Chapter 6

- 6.1 The place of model checking in applied Bayesian statistics
- 6.2 Do the inferences from the model make sense?
- 6.3 Posterior predictive checking
- 6.4 Graphical posterior predictive checks
 - this can be skimmed, see instead the paper
 Gabry et al. (2019). Visualization in Bayesian workflow
- 6.5 Model checking for the educational testing example

Model checking

- demo6_1: Posterior predictive checking light speed
- demo6_2: Posterior predictive checking sequential dependence
- demo6_3: Posterior predictive checking poor test statistic
- demo6_4: Posterior predictive checking marginal predictive p-value

Model checking - overview

- Sensibility with respect to additional information not used in modeling
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- Internal validation
 - posterior predictive checking
 - cross-validation predictive checking

Simon Newcomb's light of speed experiment in 1882

Newcomb measured (n = 66) the time required for light to travel from his laboratory on the Potomac River to a mirror at the base of the Washington Monument and back, a total distance of 7422 meters.

- Newcomb's speed of light measurements
 - model $y \sim \text{normal}(\mu, \sigma)$ with prior $(\mu, \log \sigma) \propto 1$

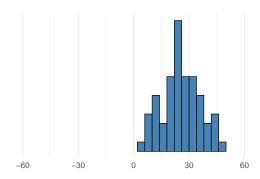
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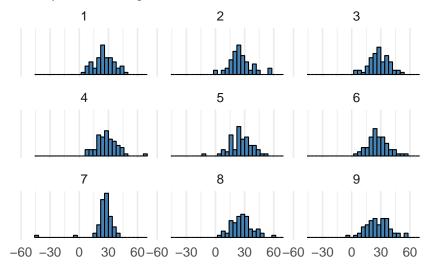
Replicates vs. future observation

Predictive ỹ is the next not yet observed possible observation.
 y^{rep} refers to replicating the whole experiment (potentially with same values of x) and obtaining as many replicated observations as in the original data.

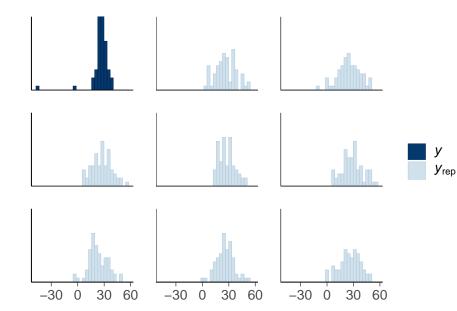
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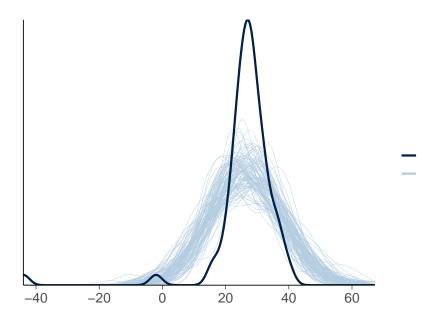


Posterior predictive checking – bayesplot ppc_hist(y, yrep[1:8,])

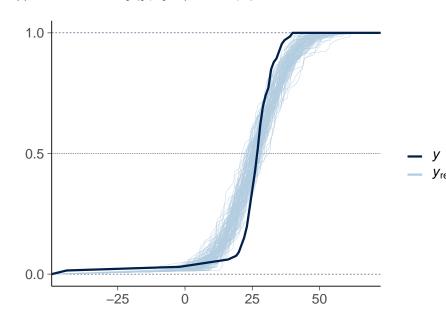


Posterior predictive checking – bayesplot

ppc_dens_overlay(y, yrep[1:100,])

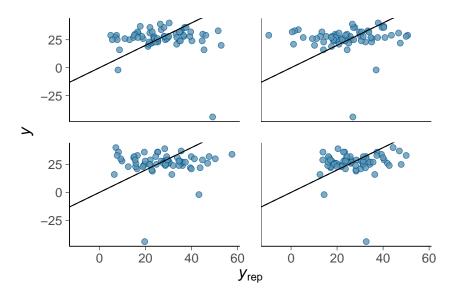


Posterior predictive checking – bayesplot ppc_ecdf_overlay(y, yrep[1:100,])



Posterior predictive checking – bayesplot

ppc_scatter(y, yrep[1:4,]) + geom_abline()



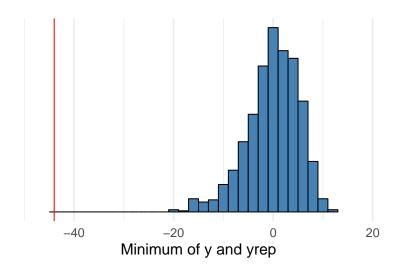
Posterior predictive checking with test statistic

- Replicated data sets y^{rep}
- Test quantity (or discrepancy measure) $T(y, \theta)$
 - summary quantity for the observed data $T(y, \theta)$
 - summary quantity for a replicated data $T(y^{rep}, \theta)$
 - can be easier to compare summary quantities than data sets

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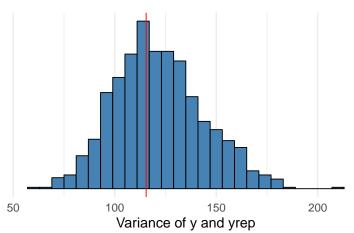
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Posterior predictive checking

Posterior predictive p-value

$$p = \Pr(T(y^{\text{rep}}, \theta) \ge T(y, \theta)|y)$$
$$= \int \int I_{T(y^{\text{rep}}, \theta) \ge T(y, \theta)} p(y^{\text{rep}}|\theta) p(\theta|y) dy^{\text{rep}} d\theta$$

where I is an indicator function

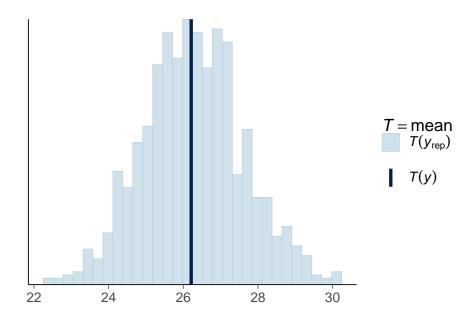
• having $(y^{\text{rep }(s)}, \theta^{(s)})$ from the posterior predictive distribution, easy to compute

$$T(y^{\text{rep}(s)}, \theta^{(s)}) \ge T(y, \theta^{(s)}), \quad s = 1, \dots, S$$

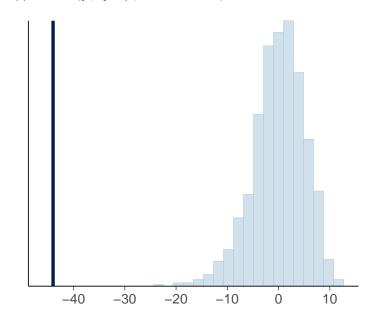
- Posterior predictive p-value (ppp-value) estimates whether difference between the model and data could arise by chance
- Not commonly used, as
 - not calibrated in case of non-ancillary statistic
 - the distribution of test statistic has more information

Posterior predictive checking – bayesplot

ppc_stat(y, yrep), the default statistic "mean" is usually bad



Posterior predictive checking – bayesplot ppc_stat(y, yrep, stat="min")

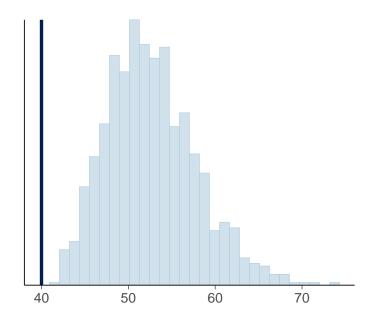


 $T = \min_{T(y_{rep})}$

T(y)

Posterior predictive checking – bayesplot

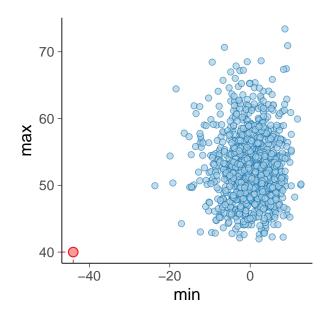
ppc_stat(y, yrep, stat="max")



 $T = \max_{T(y_{rep})}$

T(y)

Posterior predictive checking — bayesplot ppc_stat2d(y, yrep, stat=c("min", "max"))

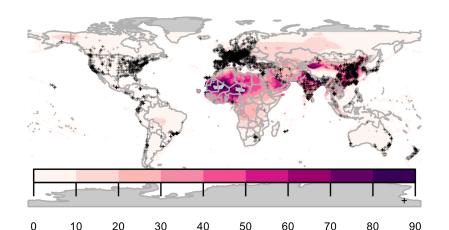


 $T = (\min, \max)$

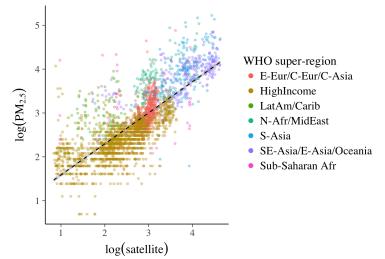


- Example from Jonah Gabry, Daniel Simpson, Aki Vehtari,
 Michael Betancourt, and Andrew Gelman (2019). Visualization in Bayesian workflow. https://doi.org/10.1111/rssa.12378
- Estimation of human exposure to air pollution from particulate matter measuring less than 2.5 microns in diameter (PM_{2.5})
 - Exposure to PM_{2.5} is linked to a number of poor health outcomes and a recent report estimated that PM_{2.5} is responsible for three million deaths worldwide each year (Shaddick et al., 2017)
 - In order to estimate the public health effect of ambient PM_{2.5}, we need a good estimate of the PM_{2.5} concentration at the same spatial resolution as our population estimates.

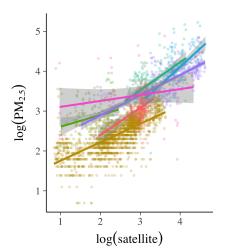
- Direct measurements of PM 2.5 from ground monitors at 2980 locations
- · High-resolution satellite data of aerosol optical depth



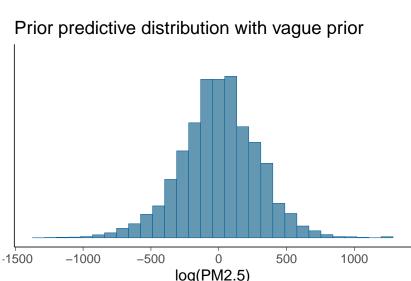
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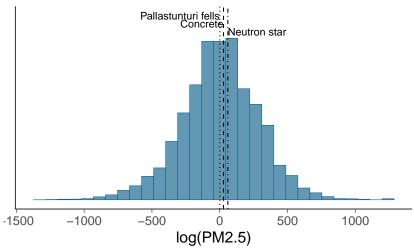


Prior predictive checking



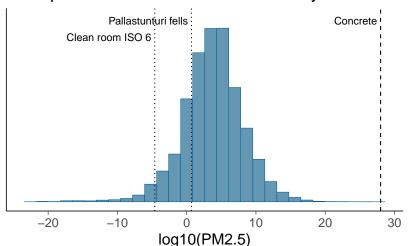
Prior predictive checking

Prior predictive distribution with vague prior

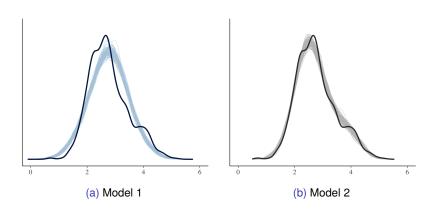


Prior predictive checking

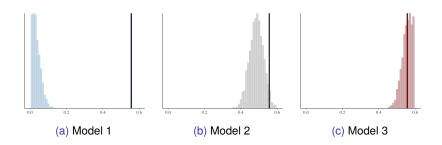
Prior predictive distribution with weakly informative



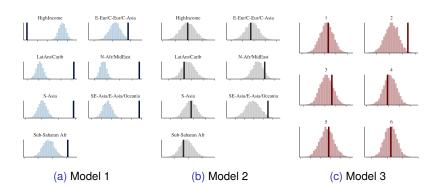
Posterior predictive checking – marginal predictive distributions



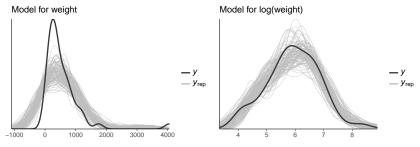
Posterior predictive checking – test statistic (skewness)



Posterior predictive checking – test statistic (median for groups)



Positive target



Predicting the yields of mesquite bushes.

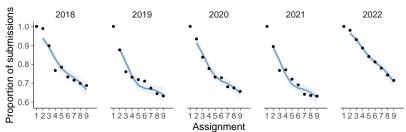
Gelman, Hill & Vehtari (2020): Regression and Other Stories, Chapter 11.

Student retention

Latent hierarchical linear + spline

```
nstudents \ | \ trials (nstudents1) \ \sim \ (assignment \ | \ year) \ + \\ s(assignment, \ k=4), \ family=binomial()
```

Latent functions + posterior uncertainty



Student retention

1. Latent hierarchical linear model

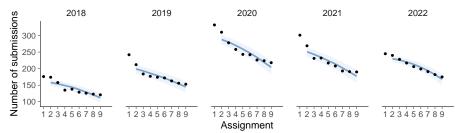
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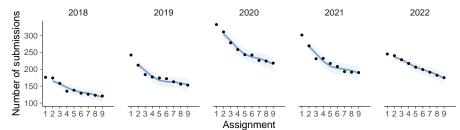
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Student retention – Posterior predictive distributions

with tidybayes



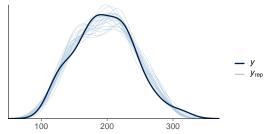


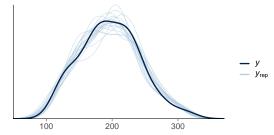


Student retention – Marginal PPC

pp_check(fit, ndraws=100)

Latent hierarchical linear model

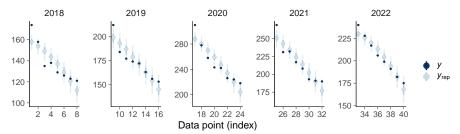


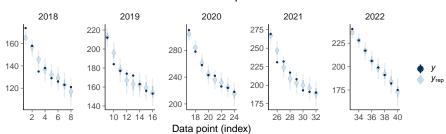


Student retention – Posterior predictive intervals

pp_check(fit, type = "intervals_grouped", group="year")

Latent hierarchical linear model

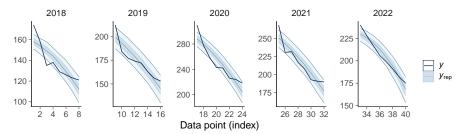


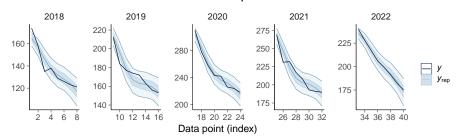


Student retention – Posterior predictive ribbon

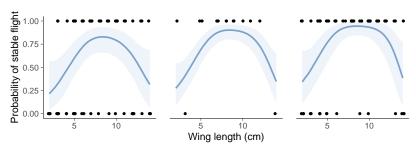
pp_check(fit, type = "ribbon_grouped", group="year")

Latent hierarchical linear model

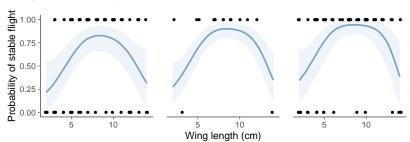


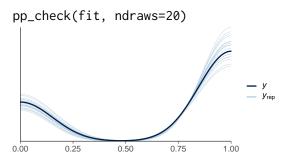


```
stable\_flight ~ s(wing\_length) ~ + ~ s(wing\_length, ~ by = nclips) \,, \\ family ~ = ~ bernoulli()
```

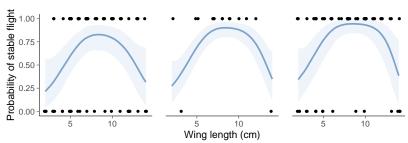


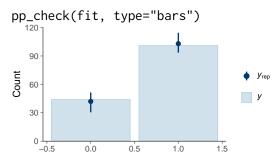
```
stable_flight \sim s(wing_length) + s(wing_length, by = nclips), family = bernoulli()
```



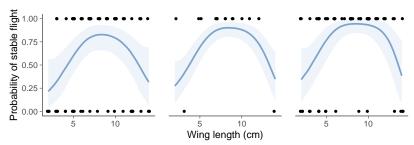


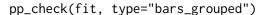
 $stable_flight ~ s(wing_length) ~ + ~ s(wing_length, ~ by = nclips) \,, \\ family ~ = ~ bernoulli()$

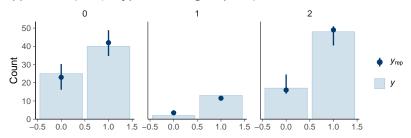




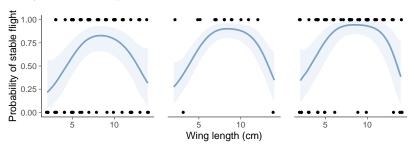
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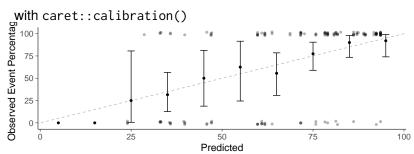




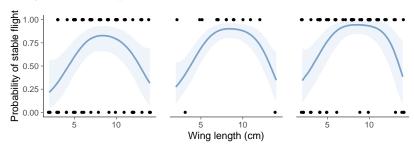


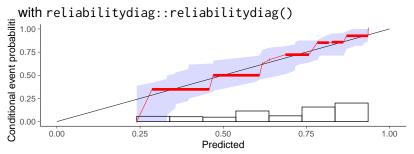
 $stable_flight \sim s(wing_length) + s(wing_length, by = nclips), family = bernoulli()$





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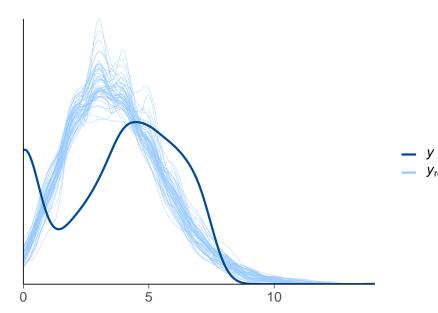
Posterior predictive checking - Stan code

demo demos_rstan/ppc/poisson-ppc.Rmd

```
data
  int < lower = 1 > N:
  int <lower=0> v[N];
parameters {
  real < lower = 0 > lambda:
model {
  lambda ~ exponential (0.2);
  v ~ poisson(lambda);
generated quantities {
  real log_lik[N];
  int y_rep[N];
  for (n in 1:N) {
    y_rep[n] = poisson_rng(lambda);
    log lik[n] = poisson lpmf(y[n] | lambda);
```

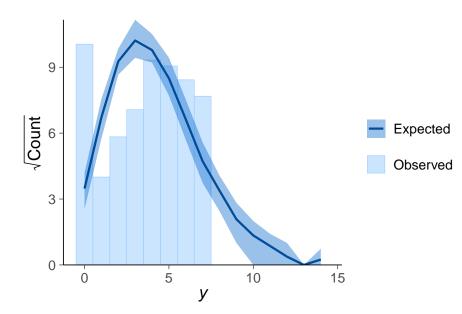
PPC for count data – Poisson model

ppc_dens_overlay(y, yrep[1:50,])



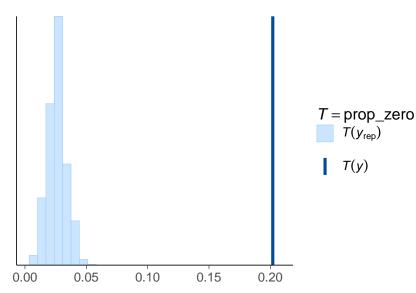
PPC for count data - Poisson model

ppc_rootogram(y, yrep)

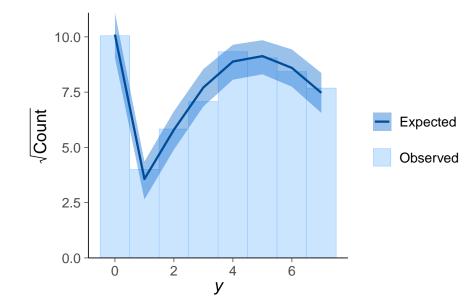


PPC for count data - Poisson model

```
prop_zero <- function(x) mean(x == 0)
ppc_stat(y, yrep, stat = "prop_zero")</pre>
```

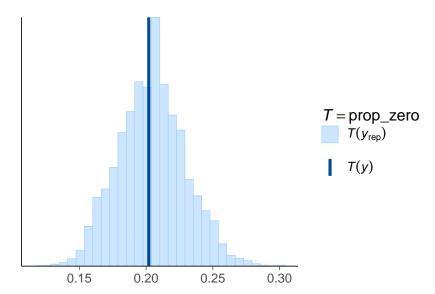


PPC for count data – hurdle truncated Poisson model ppc_rootogram(y, yrep2)



PPC for count data - hurdle truncated Poisson model

```
prop_zero <- function(x) mean(x == 0)
ppc_stat(y, yrep2, stat = "prop_zero")</pre>
```



Further reading and examples

- Gabry, Simpson, Vehtari, Betancourt, and Gelman (2019).
 Visualization in Bayesian workflow.
 https://doi.org/10.1111/rssa.12378.
- Graphical posterior predictive checks using the bayesplot package http://mc-stan.org/bayesplot/articles/graphical-ppcs.html
- Another demo demos_rstan/ppc/poisson-ppc.Rmd

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- Compare sensitivity of essential inference quantities
 - extreme quantiles are more sensitive than means and medians
 - extrapolation is more sensitive than interpolation