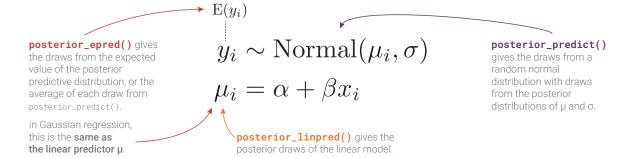
# Posterior predictions, linear predictions, and expectation of posterior predictions cheat sheet

v1.0 (2022-09-26) Andrew Heiss (https://www.andrewheiss.com; @andrewheiss)

#### **Normal Gaussian models**

Normal or Gaussian models are roughly equivalent to standard frequentist ordinary least squares (OLS) models and are generally interpreted the same way.

All the Bayesian prediction functions return values on the original scale of the outcome. In Gaussian models, the results from posterior\_epred() and posterior\_linpred() are identical. The draws from posterior\_predict() will generally have the same mean as the linear predictor, but with more variance since the overall model  $\sigma$  is incorporated into the predictions.



#### Generalized linear models with link transformations

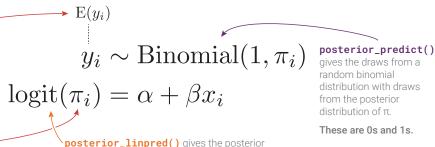
Generalized linear models (e.g., logistic, probit, ordered logistic, exponential, Poisson, negative binomial, etc.) use special link functions (e.g. logit, log, etc.) to transform the outcome variable into a scale that is more amenable to linear regression.

Estimates from these models can be used in their transformed scales (e.g., log odds in logistic regression) or can be back-transformed into their original scale (e.g., probabilities in logistic regression).

When working with links, the various Bayesian prediction functions return values on different scales, each corresponding to different parts of the model.

posterior\_epred() gives the draws from the expected value of the posterior predictive distribution, or the average of each draw from posterior\_predict().

In logistic regression, this is  $\pi$  on the probability scale (or inverse logit).



draws of  $\pi$  on the logit or log odds scale.

These are 0s and 1s.

#### Distributional models with link transformations

Regression models often focus solely on the location parameter of the model (e.g.,  $\mu$  in Normal( $\mu$ ,  $\sigma$ );  $\pi$  in Binomial(n,  $\pi$ )). However, it is also possible to specify separate predictors for the scale or shape parameters of models (e.g.,  $\sigma$  in Normal( $\mu$ ,  $\sigma$ ),  $\phi$  in Beta( $\mu$ ,  $\phi$ ))

More complex models use a collection of distributional parameters (e.g., zero-inflated beta models estimate a mean μ, precision φ, and a zero-inflated parameter zi; hurdle lognormal models estimate a mean  $\mu$ , scale  $\sigma$ , and a hurdle parameter hu).

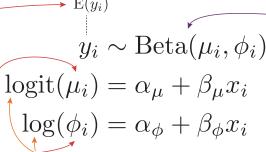
Even plain old Gaussian models become distributional models when a set of predictors is specified for  $\sigma$ (e.g.  $brm(y \sim x1 + x2, sigma \sim x2 + x3)$ )

When working with extra distributional parameters (dpar), the various Bayesian prediction functions return values on different scales for different components of the model.

posterior\_epred() gives the draws from the expected value of the posterior predictive distribution, or the average of each draw from posterior\_predict().

In beta regression, this is μ on the proportion or probability scale (or inverse logit).

posterior\_epred(dpar = "phi") gives the posterior draws of φ on the unloaged scale.



- posterior\_linpred() gives the posterior draws of  $\mu$  on the logit or log odds scale.

posterior\_linpred(dpar = "phi") gives the posterior draws of  $\varphi$  on the log scale.

### posterior\_predict()

gives the draws from a random beta distribution with draws from the posterior distribution of  $\pi$ .

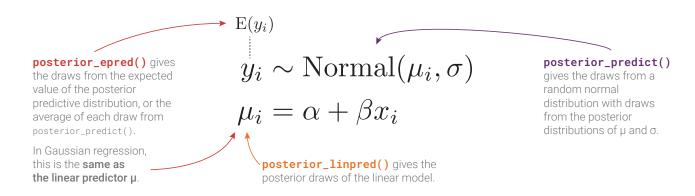
These are proportions or probabilities between 0-1.

#### Gaussian regression example

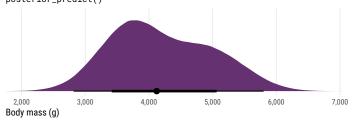
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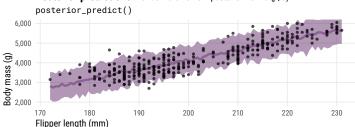
```
model_normal <- brm(
  bf(body_mass_g ~ flipper_length_mm),
  family = gaussian(),
  data = penguins
)</pre>
```



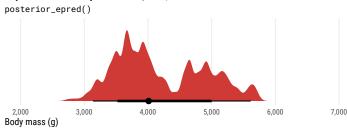
# **Posterior predictions** Random draws from posterior Normal( $\mu$ , $\sigma$ ) posterior\_predict()



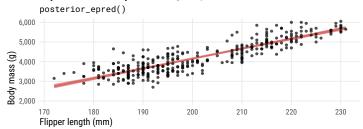
## **Posterior predictions** Random draws from posterior Normal( $\mu$ , $\sigma$ )



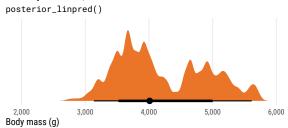
#### **Expectation of the posterior** E[y] and $\mu$ in the model



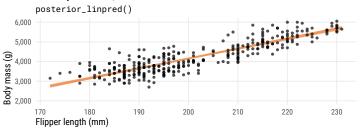
#### **Expectation of the posterior** E[y] and $\mu$ in the model



#### **Linear predictor** $\mu$ in the model



#### **Linear predictor** $\mu$ in the model



#### **Logistic regression example** (generalized linear model with link transformation)

Generalized linear models (e.g., logistic, probit, ordered logistic, exponential, Poisson, negative binomial, etc.) use special link functions (e.g. logit, log, etc.) to transform the outcome variable into a scale that is more amenable to linear regression.

Estimates from these models can be used in their transformed scales (e.g., log odds in logistic regression) or can be back-transformed into their original scale (e.g., probabilities in logistic regression).

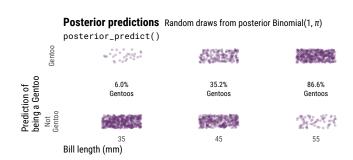
When working with links, the various Bayesian prediction functions return values on different scales, each corresponding to different parts of the model.

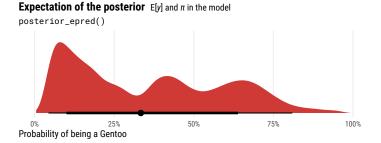
```
penguins <- penguins |>
  mutate(is_gentoo = species == "Gentoo")

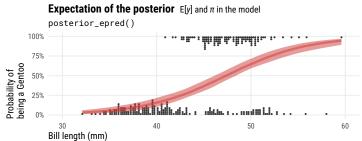
model_logit <- brm(
  bf(is_gentoo ~ bill_length_mm),
  family = bernoulli(link = "logit"),
  data = penguins
)</pre>
```

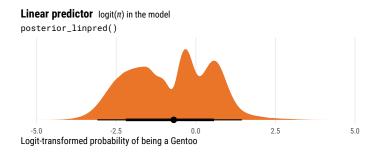
 $\rightarrow \mathrm{E}(y_i)$  $y_i \sim \text{Binomial}(1, \pi_i)$ posterior\_epred() gives posterior\_predict() the draws from the expected gives the draws from a random binomial value of the posterior  $logit(\pi_i) = \alpha + \beta x_i$ predictive distribution, or the distribution with draws average of each draw from from the posterior posterior\_predict(). distribution of  $\pi$ . In logistic regression, this is These are 0s and 1s.  $\pi$  on the probability scale posterior\_linpred() gives the posterior (or inverse logit). draws of  $\pi$  on the logit or log odds scale.

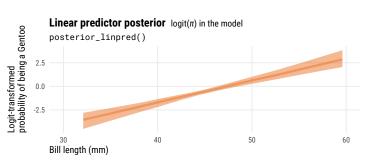
# 











#### **Beta regression example** (distributional model with multiple link transformations)

Regression models often focus solely on the location parameter of the model (e.g.,  $\mu$  in Normal( $\mu$ ,  $\sigma$ );  $\pi$  in Binomial(n,  $\pi$ )). However, it is also possible to specify separate predictors for the scale or shape parameters of models (e.g.,  $\sigma$  in Normal( $\mu$ ,  $\sigma$ ),  $\phi$  in Beta( $\mu$ ,  $\phi$ )).

More complex models use a collection of distributional parameters (e.g., zero-inflated beta models estimate a mean μ, precision φ, and a zero-inflated parameter zi; hurdle lognormal models estimate a mean  $\mu$ , scale  $\sigma$ , and a hurdle parameter hu).

Even plain old Gaussian models become distributional models when a set of predictors is specified for  $\boldsymbol{\sigma}$ (e.g.  $brm(y \sim x1 + x2, sigma \sim x2 + x3)$ )

When working with extra distributional parameters (dpar), the various Bayesian prediction functions return values on different scales for different components of the model.

```
penguins <- penguins |>
  mutate(bill_ratio = bill_depth_mm /
                        bill_length_mm)
model_beta <- brm(</pre>
  bf(bill_ratio ~ flipper_length_mm,
     phi ~ flipper_length_mm),
  family = Beta(),
  data = penguins
```

posterior\_epred() gives the draws from the expected value of the posterior predictive distribution, or the average of each draw from posterior\_predict().

In beta regression, this is μ on the proportion or probability scale (or inverse logit).

posterior\_epred(dpar = "phi") gives the posterior draws of φ on the unlogged scale.

 $\log(\phi_i) = \alpha_\phi + \beta_\phi x_i$ 

posterior\_linpred() gives the posterior draws of  $\mu$  on the logit or log odds scale.

posterior linpred(dpar = "phi") gives the posterior draws of  $\varphi$  on the log scale.

