Outline

Last week

- What is cross-validation
- LOO-PIT checking
- Fast cross-validation with PSIS
- LOO model comparison and selection (elpd_diff, se)

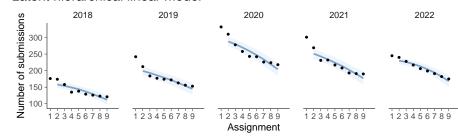
This week

- Model comparison with LOO-CV
- When is cross-validation applicable?
- K-fold cross-validation
- Related methods (WAIC, *IC, BF)
- Hypothesis testing
- Potential overfitting

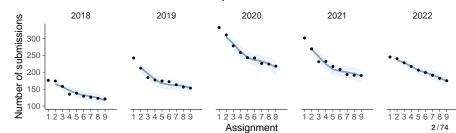
Student retention – Posterior predictive distributions

with tidybayes

Latent hierarchical linear model



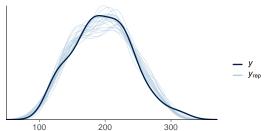
Latent hierarchical linear model + spline



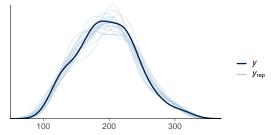
Student retention - Marginal PPC

pp_check(fit, ndraws=100)

Latent hierarchical linear model

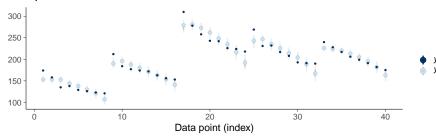


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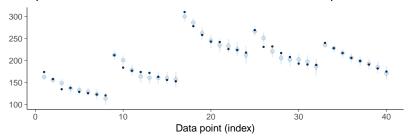


Student retention - LOO intervals

LOO predictive intervals - latent hierarchical linear



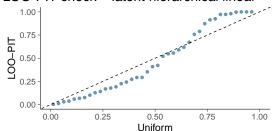
LOO predictive intervals - latent hierarchical linear + spline



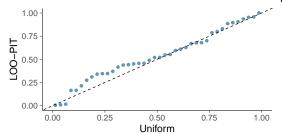
Student retention – LOO-PIT checking

pp_check(fit, type = "loo_pit_qq", ndraws=4000)

LOO-PIT check - latent hierarchical linear



LOO-PIT check - latent hierarchical linear + spline



Student retention – R^2

Latent hierarchical linear vs. latent hierarchical linear + spline

```
> loo_R2(fit4) |> round(digits=2)
    Estimate Est.Error Q2.5 Q97.5
R2     0.92     0.02 0.88     0.95
> loo_R2(fit6) |> round(digits=2)
    Estimate Est.Error Q2.5 Q97.5
R2     0.97     0.01 0.95     0.98
```

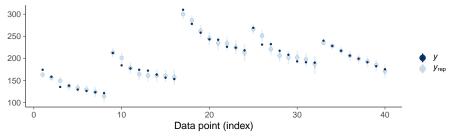
 R^2 measures the goodness of the mean of the predictive distribution

Gelman, Goodrich, Gabry, and Vehtari (2019). R-squared for Bayesian regression models. *The American Statistician*, 73(3):307-309.

- information theoretical goodness of the whole distribution
- elpd = expected log predictive density (probability)
- elpd_loo = estimated with LOO predictive densities / probs $\sum_{n=1}^{N} \log p(y_i|x_i,x_{-i},y_{-i})$

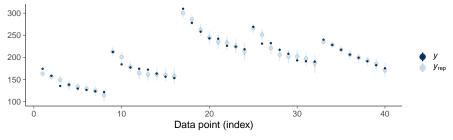
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LOO predictive intervals - latent hierarchical linear + spline



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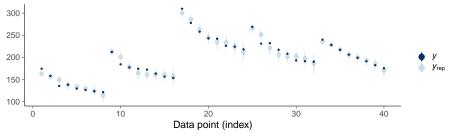
LOO predictive intervals - latent hierarchical linear + spline



-8.4 -5.6 -2.9 -2.9 -2.8 -3.0 -4.0 -3.2 -3.9 -3.2 -3.4 -3.2 -2.9 -3.9 -3.4 -3.4 -3.2 -2.7 -2.8 -3.1 -2.5 -2.8 -2.9 -3.4 -5.4 -3.7 -3.1 -3.3 -3.5 -3.2 -3.5 -3.5 -6.6 -3.8 -3.7 -3.4 -2.5 -2.8 -2.9 -3.3

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LOO predictive intervals - latent hierarchical linear + spline



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Latent hierarchical linear + spline

```
> loo(fit6)
```

Computed from 4000 by 40 log-likelihood matrix

```
Estimate SE elpd_loo -141.7 7.2 p_loo 10.9 2.5
```

Latent hierarchical linear + spline

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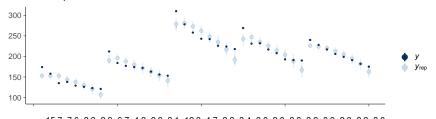
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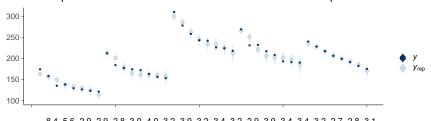
```
Estimate SE elpd_loo -184.3 17.3 p_loo 24.3 5.8
```

LOO predictive intervals - latent hierarchical linear

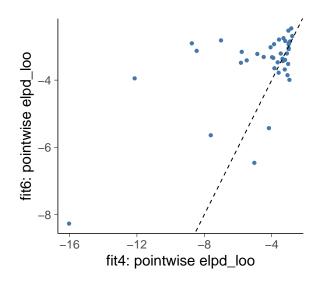


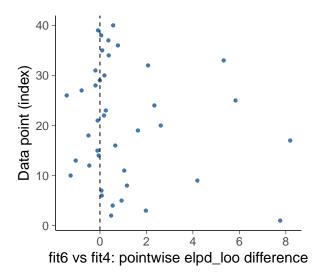
 $-2.9 - 3.3 - 3.0 - 4.6 - 4.3 - 3.3 - 3.0 - 4.0 - 3.0 - 5.6 - 3.6 - 5.4 - 4.9 - 3.6 - 3.9 - 5.2 - 2.7 - 3.7 - 3.0 - 4.1 \quad \sum = -184.3$

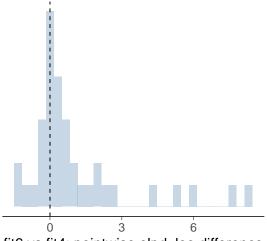




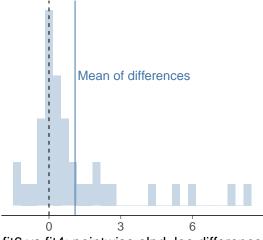
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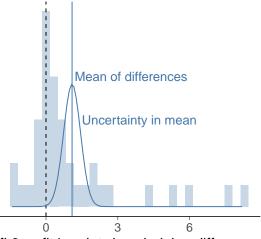




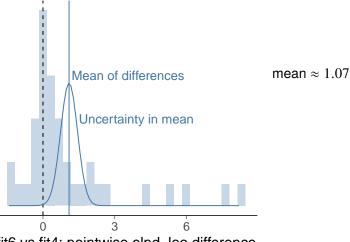
fit6 vs fit4: pointwise elpd_loo difference



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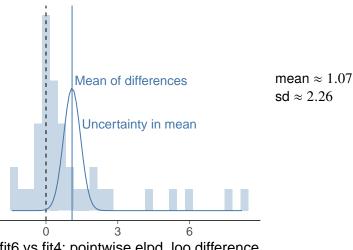


fit6 vs fit4: pointwise elpd_loo difference



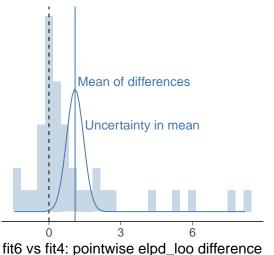
fit6 vs fit4: pointwise elpd_loo difference

Latent hierarchical linear (fit4) vs latent hierarchical linear + spline (fit6)



fit6 vs fit4: pointwise elpd_loo difference

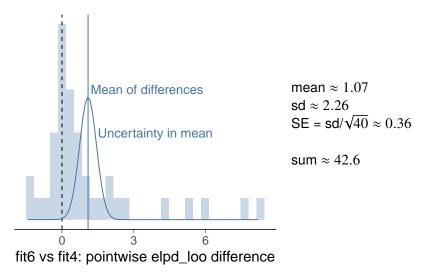
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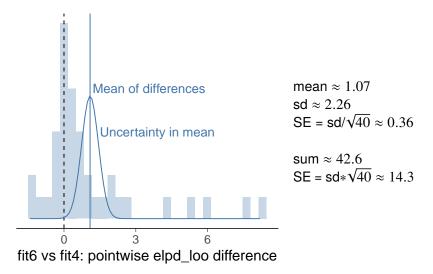


SE = $sd/\sqrt{40} \approx 0.36$

mean ≈ 1.07

 $sd \approx 2.26$





Latent hierarchical linear + spline

Latent hierarchical linear

1. The models make very similar predictions

2. The models are misspecified with outliers in the data

3. The number of observations is small

Sivula, Magnusson, Matamoros, and Vehtari (2022). Uncertainty in Bayesian leave-one-out cross-validation based model comparison. *arXiv:2008.10296v3*.

- 1. The models make very similar predictions
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 - in nested case, the skewness favors the simpler model
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- 3. The number of observations is small
 - in nested case the skewness favors the simpler model
 - any inference with small n is difficult
 - if |elpd_loo| > 4, model is well specified, and *n* > 100 then the normal approximation is good

Sivula, Magnusson, Matamoros, and Vehtari (2022). Uncertainty in Bayesian leave-one-out cross-validation based model comparison. *arXiv:2008.10296v3*.

- Log score is not easily interpretable
- but is information theoretically good utility for the goodness of the whole distribution
- and thus is useful in model comparison

- Interpretation in discrete case
 - log probability

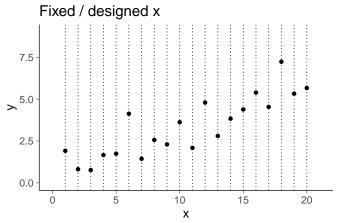
- Interpretation in discrete case
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 - compare to guessing uniformly from the data range [121,310] having $1/(310-121+1)\approx 0.5\%$ probability (log score -210)
- Interpretation in continuous case
 - can be compared to a simple reference distribution

Assumptions about the future observations

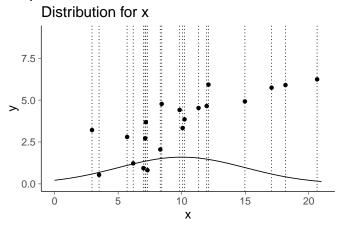


elpd_loo =
$$\sum_{i=1}^{20} \log p(y_i \mid x_i, x_{-i}, y_{-i}) \approx -29.5$$

SE = $\operatorname{sd}(\log p(y_i \mid x_i, x_{-i}, y_{-i})) \cdot \sqrt{20} \approx 3.3$

LOO is ok for fixed / designed x. SE is uncertainty about $y \mid x$.

Assumptions about the future observations

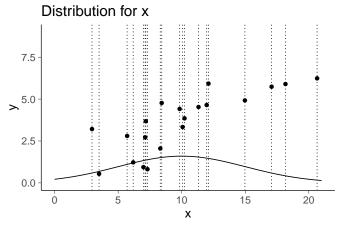


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LOO is ok for random x. SE is uncertainty about $y \mid x$ and x.

Assumptions about the future observations

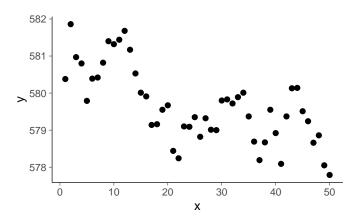


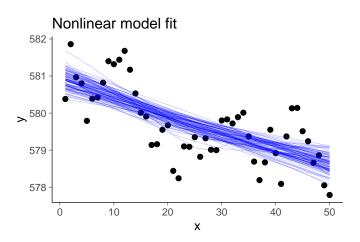
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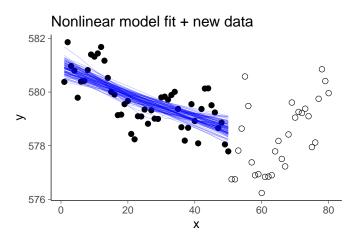
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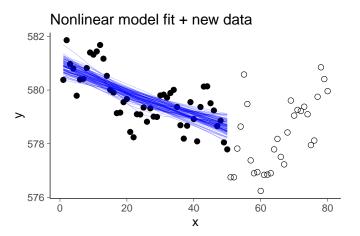
LOO is ok for random x. SE is uncertainty about $y \mid x$ and x.

Covariate shift handled with importance weighting or modelling see Vehtari & Ojanen (2012) and CV-FAQ

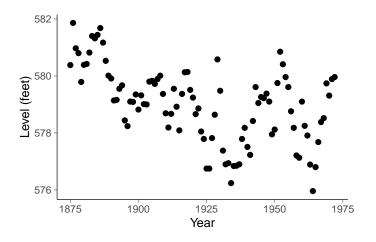




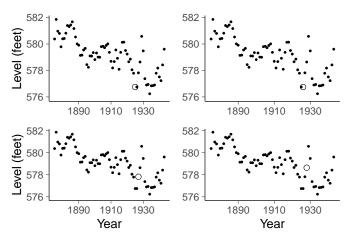




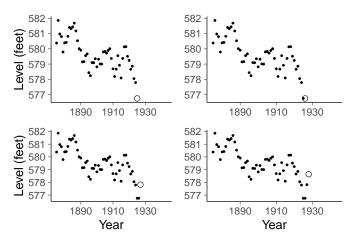
 Extrapolation is more difficult item<5> In high dimensional case mostly extrapolation



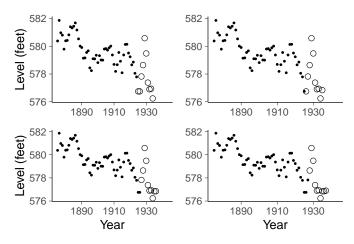
Can LOO or other cross-validation be used with time series?



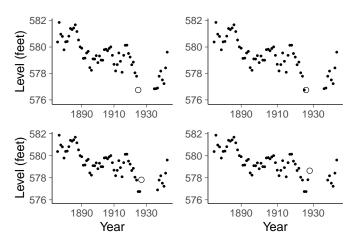
Leave-one-out cross-validation is ok for assessing conditional model



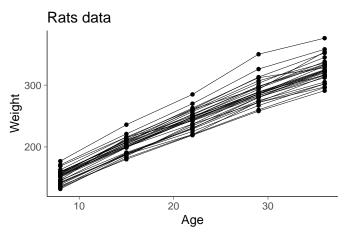
Leave-future-out (LFO) cross-validation is better for predicting future



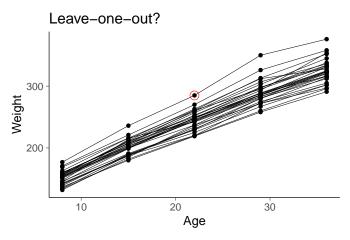
m-step-ahead cross-validation is better for predicting further future

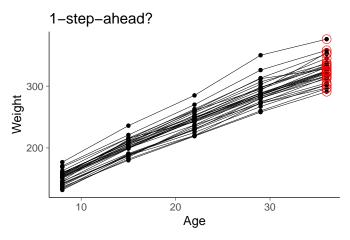


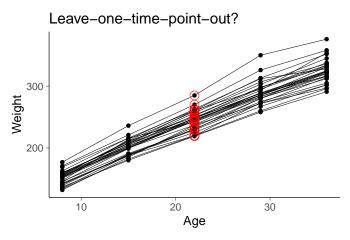
m-step-ahead leave-a-block-out cross-validation

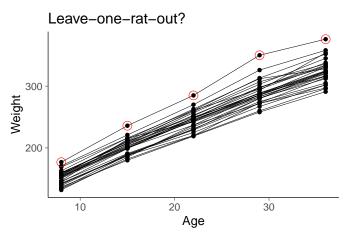


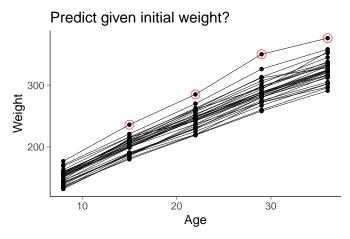
Can LOO or other cross-validation be used with hierarchical data?











Summary of data generating mechanisms and prediction tasks

- You have to make some assumptions on data generating mechanism
- · Use the knowledge of the prediction task if available
- Cross-validation can be used to analyse different parts, even if there is no clear prediction task

Pareto smoothed importance sampling CV variants

- PSIS-LOO for hierarchical models
 - leave-one-group out is challenging for PSIS-LOO
 - Stan demo of the challenges and integrated LOO at https://users.aalto.fi/~ave/modelselection/roaches.html
 - see also Merkel, Furr and Rabe-Hesketh (2018)

Pareto smoothed importance sampling CV variants

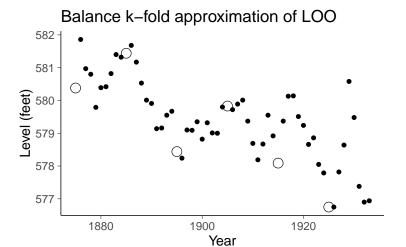
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 - mc-stan.org/loo/articles/loo2-non-factorizable.html

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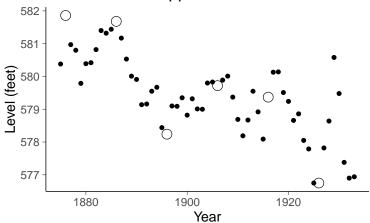
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- PSIS-LOO for non-factorized models
 - mc-stan.org/loo/articles/loo2-non-factorizable.html
- PSIS-LOO for time series
 - Approximate leave-future-out cross-validation (LFO-CV) mc-stan.org/loo/articles/loo2-lfo.html

K-fold cross-validation

- K-fold cross-validation can approximate LOO
 - the same use cases as with LOO
- K-fold cross-validation can be used for hierarchical models
 - good for leave-one-group-out
- K-fold cross-validation can be used for time series
 - with leave-block-out



Balance k-fold approximation of LOO



Random k-fold approximation of LOO 582 -

1900

Year

1920

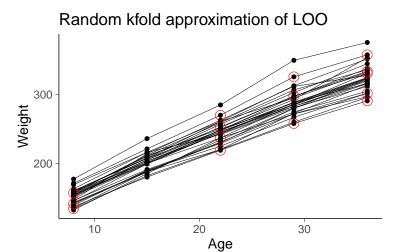
581

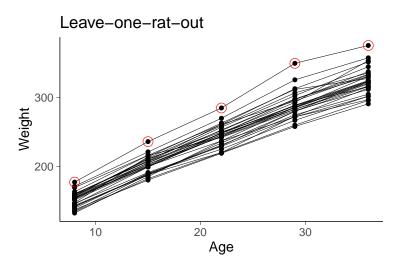
Tevel (feet) 24.00 (feet) 25.00 (feet) 25.00

578

577

1880





K-fold-CV code

- RStan, CmdStanR
 See vignette http://mc-stan.org/loo/articles/loo2-elpd.html
- RStanARM, brms kfold(fit)
- Alternative data divisions

```
kfold_split_random()
kfold_split_balanced()
kfold_split_stratified()
```

looic?

```
> loo(fit6)
```

Computed from 4000 by 40 log-likelihood matrix

```
Estimate SE
elpd_loo -141.7 7.2
p_loo 10.9 2.5
looic 283.4 14.4
```

Monte Carlo SE of elpd_loo is 0.1.

- loo output shows also looic
- for historical non-Bayesian reasons it's -2 * elpd_loo
 - connection to deviance and information criteria
 - you can just ignore it (I'd prefer it would not be shown)

Information criteria estimate predictive performance, too

AIC uses maximum likelihood estimate for prediction

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- AIC uses maximum likelihood estimate for prediction
- DIC uses posterior mean for prediction
- BIC is a simple approximation for marginal likelihood
- TIC, NIC, RIC, PIC, BPIC, QIC, AICc, ...
- WAIC is the only Bayesian information criterion

WAIC vs PSIS-LOO

WAIC has the same target and assumptions as LOO

Vehtari, Gelman and Gabry (2017). Practical Bayesian model evaluation using leave-one-out cross-validation and WAIC. *Statistics and Computing*, 27(5):1413–1432

WAIC vs PSIS-LOO

- WAIC has the same target and assumptions as LOO
- PSIS-LOO is more accurate

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- Multiplying by -2 doesn't give any benefit (Watanabe didn't multiply by -2)

Vehtari, Gelman and Gabry (2017). Practical Bayesian model evaluation using leave-one-out cross-validation and WAIC. *Statistics and Computing*, 27(5):1413–1432

Bayes Factor
$$\frac{p(y|M_1)}{p(y|M_2)}$$

Marginal likelihood
$$p(y|M_1) = \int p(y|\theta, M_1)p(\theta|M_1)d\theta$$

Bayes Factor $\frac{p(y|M_1)}{p(y|M_2)}$

Marginal likelihood $p(y|M_1) = \int p(y|\theta, M_1)p(\theta|M_1)d\theta$

Marginal likelihood with chain rule:

$$p(y|M_1) = p(y_1|M_1)p(y_2|y_1, M_1), \dots, p(y_n|y_1, \dots, y_{n-1}, M_1)$$

Bayes Factor
$$\frac{p(y|M_1)}{p(y|M_2)}$$

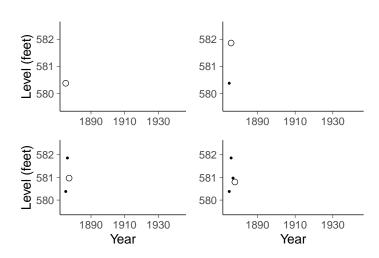
Marginal likelihood
$$p(y|M_1) = \int p(y|\theta, M_1)p(\theta|M_1)d\theta$$

Marginal likelihood with chain rule:

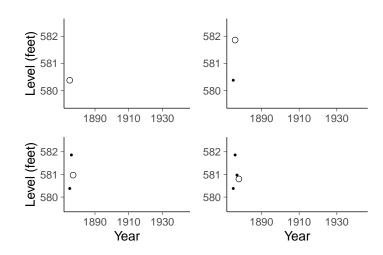
$$\begin{split} p(y|M_1) &= p(y_1|M_1)p(y_2|y_1,M_1), \dots, p(y_n|y_1,\dots,y_{n-1},M_1) \\ \text{where} \\ p(y_1|M_1) &= \int p(y_1|\theta,M_1)p(\theta|M_1)d\theta \\ p(y_2|y_1,M_1) &= \int p(y_2|\theta,M_1)p(\theta|y_1,M_1)d\theta \\ \dots \\ p(y_n|y_1,\dots,y_{n-1},M_1) &= \int p(y_n|\theta,M_1)p(\theta|y_1,\dots,y_{n-1},M_1)d\theta \end{split}$$

 Like leave-future-out 1-step-ahead cross-validation but starting with 0 observations

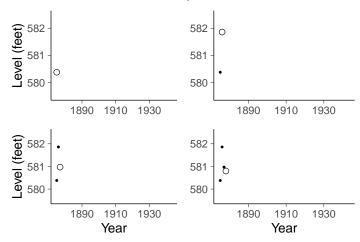
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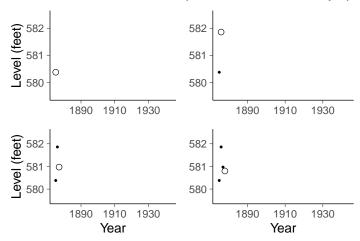
- Like leave-future-out 1-step-ahead cross-validation but starting with 0 observations
 - which makes it very sensitive to prior



- Like leave-future-out 1-step-ahead cross-validation but starting with 0 observations
 - which makes it very sensitive to prior and
 - unstable in case of misspecified models



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- Oelrich, Ding, Magnusson, Vehtari, and Villani (2020). When are Bayesian model probabilities overconfident? arXiv:2003.04026.

Predictive model selection

- Student retention
 - latent hierarchical linear vs.
 - latent hierarchical linear + spline

is a good example where predictive model selection is useful

Sometimes cross-validation is not needed

- In a simple nested case, often easier and more accurate to analyze posterior distribution of more complex model directly
 - instead of comparing Model 1: $y \sim \operatorname{normal}(\alpha, \sigma)$ vs Model 2: $y \sim \operatorname{normal}(\alpha + \beta x, \sigma)$ look at the posterior of β directly

Common statistical tests as Bayesian models

Most common statistical tests are linear models

test	model	formula
t-test	mean of data	y ~ 1
paired t-test	mean of diffs	$(y1 - y2) \sim 1$
Pearson correl.	linear model	y ~ 1 + x
two-sample t-test	group means	y ~ 1 + gid
ANOVA	hier. model	$y \sim 1 + (1 gid)$

. . .

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 - and go beyond named tests

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- Possible to extend, e.g., with group specific variances and and different distributions such t- or Poisson distribution
 - and go beyond named tests
- See longer list and illustrations (with lm) at https://lindeloev.github.io/tests-as-linear/ and with rstanarm in Regression and other stories

Beta blockers

- An experiment was performed to estimate the effect of beta-blockers on mortality of cardiac patients
- A group of patients were randomly assigned to treatment and control groups:
 - out of 674 patients receiving the control, 39 died
 - out of 680 receiving the treatment, 22 died

Beta blockers

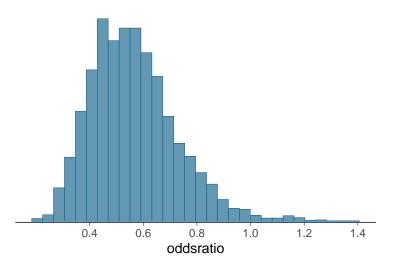
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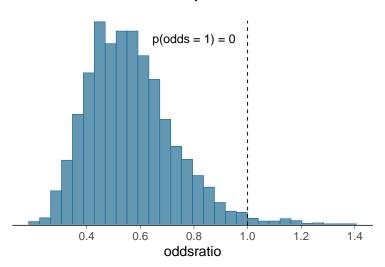
Posterior inference

- Instead of model selection, report full posterior and
 - · compare to expert information
 - · combine with utility/cost function



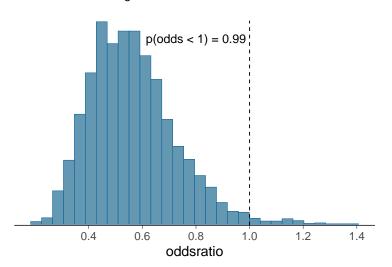
Posterior inference

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



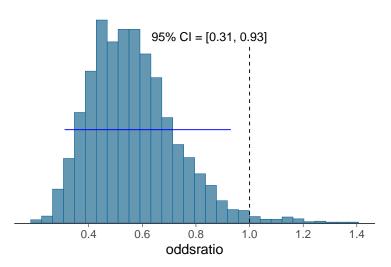
Posterior inference

- Instead of model selection, report full posterior
 - for continuous posterior we could report the probability that we know the sign of the effect

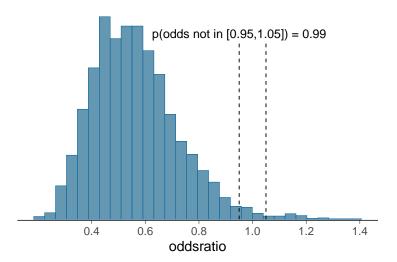


- Sometimes people want to make a dichotomous choice
 - model selection
 - hypothesis testing

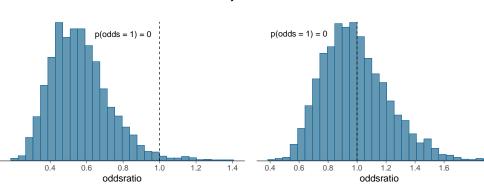
- Instead of model selection, report full posterior and
 - for continuous posterior some people compare whether posterior interval includes null case



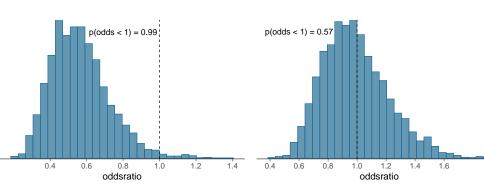
- Equivalence testing (region of practical equivalence)
 - what is the probability that the effect is closer than ϵ to null, where ϵ is based on what is practically useful effect size



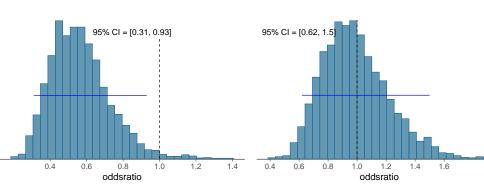
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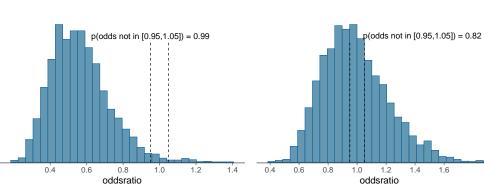
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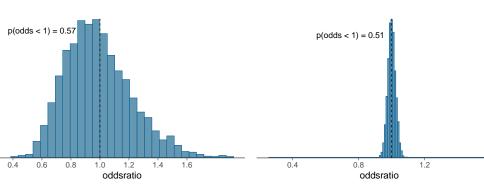


- Instead of hypothesis testing, report full posterior
 - region of practical equivalence (ROPE)

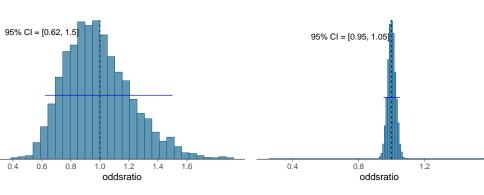


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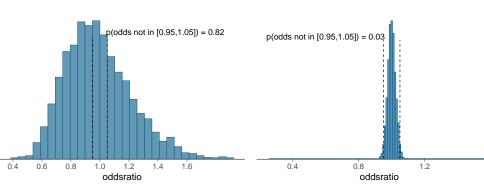
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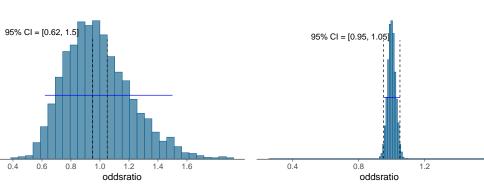
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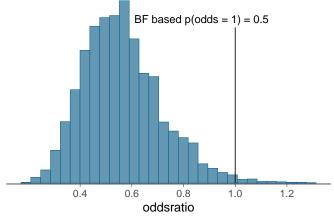
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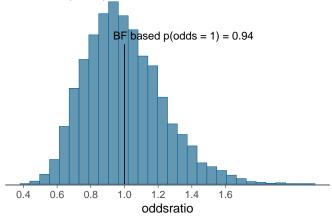
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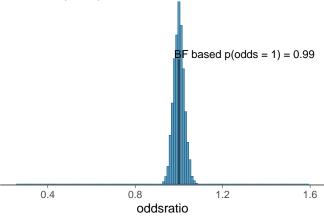
- Bayes factor
 - null model has, e.g., the treatment effect fixed to 0
 - assumes that there is non-zero probability that the treatment effect can be exactly zero (point mass)
 - requires posterior inference for the null model, too



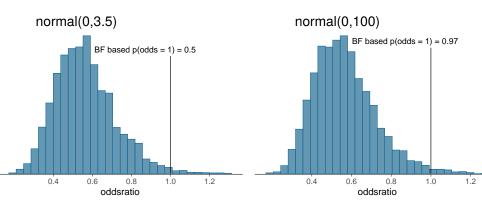
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- Bayes factor
 - sensitive to the prior choice even when the posterior is not



with bridgesampling package, see also BDA3 13.10

- 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

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In the beta blockers example

 Leave-one-person-out works, but is less efficient than looking at the posterior (see https://users.aalto.fi/~ave/modelselection/betablockers.html)

```
> loo_compare(loo(fitb1),loo(fitb2))
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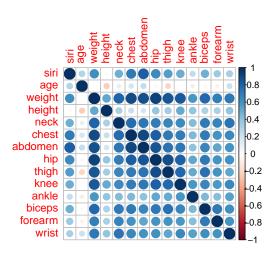
 For another similar, but more elaborate example, see https: //users.aalto.fi/~ave/casestudies/Nabiximols/nabiximols.html

Bodyfat: many predictors

- Predict bodyfat percentage
- The reference value (siri) is obtained by immersing person in water. n = 251.
- Which measurements to use in the future?

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Prediction

· Goal: prediction

Prediction

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- Use all the predictors and sensible prior

Prediction

- Goal: prediction
- Use all the predictors and sensible prior
 - no model selection needed

Predictive performance based variable selection

- Goal:
 - minimize future measurement cost
 - easier explainability of the model

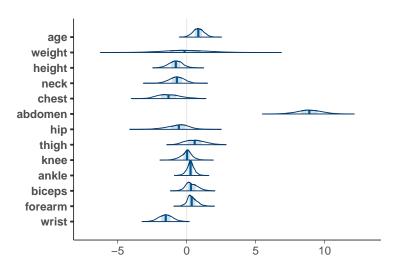
Predictive performance based variable selection

- Goal:
 - minimize future measurement cost
 - easier explainability of the model
- Select the minimal number of covariates with similar predictive performance as the full model

Hypothesis testing and posterior dependencies

Looking at the marginal posterior $p(\beta < 0)$ can be misleading when there are many parameters

Marginal posteriors of coefficients in bodyfat example

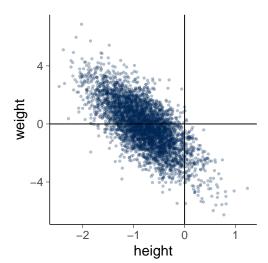


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Hypothesis testing and posterior dependencies

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

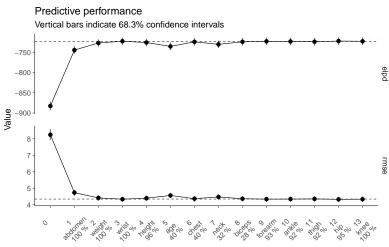
- BF in favor of removing weight (p=0.92)
- LOO in favor of removing weight (p=0.99)

In bodyfat example, starting from model y ~ abdomen

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

Predictive performance based variable selection

Projection predictive variable selection selects the minimal set of variables with similar predictive performance as the full model



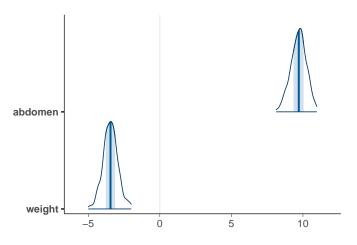
Submodel size (number of predictor terms)

Corresponding predictor from full–data predictor ranking

Corresponding main diagonal element from CV ranking proportions matrix

Projected posterior

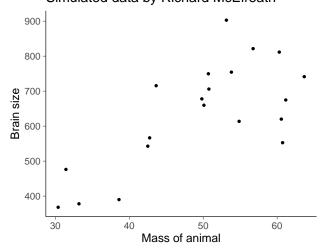
Projection predictive variable selection selects the minimal set of variables with similar predictive performance as the full model



More about projpred in the end of the course

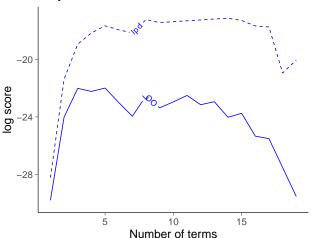
- Classic example is polynomial model with increasing number of components
 - overfits also with Bayesian inference and weak priors

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Polynomial basis functions



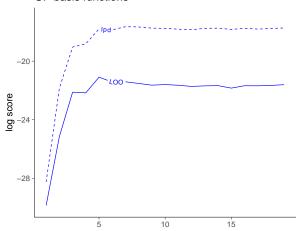
- Gaussian process can be used as a prior on function space
 - GP can be approximated with basis functions

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 - more basis functions makes the approximation more accurate, but doesn't inflate the prior on function space

Model is not needed to avoid overfitting

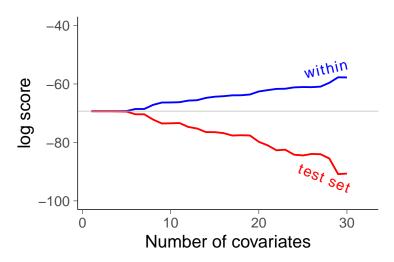
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GP basis functions

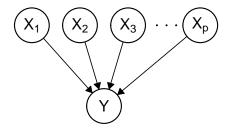


logistic regression: 30 **completely irrelevant** variables, 100 observations

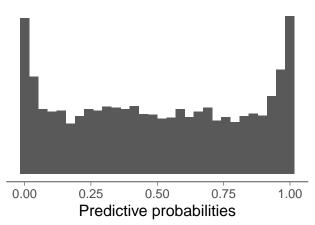
logistic regression: 30 **completely irrelevant** variables, 100 observations



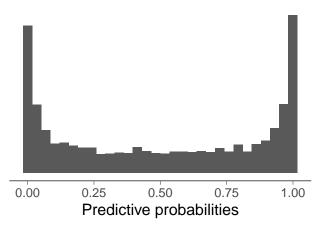
N(0,3) prior on each coefficient



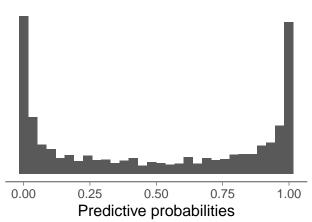
N(0,3) prior on each coefficient 1 variable



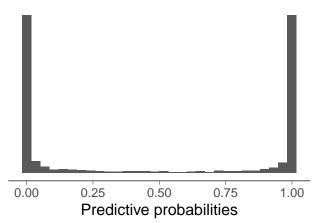
N(0,3) prior on each coefficient 2 variables



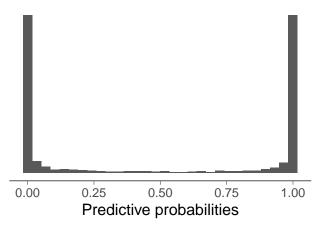
N(0,3) prior on each coefficient 3 variables



N(0,3) prior on each coefficient 30 variables



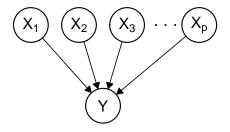
N(0,3) prior on each coefficient 30 variables



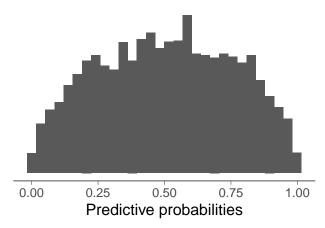
A weak prior on parameters can be a strong prior on predictions that favors overfitting

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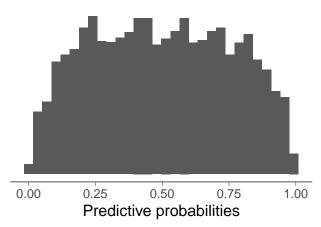
 $N(0,\frac{1}{\sqrt{p}})$ prior on each coefficient



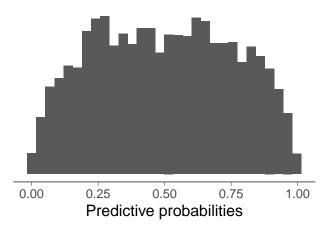
 $N(0, \frac{1}{\sqrt{p}})$ prior on each coefficient 1 variable



 $N(0, \frac{1}{\sqrt{p}})$ prior on each coefficient 2 variables

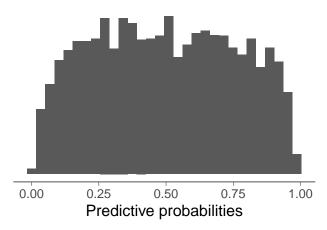


 $N(0, \frac{1}{\sqrt{p}})$ prior on each coefficient 3 variables



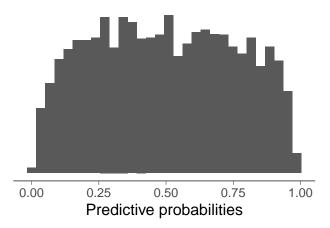
Better priors

 $N(0, \frac{1}{\sqrt{p}})$ prior on each coefficient 30 variables



Better priors

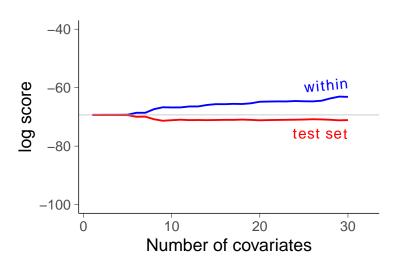
 $N(0, \frac{1}{\sqrt{p}})$ prior on each coefficient 30 variables



Prior on predictions (almost) fixed when the model gets bigger

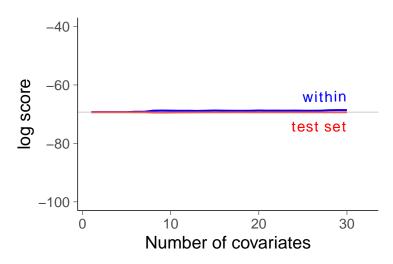
Better priors, no overfitting

logistic regression: 30 **completely irrelevant** variables, 100 observations, $N(0, \frac{1}{\sqrt{p}})$ prior



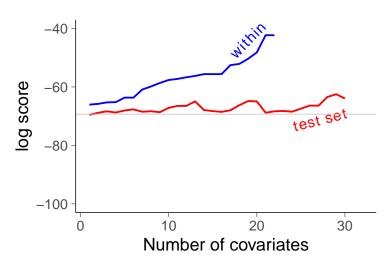
Better priors, no overfitting

logistic regression: 30 **completely irrelevant** variables, 100 observations, regularized horseshoe prior



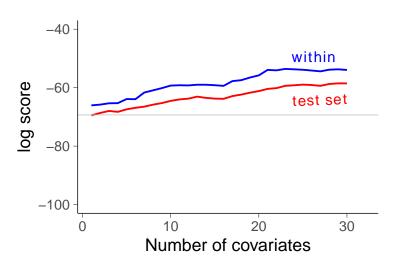
Many weak effects, wide prior on parameters

logistic regression: 30 **weakly relevant** variables, 100 observations, N(0,3) prior



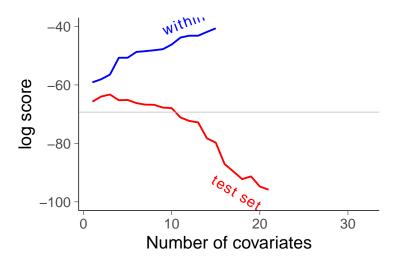
Many weak effects, better prior

logistic regression: 30 **weakly relevant** variables, 100 observations, $N(0, \frac{1}{\sqrt{p}})$ prior



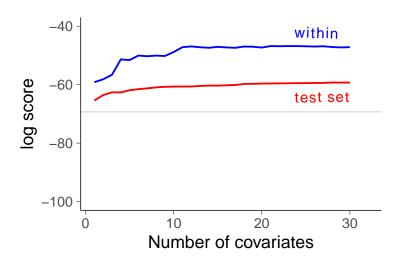
Correlating variables, wide prior on parameters

logistic regression: 30 **correlating relevant** variables, 100 observations, N(0,3) prior



Correlating variables, better prior

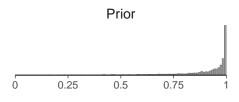
logistic regression: 30 **correlating relevant** variables, 100 observations $N(0, \frac{1}{\sqrt{p}})$ prior

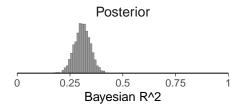


Implied prior on R^2

Regression and Other Stories, Section 12.7 Models for regression coefficients:

Wide prior on coefficients favors overfitting

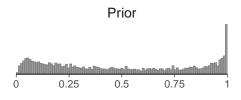


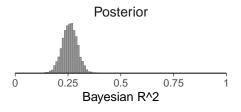


Implied prior on R^2

Regression and Other Stories, Section 12.7 Models for regression coefficients:

Scaled prior on coefficients

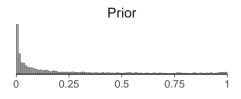


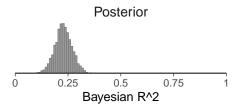


Implied prior on R^2

Regression and Other Stories, Section 12.7 Models for regression coefficients:

Regularized horseshoe prior on coefficients





Better priors

For example:

- · scaled: many weak effects
- regularized horseshoe, R2-D2: only some relevant
- R2-D2: defined directly for R²
- PCA-type: highly correlating variables

$p \gg n$

- With good priors, possible to have more variables than observations
- e.g. p = 22283, n = 85 demonstrated by Piironen, Paasiniemi, Vehtari (2020)

Variable selection

Variable selection

- 1. is not needed to avoid overfitting
- 2. can be used to reduce costs and improve explainability

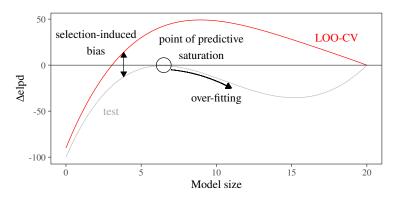
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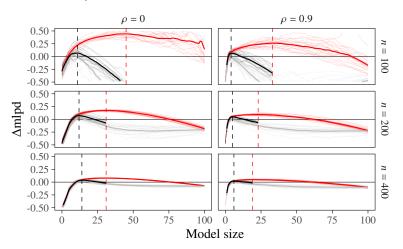
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- Bigger problem if there is a large number of models as in covariate selection

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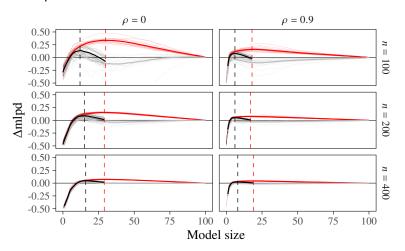
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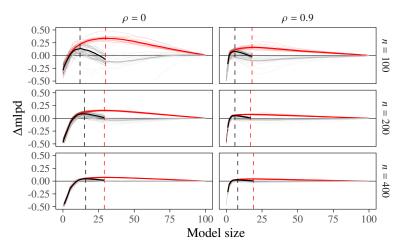
Wide normal prior



R2D2 prior reduces overfit in model selection



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Reminder: variable selection is not needed with good priors to get good predictive performance, but may be useful for other purposes

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- Bayesian stacking may work better than BMA in case of misspecified models or small data
 - Yao, Vehtari, Simpson, and Gelman (2018). Using stacking to average Bayesian predictive distributions (with discussion). Bayesian Analysis, 13(3):917-1003

Cross-validation and model selection

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- Overfitting in selection process is not unique for cross-validation

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