predator-induced hatching responses or on the short-term consequences of hatching plasticity. Short-term studies may not allow sufficient time for hatchlings to exhibit compensatory responses, which may extend to subsequent life stages and could alter the nature of the trade-offs associated with hatching plasticity. To address this issue, we conducted a

Figure: Article

THEY ARE ALL THE SAME!

Frogs hatched in different tanks (using same probability):

$$\begin{aligned}y_i &\sim \text{Bin}(n_i, p_i), \\g(p_i) &= \alpha, \\ \alpha &\sim N(0, 10).\end{aligned}$$

Hierarchical model DAG

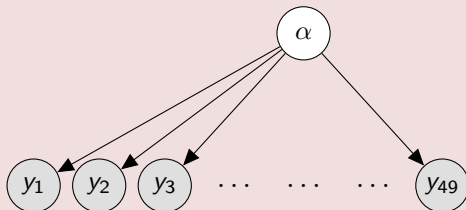


Figure: A DAG (directed acyclic graph) describing the model.

STAN simple

```
data{
  int<lower=0> N;
  int<lower=0> y[N];
  int<lower=1> n[N];
}
parameters{
  real alpha;
}
model{
  alpha ~ normal(0, 10);
  y ~ binomial_logit(n, alpha);
}
generated quantities {
  real predict_p;
  predict_p = exp(alpha)/(1 + exp(alpha));
}
```

THEY ARE ALL UNIQUE

Frogs hatched in different tanks:

$$\begin{aligned}y_i &\sim \text{Bin}(n_i, p_i), \\g(p_i) &= \alpha_i, \\ \alpha_i &\sim N(0, 10).\end{aligned}$$

STAN simple

```
data{
  int<lower=0> N;
  int<lower=0> y[N];
  int<lower=1> n[N];
  int<lower=1> ntank;
  int<lower=0> tank[N];
}
parameters{
  real a[ntank];
}
model{
  vector[N] mu;
  for(i in 1:N)
    mu[i] = a[tank[i]];

  a ~ normal(0, 10);
  y ~ binomial_logit(n, mu);
}
generated quantities {
  real predict_p[ntank];
  for(i in 1:ntank)
    predict_p[i] = exp( a[i])/(1 + exp(a[i]));
}
```

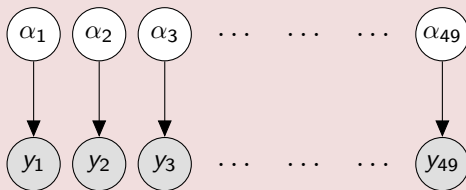


Figure: A DAG (directed acyclic graph) describing the model.

Simple binomial model

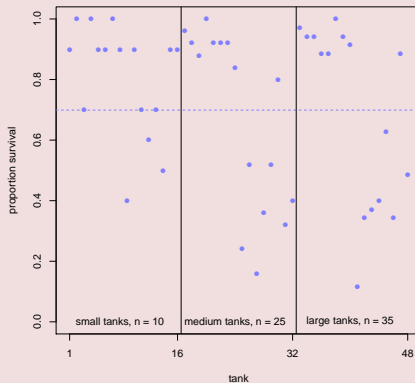
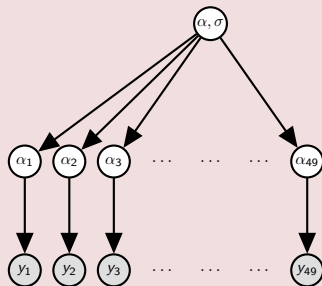


Figure: Posterior mean for each tank

Multilevel model:

$$\begin{aligned}y_i &\sim \text{Bin}(n_i, p_i), \\g(p_i) &= \alpha_i, \\ \alpha_i &\sim N(\alpha, \sigma), \\ \alpha &\sim N(0, 10), \\ \sigma &\sim \text{HC}(0, 1).\end{aligned}$$

Hierarchical model DAG




```
data{
  int<lower=0> N;
  int<lower=0> y[N];
  int<lower=1> n[N];
  int<lower=1> ntank;
  int<lower=0> tank[N];
}
parameters{
  real a[ntank];
  real alpha;
  real<lower=0> sigma;
}
model{
  vector[N] mu;
  for(i in 1:N)
    mu[i] = a[tank[i]];

  sigma ~ cauchy(0,1);
  alpha ~ normal(0, 20);
  a ~ normal(alpha, sigma);
  y ~ binomial_logit(n, mu);
}
generated quantities {
  real predict_p[ntank];
  for(i in 1:ntank)
    predict_p[i] = exp( a[i])/(1 + exp(a[i]));
}
```

Multilevel binomial model

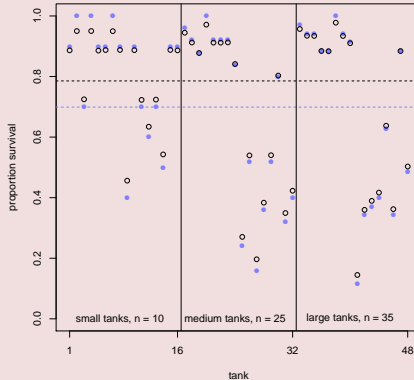


Figure: Posterior mean for each tank

Lets examine, $y_2 = 10, n_2 = 10$

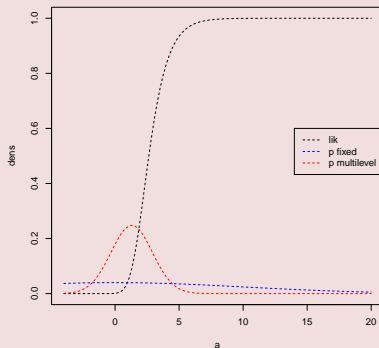


Figure: Prior + likelihood for a .

Lets examine, $y_2 = 10, n_2 = 10$

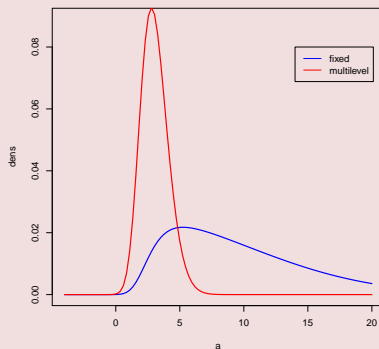
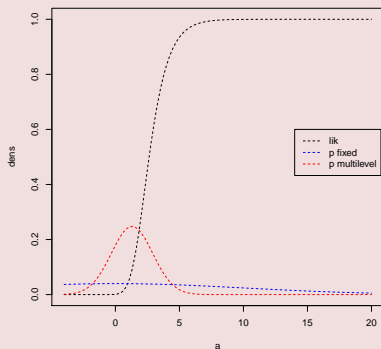


Figure: Posterior for a .

Lets examine, $y_2 = 10, n_2 = 10$

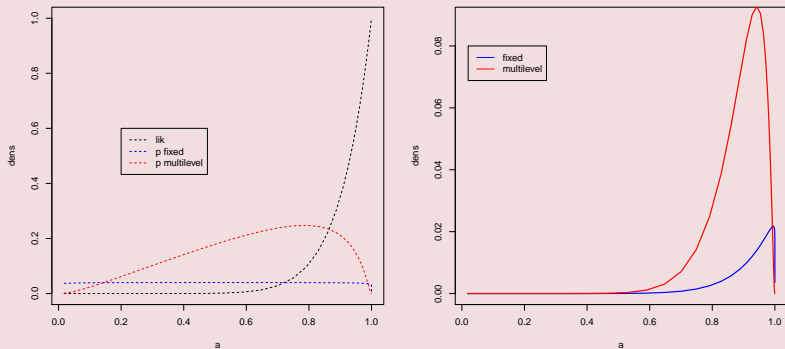
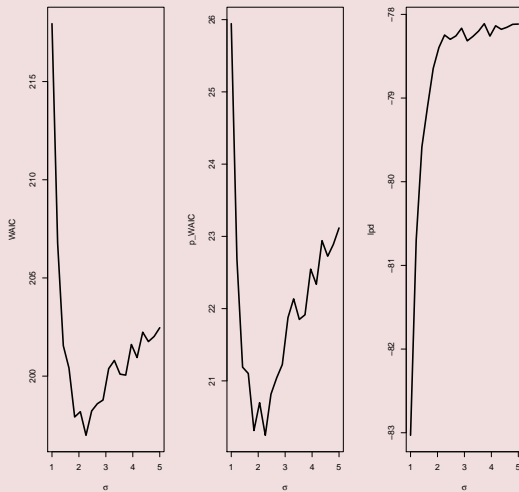
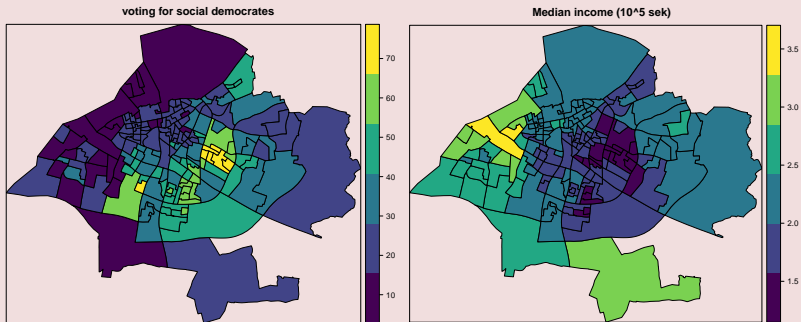


Figure: Figure on the left prior + likelihood for p . Figure on the right posterior p .

WAIC for varying σ



Voting in Malmö, data



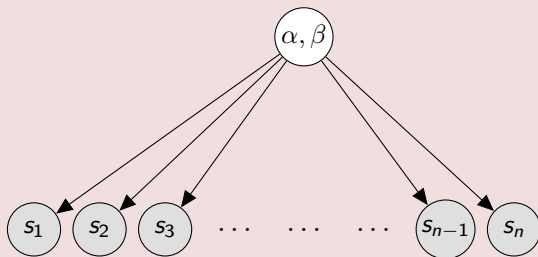
THEY ARE ALL THE UNIQUE!

$$\begin{aligned}s_i &\sim \text{bin}(n_i, p_i), \\ g(p_i) &= \alpha_i + \text{med}_i \beta_i, \\ \alpha_i &\sim N(0, 10) \\ \beta_i &\sim N(0, 10)\end{aligned}$$

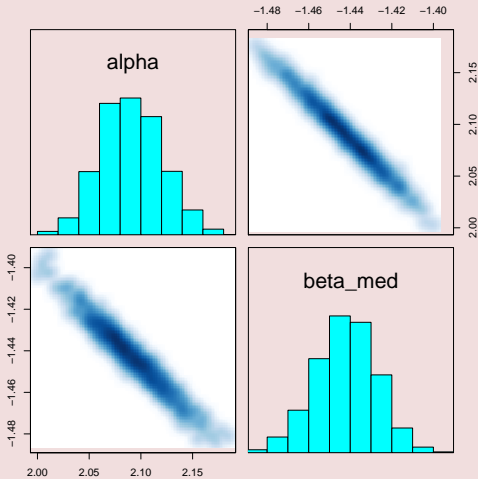
THEY ARE ALL THE SAME!

$$\begin{aligned}s_i &\sim \text{bin}(n_i, p_i), \\ g(p_i) &= \alpha + \text{med}_i \beta, \\ \alpha &\sim N(0, 10) \\ \beta &\sim N(0, 10)\end{aligned}$$

Independent model DAG



Posterior parameter



- By the model the prediction given the data is

$$\hat{Y}_i \sim \text{Bin}(n_i, p_i),$$

$$p_i \sim p(\cdot | y_1, y_2, \dots, y_n)$$

- By the model the prediction given the data is

$$\hat{Y}_i \sim \text{Bin}(n_i, p_i),$$
$$p_i \sim p(\cdot | y_1, y_2, \dots, y_n)$$

- The variance is:

$$V[\hat{Y} | p_i, n_i] = n_i(1 - p_i)p_i$$
$$V\left[\frac{\hat{Y}}{n_i} | p_i, n_i\right] = \frac{(1 - p_i)p_i}{n_i}$$

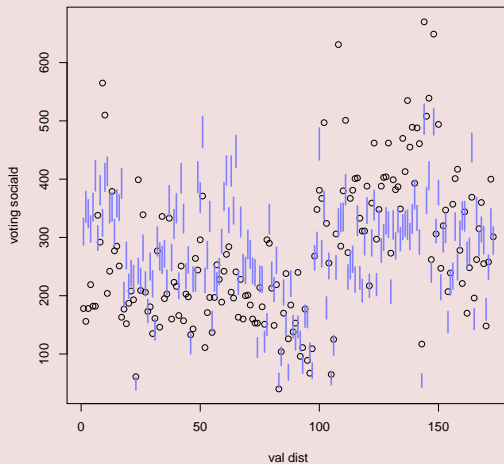


Figure: Prediction by district

- Both binomial and Poisson has only one parameter.
- These models are extremely sensitivity to incorrect parameter.
- They can not adjust it variance to the data.

- This is typically solved by overdispersion model. Like Beta-binomial.

- This is typically solved by overdispersion model. Like Beta-binomial.
- For each observation one adds a random non-negative parameter:

$$p(y_i | n_i) = \int \text{Bin}(y_i | n_i, p_i) h(p_i | p, \theta) p(p, \theta) dp_i dp d\theta,$$

Then one puts covariates on p not p_i .

- This is typically solved by overdispersion model. Like Beta-binomial.
- For each observation one adds a random non-negative parameter:

$$p(y_i | n_i) = \int \text{Bin}(y_i | n_i, p_i) h(p_i | p, \theta) p(p, \theta) dp_i dp d\theta,$$

Then one puts covariates on p not p_i .

- overdispersion is typically a multilevel model.

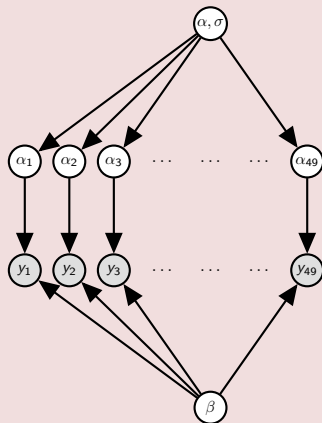
$$\begin{aligned}y_i &\sim \text{Bin}(n_i, p_i) \\g(p_i) &\sim \alpha_0 + \text{med}_i\beta + Z_i \\Z_i &\sim N(0, \sigma) \\\alpha_0 &\sim N(0, 10) \\\sigma &\sim \text{HC}(0, 5).\end{aligned}$$

$$\begin{aligned}
 y_i &\sim \text{Bin}(n_i, p_i) \\
 g(p_i) &\sim \alpha_0 + \text{med}_i \beta + Z_i \\
 Z_i &\sim N(0, \sigma) \\
 \alpha_0 &\sim N(0, 10) \\
 \sigma &\sim \text{HC}(0, 5).
 \end{aligned}$$

or equivalently

$$\begin{aligned}
 y_i &\sim \text{Bin}(n_i, p_i) \\
 g(p_i) &\sim \alpha_i + \text{med}_i \beta \\
 \alpha_i &\sim N(\alpha_0, \sigma) \\
 \alpha_0 &\sim N(0, 10) \\
 \sigma &\sim \text{HC}(0, 5).
 \end{aligned}$$

Hierarchical model DAG



```
data {  
  int<lower = 1> N;  
  int<lower = 1> voters[N];  
  int<lower = 0> soc[N];  
  vector[N] med;  
}  
  
parameters{  
  real alphas[N];  
  real alpha0;  
  real<lower=0> sigma;  
  real beta_med;  
}  
  
model{  
  vector[N] mu;  
  sigma ~ cauchy(0, 1);  
  alpha0 ~ normal(0, 10);  
  beta_med ~ normal(0, 10);  
  alphas ~ normal(alpha0, sigma);  
  for(i in 1:N)  
    mu[i] = alphas[i] + med[i] * beta_med;  
  soc ~ binomial_logit(voters, mu);  
}
```

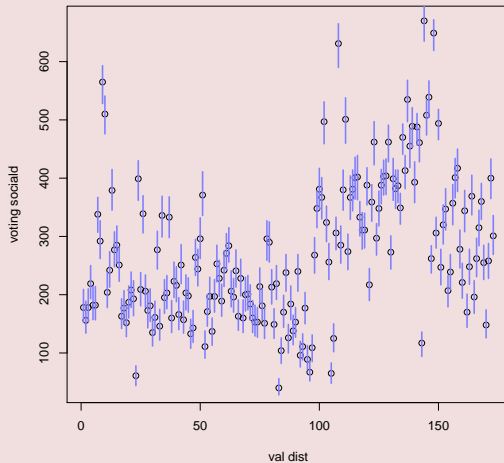


Figure: Prediction by district multilevel

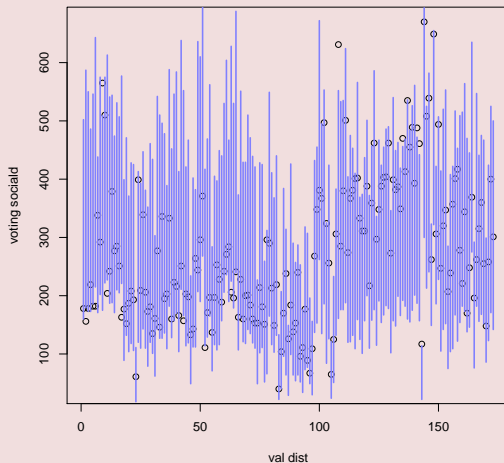


Figure: Prediction unconditional by district multilevel

- The variance is:

$$V[\hat{Y}|n_i] \approx n_i(1 - \hat{p}_i)\hat{p}_i + n_i^2\tilde{\sigma}$$
$$V[\frac{\hat{Y}}{n_i}|n_i] \approx \frac{(1 - \hat{p}_i)\hat{p}_i}{n_i} + \tilde{\sigma}$$

Where $\tilde{\sigma}$ is the variation from

PI for multilevel without cheating

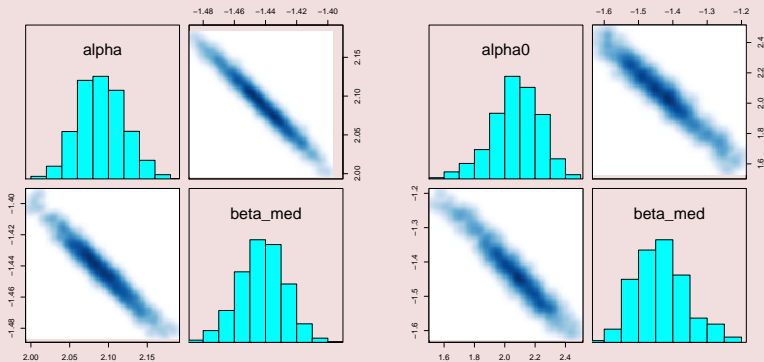


Figure: Look at parameter certainty

Multilevel binomial model

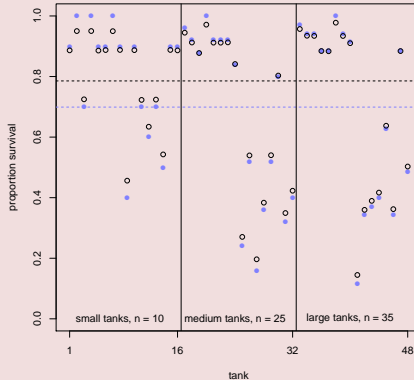


Figure: Posterior mean for each tank

Histogram of the α_i

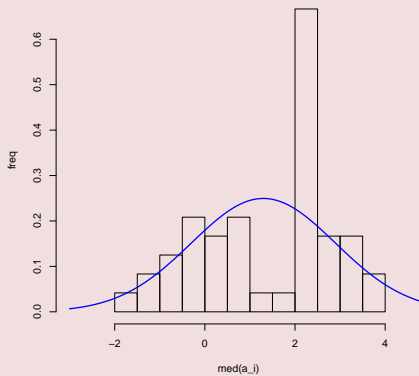


Figure: Histogram of the α_i

Multilevel mixture model:

$$y_i \sim \text{Bin}(n_i, p_i),$$

$$g(p_i) = \alpha_i,$$

$$\alpha_i \sim \theta N(\mu_1, \sigma_1) + (1 - \theta) N(\mu_2, \sigma_2),$$

$$\mu_1 \sim N(0, 10),$$

$$\mu_2 \sim N(0, 10),$$

$$\sigma_1 \sim \text{HC}(0, 1).$$

$$\sigma_2 \sim \text{HC}(0, 1).$$

$$\theta \sim B(2, 2).$$

```
parameters{
  ordered[2] a0;
  real a[ntank];
  real<lower=0> sigma_0;
  real<lower=0> sigma_1;
  real<lower=0, upper=1> theta;
}
model{
  vector[N] mu;
  theta ~ beta(2, 2);
  a0 ~ normal(0,10);
  sigma_0 ~ cauchy(0,1);
  sigma_1 ~ cauchy(0,1);
  for(i in 1:ntank){
    target += log_sum_exp(
      bernoulli_lpmf(1|theta) + normal_lpdf(a[i]| a0[1],sigma_0),
      bernoulli_lpmf(0|theta) + normal_lpdf(a[i]| a0[2],sigma_1)
    );
  }
  for(i in 1:N)
    mu[i] = a[tank[i]];

  y ~ binomial_logit(n, mu);
}
```

Multilevel mixture binomial model

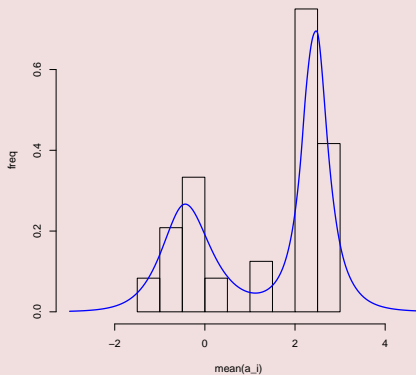


Figure: Histogram of the α_i

Multilevel mixture binomial model

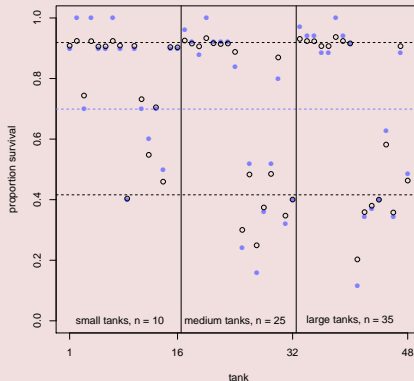


Figure: Posterior mean for each tank