

- Model assessment and selection
 - Measures of predictive accuracy
 - Model selection
- Cross-validation
 - When is LOO applicable
 - PSIS LOO-CV
 - K-fold CVModel Comparison
- Information criteria
- Model averaging and Stacking
- Summary

Bayesian Statistics and Data Analysis Lecture 8b

Måns Magnusson Department of Statistics, Uppsala University Thanks to Aki Vehtari, Aalto University



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Predictive performance

Modeling complex phenomena with models that are simplified

All models are wrong... but some are useful.



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Predictive performance

- Modeling complex phenomena with models that are simplified
 - All models are wrong... but some are useful.
- True predictive performance is found out by using it to make predictions and comparing predictions to true observations
 - external validation



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Predictive performance

- Modeling complex phenomena with models that are simplified
 - All models are wrong... but some are useful.
- True predictive performance is found out by using it to make predictions and comparing predictions to true observations
 - external validation
- Expected predictive performance
 - approximates the external validation



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Goal of model evaluation

• Model choice is a (model-)decision-theoretic problem



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Goal of model evaluation

- Model choice is a (model-)decision-theoretic problem
- Evaluate the utility of a model M for new unseen data \tilde{y} :

$$U = \int u(\tilde{y}) p_{\mathsf{true}}(\tilde{y}) d\tilde{y} \,,$$

where \tilde{y} is an unseen observation generated from the true data generating process $p_{\text{text}}(\tilde{y})$, and y are observed data and $u(\tilde{y})$ is a utility function.



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• The expectation is with respect to p_{true} (f in BDA3)



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- The expectation is with respect to p_{true} (f in BDA3)
- Choose the model function to maximize our utility



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Model choice utility

- Application specific utility/cost functions are important
 - eg. money, life years, quality adjusted life years, etc.



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 - eg. money, life years, quality adjusted life years, etc.
- General utility: overall in the goodness of the predictive distribution
 - we don't know (yet) the application specific utility then good information theoretically justified choice is log-score for model M

 $\log p_M(y^{\mathsf{rep}}|y)$



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$$\log p_M(y^{\text{rep}}|y)$$

• We want the "best" model to explain the data



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Measures of predictive accuracy



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(Bayesian) Points Prediction Accuracy

Point residuals

$$e_i = y_i - E(\tilde{y}_i|y)$$
,

where

$$E(\tilde{y}|y) = \int \tilde{y}_i p(\tilde{y}_i|y) d\tilde{y},$$

i.e. the expected predicted value



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i.e. the expected predicted value

Mean squared (prediction) error (MSE)

$$MSE = \frac{1}{n} \sum_{i=1}^{n} [y_i - E(\tilde{y}_i|y)]^2.$$



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Probabilistic predictions

• The log score (a local and proper scoring rule)* $\log p(y|\theta)$



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Probabilistic predictions

• The log score (a local and proper scoring rule)*

$$\log p(y|\theta)$$

The log predictive density (lpd)

$$lpd = log p(y|y)$$
$$= log \int p(y|\theta)p(\theta|y)d\theta$$



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Probabilistic predictions

 The lpd is usually approximated with the log point predictive density (lppd or just lpd)

$$\mathsf{lppd} = \sum_{i}^{n} \log p(y_{i}|y)$$

$$\approx \log p(y|y)$$



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Probabilistic predictions

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$$\mathsf{lppd} = \sum_{i}^{n} \mathsf{log} \ p(y_{i}|y)$$
$$\approx \mathsf{log} \ p(y|y)$$

Estimation using MCMC

$$\mathsf{lppd} = \sum_{i}^{n} \log \left(\frac{1}{S} \sum_{s}^{S} p(y_{i} | \theta_{s})) \right)$$



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 Evaluate how model M generalizes to unseen data ỹ (the expected log predictive density):

$$\mathsf{elpd}_{M} = \int \mathsf{log}\, p_{M}(\tilde{oldsymbol{y}}|oldsymbol{y}) p_{\mathsf{true}}(\tilde{oldsymbol{y}}) d ilde{oldsymbol{y}}\,,$$

where \tilde{y} is an unseen observation generated from the true data generating process $p_{\text{true}}(\tilde{y})$, and y are observed data.

• $\log p_M(\tilde{y}|y)$ is the \log score (the utility of the model)



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- The expectation is with respect to p_{true}



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- The expectation is with respect to p_{true}
- p_{true} is (almost always) unknown



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- $\log p_M(\tilde{y}|y)$ is the \log score (the utility of the model)
- The expectation is with respect to p_{true}
- p_{true} is (almost always) unknown
- The utility function is the log scoring rule.



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Leave-one-out cross-validation (LOO-CV)

 Evaluate how model M generalizes to unseen data (the expected log predictive density):

$$\mathsf{elpd}_M = \int \mathsf{log}\, p_M(ilde{y_i}|y) p_\mathsf{true}(ilde{y_i}) d ilde{y_i}\,,$$

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where \tilde{y}_i is an unseen observation generated from the true data generating process $p_{\text{true}}(\tilde{y}_i)$, and y are observed data.

• Can we approximate $p_{\text{true}}(\tilde{y}_i)$?



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Leave-one-out cross-validation (LOO-CV)

• Approximate $p_{\mathsf{true}}(\tilde{y}_i)$ with data y



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Leave-one-out cross-validation (LOO-CV)

- Approximate $p_{\text{true}}(\tilde{y}_i)$ with data y
- Hold out observation i and try to predict y_i based on \mathbf{y}_{-i}
- Estimation of elpd_M using leave-one-out cross-validation

$$\begin{aligned} \mathsf{elpd}_{\mathsf{loo}} &= \sum_{i=1}^{n} \mathsf{log} \, p_{M}(y_{i} | \mathbf{y}_{-i}) \\ &= \sum_{i=1}^{n} \mathsf{log} \int p_{M}(y_{i} | \theta) p(\theta | \mathbf{y}_{-i}) d\theta \end{aligned}$$



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- Analogy: Monte Carlo approximation using our data
- Similar to jack-knife resampling



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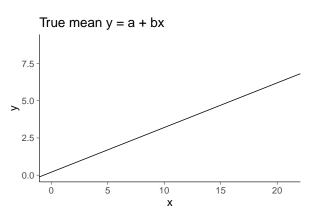
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- Analogy: Monte Carlo approximation using our data
- Similar to jack-knife resampling
- The elpd, lpd and efficient number of parameters (p_{loo})

$$elpd_{loo} = lpd - p_{loo}$$

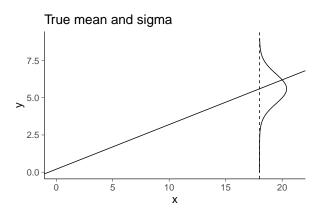


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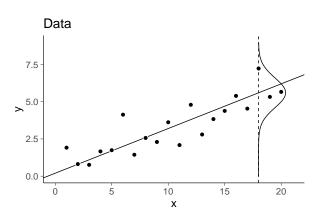


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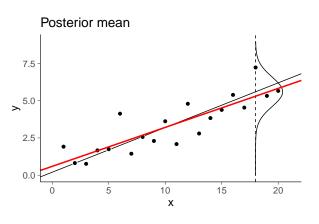


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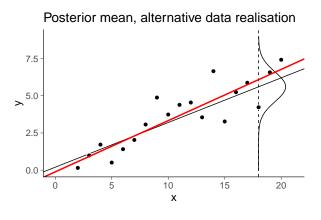


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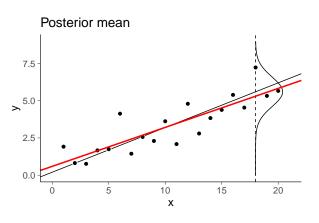


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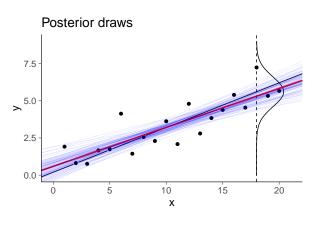


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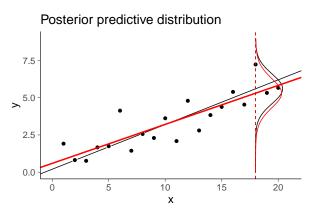


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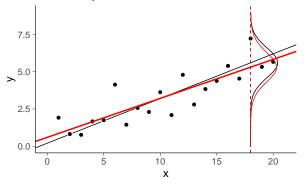




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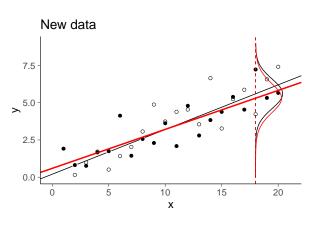
Posterior predictive distribution



$$p(\tilde{y}|\tilde{x}=18,x,y) = \int p(\tilde{y}|\tilde{x}=18,\theta)p(\theta|x,y)d\theta$$

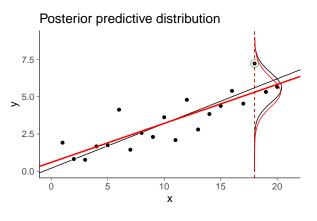


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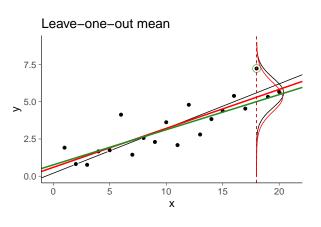


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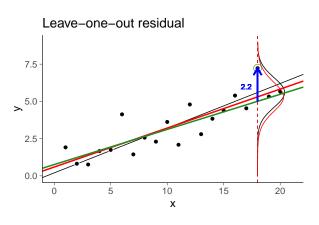


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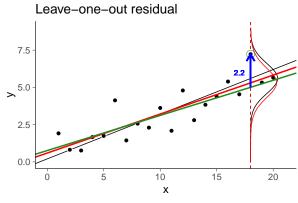


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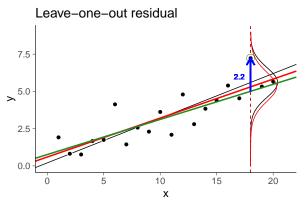
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$$y_{18} - E[p(\tilde{y}|\tilde{x}=18, x_{-18}, y_{-18})]$$



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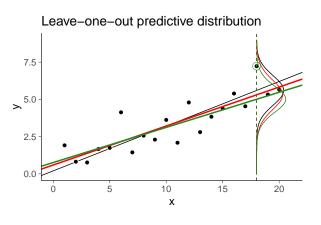


$$y_{18} - E[p(\tilde{y}|\tilde{x}=18, x_{-18}, y_{-18})]$$

Can be use to compute, e.g., RMSE, R², 90% error

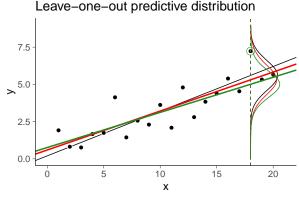


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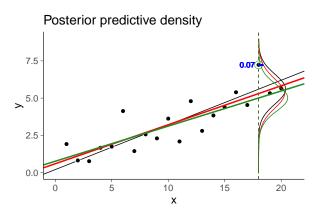
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$$p(\tilde{y}|\tilde{x} = 18, x_{-18}, y_{-18}) = \int p(\tilde{y}|\tilde{x} = 18, \theta)p(\theta|x_{-18}, y_{-18})d\theta$$

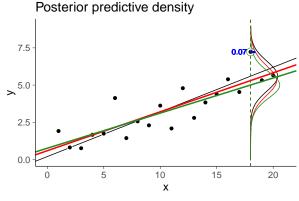


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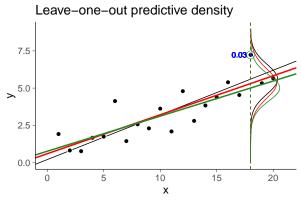


$$p(\tilde{y}=y_{18}|\tilde{x}=18,x,y)\approx 0.07$$



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 - PSIS LOO-CV

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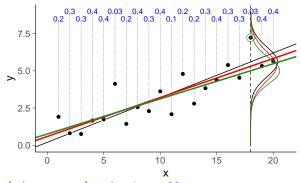


$$p(\tilde{y} = y_{18}|\tilde{x} = 18, x, y) \approx 0.07$$

$$p(\tilde{y} = y_{18}|\tilde{x} = 18, x_{-18}, y_{-18}) \approx 0.03$$



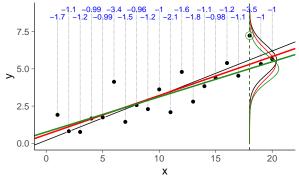
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$$p(y_i|x_i,x_{-i},y_{-i}), \quad i=1,\ldots,20$$



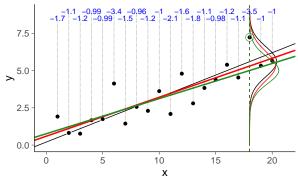
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$$\log p(y_i|x_i,x_{-i},y_{-i}), \quad i=1,\ldots,20$$



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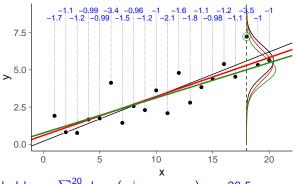


$$\sum_{i=1}^{20} \log p(y_i|x_i, x_{-i}, y_{-i}) \approx -29.5$$



- Measures of predictive accuracy
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Leave-one-out log predictive densities

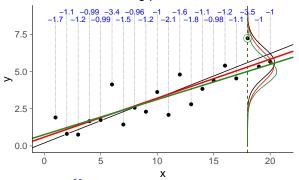


elpd_loo = $\sum_{i=1}^{20} \log p(y_i|x_i, x_{-i}, y_{-i}) \approx -29.5$



- Measures of predictive accuracy
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Leave-one-out log predictive densities

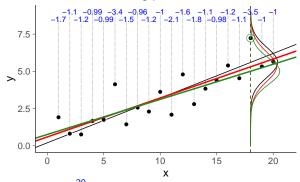


elpd_loo = $\sum_{i=1}^{20} \log p(y_i|x_i,x_{-i},y_{-i}) \approx -29.5$ unbiased estimate of log posterior pred. density for new data



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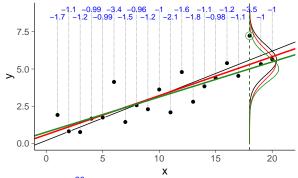
elpd_loo =
$$\sum_{i=1}^{20} \log p(y_i|x_i, x_{-i}, y_{-i}) \approx -29.5$$

$$lpd = \sum_{i=1}^{20} log p(y_i|x_i, x, y) \approx -26.8$$



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- Model assessment and selection
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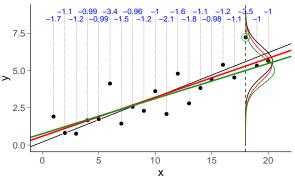
elpd_loo =
$$\sum_{i=1}^{20} \log p(y_i|x_i, x_{-i}, y_{-i}) \approx -29.5$$

lpd = $\sum_{i=1}^{20} \log p(y_i|x_i, x, y) \approx -26.8$
p_loo = lpd - elpd_loo ≈ 2.7



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

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



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Stan and loo package

Computed from 4000 by 20 log-likelihood matrix

 $\begin{array}{ccc} & \text{Estimate} & \text{SE} \\ \text{elpd_loo} & -29.5 & 3.3 \\ \text{p_loo} & 2.7 & 1.0 \end{array}$

Monte Carlo SE of elpd_loo is 0.1.

Pareto k diagnostic values:

		Count	Pct.	Min.	n_eff
(-Inf, 0.5]	(good)	18	90.0%	899	
	(ok)	2	10.0%	459	
(0.7, 1]	(bad)	0	0.0%	<NA $>$	
	(very bad)	0	0.0%	<NA $>$	

All Pareto k estimates are ok (k < 0.7). See help('pareto-k-diagnostic') for details.



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Subsection 1

When is LOO applicable



- Measures of predictive accuracy
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Pro and cons with LOO-CV

- + Intuitive
- + Robust
- + Good theoretical properties



- Measures of predictive
- accuracy
- Model selection

Cross-validation

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Pro and cons with LOO-CV

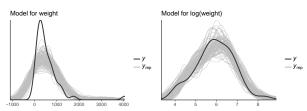
- + Intuitive
- + Robust
- + Good theoretical properties
 - Can be costly (naive LOO-CV mean n posterior computations)



- Measures of predictive accuracy
- Model selection
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Sometimes cross-validation is not needed

Posterior predictive checking can be sufficient



Predicting the yields of mesquite bushes.

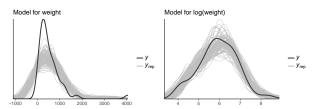
Gelman, Hill & Vehtari (2020, Ch. 11)



- Measures of predictive accuracy
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Sometimes cross-validation is not needed

Posterior predictive checking can be sufficient



Predicting the yields of mesquite bushes.

Gelman, Hill & Vehtari (2020, Ch. 11)

 In nested case, often easier and more accurate to analyse posterior distribution of more complex model directly



- Measures of predictive accuracy
- Model selection
- Iviodel selection

Cross-validation When is LOO applicable

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Data generating mechanisms and prediction tasks

1. You have to make some assumptions on data generating mechanism p_{true}

$$\mathsf{elpd}_M = \int \mathsf{log}\, p_M(\boldsymbol{\tilde{y}}|y) p_\mathsf{true}(\boldsymbol{\tilde{y}}) d\boldsymbol{\tilde{y}}\,,$$

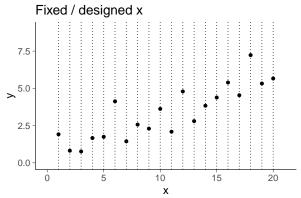
- 2. Use the knowledge of the prediction task if available
- 3. Cross-validation can be used to analyse different parts, even if there is no clear prediction task



- Measures of predictive accuracy
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$x \text{ in } p_{\text{true}}$



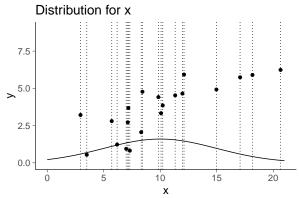
LOO is ok for fixed / designed x: $p_{true}(y|x)$



- Measures of predictive accuracy
- Model selection
- Cross-validation
 When is LOO applicable
 - When is LOO:

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$x \text{ in } p_{\text{true}}$



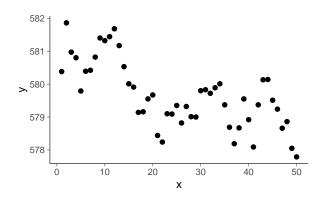
LOO is ok for random x. $p_{true}(y, x)$



- Measures of predictive
- accuracy

 Model selection
- Model selection
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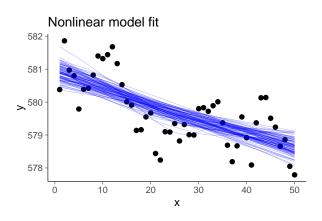
p_{true} extrapolation





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p_{true} extrapolation

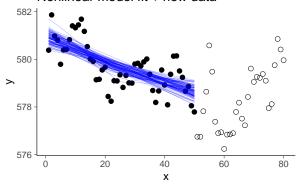




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p_{true} extrapolation





Extrapolation is more difficult



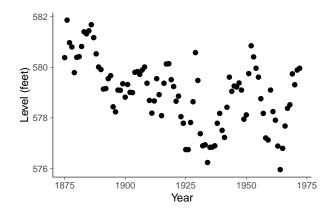
- Measures of predictive
- accuracy

 Model selection
- Woder Selection

Cross-validation

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LOO for time series data



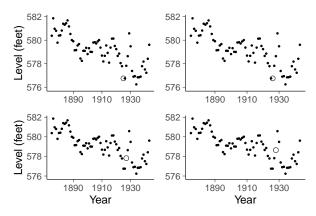
 p_{true} for time series?



- Measures of predictive
- accuracy

 Model selection
- ...
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LOO for time series



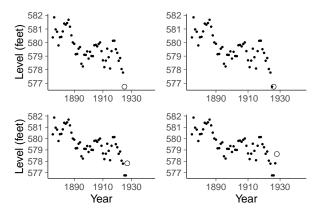
Leave-one-out cross-validation is ok for assessing conditional model



- Model assessment and selection
 - Measures of predictive
 - accuracy

 Model selection
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LOO for time series

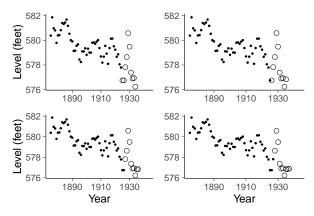


Leave-future-out cross-validation is better for predicting future



- Measures of predictive accuracy
- Model selection
-
- Cross-validation
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LOO for time series

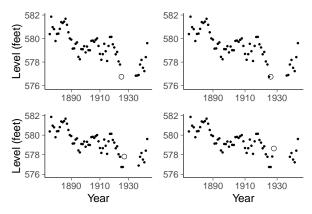


m-step-ahead cross-validation is better for predicting further future



- Measures of predictive accuracy
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LOO for time series



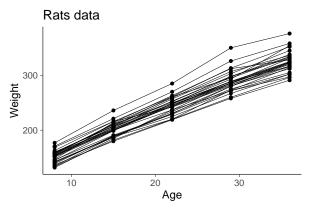
m-step-ahead leave-a-block-out cross-validation



- Model assessment and selection
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LOO for hiearchical data

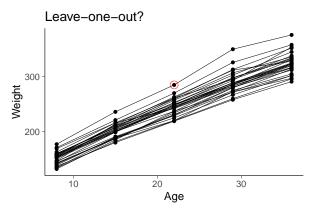


Can LOO or other cross-validation be used with hierarchical data?



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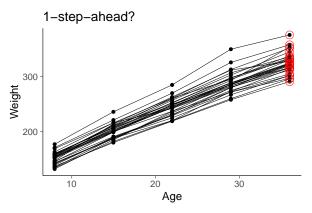
LOO for hiearchical data





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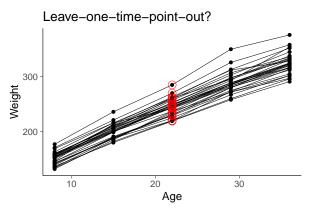




- Measures of predictive
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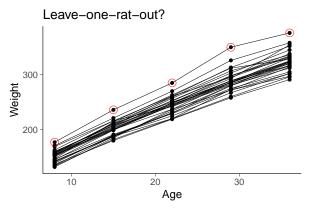




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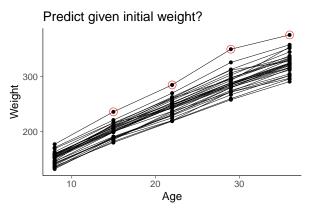
LOO for hiearchical data





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Fast cross-validation

1. Pareto smoothed importance sampling LOO (PSIS-LOO)



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Fast cross-validation

- 1. Pareto smoothed importance sampling LOO (PSIS-LOO)
- 2. K-fold cross-validation



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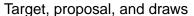
PSIS LOO-CV

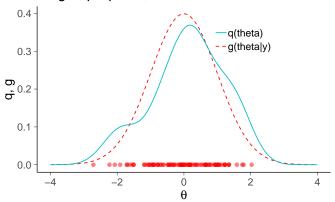


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Recap: Importance sampling



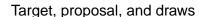


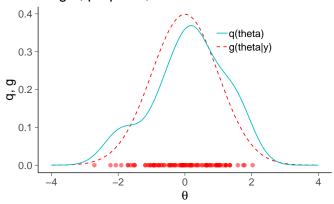


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Recap: Importance sampling





$$\mathrm{E}[f(heta)] pprox rac{\sum_s w_s f(heta^{(s)})}{\sum_s w_s}, \qquad ext{where} \quad w_s = rac{q(heta^{(s)})}{g(heta^{(s)})}$$

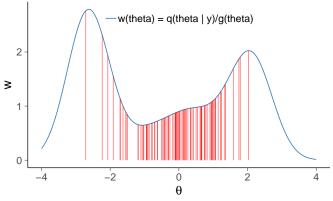


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Recap: Importance sampling

Draws and importance weights



$$\mathrm{E}[f(heta)] pprox rac{\sum_s w_s f(heta^{(s)})}{\sum_s w_s}, \qquad ext{where} \quad w_s = rac{q(heta^{(s)})}{g(heta^{(s)})}$$



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Importance sampling for LOO-CV

• Let $f = p(\tilde{y}_i|y_{-i})$, $g = p(\theta|y)$, and $q = p(\theta|y_{-i})$



- Measures of predictive accuracy
- Model selection

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Importance sampling for LOO-CV

- Let $f = p(\tilde{y}_i|y_{-i})$, $g = p(\theta|y)$, and $q = p(\theta|y_{-i})$
- Having samples θ^s from $p(\theta^s|y)$

$$p(\tilde{y}_i|y_{-i}) \approx \frac{\sum_{s=1}^{S} p(\tilde{y}_i|\theta^s)w_i^s}{\sum_{s=1}^{S} w_i^s},$$

where w_i^s are importance weights and

$$w_i^s = \frac{p(\theta^s|y_{-i})}{p(\theta^s|y)}.$$



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Importance sampling for LOO-CV

• We get a nice solution

$$w_i^s = \frac{p(\theta^s|y_{-i})}{p(\theta^s|y)} = \frac{\frac{p(\theta^s)\prod_{j\neq i}p(y_j|\theta^s)}{p(y_{-i})}}{\frac{p(\theta^s)\prod_jp(y_j|\theta^s)}{p(y)}} \propto \frac{1}{\frac{p(y_i|\theta^s)}{p(y)}}.$$



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Importance sampling for LOO-CV

• We get a nice solution

$$w_i^s = \frac{p(\theta^s|y_{-i})}{p(\theta^s|y)} = \frac{\frac{p(\theta^s)\prod_{j\neq i}p(y_j|\theta^s)}{p(y_{-i})}}{\frac{p(\theta^s)\prod_{j}p(y_j|\theta^s)}{p(y)}} \propto \frac{1}{\frac{p(y_i|\theta^s)}{p(y)}}.$$

• If evaluated with $\tilde{y}_i = y_i$

$$p(\tilde{y}_i|y_{-i}) \approx \frac{\sum_{s=1}^{S} p(\tilde{y}_i|\theta^s) w_i^s}{\sum_{s=1}^{S} w_i^s} = \frac{1}{\sum_{s=1}^{S} \frac{1}{p(y_i|\theta^s)}},$$

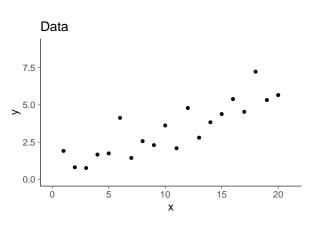
and

$$\log w_i^{(s)} = -\log p(y_i|\theta^{(s)})$$



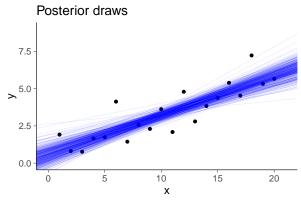
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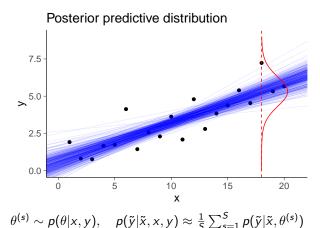
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$$\theta^{(s)} \sim p(\theta|x,y)$$

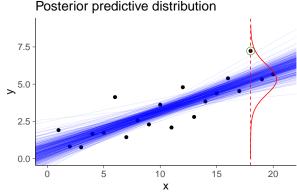


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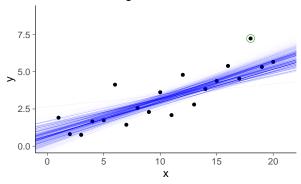


$$\theta^{(s)} \sim p(\theta|x,y), \quad p(\tilde{y}|\tilde{x},x,y) \approx \frac{1}{S} \sum_{s=1}^{S} p(\tilde{y}|\tilde{x},\theta^{(s)})$$



- Measures of predictive accuracy
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PSIS-LOO weighted draws

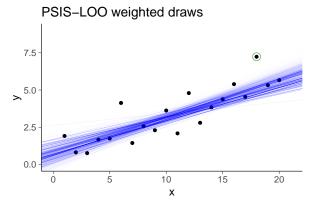


$$\theta^{(s)} \sim p(\theta|x, y)$$

$$w_i^{(s)} = p(\theta^{(s)}|x_{-i}, y_{-i})/p(\theta^{(s)}|x, y)$$



- Measures of predictive accuracy
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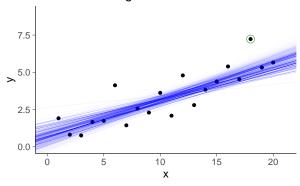
$$\theta^{(s)} \sim p(\theta|x,y)$$

$$w_i^{(s)} = p(\theta^{(s)}|x_{-i},y_{-i})/p(\theta^{(s)}|x,y) \propto 1/p(y_i|x_i,\theta^{(s)})$$



- Measures of predictive accuracy
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PSIS-LOO weighted draws



$$\theta^{(s)} \sim p(\theta|x, y)$$

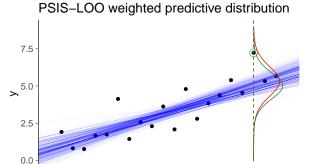
$$w_i^{(s)} = p(\theta^{(s)}|x_{-i}, y_{-i})/p(\theta^{(s)}|x, y) \propto 1/p(y_i|x_i, \theta^{(s)})$$

$$\log(1/p(y_i|x_i, \theta^{(s)})) = -\log_{-lik}[i]$$



- Measures of predictive accuracy
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$$\theta^{(s)} \sim p(\theta|x, y)$$

 $w_i^{(s)} = p(\theta^{(s)}|x_{-i}, y_{-i})/p(\theta^{(s)}|x, y) \propto 1/p(y_i|x_i, \theta^{(s)})$

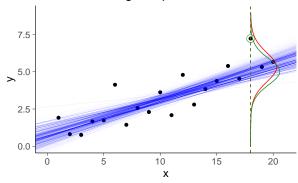
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PSIS-LOO weighted predictive distribution



$$\theta^{(s)} \sim p(\theta|x, y)$$

$$w_i^{(s)} = p(\theta^{(s)}|x_{-i}, y_{-i})/p(\theta^{(s)}|x, y) \propto 1/p(y_i|x_i, \theta^{(s)})$$

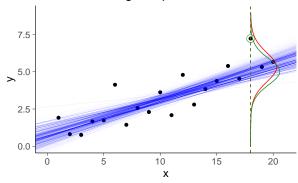
$$p(y_i|x_i, x_{-i}, y_{-i}) \approx \sum_{s=1}^{S} [w_i^{(s)} p(y_i|x_i, \theta^{(s)})]$$



- Measures of predictive accuracy
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PSIS-LOO weighted predictive distribution



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$$p(y_i|x_i, x_{-i}, y_{-i}) \approx \sum_{s=1}^{S} [w_i^{(s)} p(y_i|x_i, \theta^{(s)})]$$



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Recap: Pareto smoothed importance sampling

- Pareto-Smoothed Importance sampling smooth the weights according to a Generalized Pareto(k) distribution
- Pareto-k diagnostic estimate the number of existing moments (|1/k|)



- Measures of predictive accuracy
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Recap: Pareto smoothed importance sampling

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- Finite variance and central limit theorem for k < 1/2



- Model assessment and selection
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- Finite variance and central limit theorem for k < 1/2
- Finite mean and generalized central limit theorem for k < 1, but pre-asymptotic constant grows impractically large for k > 0.7



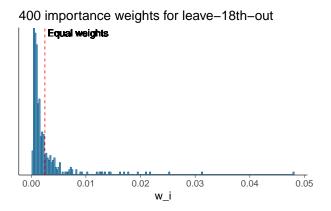
- Measures of predictive accuracy
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Recap: Pareto smoothed importance sampling

- Pareto-Smoothed Importance sampling smooth the weights according to a Generalized Pareto(k) distribution
- Pareto-k diagnostic estimate the number of existing moments (|1/k|)
- Finite variance and central limit theorem for k < 1/2
- Finite mean and generalized central limit theorem for k < 1, but pre-asymptotic constant grows impractically large for k > 0.7
- Hence k > 0.7 can be used to indicate what observation that can "hold-out" using IS

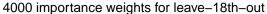


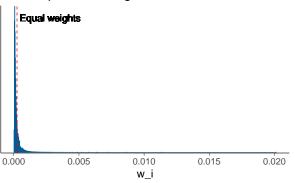
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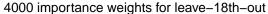
- Model assessment and selection
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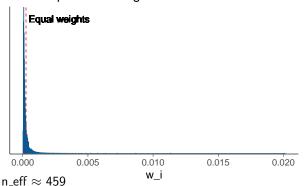






- Model assessment and selection
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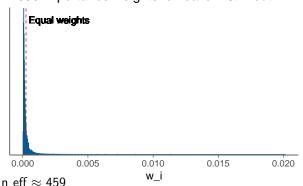




- Model assessment and selection
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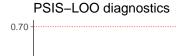
4000 importance weights for leave-18th-out

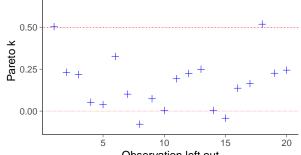


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- Measures of predictive accuracy
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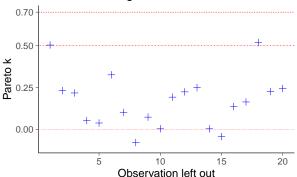


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- Model assessment and selection
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PSIS-LOO diagnostics



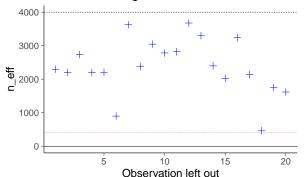
Pareto k diagnostic values: Count Pct. Min. n_eff (-Inf, 0.5] (0.5, 0.7] 90.0% (good) 18 899 (ok) 10.0% 459 (bad) 0.0% <NA>(verv bad) 0.0% < NA >



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PSIS-LOO diagnostics



Pareto k diagnostic values:

	_	Count	Pct.	Min.	n_eff
(-Inf, 0.5]	(good)	18	90.0%	899	
(0.5, 0.7]	(ok)	2	10.0%	459	
(0.7, 1]	(bad)	0	0.0%	<NA $>$	
(1 Inf)	(very had)	0	0.0%	∠NIΔ\	



- Model assessment and selection
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Stan code

$$\log(w_i^{(s)}) = \log(1/p(y_i|x_i,\theta^{(s)})) = -\log_{-1}[ik[i]]$$



- Model assessment and selection
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Stan code

```
\log(w_i^{(s)}) = \log(1/p(y_i|x_i,\theta^{(s)})) = -\log_{-\text{lik}}[i]
model {
  alpha " normal(pmualpha, psalpha);
  beta ~ normal(pmubeta, psbeta);
  v ~ normal(mu, sigma);
generated quantities {
  vector[N] log_lik;
  for (i in 1:N)
    log_lik[i] = normal_lpdf(y[i] | mu[i], sigma);
```



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- Model assessment and selection
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Stan and loo package

Computed from 4000 by 20 log-likelihood matrix

 $\begin{array}{ccc} & \text{Estimate} & \text{SE} \\ \text{elpd_loo} & -29.5 & 3.3 \\ \text{p_loo} & 2.7 & 1.0 \end{array}$

Monte Carlo SE of elpd_loo is 0.1.

Pareto k diagnostic values:

		Count	Pct.	Min.	n_eff
(-Inf, 0.5]	(good)	18	90.0%	899	
(0.5, 0.7]	(ok)	2	10.0%	459	
(0.7, 1]	(bad)	0	0.0%	<NA $>$	
(1, Inf)	(very bad)	0	0.0%	<NA $>$	

All Pareto k estimates are ok (k < 0.7). See help('pareto-k-diagnostic') for details.



- Measures of predictive accuracy
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PSIS-LOO

- 1. PSIS-LOO for hierarchical models
 - 1.1 leave-one-group out is challenging for PSIS-LOO



- Measures of predictive accuracy
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PSIS-LOO

- 1. PSIS-LOO for hierarchical models
 - 1.1 leave-one-group out is challenging for PSIS-LOO
- 2. PSIS-LOO for non-factorizable models



- Measures of predictive accuracy
- Model selection
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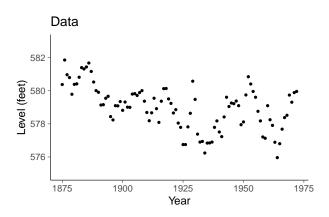
PSIS-LOO

- 1. PSIS-LOO for hierarchical models
 - 1.1 leave-one-group out is challenging for PSIS-LOO
- 2. PSIS-LOO for non-factorizable models
- 3. PSIS-LOO for time series
 - 3.1 Approximate leave-future-out cross-validation



- Model assessment and selection
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PSIS-LOO for time series

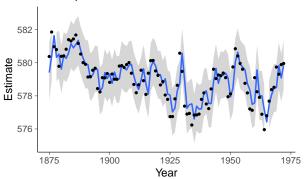




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PSIS-LOO for time series





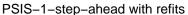


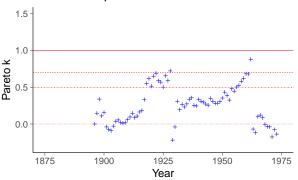
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 - PSIS LOO-CV

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PSIS-LOO for time series





mc-stan.org/loo/articles/loo2-lfo.html



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Subsection 3

K-fold CV



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K-fold cross-validation

We can approximate elpd_{loo} using K-fold CV



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K-fold cross-validation

- We can approximate elpd_{loo} using K-fold CV
 - 1. for k in K
 - 1.1 hold 1/K observations out (i.e. $y_{-v(k)}$)
 - 1.2 compute $p(\theta|y_{-y(k)})$
 - 1.3 compute lpd for $y_{-y(k)}$
 - 2. sum the lpd over *K* folds



- Measures of predictive accuracy
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K-fold cross-validation

- We can approximate elpd_{loo} using K-fold CV
 - 1. for k in K
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 - 1.2 compute $p(\theta|y_{-y(k)})$
 - 1.3 compute lpd for $y_{-y(k)}$
 - 2. sum the lpd over K folds
- We only need to estimate K models



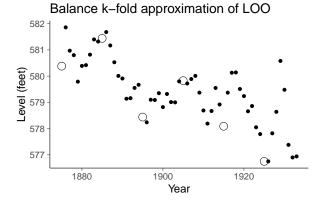
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- Cross-validation
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K-fold cross-validation

- 1. K-fold cross-validation can approximate LOO
 - 1.1 can be used all uses for LOO
- 2. K-fold cross-validation can be used for hierarchical models
 - 2.1 good for leave-one-group-out
- 3. K-fold cross-validation can be used for time series
 - 3.1 with leave-block-out

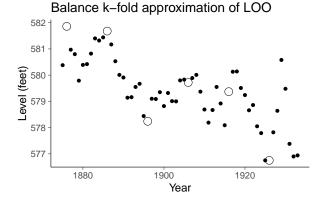


- Measures of predictive accuracy
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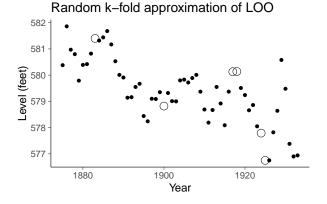


- Model assessment and selection
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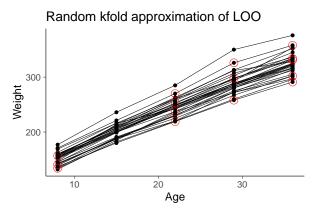


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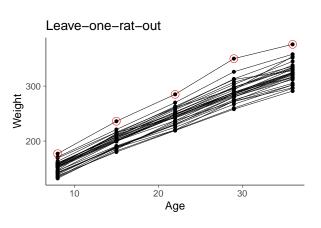


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Cross-validation for model assessment

1. CV is good for model assessment when application specific utility/cost functions are used



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Cross-validation for model assessment

- CV is good for model assessment when application specific utility/cost functions are used
- 2. Also useful in model checking in similar way as posterior predictive checking (PPC)
 - 2.1 model misspecification diagnostics (e.g. Pareto-*k* and p_loo)
 - 2.2 checking calibration of leave-one-out predictive posteriors (ppc_loo_pit in bayesplot)



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Subsection 4

Model Comparison



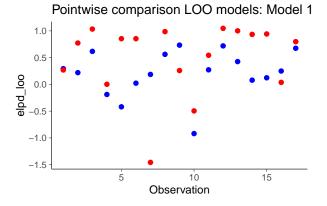
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Model comparison

- "A popular hypothesis has it that primates with larger brains produce more energetic milk, so that brains can grow quickly" (from Statistical Rethinking)
 - 1.1 Model 1: formula = kcal.per.g \sim neocortex
 - 1.2 Model 2: formula = kcal.per.g \sim neocortex + log(mass)



- Model assessment and selection
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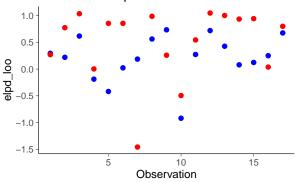


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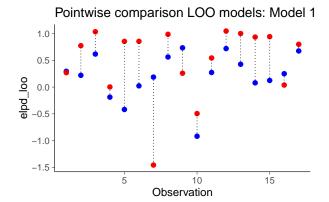


Model 1 elpd_loo \approx 3.7, SE=1.8 Model 2 elpd_loo \approx 8.4, SE=2.8



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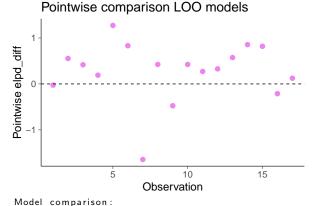
- Model assessment and selection
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Model 1 elpd_loo \approx 3.7, SE=1.8 Model 2 elpd_loo \approx 8.4, SE=2.8



- Model assessment and selection
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Arsenic well example – Model comparison

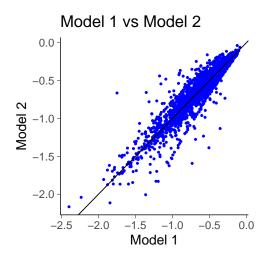
- Probability of switching well with high arsenic level in rural Bangladesh
 - 1.1 Model 1 covariates: log(arsenic) and distance
 - 1.2 Model 2 covariates: log(arsenic), distance and education level

Gelman, Hill & Vehtari (2020): Regression and Other Stories, Chapter 13.



- Measures of predictive
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Arsenic well example – Model comparison



Model 1 elpd_loo \approx -1952, SE=16 Model 2 elpd_loo \approx -1938, SE=17

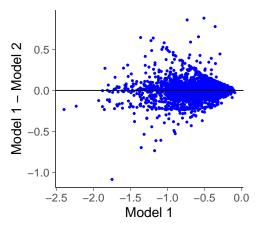


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Arsenic well example – Model comparison





> loo_compare(model1, model2) elpd_diff se_diff model2 0.0 0.0 model1 -14.4 6.1



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Arsenic well example – Model comparison

```
> loo_compare(model1, model2)
elpd_diff se_diff
model2 0.0 0.0
model1 -14.4 6.1
```

se_diff and normal approximation for the uncertainty in the difference is good only if models are well specified and the number of observations is relatively big.



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- 1. Continuous expansion including all models?
 - 1.1 and then analyse the posterior distribution directly
 - 1.2 sparse priors like regularized horseshoe prior instead of variable selection



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- 1. Continuous expansion including all models?
 - 1.1 and then analyse the posterior distribution directly
 - 1.2 sparse priors like regularized horseshoe prior instead of variable selection
- 2. Model averaging with BMA or Bayesian stacking?



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- 1. Continuous expansion including all models?
 - 1.1 and then analyse the posterior distribution directly
 - 1.2 sparse priors like regularized horseshoe prior instead of variable selection
- 2. Model averaging with BMA or Bayesian stacking?
- 3. In a nested case choose simpler if assuming some cost for extra parts?
- In a nested case choose more complex if you want to take into account all the uncertainties.



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Cross-validation and model selection

- 1. Cross-validation can be used for model selection if
 - 1.1 small number of models
 - 1.2 the difference between models is clear



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Cross-validation and model selection

- 1. Cross-validation can be used for model selection if
 - 1.1 small number of models
 - 1.2 the difference between models is clear
- Do not use cross-validation to choose from a large set of models
 - 2.1 selection process leads to overfitting



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Cross-validation and model selection

- 1. Cross-validation can be used for model selection if
 - 1.1 small number of models
 - 1.2 the difference between models is clear
- Do not use cross-validation to choose from a large set of models
 - 2.1 selection process leads to overfitting
- Overfitting in selection process is not unique for cross-validation



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Selection induced bias and overfitting

- Selection induced bias in cross-validation
 - same data is used to assess the performance and make the selection
 - the selected model fits more to the data
 - the CV estimate for the selected model is biased



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Selection induced bias and overfitting

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 - same data is used to assess the performance and make the selection
 - the selected model fits more to the data
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- Performance of the selection process itself can be assessed using two level cross-validation, but it does not help choosing better models



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Selection induced bias and overfitting

- Selection induced bias in cross-validation
 - same data is used to assess the performance and make the selection
 - the selected model fits more to the data
 - the CV estimate for the selected model is biased
- Performance of the selection process itself can be assessed using two level cross-validation, but it does not help choosing better models
- Bigger problem if there is a large number of models as in covariate selection



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Section 3

Information criteria



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Akaike Information Criteria

1. Remember the relationship between the elpd and the lpd

$$\mathsf{elpd}_{\mathsf{loo}} = \sum_{i} \mathsf{log}\, p(y_i|y) - p_{\mathsf{loo}}$$



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Akaike Information Criteria

1. Remember the relationship between the elpd and the lpd

$$elpd_{loo} = \sum_{i} log p(y_i|y) - p_{loo}$$

2. Akaike information criteria

$$\mathsf{elpd}_{\mathsf{AIC}} = \sum_{i} \log p(y_i | \hat{\theta}_{\mathsf{MLE}}) - k$$

3. AIC

$$\mathsf{AIC} = -2\mathsf{elpd}_{\mathsf{AIC}}$$



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Deviance Information Criteria

1. Deviance information criteria

$$\mathsf{elpd}_\mathsf{AIC} = \sum_i \mathsf{log}\, p(y_i|\hat{ heta}_\mathsf{Bayes}) - p_\mathsf{DIC}$$

- 2. We estimate the number of effeicient parameters with
- 3. DIC

$$\mathsf{DIC} = -2\mathsf{elpd}_{\mathsf{DIC}}$$



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WAIC vs PSIS-LOO

1. WAIC has same assumptions as LOO



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- 1. WAIC has same assumptions as LOO
- 2. PSIS-LOO is more accurate



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- 1. WAIC has same assumptions as LOO
- 2. PSIS-LOO is more accurate
- 3. PSIS-LOO has much better diagnostics



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- 1. WAIC has same assumptions as LOO
- 2. PSIS-LOO is more accurate
- 3. PSIS-LOO has much better diagnostics
- 4. LOO makes the prediction assumption more clear, which helps if K-fold-CV is needed instead



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- 1. WAIC has same assumptions as LOO
- 2. PSIS-LOO is more accurate
- 3. PSIS-LOO has much better diagnostics
- LOO makes the prediction assumption more clear, which helps if K-fold-CV is needed instead
- 5. Multiplying by -2 doesn't give any benefit (Watanabe didn't multiply by -2)



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- 1. AIC uses maximum likelihood estimate for prediction
- 2. DIC uses posterior mean for prediction
- 3. BIC is an approximation for marginal likelihood
- 4. TIC, NIC, RIC, PIC, BPIC, QIC, AICc, ...



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Model averaging and Stacking



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Special lecture!

1. Special lecture on model averaging and stacking next week!



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- 1. It's good to think predictions of observables, because observables are the only ones we can observe
- Cross-validation can simulate predicting and observing new data
- 3. Cross-validation is good if you don't trust your model
- Different variants of cross-validation are useful in different scenarios
- 5. Cross-validation has high variance, and **if** you trust your model you can beat cross-validation in accuracy



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