



Scaleable Distributional Learning

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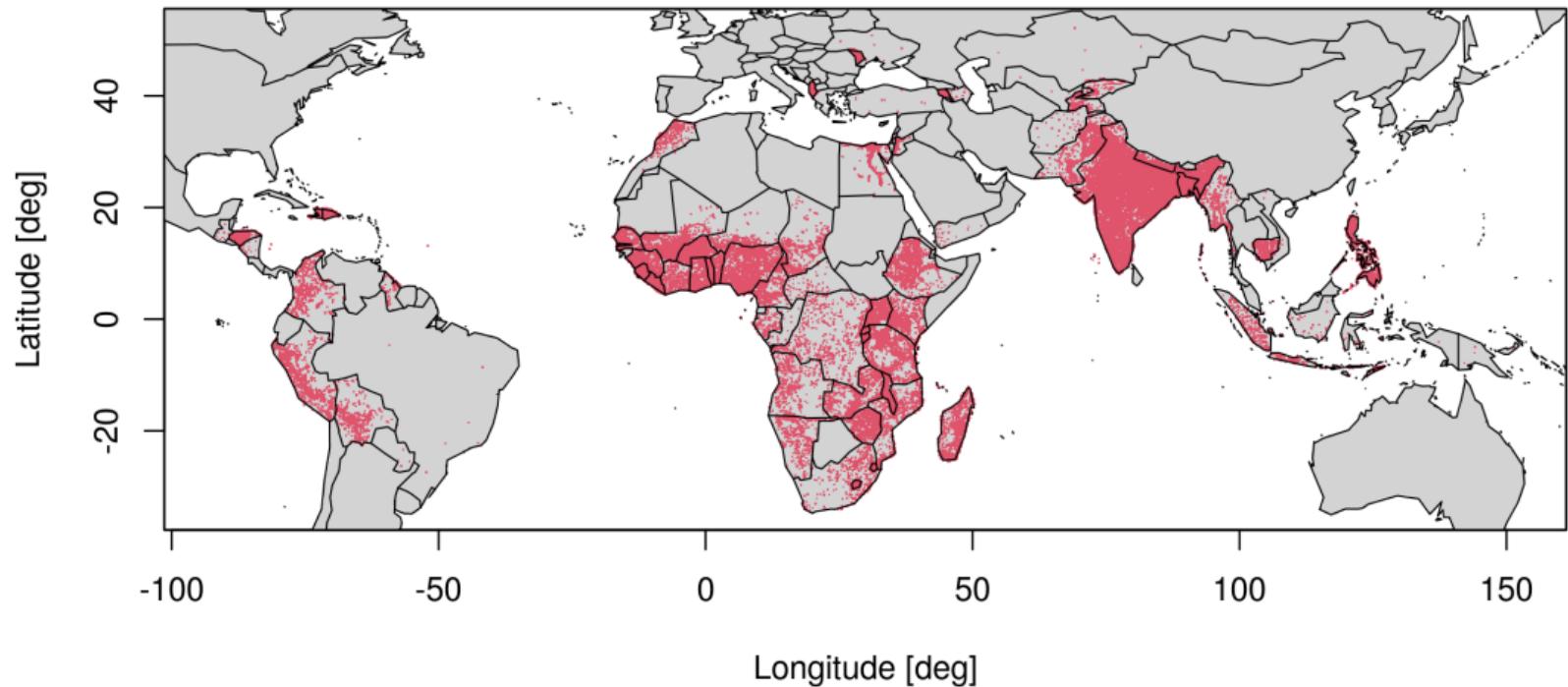
Childhood malnutrition

Joint work with:

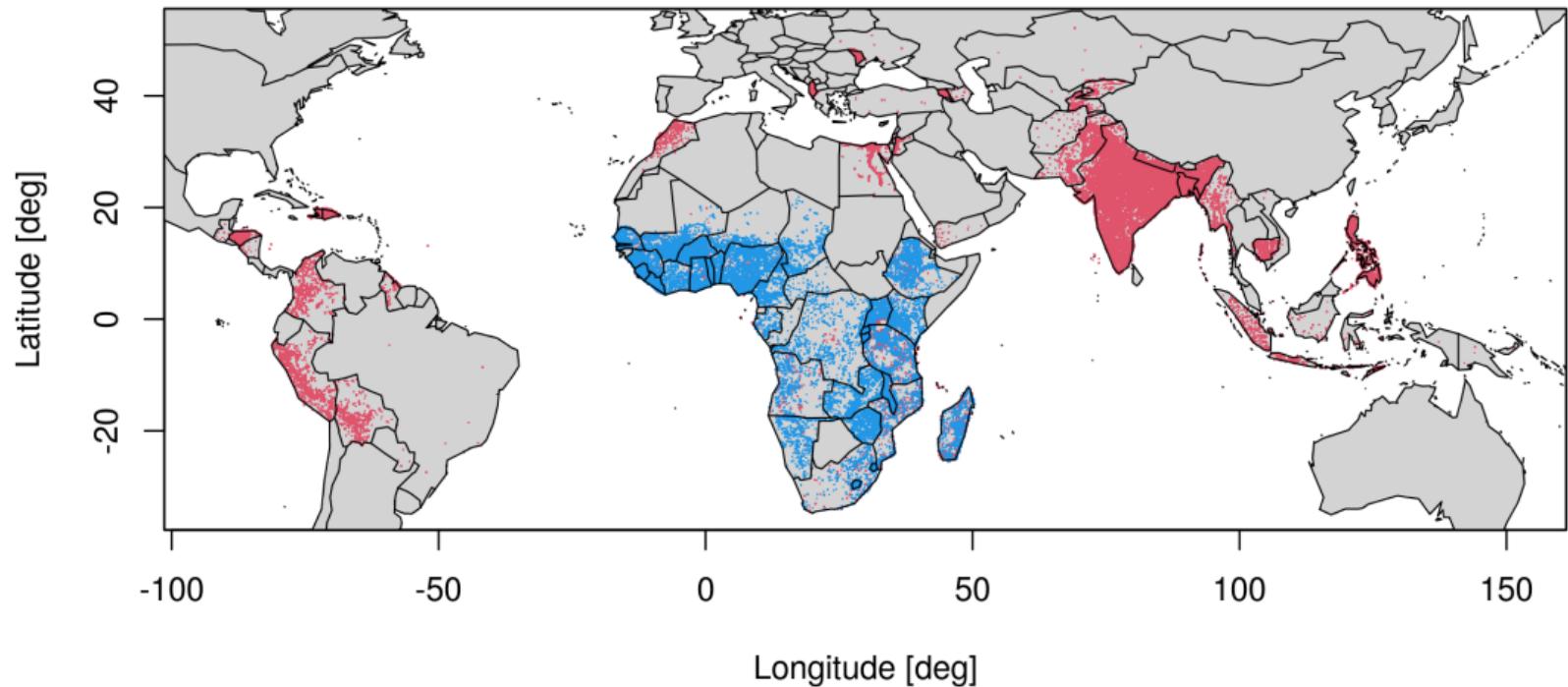
Johannes Seiler, Mattias Wetcher, Thorsten Simon & Stefan Lang.

- Project aiming to better explain the problems of childhood malnutrition in low- and middle-income countries.
- Contribute to monitoring of the Sustainable Development Goals (SGD).
- We compiled a brand new data set using DHS data.
- Data on global conflicts, topography and environmental data from satellite earth observations (NDVI), temperature and precipitation data from ERA5.
- Data from 1990–2019 with $n > 3M$ observations.

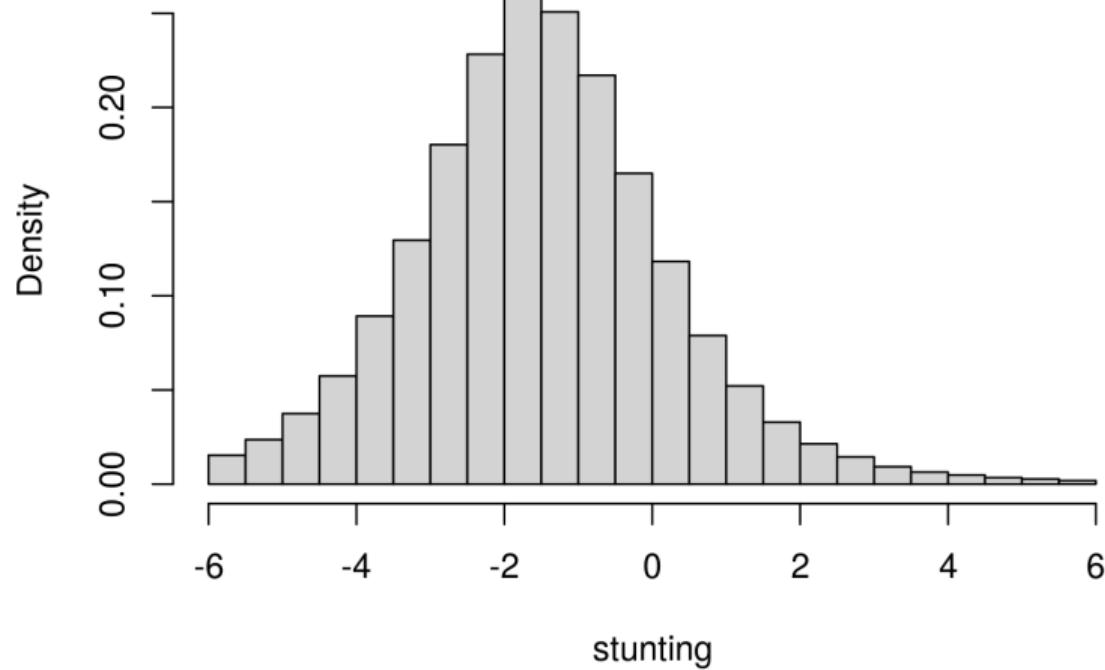
Childhood malnutrition



Childhood malnutrition



Childhood malnutrition



Computational challenges

- Distributional regression using (very) large data sets?
- Variable selection?
- Inference?

Model specification

Any parameter of a population distribution \mathcal{D} may be modeled by explanatory variables

$$y \sim \mathcal{D}(\theta_1(\mathbf{x}; \boldsymbol{\beta}_1), \dots, \theta_K(\mathbf{x}; \boldsymbol{\beta}_K)),$$



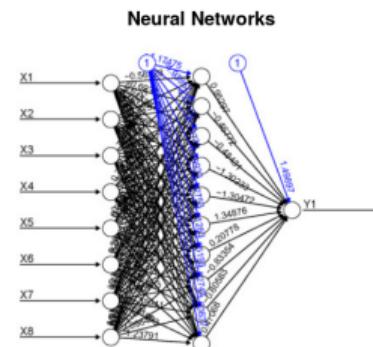
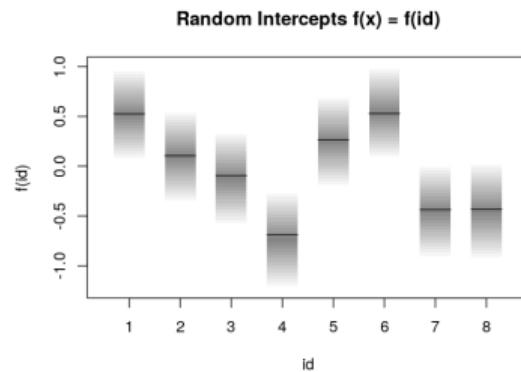
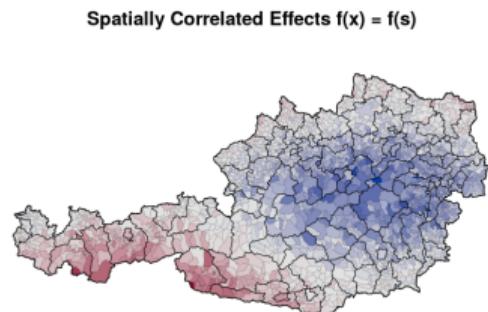
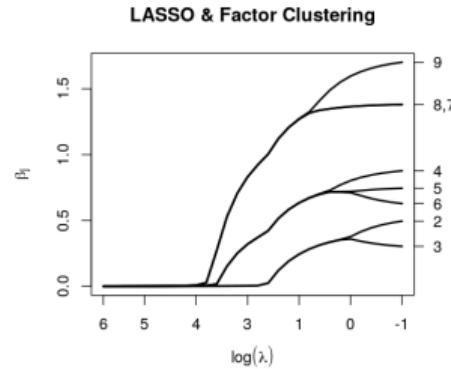
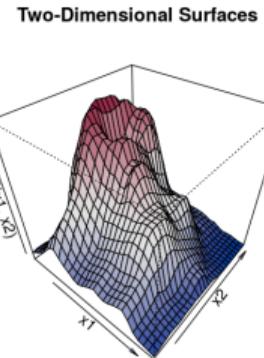
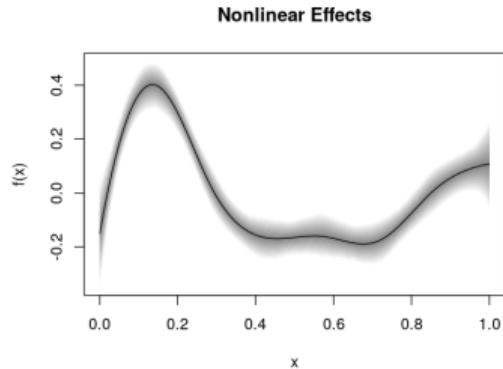
with $\boldsymbol{\beta} = (\boldsymbol{\beta}_1^\top, \dots, \boldsymbol{\beta}_K^\top)^\top$.

Each parameter is linked to a structured additive predictor

$$h_k(\theta_k(\mathbf{x}; \boldsymbol{\beta}_k)) = f_{1k}(\mathbf{x}; \beta_{1k}) + \dots + f_{J_k k}(\mathbf{x}; \beta_{J_k k}),$$

- $j = 1, \dots, J_k$ and $k = 1, \dots, K$.
- $h_k(\cdot)$: Link functions for each distribution parameter.
- $f_{jk}(\cdot)$: Model terms of one or more variables.

Model terms $f_{jk}(\cdot)$



Estimation

The main building block of regression model algorithms is the probability density function $d_y(\mathbf{y}|\boldsymbol{\theta}_1, \dots, \boldsymbol{\theta}_K)$.

Estimation typically requires to evaluate the log-likelihood

$$\ell(\boldsymbol{\beta}; \mathbf{y}, \mathbf{X}) = \sum_{i=1}^n \log d_y(y_i; \theta_1(\mathbf{x}_i; \boldsymbol{\beta}_1), \dots, \theta_K(\mathbf{x}_i; \boldsymbol{\beta}_K)),$$

with $\mathbf{X} = (\mathbf{X}_1, \dots, \mathbf{X}_K)$.

The log-posterior (frequentist penalized log-likelihood)

$$\log \pi(\boldsymbol{\beta}, \boldsymbol{\tau}; \mathbf{y}, \mathbf{X}, \boldsymbol{\alpha}) \propto \ell(\boldsymbol{\beta}; \mathbf{y}, \mathbf{X}) + \sum_{k=1}^K \sum_{j=1}^{J_k} [\log p_{jk}(\beta_{jk}; \tau_{jk}, \alpha_{jk})],$$

where $p_{jk}(\cdot)$ are priors, τ_{jk} (smoothing) variances and α_{jk} fixed hyper parameters.

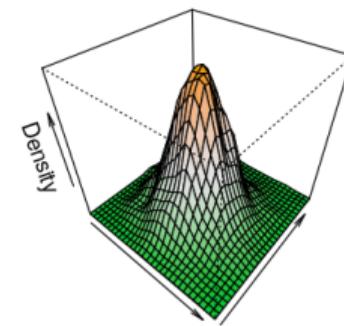
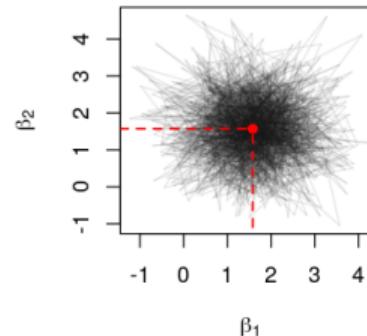
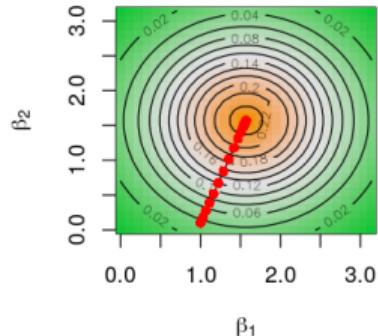
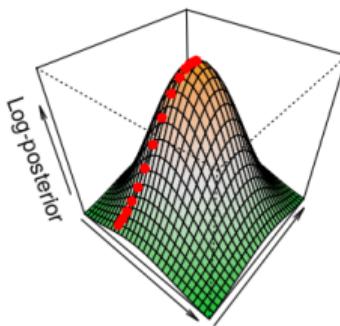
Estimation

Bayesian point estimates of parameters are obtained by:

- ① Maximization of the log-posterior for posterior mode estimation.
- ② Solving high dimensional integrals, e.g., for posterior mean or median estimation.

Problems 1 and 2 are commonly solved by computer intensive iterative algorithms of the following type:

$$(\beta^{[t+1]}, \tau^{[t+1]}) = U(\beta^{[t]}, \tau^{[t]}; \mathbf{y}, \mathbf{X}, \alpha).$$



Scaleable distributional learning

Consider the following updating scheme

$$\boldsymbol{\beta}_k^{[t+1]} = \text{U}_k(\boldsymbol{\beta}_k^{[t]}; \cdot) = \boldsymbol{\beta}_k^{[t]} - \mathbf{H}_{kk} \left(\boldsymbol{\beta}_k^{[t]} \right)^{-1} \mathbf{s} \left(\boldsymbol{\beta}_k^{[t]} \right).$$

Assuming model terms that can be written as a matrix product of a design matrix and coefficients we obtain an iteratively weighted least squares scheme given by

$$\boldsymbol{\beta}_{jk}^{[t+1]} = \text{U}_{jk}(\boldsymbol{\beta}_{jk}^{[t]}; \cdot) = (\mathbf{X}_{jk}^\top \mathbf{W}_{kk} \mathbf{X}_{jk} + \mathbf{G}_{jk}(\tau_{jk}))^{-1} \mathbf{X}_{jk}^\top \mathbf{W}_{kk} (\mathbf{z}_k - \boldsymbol{\eta}_{k,-j}^{[t+1]}),$$

with working observations $\mathbf{z}_k = \boldsymbol{\eta}_k^{[t]} + \mathbf{W}_{kk}^{-1[t]} \mathbf{u}_k^{[t]}$, working weights $\mathbf{W}_{kk}^{-1[t]}$ and score vector $\mathbf{u}_k^{[t]}$.

Scaleable distributional learning

Instead of using all observations of the data, we only use a randomly chosen **subset** denoted by the subindex $[s]$ in one updating step

$$\begin{aligned}\beta_{jk}^{[t+1]} &= \nu \cdot (\mathbf{X}_{[s],jk}^\top \mathbf{W}_{[s],kk} \mathbf{X}_{[s],jk} + \mathbf{G}_{jk}(\boldsymbol{\tau}_{jk}))^{-1} \mathbf{X}_{[s],jk}^\top \mathbf{W}_{[s],kk} (\mathbf{z}_{[s],k} - \boldsymbol{\eta}_{[s],k,-j}^{[t+1]}) + \\ &\quad (1 - \nu) \cdot \beta_{jk}^{[t]},\end{aligned}$$

where ν is a weight parameter which specifies how much the parameters at iteration $t + 1$ are influenced by parameters of the previous iteration t .

Use **flat file** format for each \mathbf{X}_{jk} , i.e., only batch $[s]$ is in memory. This way, we can estimate models with **really** large datasets!

Scaleable distributional learning

Mimics a second order **stochastic gradient descent** (SGD) algorithm

$$\boldsymbol{\beta}_{jk}^{[t+1]} = \boldsymbol{\beta}_{jk}^{[t]} + \nu \cdot (\boldsymbol{\beta}_{jk,[\mathbf{s}]} - \boldsymbol{\beta}_{jk}^{[t]}) = \boldsymbol{\beta}_{jk}^{[t]} + \nu \cdot \boldsymbol{\delta}_{jk}^{[t]},$$

and $\boldsymbol{\delta}_{jk}^{[t]}$ is composed from first and second order derivative information with

$$\begin{aligned}\boldsymbol{\delta}_{jk}^{[t]} &= \boldsymbol{\beta}_{jk,[\mathbf{s}]} - \boldsymbol{\beta}_{jk}^{[t]} \\ &= \left[\boldsymbol{\beta}_{jk}^{[t]} - \mathbf{H}_{[\mathbf{s}],kk} \left(\boldsymbol{\beta}_{jk}^{[t]} \right)^{-1} \mathbf{s}_{[\mathbf{s}]} \left(\boldsymbol{\beta}_{jk}^{[t]} \right) \right] - \boldsymbol{\beta}_{jk}^{[t]} \\ &= -\mathbf{H}_{[\mathbf{s}],kk} \left(\boldsymbol{\beta}_{jk}^{[t]} \right)^{-1} \mathbf{s}_{[\mathbf{s}]} \left(\boldsymbol{\beta}_{jk}^{[t]} \right)\end{aligned}$$

Hence, the updating step length is adaptive.

Scaleable distributional learning

The idea is to select τ_{jk} using a stepwise algorithm which is based on an **"out-of-sample" criterion**, i.e., the criterion $C(\cdot)$ is evaluated on another batch denoted by $[\tilde{s}]$, $C_{[\tilde{s}]}(\cdot)$ respectively, i.e.

$$\tau_{ljk}^{[t+1]} \leftarrow \arg \min_{\tau_{ljk}^* \in \mathcal{I}_{ljk}} C_{[\tilde{s}]}(U_{jk}(\beta_{jk}^{[t]}, \tau_{ljk}^*; \cdot)),$$

where \mathcal{I}_{ljk} is a search interval for $\tau_{ljk}^{[t+1]}$, e.g.,

$$\mathcal{I}_{ljk} = [\tau_{ljk}^{[t]} \cdot 10^{-1}, \tau_{ljk}^{[t]} \cdot 10].$$

Scaleable distributional learning

Some interesting features:

- ① Set, e.g., $\nu = 0.1$, convergence after visiting m batches [s].
- ② Only update if "out-of-sample" log-likelihood is increased.
- ③ **Boosting for variable selection:** Update only $f_{jk}(\cdot)$ with greatest contribution in "out-of-sample" log-likelihood.
- ④ **Bagging:** If $\nu = 1$, each update is so to say a "sample". Convergence similar to MCMC algorithms, i.e., if $\beta_{jk}^{[t+1]}$ start fluctuating around a certain level.
- ⑤ **Slice sample** τ_{ljk} under $C_{[\bar{s}]}(\cdot)$, much faster!

Simulation study

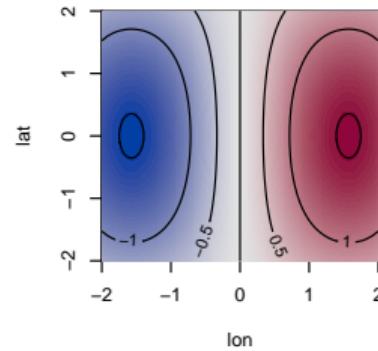
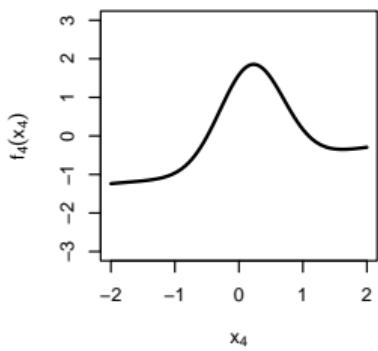
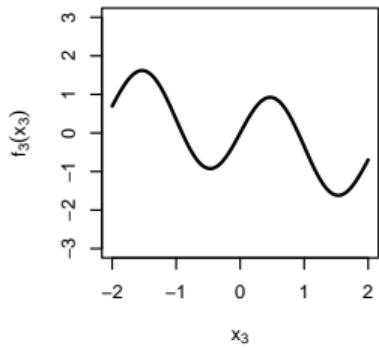
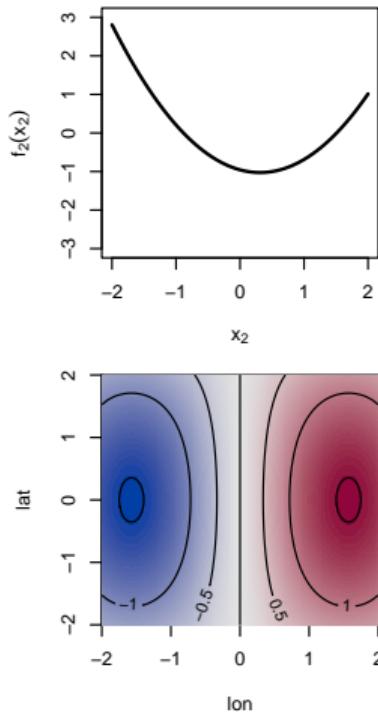
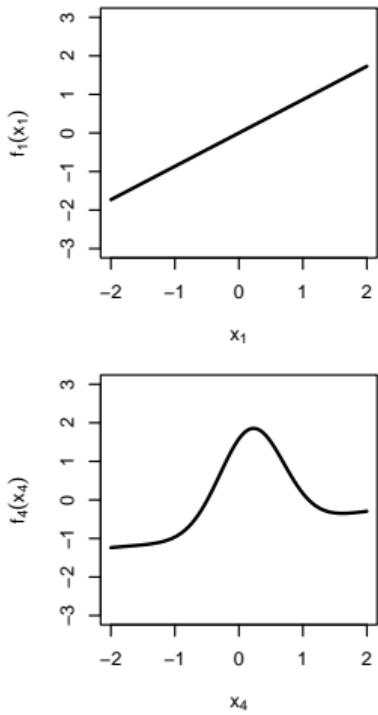
- We consider models using the following predictors:

$$\eta_\mu = f_1(x_1) + f_3(x_3) + f_{2d}(\text{lon}, \text{lat})$$

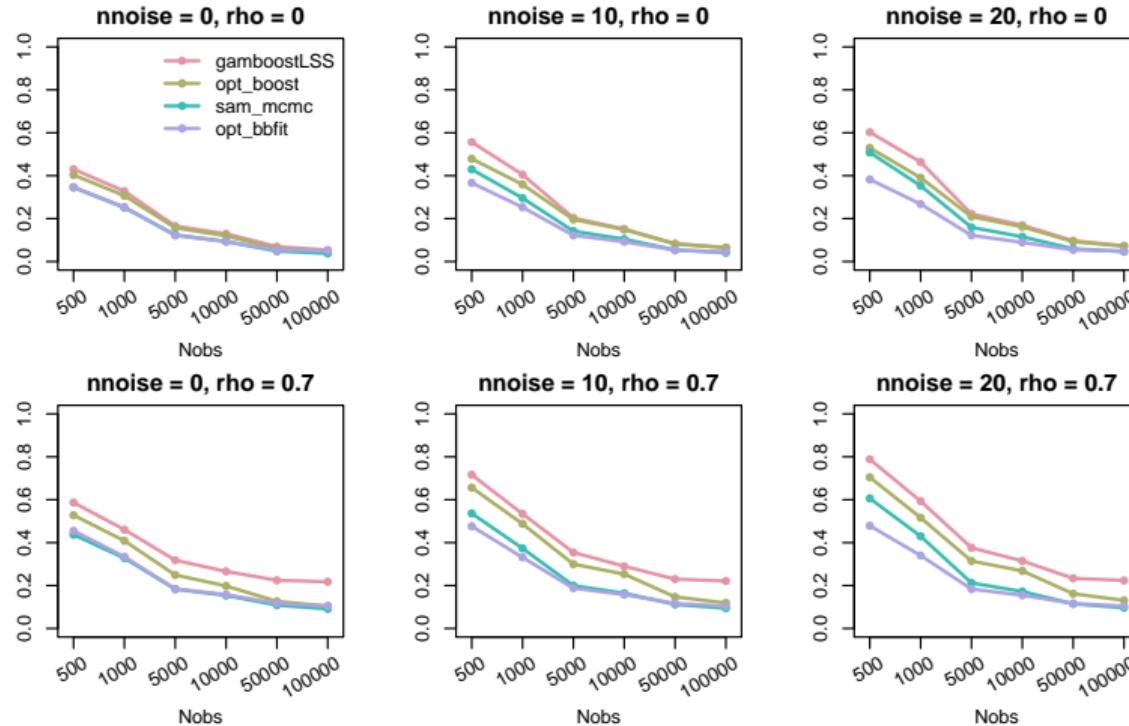
$$\eta_\sigma = f_2(x_2) + f_3(x_3) + f_4(x_4) \text{ with } x_j, \text{lon}, \text{lat} \sim U(-2, 2).$$

- We investigate performance for NO(), GA() and ZAP() distribution,
- using $n = 500; 1000; 5000; 10000; 50000; 100000$ observations.
- Moreover, we study performance using nnoise = 0, 10, 20 variables.
- Two settings for covariate data: uncorrelated and pairwise correlation of $\rho = 0.7$.
- 100 replications each, evaluation using bias, mse, (continuous-) ranked probability score ((C)RPS).
- Algorithms: *gamboostLSS*, *sam_MCMC*, *opt_boost*, *opt_bbfit*.

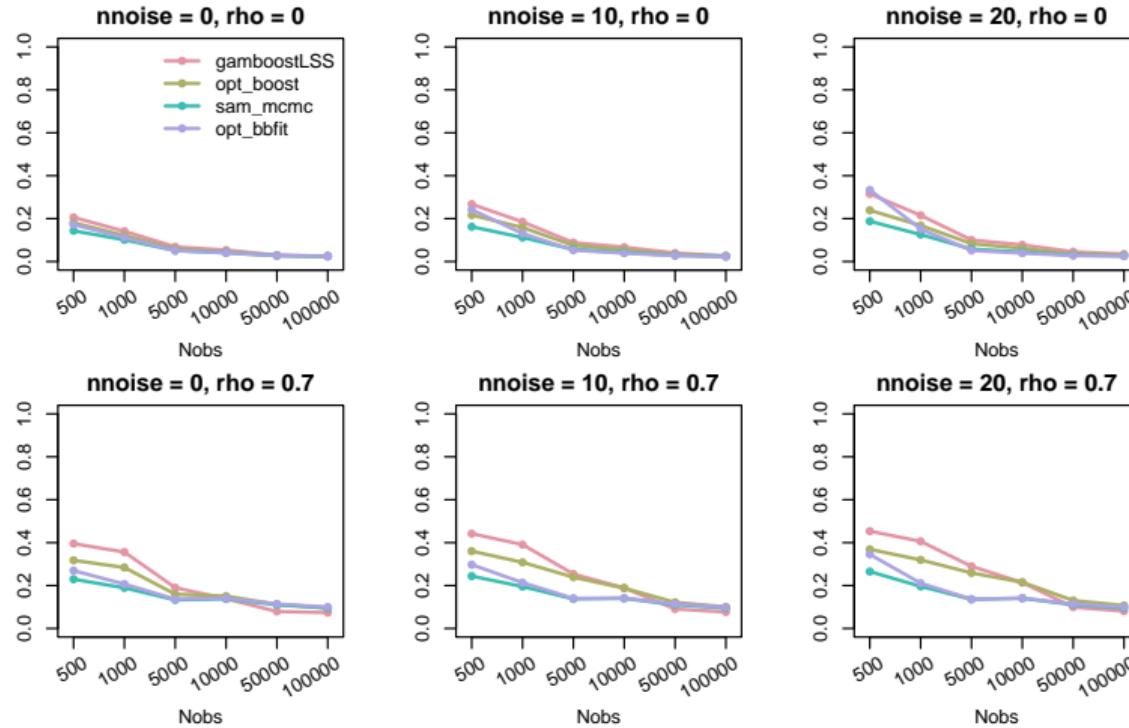
Simulation study



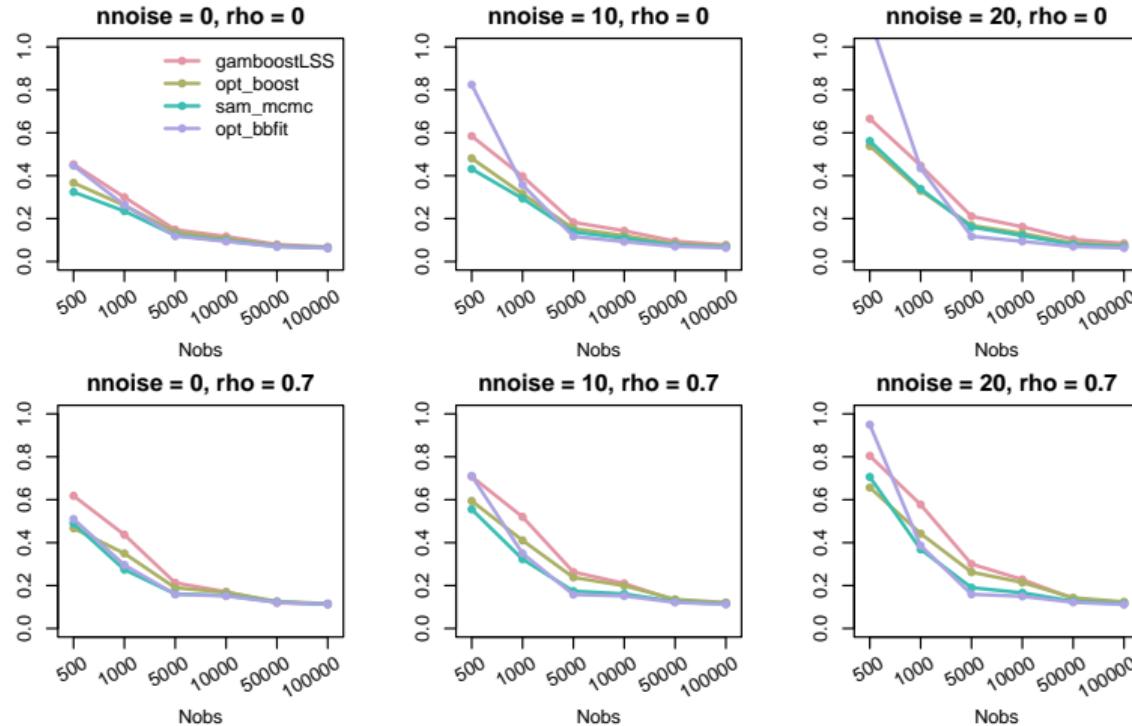
Simulation study: NO(), $\sqrt{\text{bias}}$ η_μ



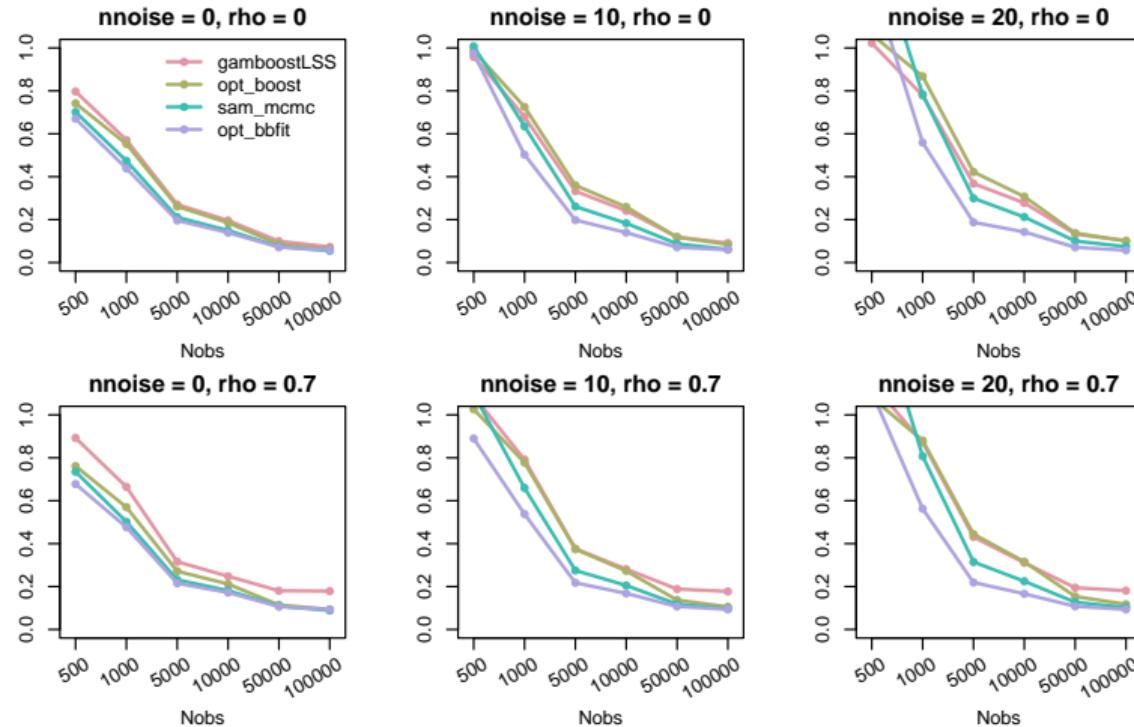
Simulation study: GA(), $\sqrt{\text{bias}}$ η_μ



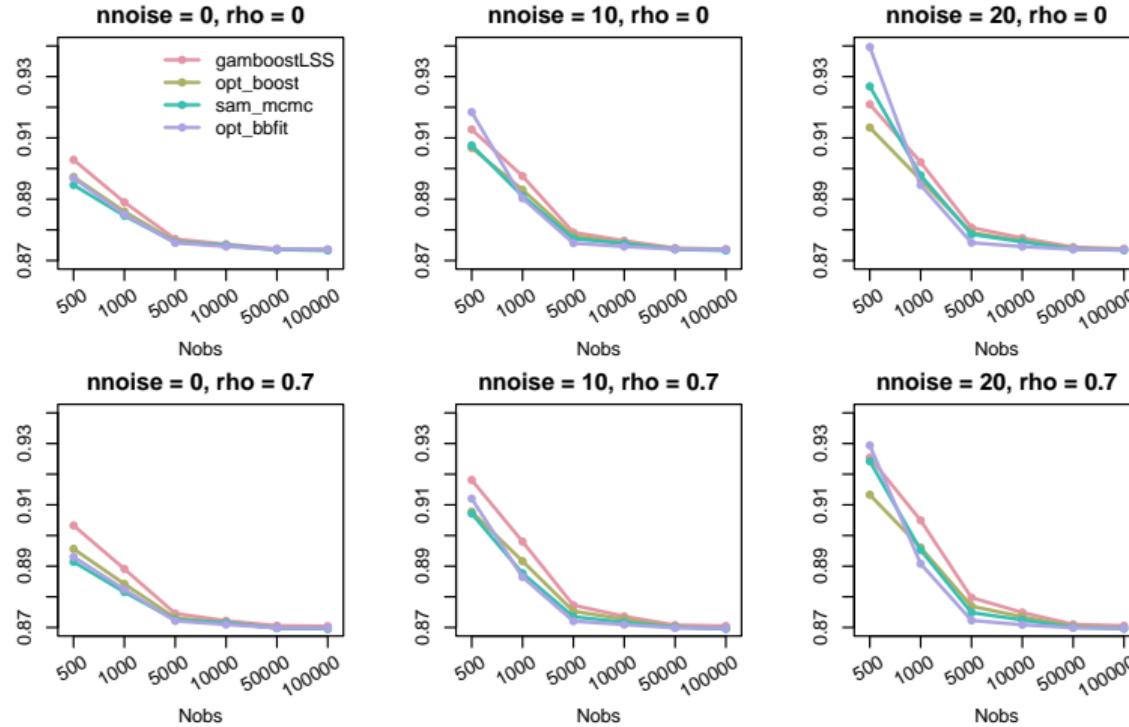
Simulation study: GA(), $\sqrt{\text{bias}}$ η_σ



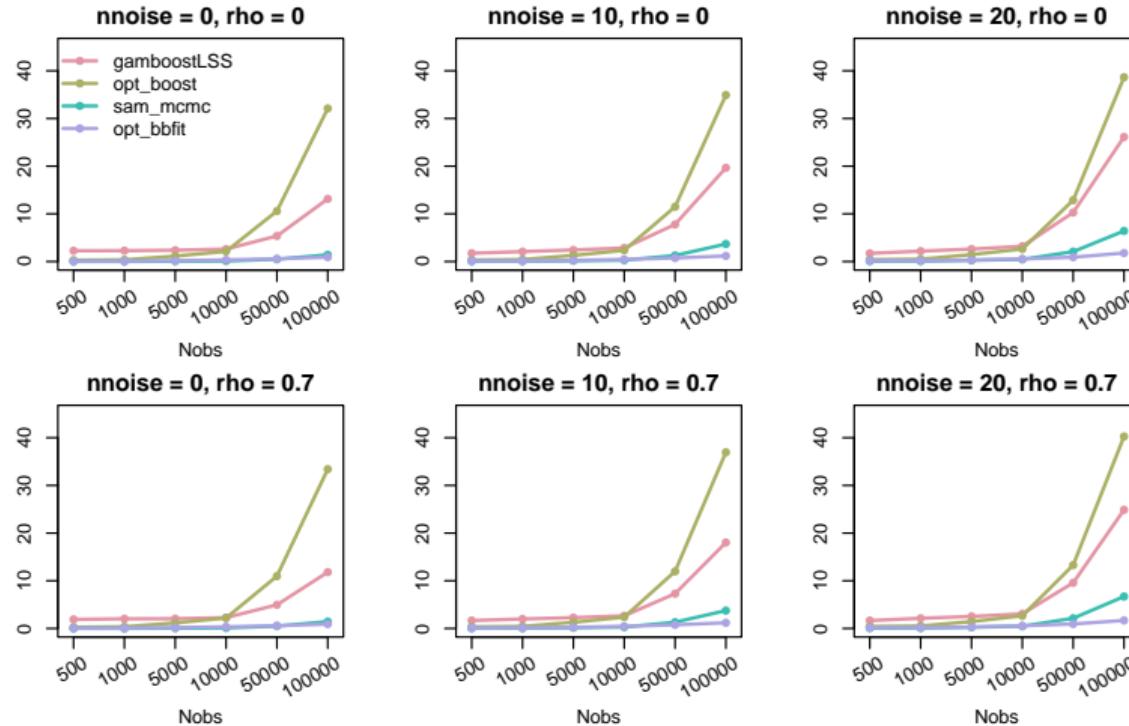
Simulation study: ZAP(), $\sqrt{\text{bias}}$ η_σ



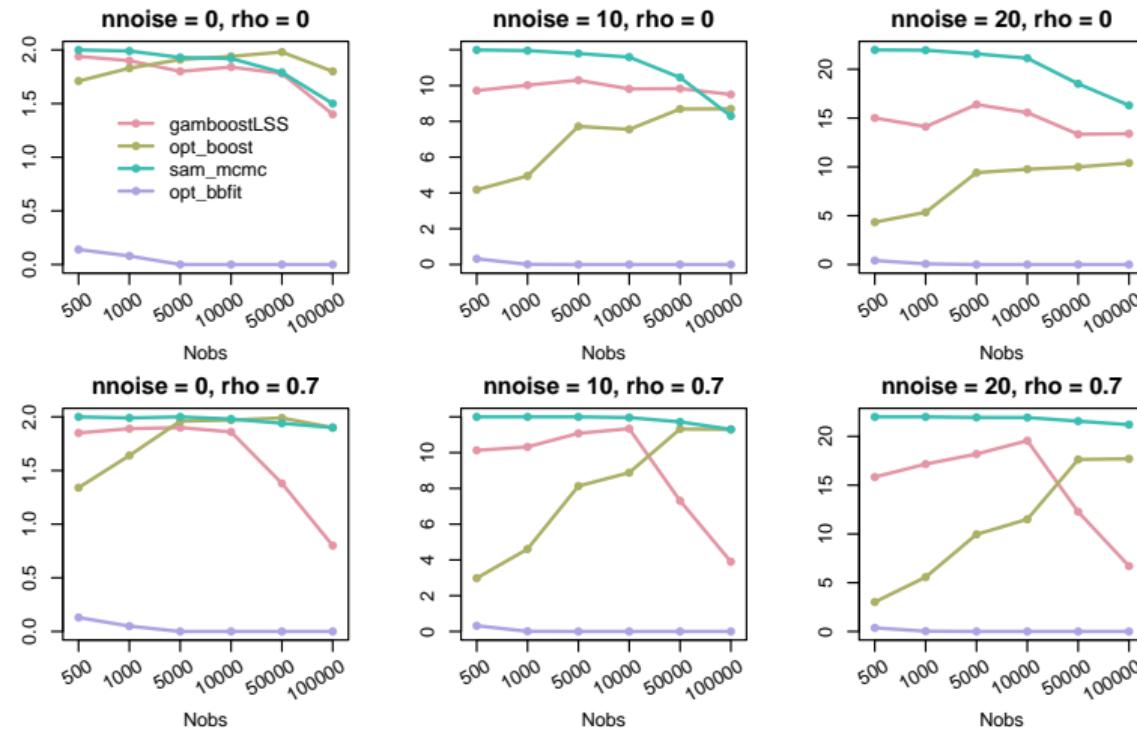
Simulation study: GA(), $\sqrt{\text{CRPS}}$



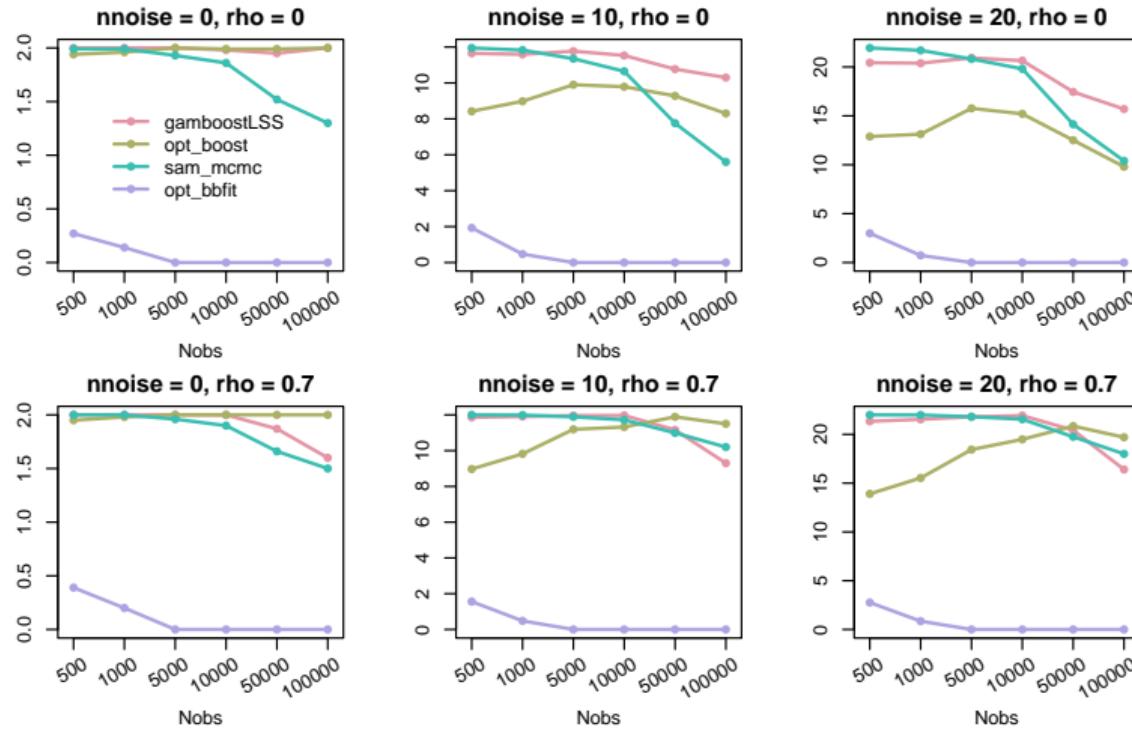
Simulation study: NO(), GA(), ZAP(); runtime [h]



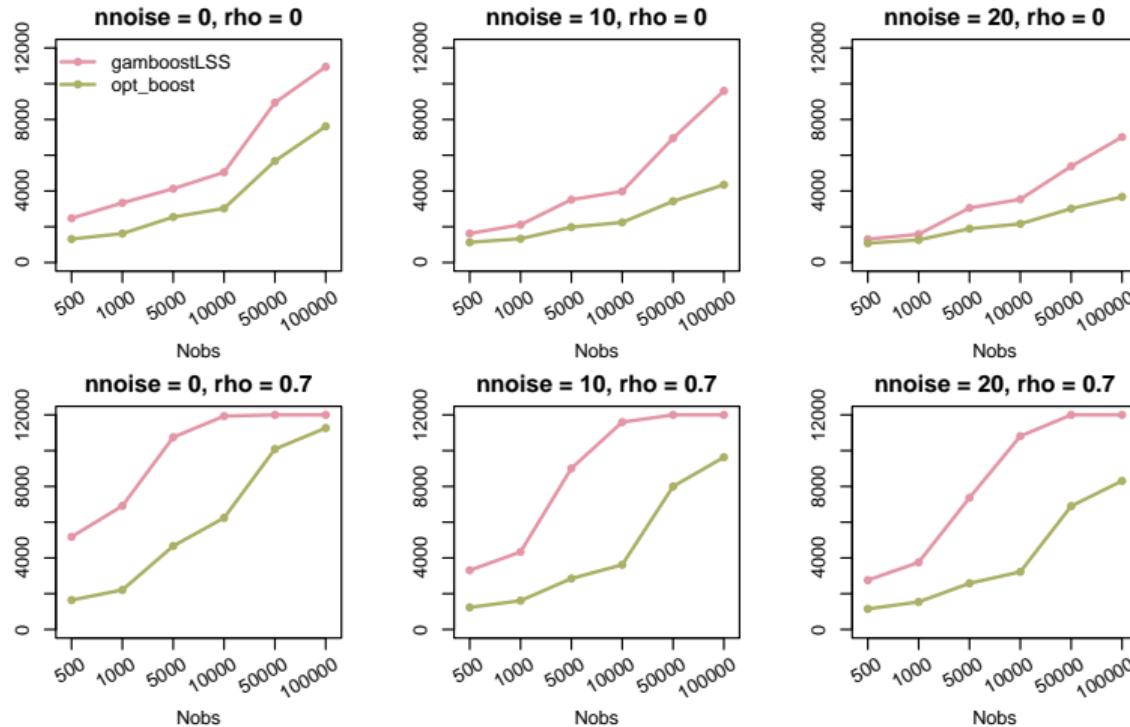
Simulation study: NO(), FP μ



Simulation study: NO(), FP σ



Simulation study: NO(); mstop



Application

Example: Search distribution.

Define the batchsize.

```
R> bs <- 2000
```

Generate batches.

```
R> batch_ids <- lapply(1:200, function(...) {  
+   sample(1:nrow(d), size = bs, replace = FALSE)  
+ })
```

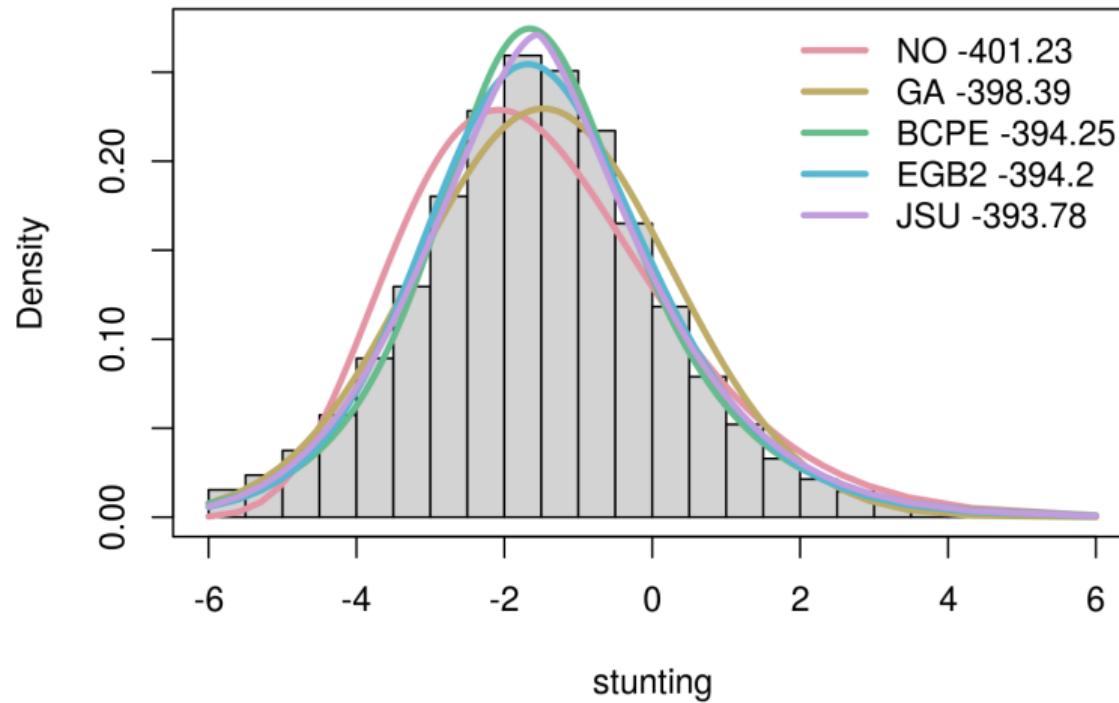
Estimate model.

```
R> b <- bamLSS(y ~ 1, data = d, family = JSU,  
+   sampler = FALSE, optimizer = opt_bbfip, slice = 10, aic = TRUE, K = 2,  
+   batch_ids = batch_ids)
```

Compute log-likelihood.

```
R> logLik(b, newdata = nd)
```

Application



Application

Example: Boosting flavour with *ff* data frame.

Set up a model formula.

```
R> f <- list(  
+   stunting ~ s(cage) + s(bord) + s(hhs) + s(x, y) + ...,  
+   ~ s(cage) + s(bord) + s(hhs) + s(x, y) + ...  
+ )
```

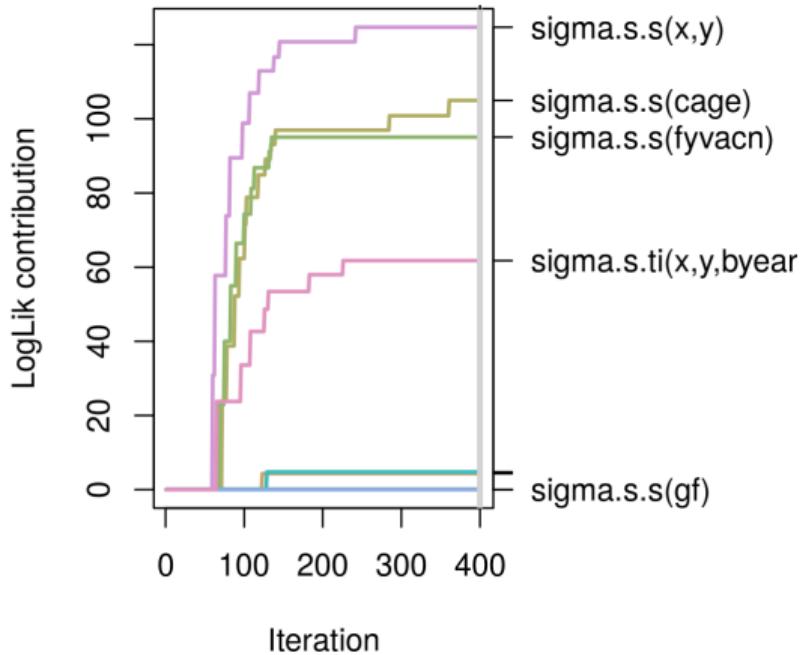
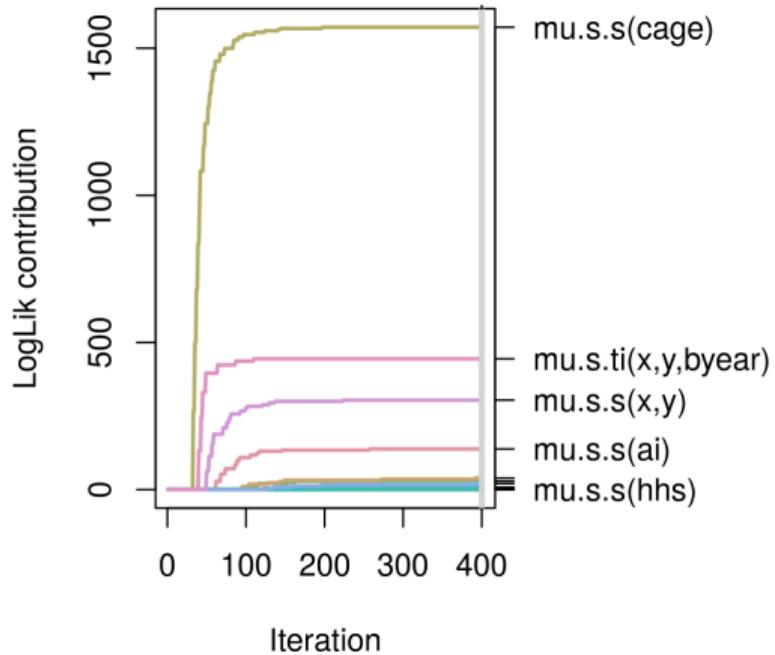
Estimate model.

```
R> b <- bamLSS(f, data = dff, family = JSU,  
+   sampler = FALSE, optimizer = opt_bbfit,  
+   batch_ids = batch_ids, select = TRUE, aic = TRUE, always = FALSE,  
+   eps_loglik = 0.001, K = 2, overwrite = TRUE, delete = FALSE,  
+   ff_name = ff_name)
```

Plot results.

```
R> contribplot(b)
```

Application



Application

Example: Bagging type flavour with slice sampling.

Extract formula.

```
R> nf <- new_formula(b)
```

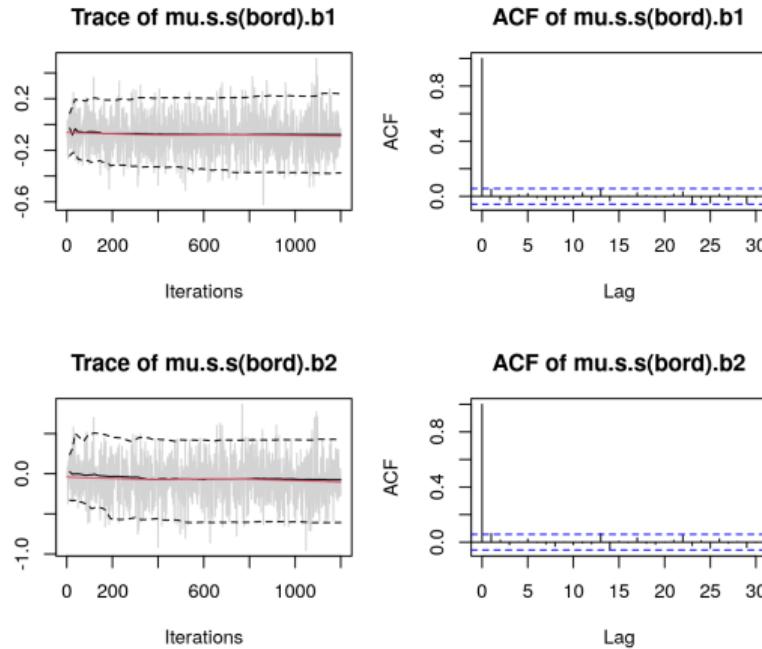
Estimate model using ff processed data.

```
R> m <- bamLSS(nf, data = dff, family = JSU,
+   sampler = FALSE, optimizer = opt_bbfitp,
+   batch_ids = batch_ids, aic = TRUE, slice = TRUE,
+   ff_name = ff_name)
```

Afterwards, all extractor functions provided by *bamLSS* can be used, e.g., `summary()`, `predict()`, `plot()`, etc.

Application

```
R> plot(m, which = "samples")
```



References

- ▶ Umlauf, Klein, and Zeileis (2018). *BAMLSS: Bayesian Additive Models for Location, Scale and Shape (and Beyond)*. Journal of Computational and Graphical Statistics, doi:10.1080/10618600.2017.1407325.
- ▶ Groll, Hambucker, Kneib, and Umlauf (2019). *LASSO-Type Penalization in the Framework of Generalized Additive Models for Location Scale and Shape*. Computational Statistics & Data Analysis, doi:10.1016/j.csda.2019.06.005.
- ▶ Umlauf et al. (2021). *bamlss: Bayesian Additive Models for Location Scale and Shape (and Beyond)*. R package version 1.1-6,
<http://CRAN.R-project.org/package=bamlss>, <http://bamlss.org>.
- ▶ Umlauf, Klein, Simon, Zeileis (2021). *bamlss: A Lego Toolbox for Flexible Bayesian Regression (and Beyond)* . Journal of Statistical Software, doi:10.18637/jss.v100.i04.