

# Bayesian Skew-Normal and Skew-t Models of Birth Weight and Food Security

Carter Allen<sup>1</sup>; Brian Neelon, PhD<sup>1</sup>; Sara E. Benjamin Neelon, PhD, JD, MPH<sup>2</sup>

<sup>1</sup>Department of Public Health Sciences, Medical University of South Carolina; <sup>2</sup>Bloomberg School of Public Health, Johns Hopkins University

## Objectives

We examine the properties of skew-normal and skew-t models from both a Bayesian and frequentist perspective, and investigate the computational tools available for fitting these models. We apply skew-normal and skew-t models to data from the Nurture study, a cohort of mothers who gave birth between 2013 and 2016, where we seek to model the effect of food security during pregnancy on birth weight.

## Introduction

In many applications of classical linear regression, the distribution of residuals exhibits non-normal qualities such as skewness or heavy tails, making the assumption of normal error terms difficult to justify. The common statistical suggestion in these cases is to implement a transformation of the response variable, but this can result in a loss of interpretability. The skew-elliptical family is a broad class of probability distributions that contain the normal distribution as a special case and allow for flexible modeling when data exhibit skewness.

## Definitions

Let  $\phi$  and  $\Phi$  be the standard normal pdf and cdf, respectively. Azzalini (1985) defined the density of a skew-normal random variable  $Z$  follows.

$$f(z; \lambda) = 2\phi(z)\Phi(\lambda z)$$

Similar to the construction of the familiar student's  $t$  random variable, we (cite) can define a skew-t random variable as the ratio of a skew normal and the square root of a  $\chi^2$  divided by its degrees of freedom. The resultant density is

$$t(x; \lambda, \nu) = 2t_0(x; \nu)T_0(\lambda x \sqrt{\frac{\nu+1}{\nu+x^2}}; \nu+1)$$

where  $t_0$  and  $T_0$  are the density and mass functions of the student's  $t$  distribution, respectively. A linear regression model with skew error terms is a modification of classical regression with the modification of either assuming  $\mathcal{SN}$  or  $\mathcal{ST}$  random errors.

## Motivation

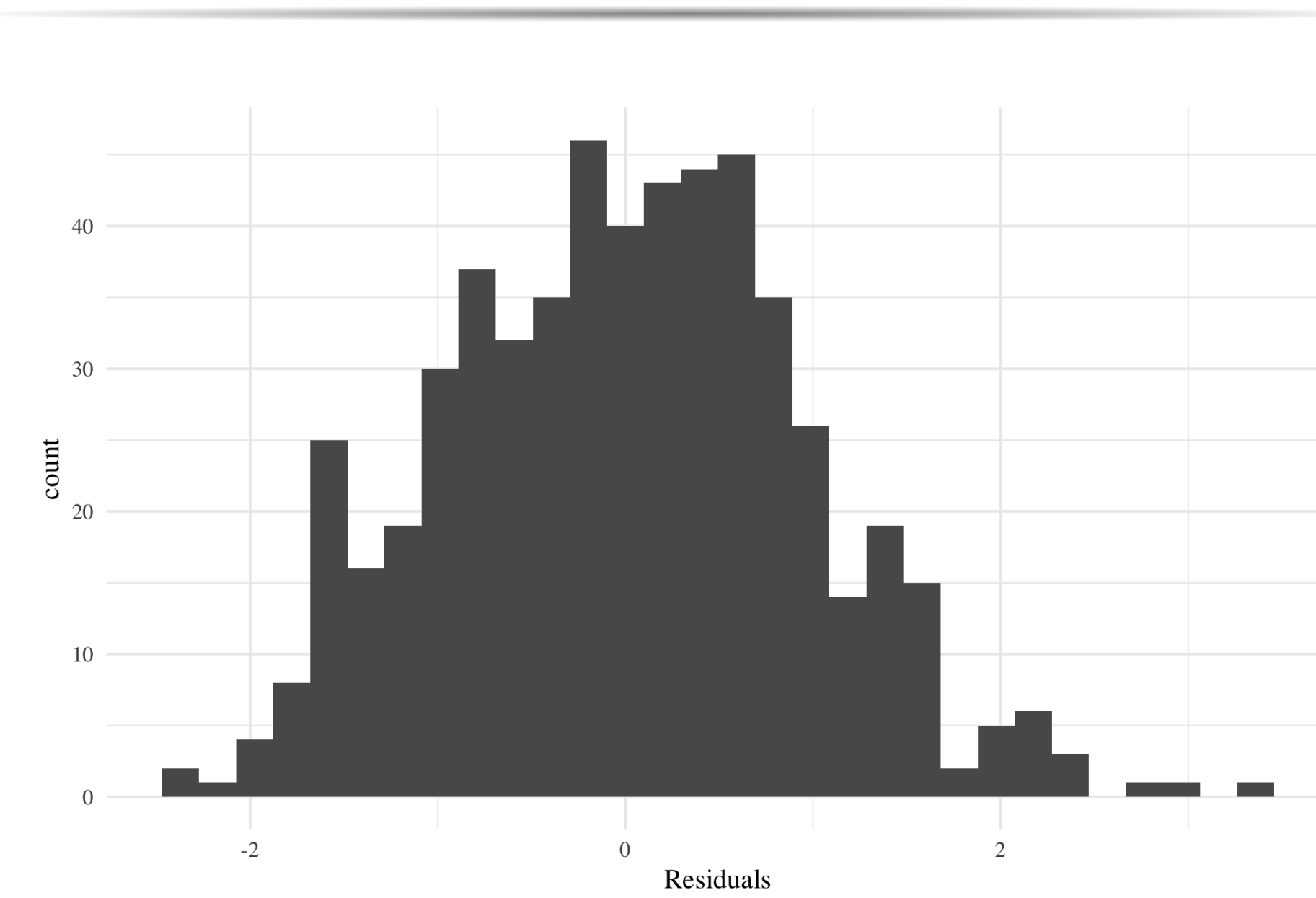


Figure 1: Distribution of residuals exhibiting skewness. Pearson's skewness = 0.17. Shapiro-Wilk p-value = 0.08.

## Full Conditionals

Let  $Y_i = \mathbf{x}_i\beta + \psi z_i + \sigma\epsilon$  with  $Z \sim N_+(0, 1)$ , and  $\epsilon \sim N(0, 1)$ . Define  $\mathbf{X}^* = [\mathbf{X}|\mathbf{z}]$ , and  $\beta^* = [\beta_0, \beta_1, \dots, \beta_p, \psi]$ . Then,

$$\beta^*|Y, \mathbf{X}^*, \tau \sim N_{p+2}\left(\frac{(T_0\beta_0 + \tau\mathbf{X}^{*T}Y)}{\tau\mathbf{X}^{*T}\mathbf{X}^* + T_0}, \tau\mathbf{X}^{*T}\mathbf{X}^* + T_0\right)$$

$$\tau|\beta^*, \mathbf{X}^*, Y \sim \Gamma(n/2 + \alpha, \frac{1}{2}(Y - \mathbf{X}^*\beta^*)^T(Y - \mathbf{X}^*\beta^*) + b)$$

$$z_i|y_i, \mathbf{x}_i, \beta, \tau \sim N_+\left(\frac{\psi\tau(y_i - \mathbf{x}_i\beta)}{\tau\psi^2 + 1}, \frac{1}{\tau\psi^2 + 1}\right)$$

## Important Results

Through simulation studies, we validated Bayesian Gibbs samplers for skew-normal and skew-t error regression models. We used these models to find an association between birth weight and food security during pregnancy using data from the Nurture study.

## Modeling Approaches

- Maximum Likelihood Estimation:** For a random sample  $Y_1, Y_2, \dots, Y_n \stackrel{iid}{\sim} \mathcal{SN}(\xi, \omega^2, \alpha)$ , where  $\xi = \mathbf{x}^T\beta$  for a collection of predictors  $x_1, x_2, \dots, x_n$  and a vector of unknown regression coefficients  $\beta \in \mathbb{R}^p$ . Azzalini (2014) describes procedures for obtaining MLE estimates of  $\beta$  and  $\alpha$ , our primary parameters of interest. Azzalini's R package **sn** contains a function **selm** for fitting regression models with  $\mathcal{SN}$  or  $\mathcal{ST}$  random errors.
- Bayesian Gibbs Sampler:** We introduce the following stochastic representation of the skew normal distribution

$$Y_i = \mathbf{x}_i\beta + \psi z_i + \sigma\epsilon$$

where  $z_i \sim N_+(0, 1)$  and  $\epsilon \sim N(0, 1)$ . The marginal density of  $Y_i$  integrating over  $z_i$  and  $\epsilon$  is  $\mathcal{SN}(\mathbf{x}_i\beta, \omega^2)$ . We use this stochastic representation to obtain full conditionals for all parameters.

## Gibbs Sampler Simulation

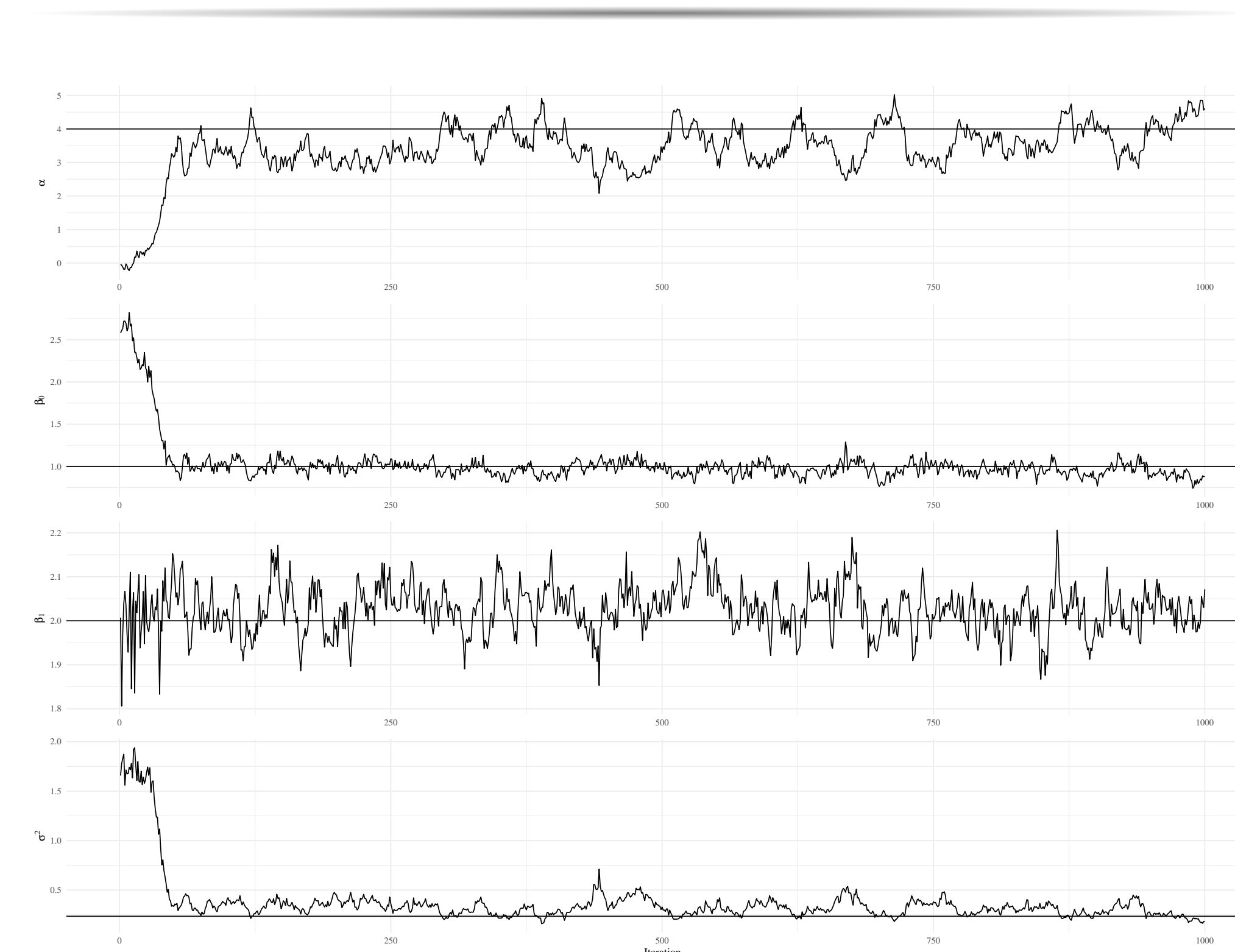


Figure 2: Trace plots for skew normal Gibbs sampler simulation

Parameter	True	MLE	Gibbs
$\alpha$	4.00	0.3928	4.016
$\beta_0$	1.00	1.013	1.009
$\beta_1$	2.00	2.006	2.01
$\sigma^2$	0.235	0.257	0.257

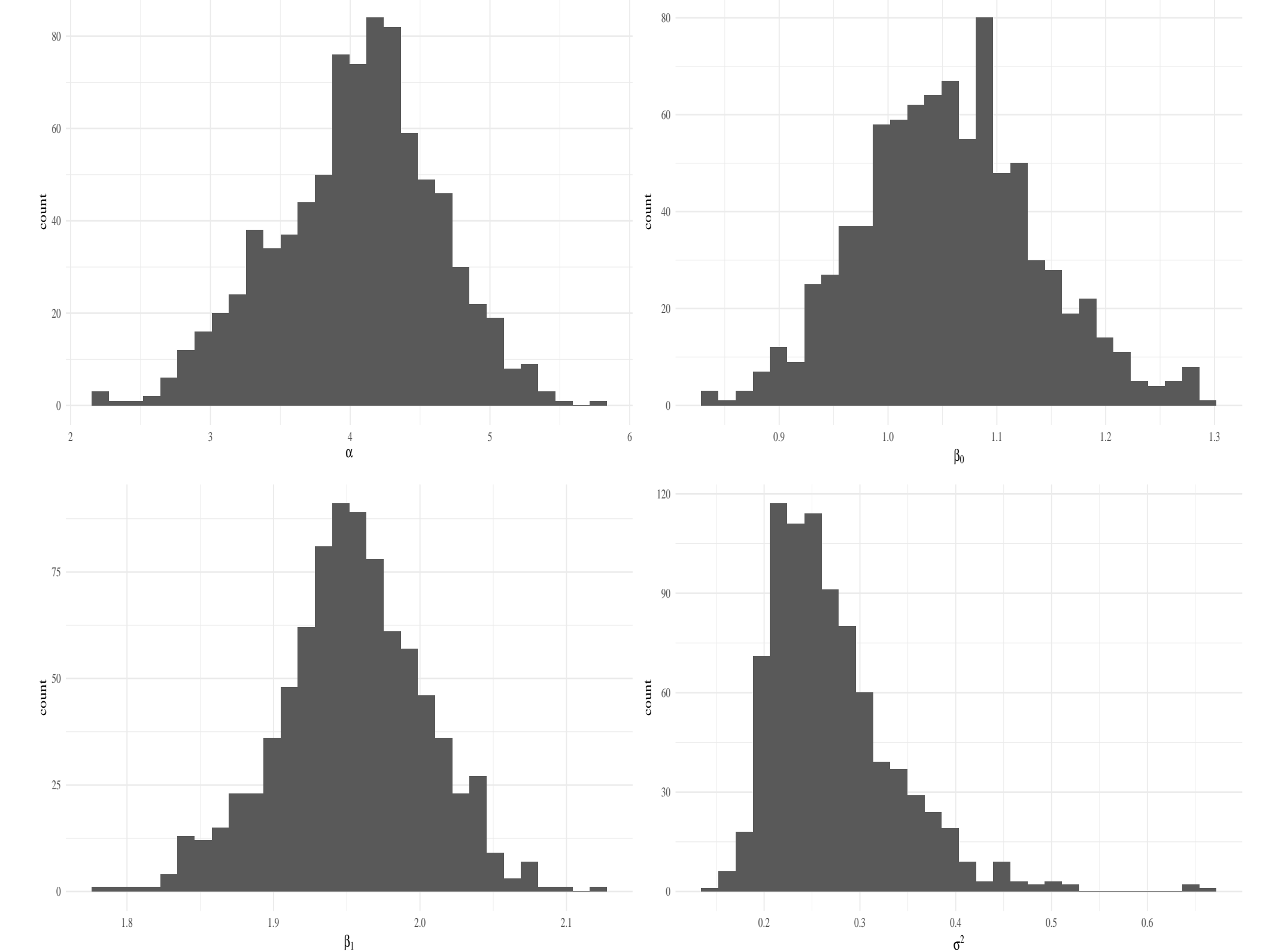


Figure 3: Simulated posterior distributions

## Modeling Results

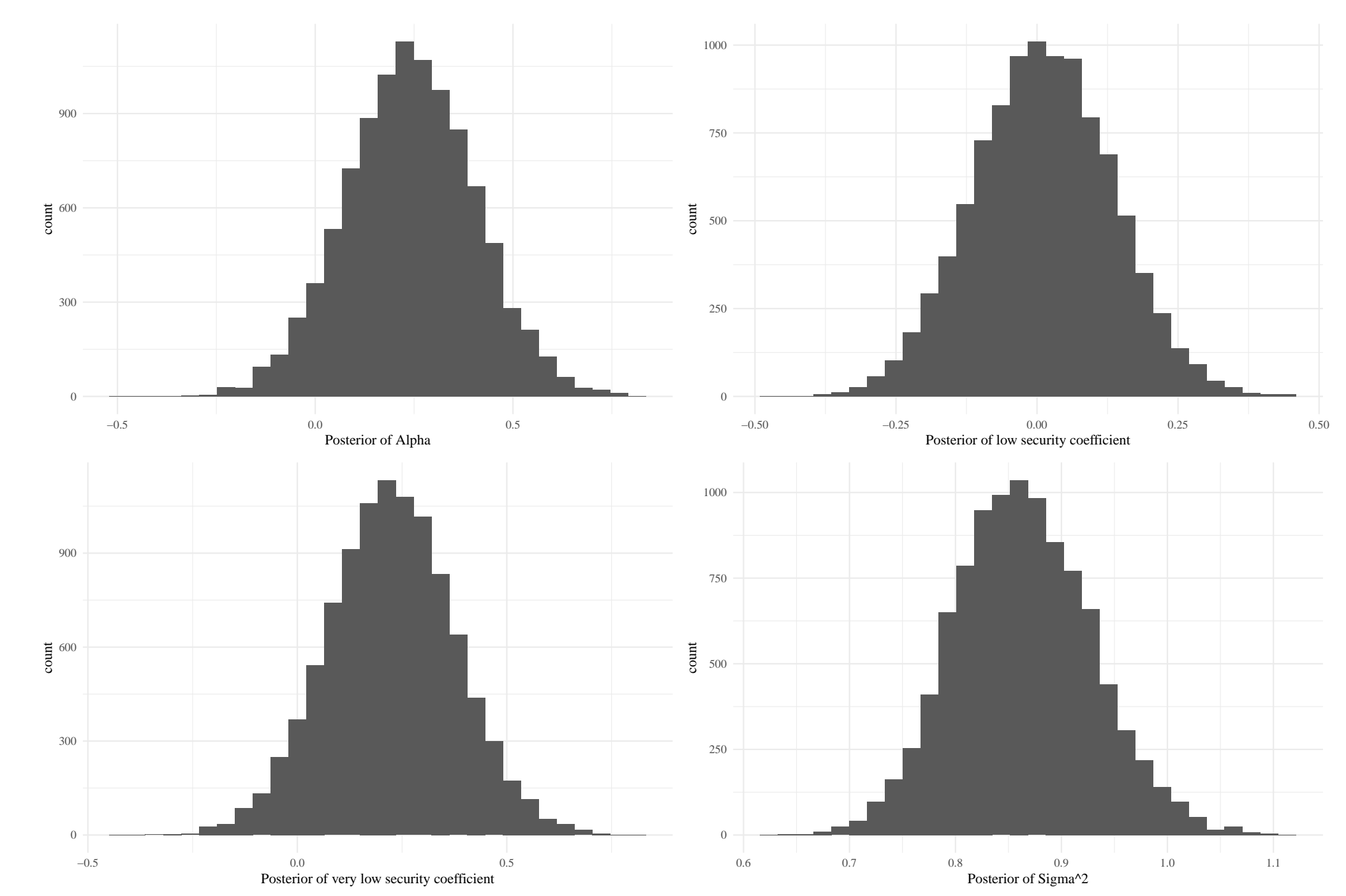


Figure 4: Posterior distributions of model parameters

## References

- (1) Azzalini, S. (1985). A class of distributions which includes the normal ones. SJS; (2) Fruhwirth-Schnatter, S and Pyne, S. (2010). Bayesian inference for finite mixtures of univariate ... Biostatistics; (3) Benjamin-Neelon SE, Ostbye T, Bennett GG, et al. Cohort profile for the Nurture Observational Study ... BMJ Open 2017; (4) Neelon, B. (2015) Bayesian two-part spatial models ... Biostatistics.

## Further Resources

<https://carter-allen.github.io/SN>

