

# Frequency evaluations

- Bayesian theory has epistemic and aleatory probabilities
- Frequency evaluations focus on frequency properties given aleatoric repetition of an observation and modeling
  - Asymptotic consistency
  - Unbiasedness
    - not that important in Bayesian inference, small error more important
  - Efficiency
    - small squared error
    - other utility/cost functions possible
  - 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.

# Frequentist statistics

- Frequentist statistics accepts only aleatory probabilities
  - Estimates are based on data
  - Uncertainty of estimates are based on all possible data sets which could have been generated by the data generating mechanism
    - inference is based also on data we did not observe
- Estimates are derived to fulfill frequency properties
  - Maximum likelihood fulfills just asymptotic frequency properties
  - Common desiderata are 1) unbiasedness, 2) minimum variance, 3) calibration of confidence interval

# Frequentist statistics

- Estimates are derived to fulfill frequency properties
  - Maximum likelihood fulfills just asymptotic frequency properties
  - Common desiderata are 1) unbiasedness, 2) minimum variance, 3) calibration of confidence interval
- Requirement of unbiasedness may lead to higher variance or silly estimates
  - 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

# Frequentist vs Bayes vs others

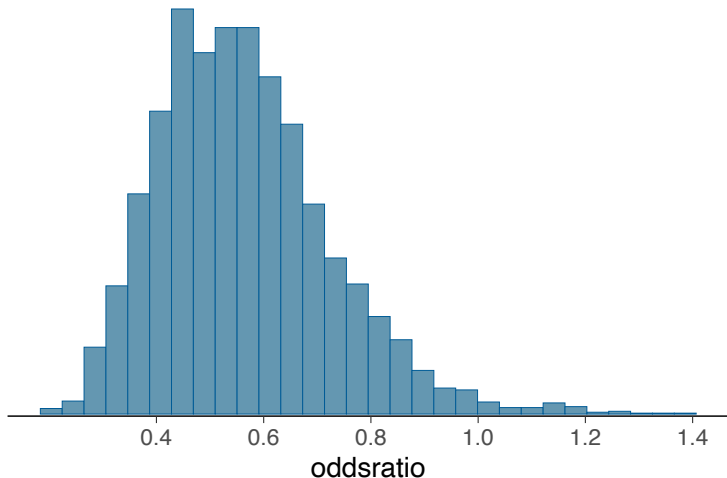
- There is a great amount of very useful frequentist statistics
  - also for simple models and lot's of data there is not much difference
- Bayesian inference
  - easier for complex, e.g. hierarchical, models
  - easier when model changes
  - a consistent way to add prior information
- Lot of machine learning is not pure frequentist or Bayesian

# Hypothesis testing

- Frequentist approach can be used to to make estimates and confidence intervals, but for some reason null hypothesis testing has a very big role
  - reporting just the null hypothesis testing result throws away lot of useful information
  - some Bayesians are also into null hypothesis testing
- 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

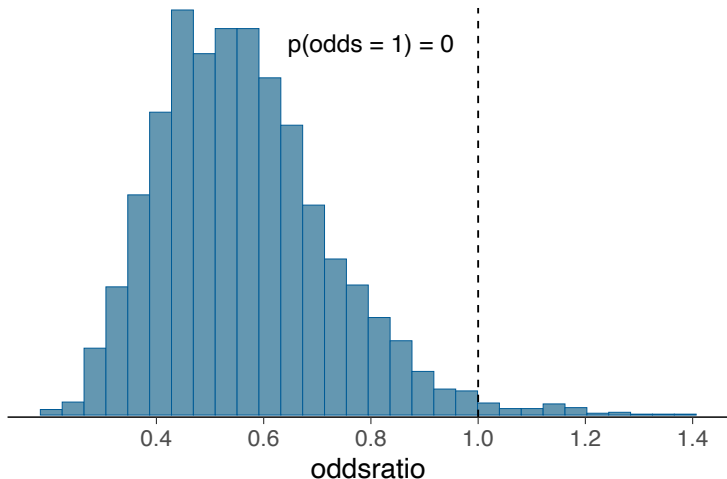
# Bayesian hypothesis testing

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



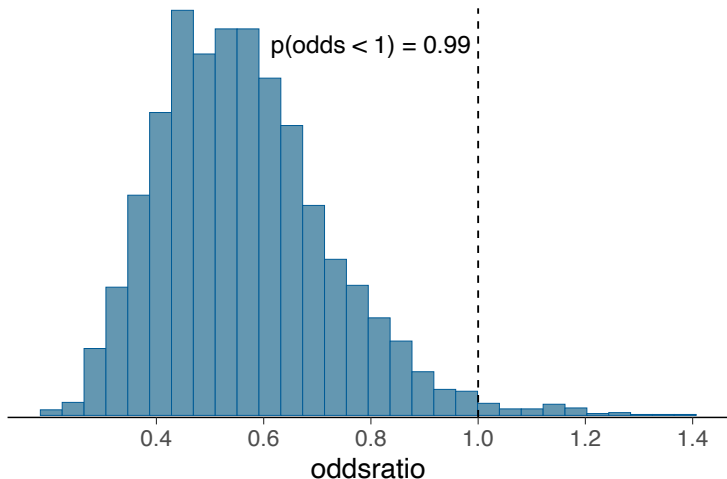
# Bayesian hypothesis testing

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



# Bayesian hypothesis testing

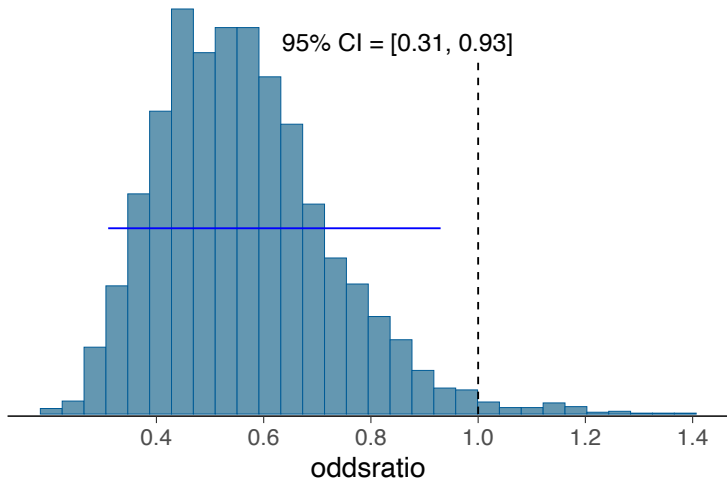
- Instead of hypothesis testing, report full posterior
  - for continuous posterior we could compute the probability that we know the sign of the effect





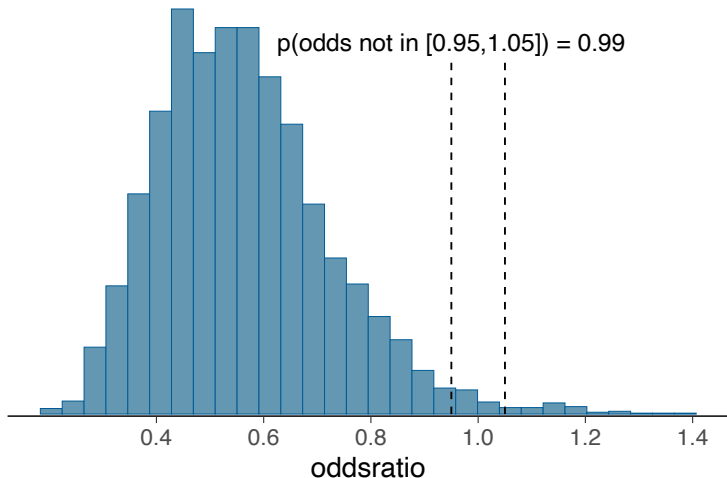
# Bayesian hypothesis testing

- Instead of hypothesis testing, report full posterior
  - for continuous posterior some people compare whether posterior interval includes null case



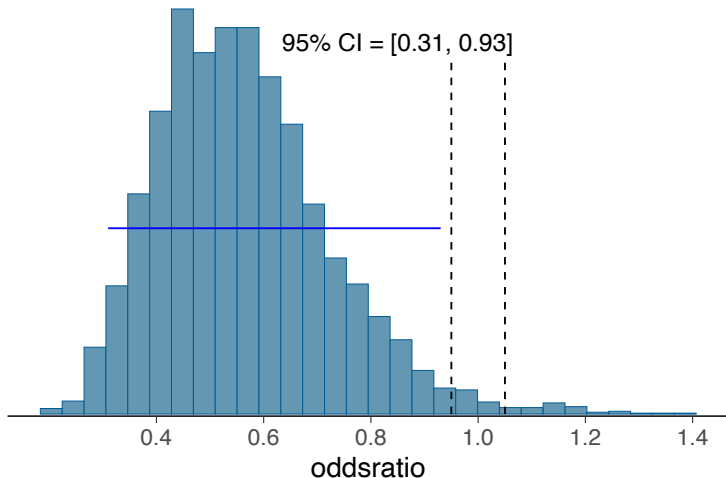
# Bayesian hypothesis testing

- 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



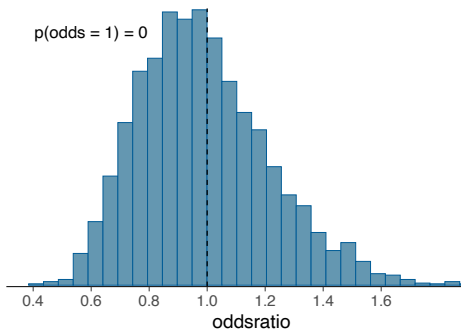
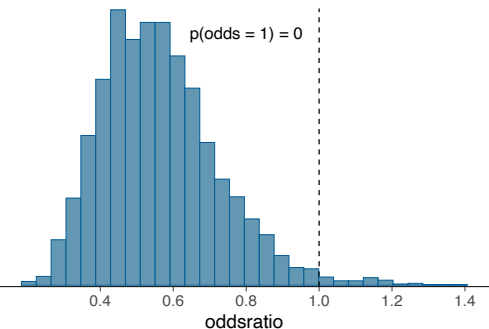
# Bayesian hypothesis testing

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



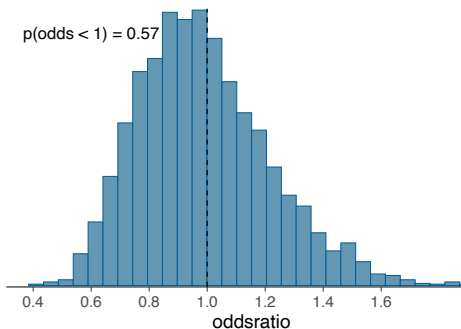
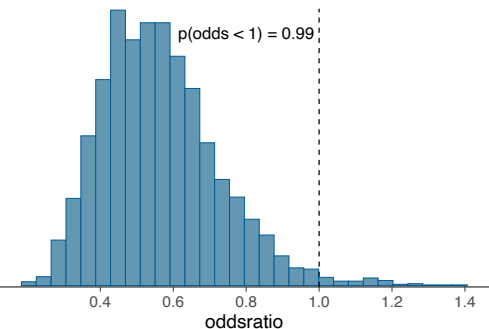
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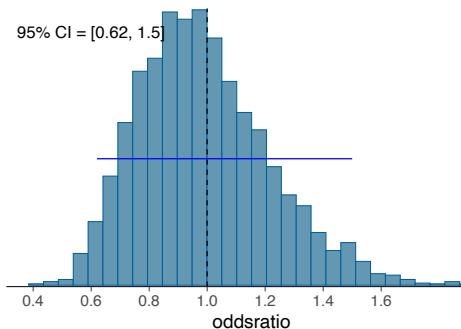
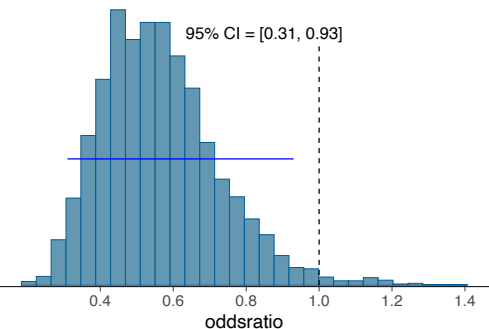
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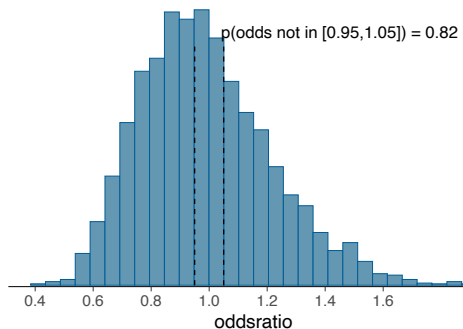
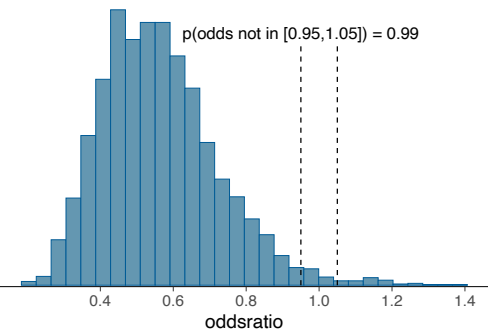
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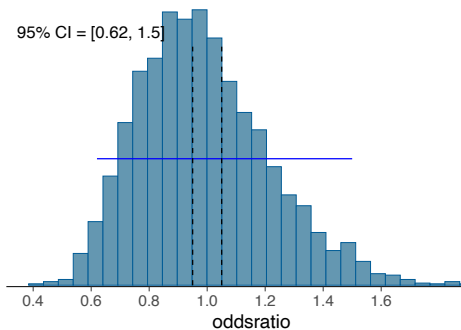
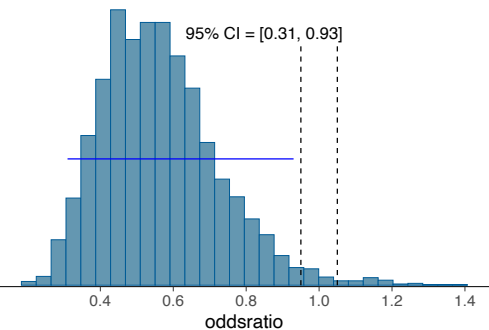
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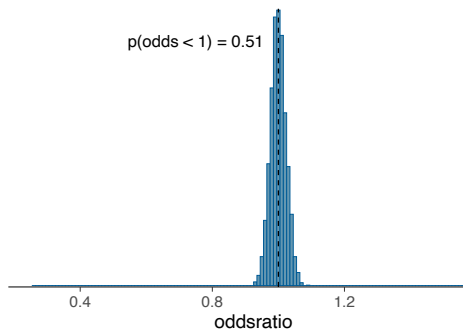
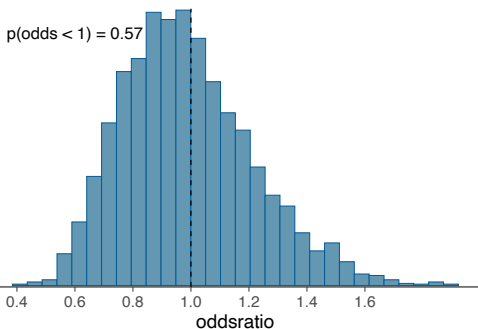


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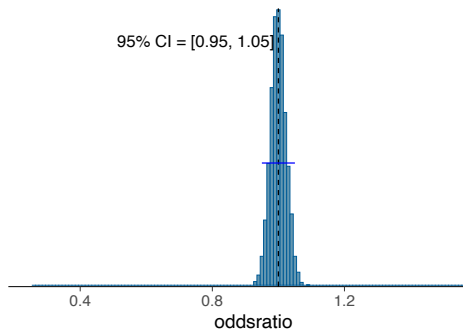
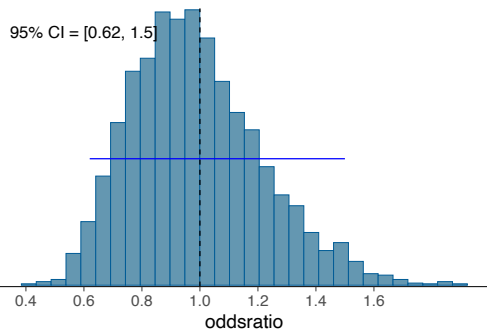
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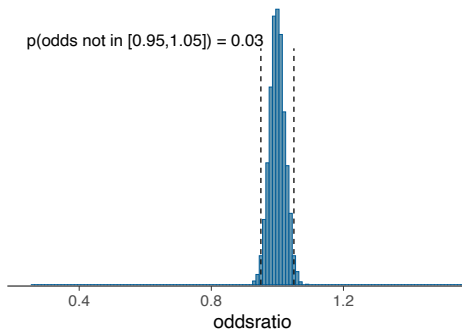
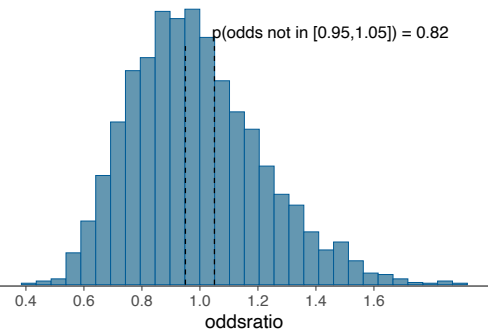
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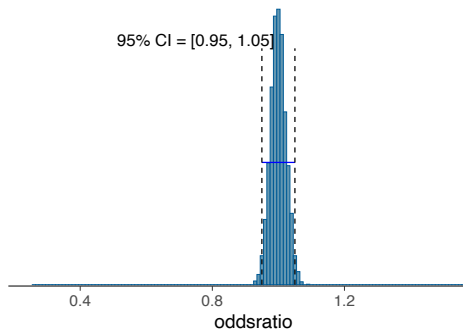
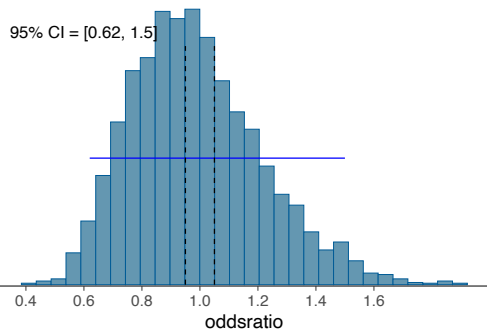
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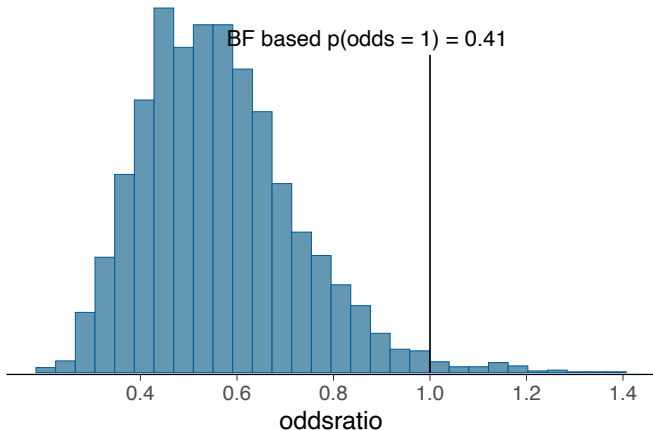
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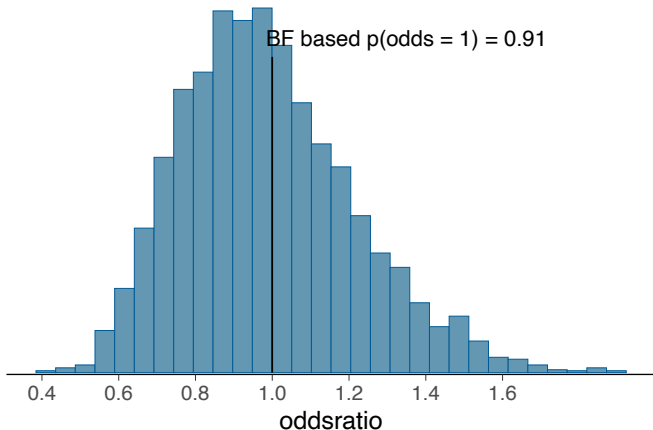
- Bayes factor
  - null model has, e.g., the treatment effect fixed to 0
  - assumes that there is non-zero probability that the treatment effect can be exactly zero
  - requires posterior inference for the null model, too



with `bridgesampling` package, see also BDA3 13.10

# Bayesian hypothesis testing

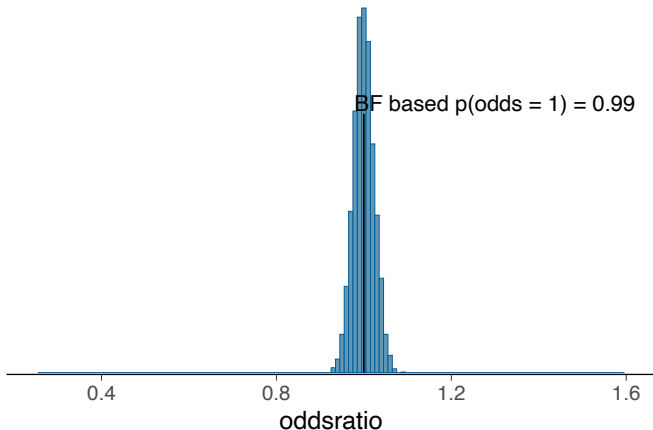
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# Bayesian hypothesis testing

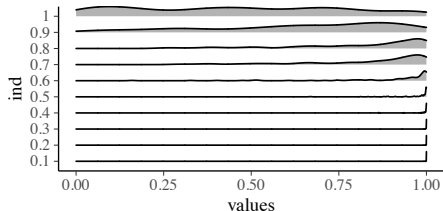
- Predictive performance
  - is there difference in predictive performance with, e.g., treatment effect fixed to zero or unknown treatment effect
  - requires posterior inference for the null model or projection from the full to null
  - looking at the posterior is better if parameters are independent

In the beta blockers example

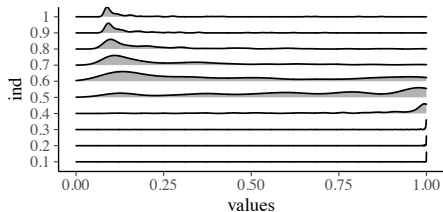
- Leave-one-group-out is not sensible as there are only two groups
- Leave-one-person-out works, but is less efficient than looking at the posterior (see <https://avehtari.github.io/modelselection/betablockers.html>)

# Simulation experiment

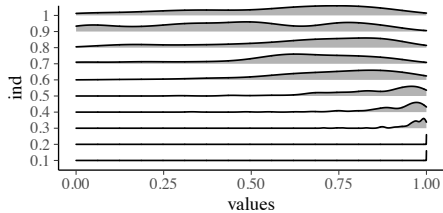
$p(\text{odds} < 1)$



Marginal likelihood  
comparison



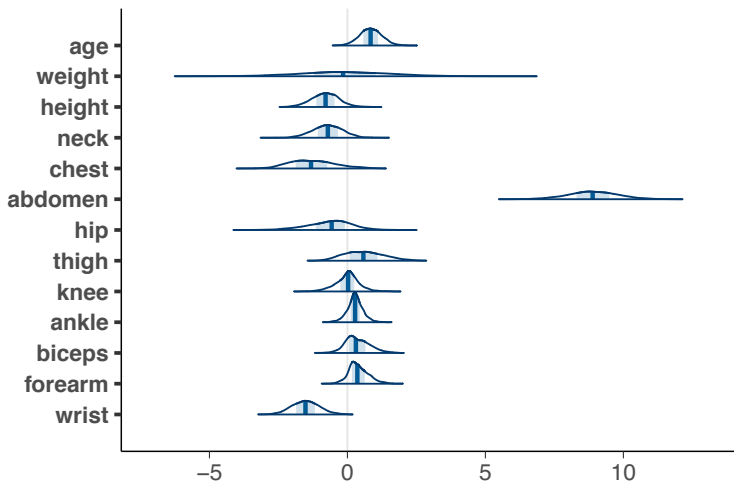
LOO comparison



# Hypothesis testing and posterior dependencies

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

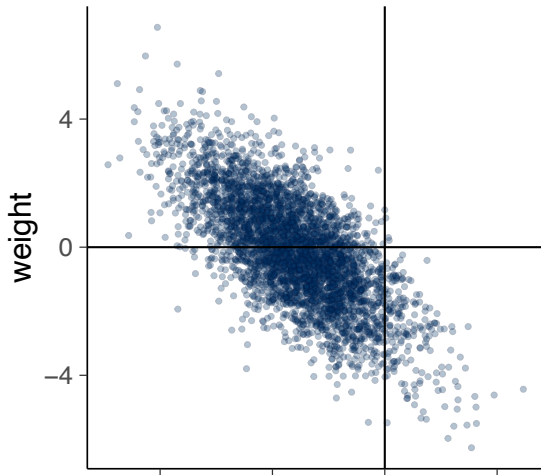
Marginal posteriors of coefficients



# Hypothesis testing and posterior dependencies

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

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 ( $p=0.99$ )

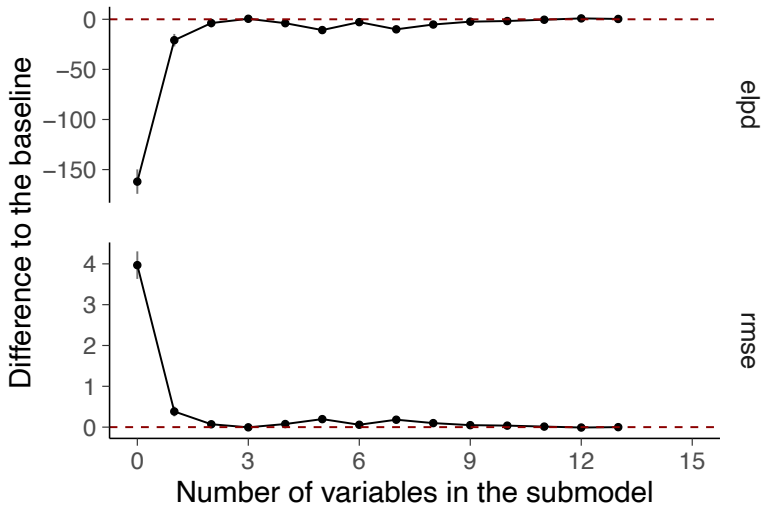
In bodyfat example, starting from model  $y \sim \text{abdomen}$

- 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>
ANOVA	hier. model	<code>stan_glm(y ~ 1 + (1   gid))</code>
...		

possible to extend, e.g., with group specific variances and and different distributions such *t*- or Poisson distribution

See longer list and illustrations (with `lm`) at

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

and

in the forthcoming *Regression and other stories* book

## Chapter 8: Modelling accounting for data collection

Highly recommended to read. Very informative, but also dense chapter.

- We need to model the data collection unless it is ignorable
- We need to know when data collection is ignorable
- Data collection
  - Sample surveys
  - Designed experiments
  - Randomization
  - Observational studies
  - Censoring and truncation



## Chapter 14: Introduction to regression models

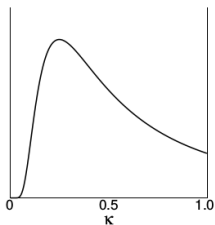
- Justification of conditional modeling
  - if joint model factorizes  $p(y, x|\theta, \phi) = p(y|x, \theta)p(x|\phi)$  we can model just  $p(y|x, \theta)$
- Gaussian linear model with conjugate prior
  - the conditional posterior is multivariate normal
  - with fixed prior on weights, the joint posterior is N-Inv- $\chi^2$
  - these properties are sometimes useful and thus good to know, but with probabilistic programming less often needed
- Bit on causal analysis (see much more in ROS)
- Assembling matrix of explanatory variables
  - identifiability, collinearity, nonlinear relations, indicator and categorical variables, interactions
  - variable selection is not much discussed (see lectures 9.2, 9.3)
- Regularization
  - not much discussed (see more in lecture 9.3 and e.g. [https://betanalpha.github.io/assets/case\\_studies/bayes\\_sparse\\_regression.html](https://betanalpha.github.io/assets/case_studies/bayes_sparse_regression.html))
- Unequal variances and correlations

## Lasso and Bayesian lasso

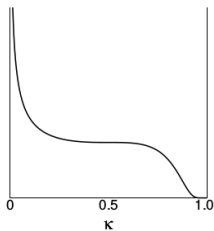
- Lasso is penalized maximum likelihood linear regression, with L1 one penalty where the amount penalty is adapted
  - penalized maximum likelihood finds the mode given the penalty parameter, and is almost the same as maximum a posteriori
  - when the amount of penalty is increased, marginal modes of weak effects go to zero first
  - when the amount of penalty is increased, also the relevant coefficients are shrunk towards zero
  - sometimes relaxed lasso is used, where after variable selection coefficients are re-estimated
- Bayesian lasso uses Laplace distribution as prior
  - Laplace prior is equivalent to L1 penalty
  - but the Bayesian inference includes distribution for parameters and that distribution doesn't shrink to a point at zero, even if the mode would be at zero
  - 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

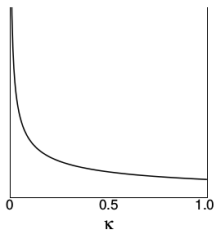
**Laplacian**



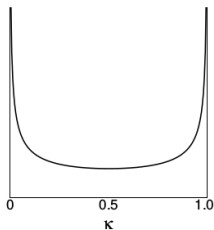
**Student-t**



**Strawderman-Berger**

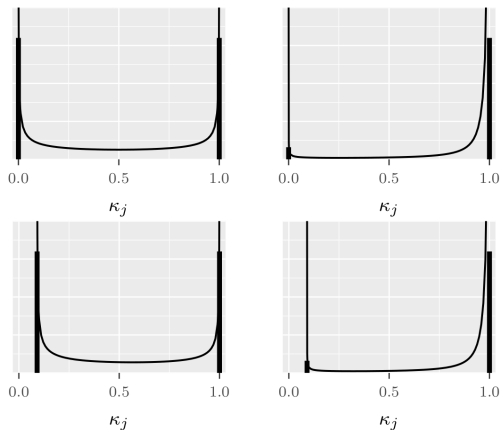


**Horseshoe**



from Carvalho, Polson, Scott (2009).

# Regularized horseshoe



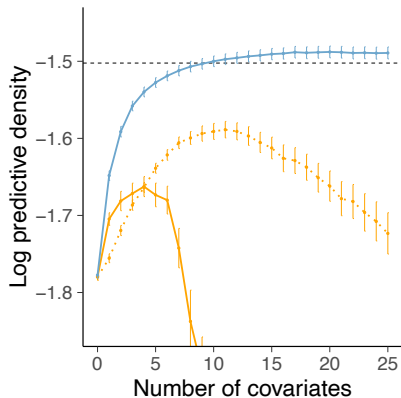
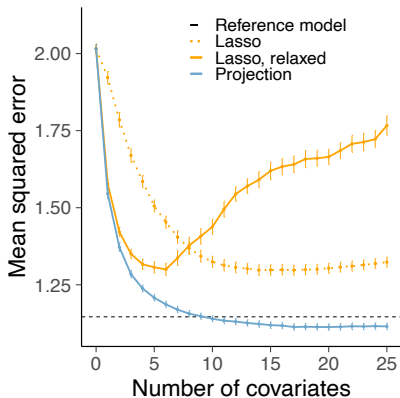
for more see

- 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

Same simulated regression data as in lecture 9,3,  
 $n = 50$ ,  $p = 500$ ,  $p_{\text{rel}} = 150$ ,  $\rho = 0.5$



## Chapter 15: Hierarchical linear models

- Since you know hierarchical models, theory is easy
- With probabilistic programming computation is also easy
  - BDA3 discusses some other computational issues
  - section on transformations for HMC is relevant (see also Stan user guide 21.7 Reparameterization)
- Fixed, random, and mixed effects models
  - we don't recommend using these terms, but they are so popular that it's useful to know them

$y \sim 1 + x$	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`)

## Chapter 16: Generalized linear models

- Bioassay model is an example of GLM
- Components:
  1. The linear predictor  $\eta = X\beta$
  2. The link function  $g(\cdot)$  and  $\mu = g^{-1}(\eta)$
  3. Outcome distribution model with location parameter  $\mu$ 
    - the distribution can also depend on dispersion parameter  $\phi$
    - originally just exponential family distributions (e.g. Poisson, binomial, negative-binomial), which all have natural location-dispersion parameterization
    - after MCMC made computation easy, GLM can refer to models where outcome distribution is not part of exponential family and dispersion parameter may have its own latent linear predictor
- 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 →  $t$ -distribution
  - Poisson → negative-binomial
  - binomial → beta-binomial
  - probit → logistic / robit
- Computation with MCMC easy
  - posterior can be multimodal
  - rstanarm doesn't have  $t$ -distribution for outcome, but brms has



## Chapter 18: Models for missing data

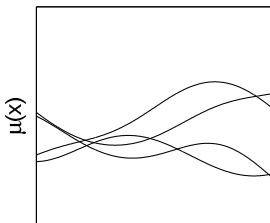
- Extends the data collection modelling from Chapter 8
- Useful terms
  - Missing completely at random (MCAR)  
missingness does not depend on missing values or other observed values (including covariates)
  - Missing at random (MAR)  
missingness does not depend on missing values but may depend on other observed values (including covariates)
  - Missing not at random (MNAR)  
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

- Gaussian process is
  - infinite dimensional extension of normal distribution
  - useful prior for non-linear functions
  - for any finite number of variables, the marginal is multivariate normal  $f_1, \dots, f_n \sim N(\mu(x_1, \dots, x_n), K(x_1, \dots, x_n))$
- Often a priori  $\mu = 0$
- Prior for smooth non-linear functions, e.g. with

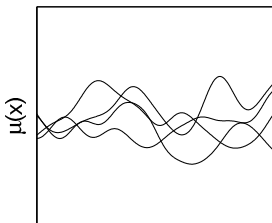
$$k(x, x') = \tau^2 \exp\left(-\frac{|x-x'|^2}{2l^2}\right)$$

$\tau=1/2, l=2$



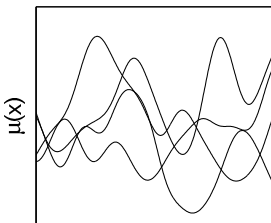
$x$

$\tau=1/4, l=1/2$



$x$

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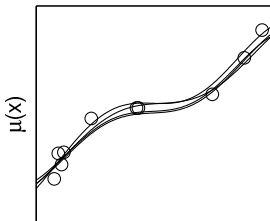
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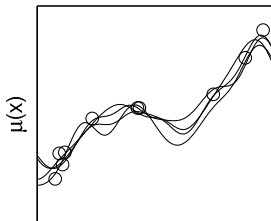
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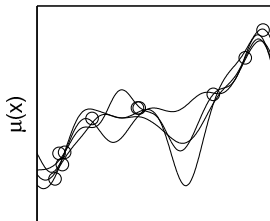
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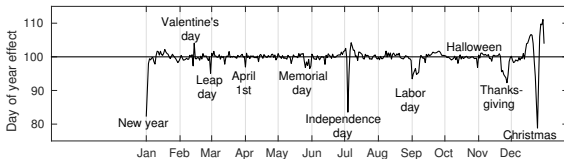
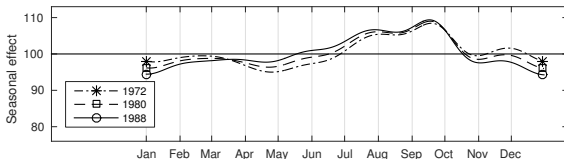
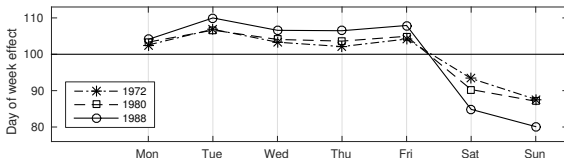
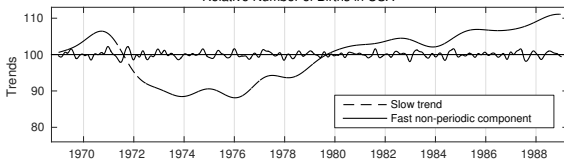
## Chapter 21: Gaussian process models

- 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

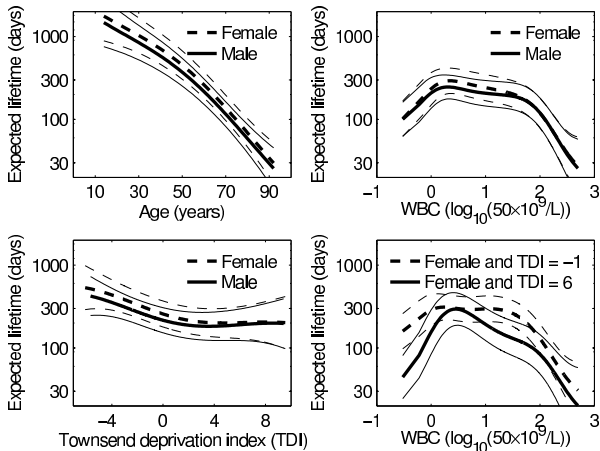
$$y_t(t) = f_1(t) + f_2(t) + f_3(t) + f_4(t) + f_5(t) + \epsilon_t$$

Relative Number of Births in USA



## 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)
- In case of non-Gaussian outcome models, sampling of latent variables can be slow (Laplace integration over the latents coming)

## GPs in Stan

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- Stan has some built-in covariance functions (and soon GPU support)
- 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"))`