

# *Introducing Brms: A R package for Bayesian Regression*

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🔗 <https://github.com/lawsofthought/intro-to-brms>

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## What is Brms?

- Brms is an R package for doing Bayesian general and generalized linear models, and general and generalized multilevel variants, etc.
- It includes far more probability models for outcome variables than almost all other regression packages: gaussian, student, binomial, bernoulli, poisson, negbinomial, geometric, Gamma, skew\_normal, lognormal, shifted\_lognormal, exgaussian, wiener, inverse.gaussian, exponential, weibull, frechet, Beta, von\_mises, asym\_laplace, gen\_extreme\_value, categorical, cumulative, cratio, sratio, acat, hurdle\_poisson, hurdle\_negbinomial, hurdle\_gamma, hurdle\_lognormal, zero\_inflated\_binomial, zero\_inflated\_beta, zero\_inflated\_negbinomial, zero\_inflated\_poisson, and zero\_one\_inflated\_beta.
- It also allows for censored data, missing data, measurement error, nonlinear regression, probabilistic mixture models, *distributional* models (whereby all parameters of the outcome variables have predictors), and so on.

## *The how and why of Brms*

- ▶ Brms writes Stan and Stan writes and compiles a MCMC sampler.
- ▶ To understand this process and its importance better, we must appreciate the following:
  1. Bayes is best. No further discussion necessary.
  2. Doing Bayesian data analysis, except for when using a prohibitively small set of models, requires Markov Chain Monte Carlo (MCMC) samplers.
  3. Writing your own MCMC is either hard or very hard.
  4. Probabilistic programming languages like Stan essentially write your MCMC sampler for any model you programmatically define.
  5. Although probabilistic programming languages reduce down the time and effort to obtain your sampler by orders of magnitude, they *still* require considerable time and effort relative to writing a single R command.
- ▶ Brms allows you to write your Bayesian model (with some restrictions) using standard R regression commands.