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  - Calibration
    - $\alpha\%$ -posterior interval has the true value in  $\alpha\%$  cases
    - $\alpha\%$ -predictive interval has the true future values in  $\alpha\%$  cases
    - approximate calibration with shorter intervals for likely true values more important than exact calibration with bad intervals for all possible values.

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  - unbiased estimate for strictly positive parameter can be negative
- Confidence interval is defined to have true value inside the interval in  $\alpha\%$  cases of repeated data generation from the data generating mechanism
  - doesn't say how likely the true value is inside the interval given the observed data
  - doesn't need be useful to have perfect calibration

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  - a consistent way to add prior information
- Lot of machine learning is not pure frequentist or Bayesian

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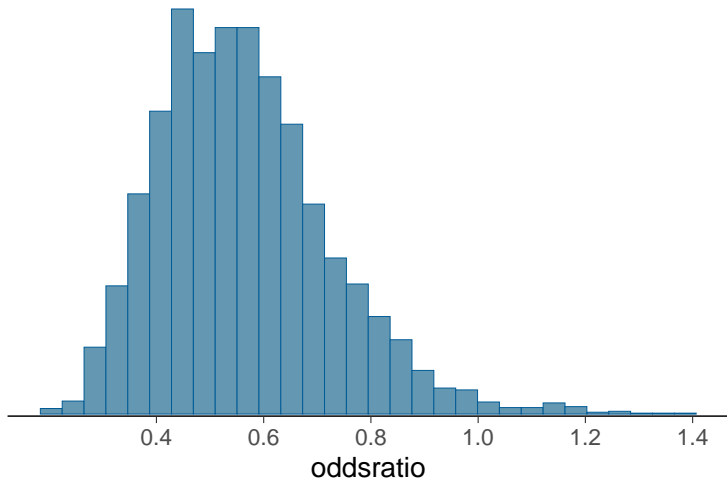


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- Frequentist null hypothesis testing
  - asks what if data is generated from the smaller model
  - doesn't tell whether the more complex model is good enough

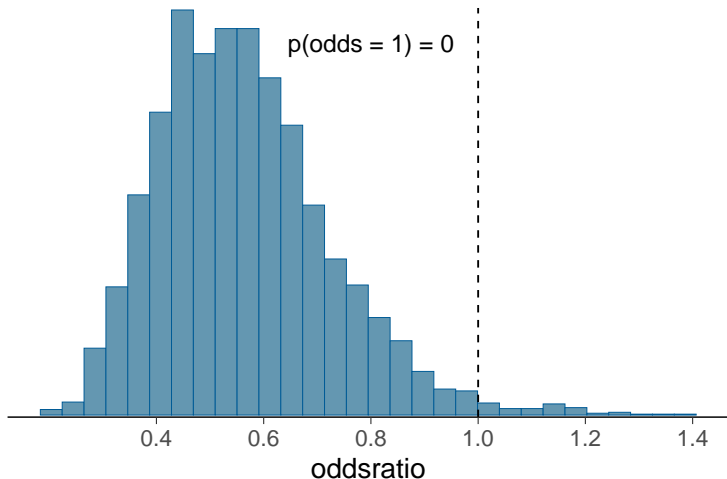
# Bayesian hypothesis testing

- Instead of hypothesis testing, report full posterior and
  - compare to expert information
  - combine with utility/cost function



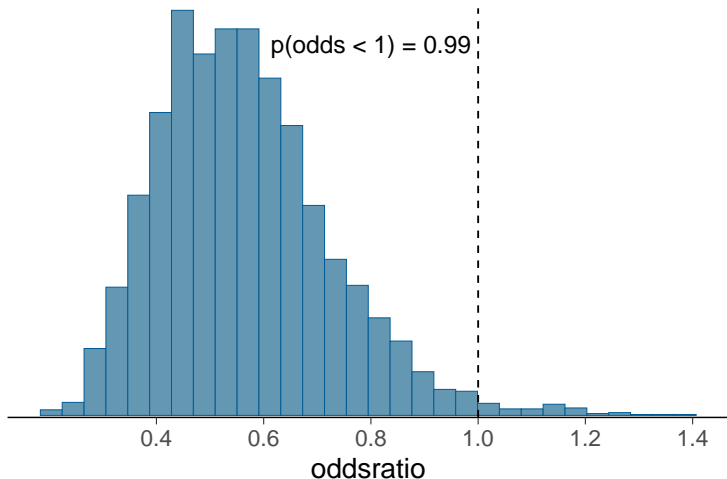
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- Instead of hypothesis testing, report full posterior
  - for continuous posterior there is zero probability that e.g. treatment effect is exactly zero



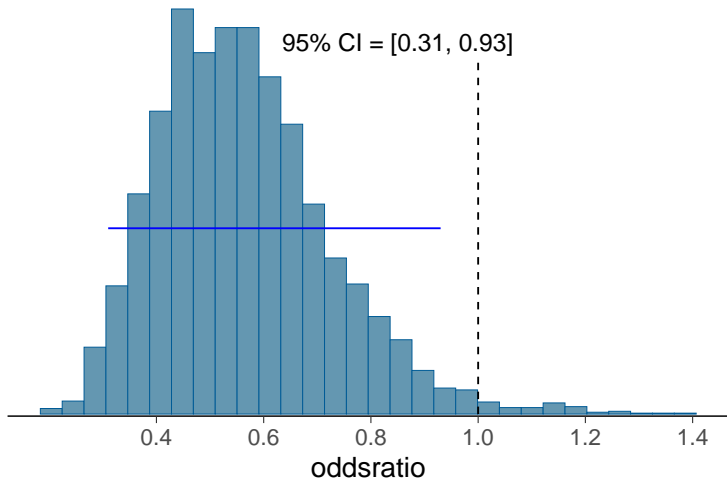
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- Instead of hypothesis testing, report full posterior
  - for continuous posterior we could compute the probability that we know the sign of the effect



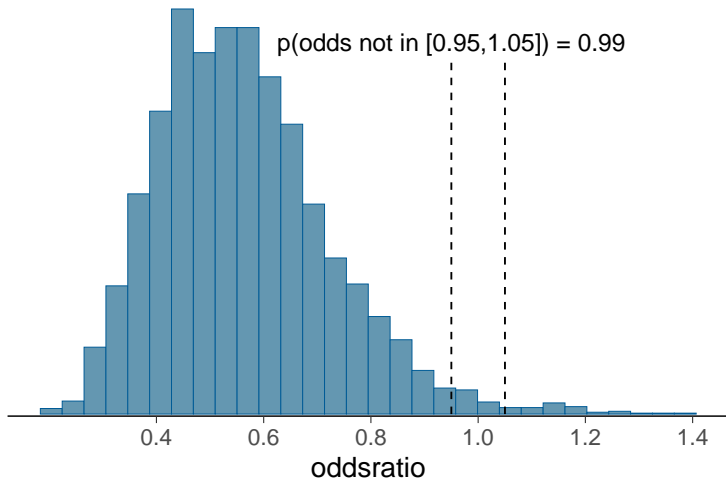
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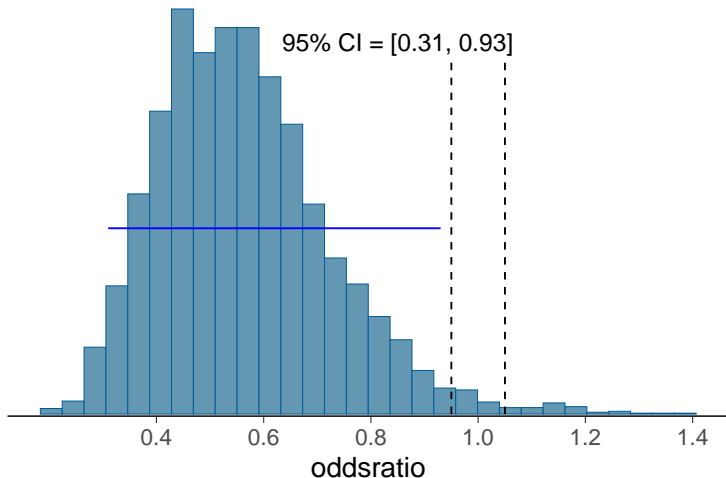
# Bayesian hypothesis testing

- Equivalence testing (region of practical equivalence)
  - what is the probability that the effect is closer than  $\epsilon$  to null, where  $\epsilon$  is based on what is practically useful effect size



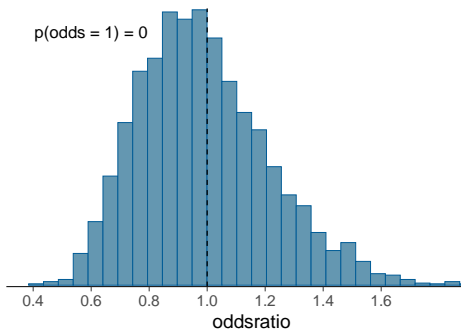
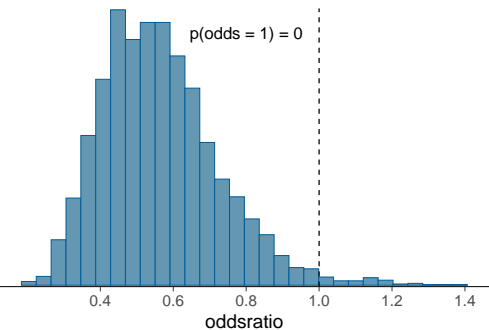
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- Equivalence testing (region of practical equivalence)
  - some people combine posterior interval and region of practical equivalence



# Bayesian hypothesis testing

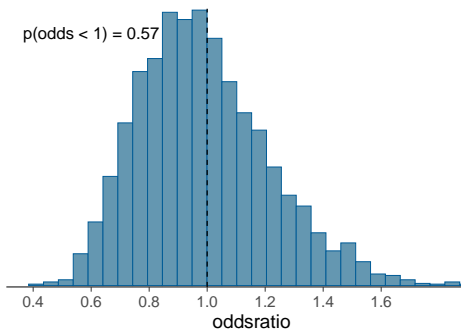
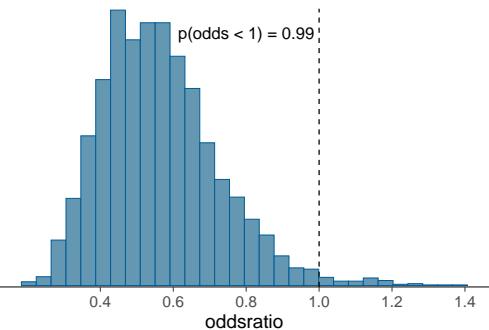
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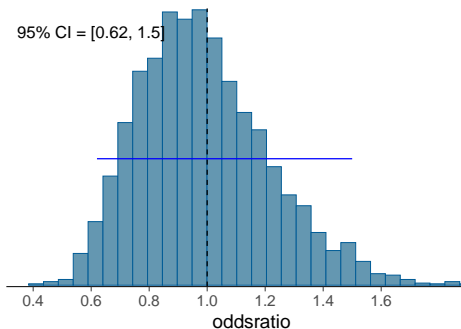
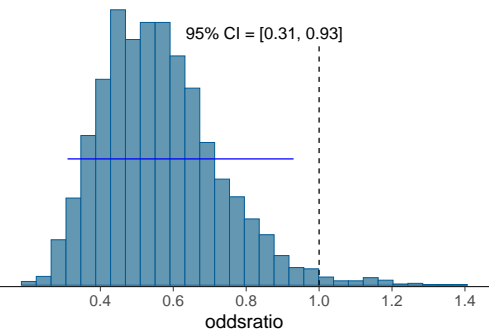
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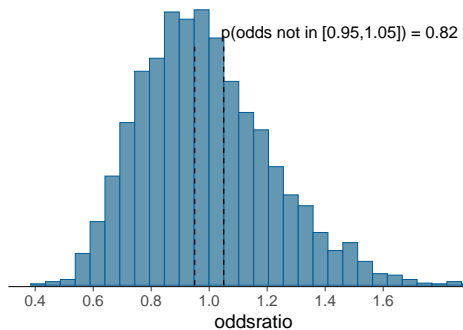
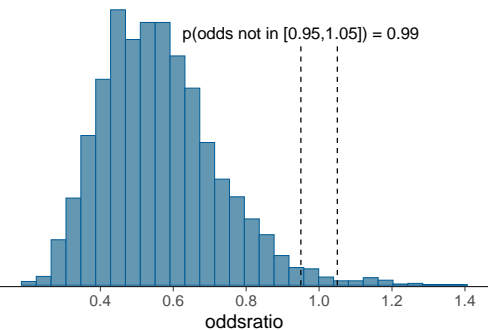
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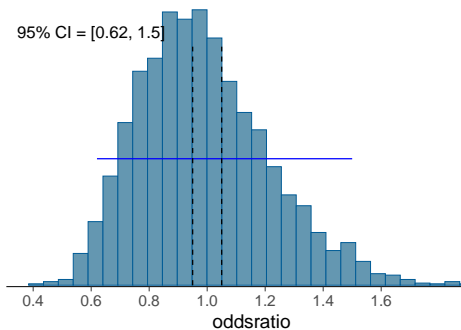
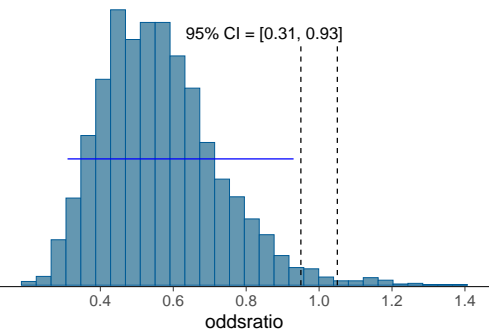
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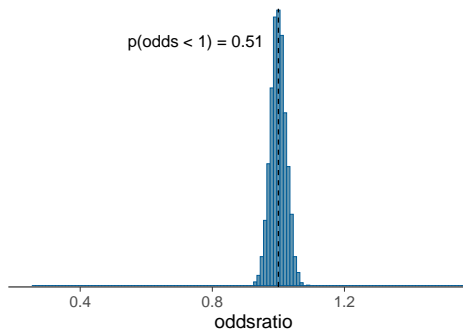
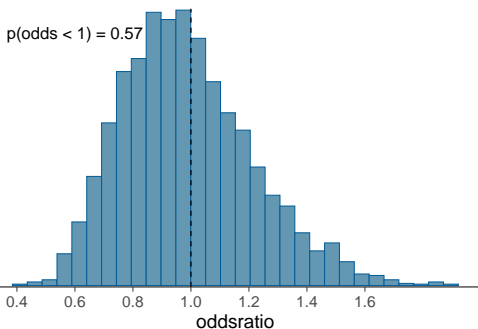


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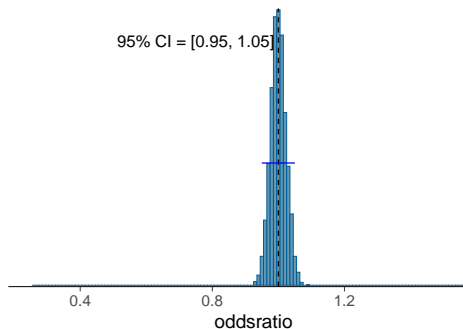
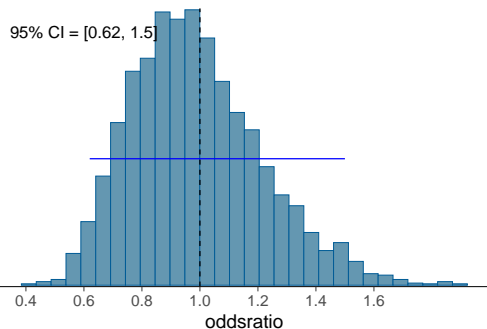
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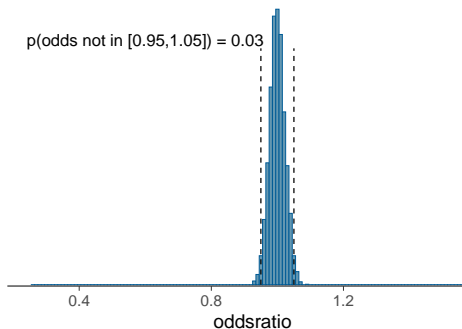
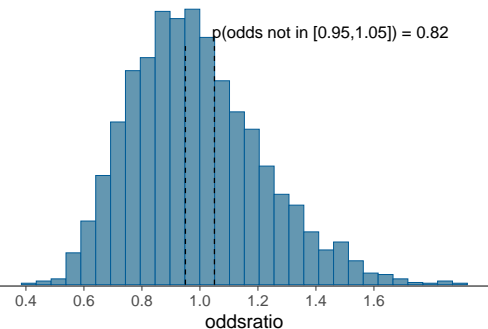
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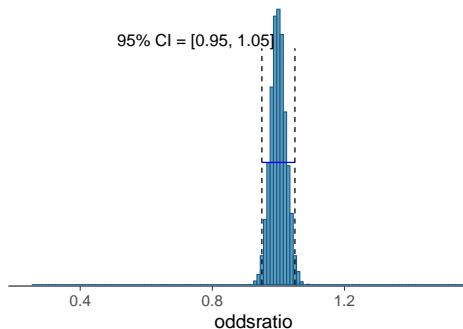
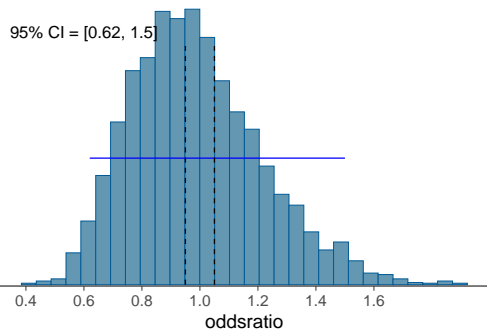
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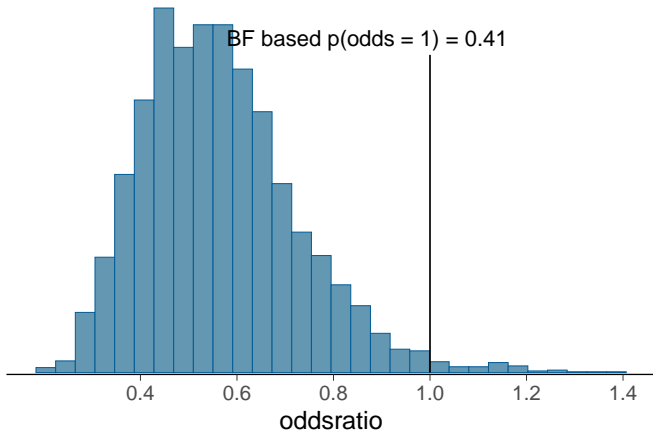
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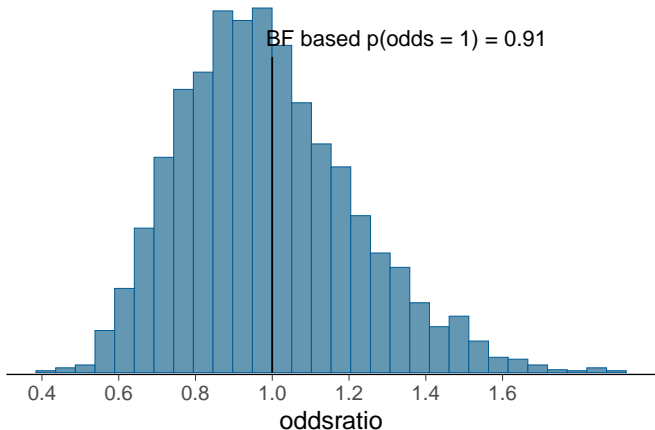
- Bayes factor
  - null model has, e.g., the treatment effect fixed to 0
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with `bridgesampling` package, see also BDA3 13.10

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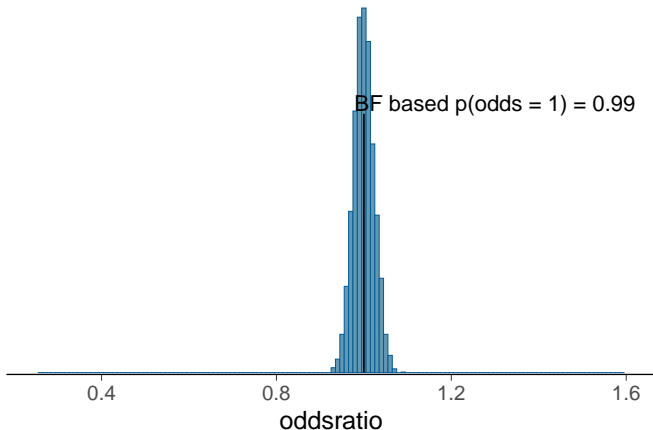
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  - is there difference in predictive performance with, e.g., treatment effect fixed to zero or unknown treatment effect
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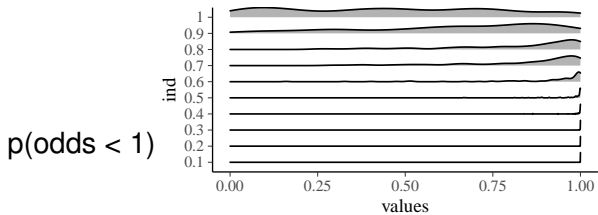
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In the beta blockers example

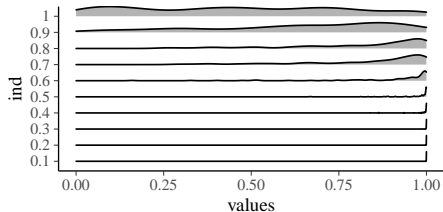
- Leave-one-group-out is not sensible as there are only two groups
- Leave-one-person-out works, but is less efficient than looking at the posterior (see <https://avehtari.github.io/modelselection/betablockers.html>)

# Simulation experiment

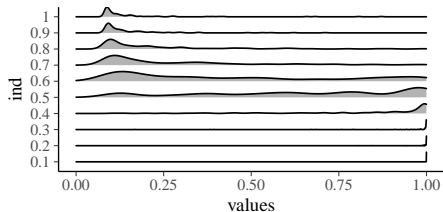


# Simulation experiment

$p(\text{odds} < 1)$



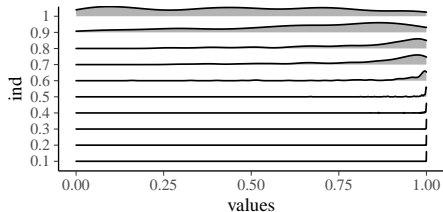
Marginal likelihood  
comparison



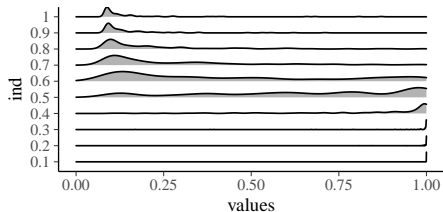


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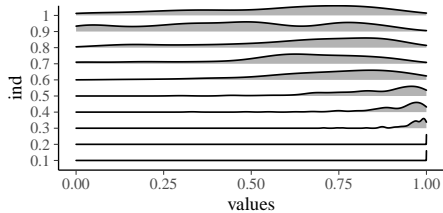
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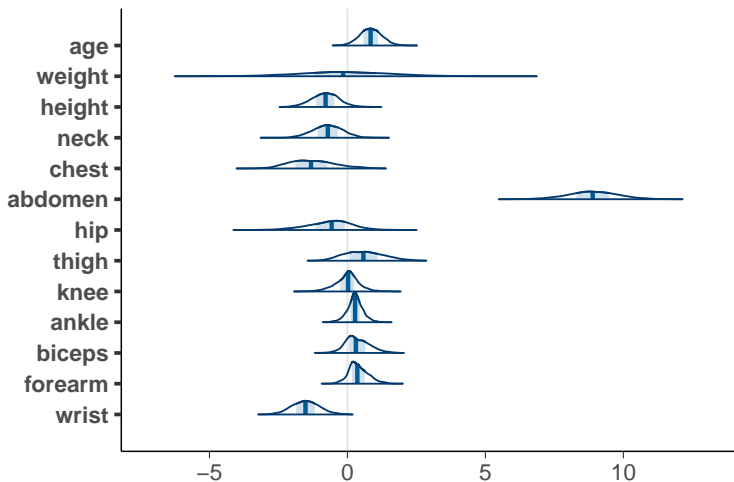
LOO comparison



# Hypothesis testing and posterior dependencies

Looking at the marginal posterior(s) can be misleading when there are many parameters

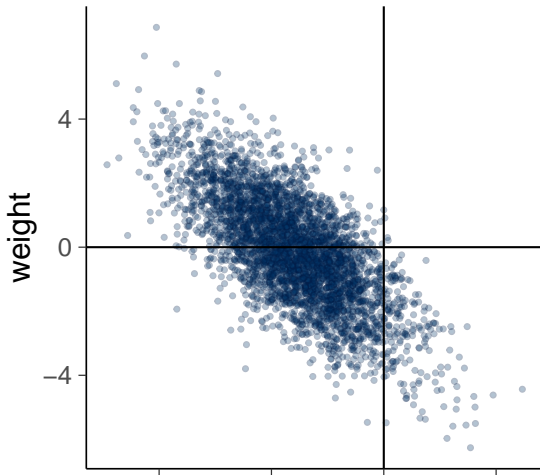
Marginal posteriors of coefficients



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Bivariate marginal of weight and height



# Hypothesis testing and posterior dependencies

In bodyfat example, starting from full model

- BF in favor of removing weight ( $p=0.92$ )
- LOO in favor of removing weight ( $p=0.99$ )

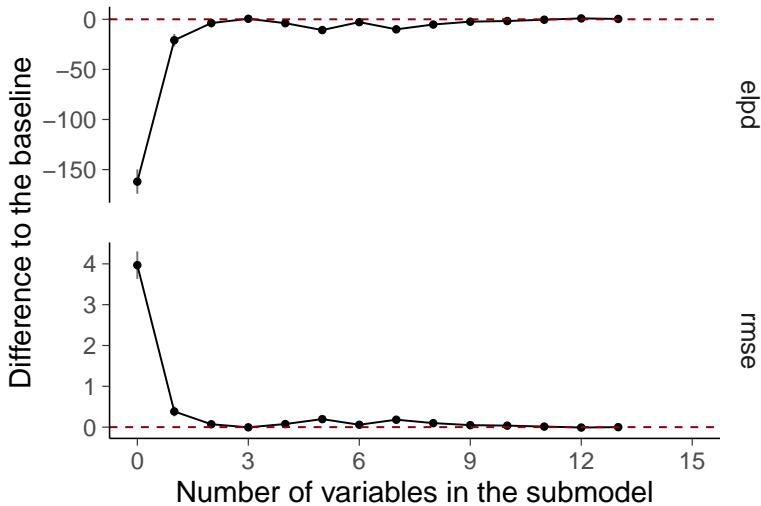
In bodyfat example, starting from model  $y \sim \text{abdomen}$

- BF in favor of adding weight ( $p=1.0$ )
- LOO in favor of adding weight ( $p=1.0$ )

## Variable selection

More elaborate approaches are needed for variable selection

See Lecture 9.3 on projection predictive variable selection



# Common statistical tests as Bayesian models

Most common statistical tests are linear models

<i>t</i> -test	mean of data	<code>stan_glm(y ~ 1)</code>
paired <i>t</i> -test	mean of diffs	<code>stan_glm((y1 - y2) ~ 1)</code>
Pearson correl.	linear model	<code>stan_glm(y ~ 1 + x)</code>
two-sample <i>t</i> -test	group means	<code>stan_glm(y ~ 1 + gid)</code>
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See longer list and illustrations (with `lm`) at

<https://lindeloev.github.io/tests-as-linear/>

and

in the forthcoming *Regression and other stories* book



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- Data collection
  - Sample surveys
  - Designed experiments
  - Randomization
  - Observational studies
  - Censoring and truncation

## Chapter 14: Introduction to regression models

- Justification of conditional modeling
  - if joint model factorizes  $p(y, x|\theta, \phi) = p(y|x, \theta)p(x|\phi)$   
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- Unequal variances and correlations

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  - sometimes relaxed lasso is used, where after variable selection coefficients are re-estimated

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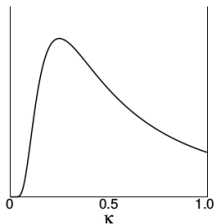
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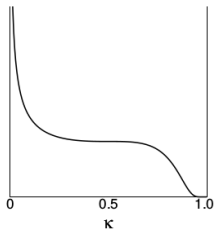
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  - empirically better results obtained with more sparse priors
  - it's best to separate selection of sensible prior, good posterior inference, and the decision analysis of which variables are important

# Sparse priors

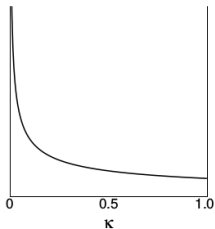
**Laplacian**



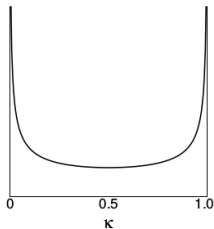
**Student-t**



**Strawderman-Berger**

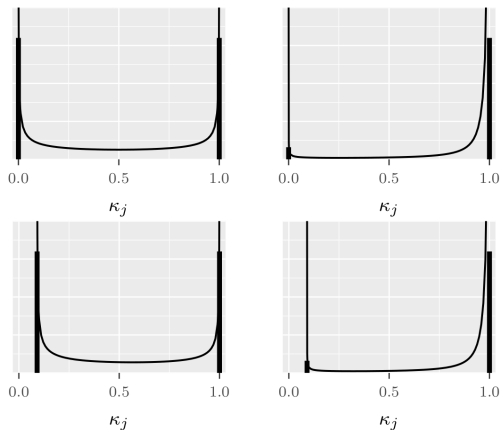


**Horseshoe**



from Carvalho, Polson, Scott (2009).

# Regularized horseshoe



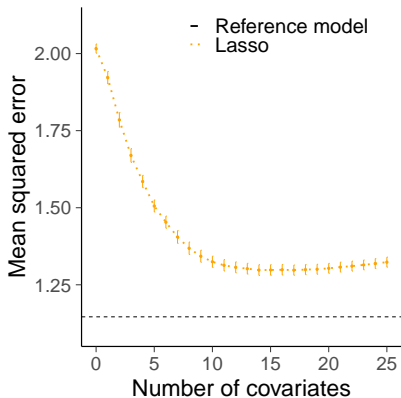
for more see

- Piironen and Vehtari (2017). Sparsity information and regularization in the horseshoe and other shrinkage priors. In Electronic Journal of Statistics, 11(2):5018-5051. [Online](#)
- [https://betanalpha.github.io/assets/case\\_studies/bayes\\_sparse\\_regression.html](https://betanalpha.github.io/assets/case_studies/bayes_sparse_regression.html)

# Projpred selection vs. Lasso

See projpred in lecture 9.3

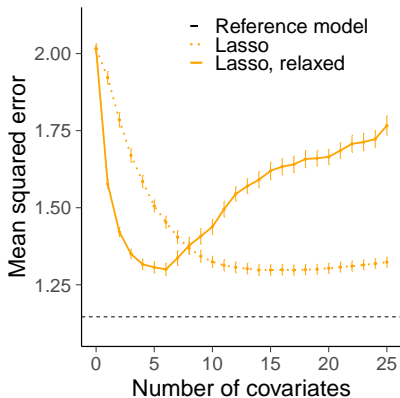
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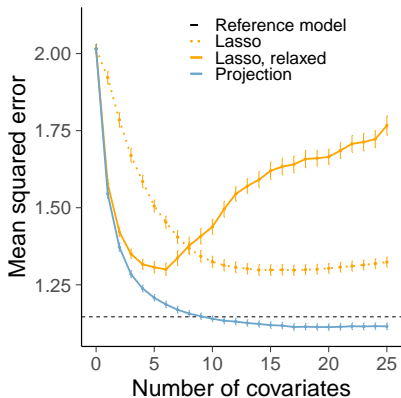
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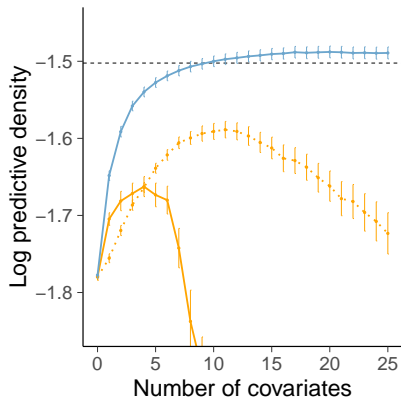
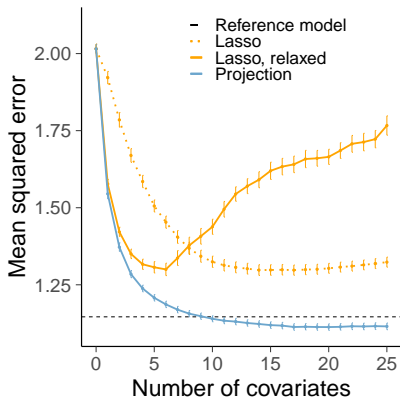
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## Chapter 15: Hierarchical linear models

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  - section on transformations for HMC is relevant  
(see also Stan user guide 21.7 Reparameterization)

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$y \sim 1 + x$	fixed / population effect; pooled model
$y \sim 1 + (0 + x \mid g)$	random / group effects
$y \sim 1 + x + (1 + x \mid g)$	mixed effects; hierarchical model

- ANOVA in section 15.6 (see also `stan_aov`)

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- Hierarchical GLM natural extension
- 16.3 Weakly informative priors section is excellent although the recommendation on using Cauchy has changed (see <https://github.com/stan-dev/stan/wiki/Prior-Choice-Recommendations>)

## Chapter 17: Models for robust inference

- For example

normal       $\rightarrow$      $t$ -distribution

Poisson      $\rightarrow$     negative-binomial

binomial     $\rightarrow$     beta-binomial

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  - rstanarm doesn't have  $t$ -distribution for outcome, but brms has

## Chapter 18: Models for missing data

- Extends the data collection modelling from Chapter 8
- Useful terms



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missingness depends on missing values
- Multiple imputation
  1. make a model predicting missing data
  2. sample repeatedly from the missing data model to generate multiple imputed data sets
  3. make usual inference for each imputed data set
  4. combine results

## Chapter 21: Gaussian process models

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  - infinite dimensional extension of normal distribution
  - useful prior for non-linear functions
  - for any finite number of variables, the marginal is multivariate normal  $f_1, \dots, f_n \sim \mathcal{N}(\mu(x_1, \dots, x_n), K(x_1, \dots, x_n))$

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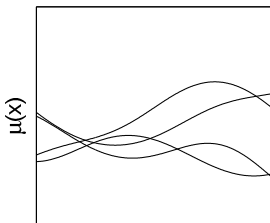
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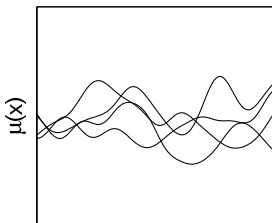
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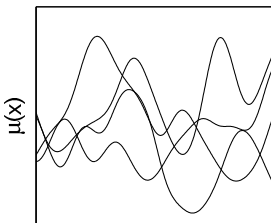
$x$

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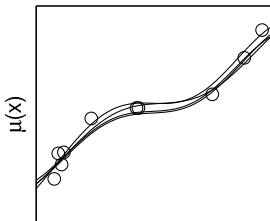


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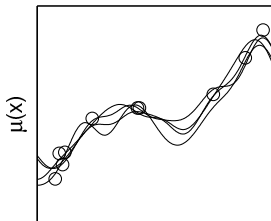
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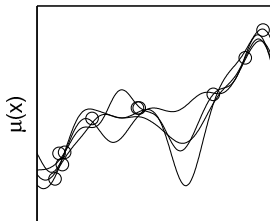
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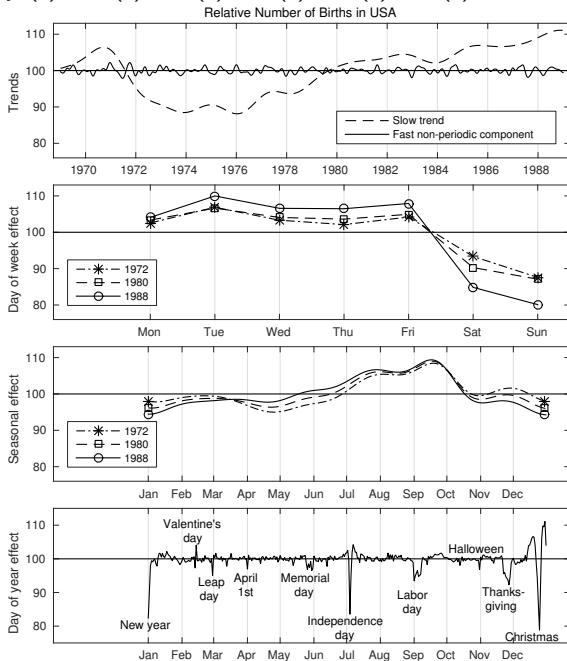
$x$

## Chapter 21: Gaussian process models

- Conditional on covariance function parameter the posterior is just multivariate normal
  - need to make inference for covariance function parameters given the marginal likelihood
  - the exact computation of the marginal likelihood scales  $O(N^3)$

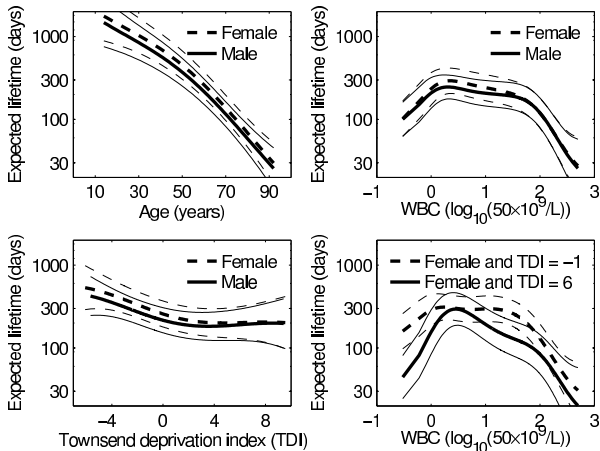
• Easy to make additive models

$$y_t(t) = f_1(t) + f_2(t) + f_3(t) + f_4(t) + f_5(t) + \epsilon_t$$



## Chapter 21: Gaussian process models

- For non-Gaussian outcome models similar extension as GLMs
- Survival model example:



## GPs in Stan

- GP specific software (e.g. GPy, GPflow, GPyTorch) scale computationally better for GPs than Stan
- Stan has some built-in covariance functions (and soon GPU support)
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- In case of non-Gaussian outcome models, sampling of latent variables can be slow (Laplace integration over the latents coming)
- Instead of covariance matrix based approach, for low dimensional cases faster to use basis function representation
  - e.g. `stan_glm(y ~ s(x, bs="gp"))`