

Choosing the threshold in extreme value analysis

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Motivation: peaks over threshold

In the simplest applications:

1. we choose a high threshold u
 - equivalently, choose the number of upper order statistics n_u with threshold $u = X_{(n-n_u)}$.
2. we fit the limiting generalized Pareto distribution with scale σ_u and shape ξ to exceedances $X - u$ over threshold u .
 - if the shape $\xi > 0$, we can use Hill's estimator
3. we use the resulting model for extrapolation beyond u .

Threshold selection

Bias and variance trade-off:

- taking u too high will mean that the number of exceedances n_u is small (increase uncertainty)
- taking u too low will increase n_u , at the risk of a biased extrapolation.

Since the limit holds as u increases to the upper support point of X , we must use a so-called intermediate sequence $n_u/n \rightarrow 0$ as $n_u \rightarrow \infty$ for consistency.

Limiting distribution and threshold stability

- Limiting distribution of threshold exceedances is generalized Pareto. If the limit was exact and $X - u \sim \text{GPD}(\sigma_u, \xi)$

- For higher threshold $v > u$, then

$$X - v \mid X > v \sim \text{GPD}\{\sigma_v = \sigma_u + \xi(v - u), \xi\}.$$

- Provided $\xi > -1$, the expected value of exceedances is

$$\mathbf{E}(X - u) = \sigma_u / (1 + \xi).$$

Literature review

There are earlier reviews of the topic, but the literature keeps increasing...

- The most comprehensive reviews are Scarrott and MacDonald (2012), Caeiro and Gomes (2016) and Langousis et al. (2016).
- Selective numerical comparisons in Gomes and Oliveira (2001), Murphy, Tawn, and Varty (2025) and Schneider, Krajina, and Krivobokova (2021), among others.

Some problems with (most methods)

Consider selecting a threshold among candidates $u_1 < \dots < u_k$.

- Sample overlap: data above different thresholds are overlapping (so dependence between estimates or test statistics and P -values).
- Non-nested models: we only model exceedances above u_j (so most models are not nested).
- Multiple testing problem.
- Potential for automation.
- Non-stationarity (dependence on covariate, change over time).

Objective

We provide an extensive review of threshold selection mechanisms for peak over threshold analysis, including

- semiparametric methods based on Hill's estimator (not covered in this presentation for the sake of time),
- visual diagnostics,
- goodness-of-fit tests,
- extended generalized Pareto models,
- etc.

How to benchmark methods?

What makes a threshold procedure good? In practice, we care about the extrapolation, often

- a high quantile (return level), or
- a probability of exceedance.

Benchmarking the method based on proximity with the asymptotic shape parameter is **not** a good point of reference.

Scale and shape parameters are negatively correlated.

Penultimate effects

Smith (1987) show that a better approximation is obtained by letting the shape vary with u , with $\xi_t = r'(u)$, where

$$r(t) = \{1 - F(t)\} / f(t)$$

is the reciprocal hazard function.

mev package: `smith.penult(family = "norm", method = "pot", qu = c(0.9, 0.95, 0.99))`.

Illustration of penultimate effects: normal example

sampling distribution of shape for MLE based on 1000 exceedances of standard normal

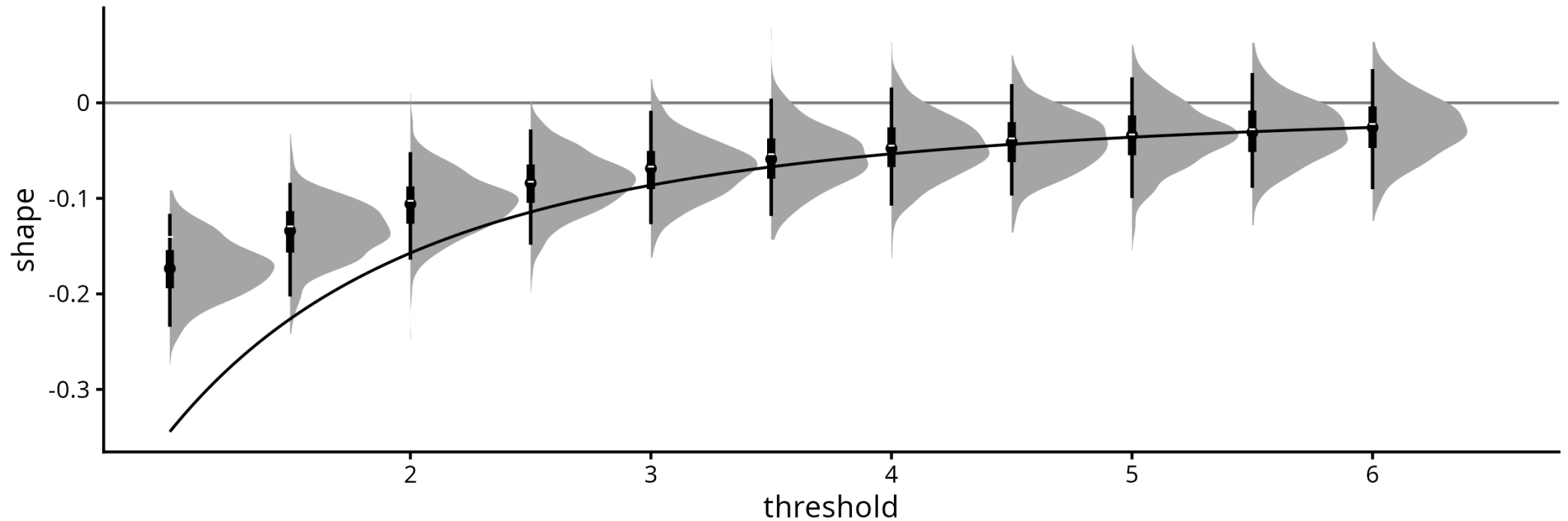
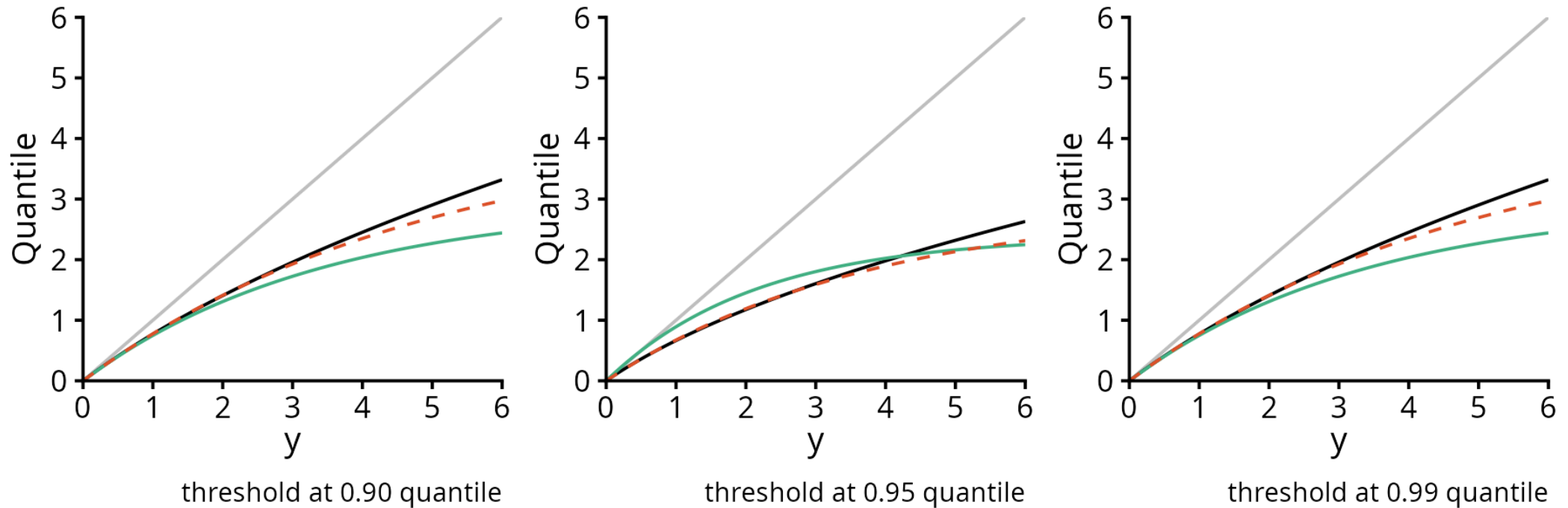


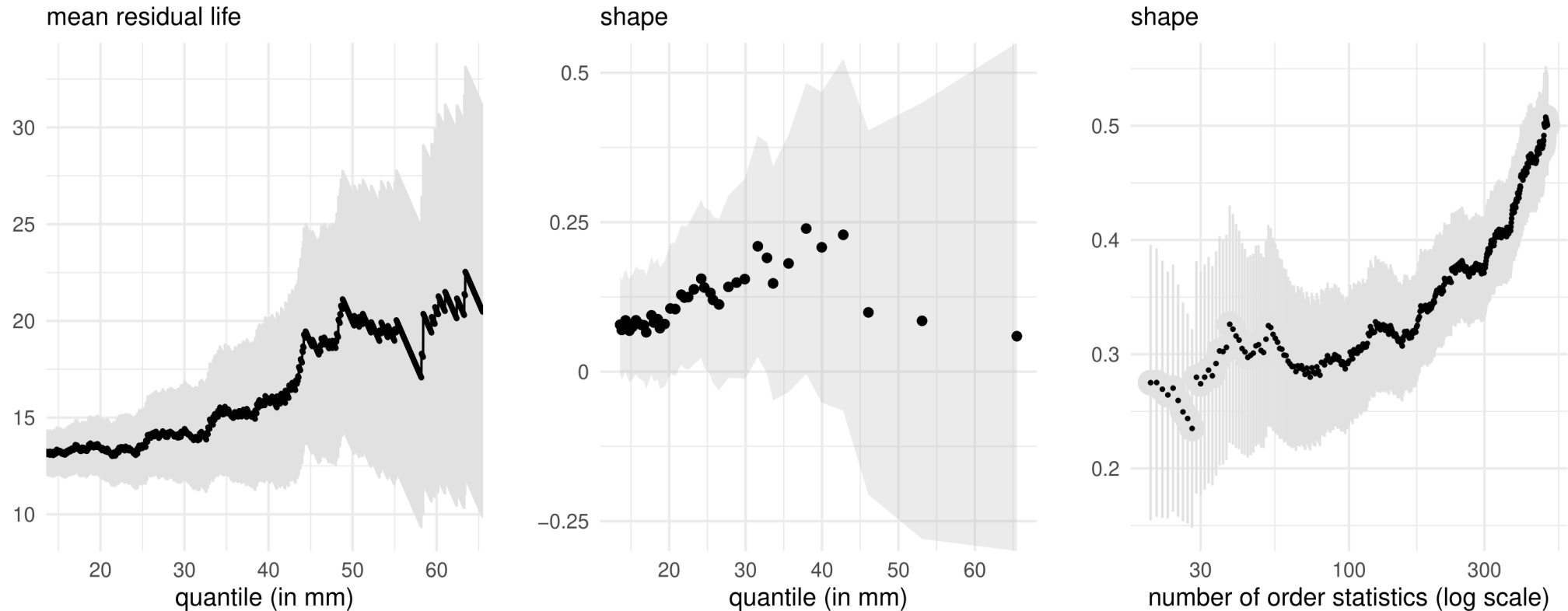
Illustration of penultimate effects: normal quantiles



Conditional quantiles above threshold, rescaled. 45° line shows limiting exponential with true conditional (black), fitted GPD with 1M observations (dashed red) and penultimate (green).

Visual diagnostics on Padova data

Mean residual life ([Davison and Smith 1990](#)), parameter stability and Hill plots.

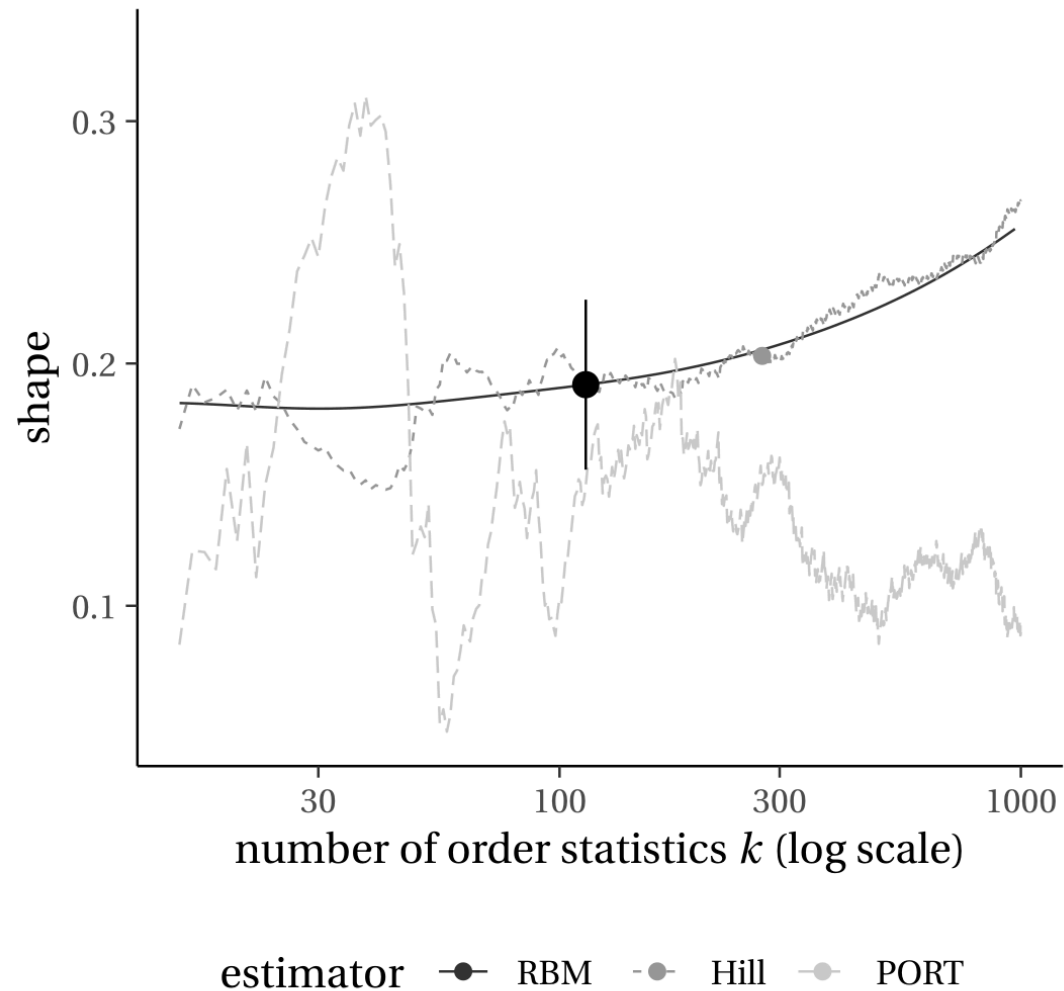


Which threshold would you choose?

Caveats

- Difficulty in automating selection (need visual inspection); proposals in Langousis et al. (2016) or Danielsson et al. (2019), but both fall short.
- Problems: sample overlap leads to dependence between estimates (pointwise confidence intervals).
- Underlying assumptions affected by penultimate effects.
- The plots say nothing about goodness-of-fit!

Stability?



Hill, PORT and random block maxima ([Wager 2014](#)) estimates as a function of n_u .

Generalized Pareto model extensions

Build extended models with additional parameters (with continuity constraints) and test for equality of shape.

Some model are inspired by penultimate approximations (GPD-like, but with varying shape).

Northrop and Coleman (2014): piecewise generalized Pareto model coupled with score tests.

- tests are dependent because the data are re-used; no control of overall error rate.
- the power of the test depends on the choice of thresholds and particularly on the largest threshold u_K .

Extended generalized Pareto models

Embed generalized Pareto $F(x; \sigma, \xi)$ in a more flexible model using a continuous distribution function G_κ on $[0, 1]$. The EGP(σ, ξ, G_κ) distribution function is then

$$\Pr(X \leq x) = G_\kappa\{F(x; \sigma, \xi)\}.$$

Choose G to keep the tail properties. See the chapter of P. Naveau ([2025](#)) for a recent review.

- Papastathopoulos and Tawn ([2013](#)): models imply the density at the origin is zero.
- Gamet and Jalbert ([2022](#)) propose two models, but one leads to non-regular asymptotics (restriction on boundary of the parameter space).
- Philippe Naveau et al. ([2016](#)) additional models with two parameters.

Extended generalized Pareto models

Test for restriction to generalized Pareto sub-model using likelihood ratio tests (profile).

- Could allow for a bit more data to be included, at the expense of additional parameters to estimate (and potentially more variability).
- Same problems as parameter stability plots (sequential tests, overlapping data).

Splicing models

Glue a distribution for the bulk with one for the tail using a mixture of disjoint components below u (bulk) and above u (generalized Pareto). See Scarrott and MacDonald (2012) and Hu and Scarrott (2018).

- Leads to sample contamination, fit may be driven by bulk.
- profile likelihood for threshold u is non-monotone.
- Bayesian version with random threshold (Nascimento, Gamerman, and Lopes 2012) provides threshold selection uncertainty.

Goodness-of-fit measures

1. Fit a generalized Pareto distribution at each candidate threshold.
2. Compute either
 - a suitable statistic which indicates departure from the posulated distribution.
 - a measure of discrepancy between empirical distribution of exceedances above u and generalized Pareto model (via Kolmogorov–Smirnov, Cramér–von Mises, etc.)
3. Perform tests sequentially until rejection, or select the “best” threshold according to the criterion.

Some proposals

- Idea dates back to Pickands ([1975](#)).
- Choulakian and Stephens ([2001](#)), Bader, Yan, and Zhang ([2018](#)) (using ForwardStop).
- Recent proposals using L -moment estimators including Kiran and Srinivas ([2021](#)), Solari et al. ([2017](#)), Silva Lomba and Fraga Alves ([2020](#)).

Note: goodness-of-fit tests null distributions require adjustment for rounded values (estimate null via Monte Carlo).

Testing using maximum likelihood distribution

Thompson et al. (2009) propose using constant values

$$\tau_j = \hat{\sigma}_j - \hat{\xi}_j u_j$$

and performing Pearson's test of normality for the differences $\tau_{j+1} - \tau_j$ ($j = 1, \dots, k - 1$), stopping whenever the hypothesis is rejected at level $\alpha = 0.2$.

Ad hoc proposal... but works well in simulations.

Sequential analysis

Wadsworth (2016) obtains asymptotic joint distribution of MLE from a superposition of Poisson processes.

Build independent increments of shape to form a white noise sequence $\xi_i^* = (\hat{\xi}_{u_{i+1}} - \hat{\xi}_{u_i}) / \{(I_{u_{i+1}}^{-1} - I_{u_i}^{-1})_{\xi, \xi}^{1/2}\}$.

- Parameter stability plots with simultaneous confidence intervals!
- White noise test for ξ_i^* , with alternative hypothesis $\mathcal{H}_a : \xi_i^* \sim \text{Normal}(\beta, \sigma) (i = 1, \dots, j - 1)$ and $\xi_i^* \sim \text{Normal}(0, 1)$ for $j, \dots, k - 1$ motivated by results on model misspecification (White 1982).

Comments on Wadsworth (2016)

Fails 17% of the time in our simulations with equally spaced quantiles.

- Problems with positive definiteness of information matrices for shape increments.
- Method sensitive to rounding (add noise?)
- Must choose the threshold sequence carefully.

Predictive distribution

Northrop, Attalides, and Jonathan (2017) propose a Bayesian method based on leave-one-out cross validation with a binomial-generalized Pareto (BGP) model and a single validation threshold $v > u_k$ above which we assess the model performance.

The measure of goodness-of-fit proposed is an estimate of the negated Kullback–Leibler divergence,

$$\hat{T}_v(u_i) = \sum_{i=1}^n \log\{\hat{f}_v(x_r \mid \mathbf{x}_{-r}, u_i)\}.$$

The selected threshold is the one among the candidates maximizing this diagnostic.

Further propose using Bayesian model averaging to account for the uncertainty originating from threshold selection.

Metric-based diagnostic

Build quantile-quantile (QQ-) plot, with pointwise confidence intervals can be obtained using a parametric bootstrap.

Each bootstrap sample $b = 1, \dots, B$, estimate the empirical quantile function F_b , which is evaluated at the plotting position $p_i = i/(n + 1)$ for $i = 1, \dots, n$.

Varty et al. (2021) and Murphy, Tawn, and Varty (2025) propose repeating this with simulated iid data from F_0 (tolerance intervals).

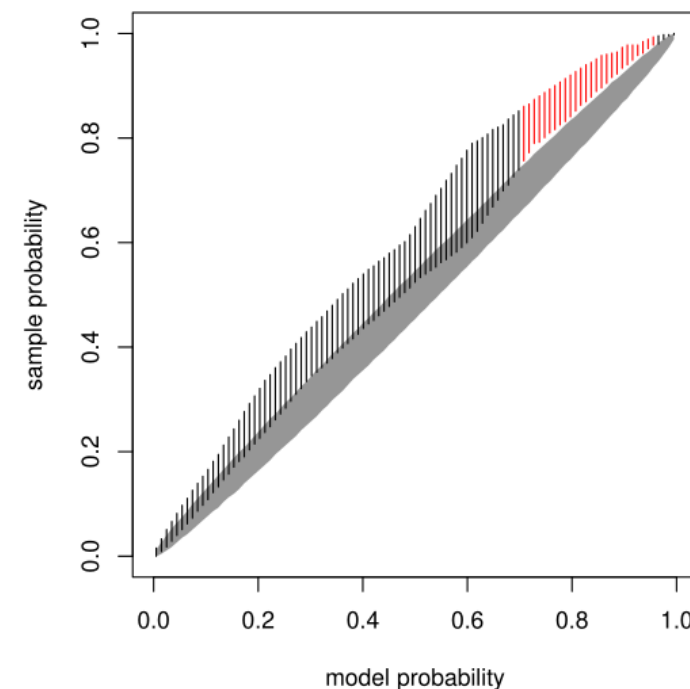


Figure from Varty et al. (2021)

Metric-based adjustment

Build metric based on exponential (or generalized Pareto quantile $F_0^{-1}\{p_i\}$ against $F_b^{-1}(p_i)$ with mean absolute difference or mean squared difference.

Pick the threshold with the smallest average distance is chosen.

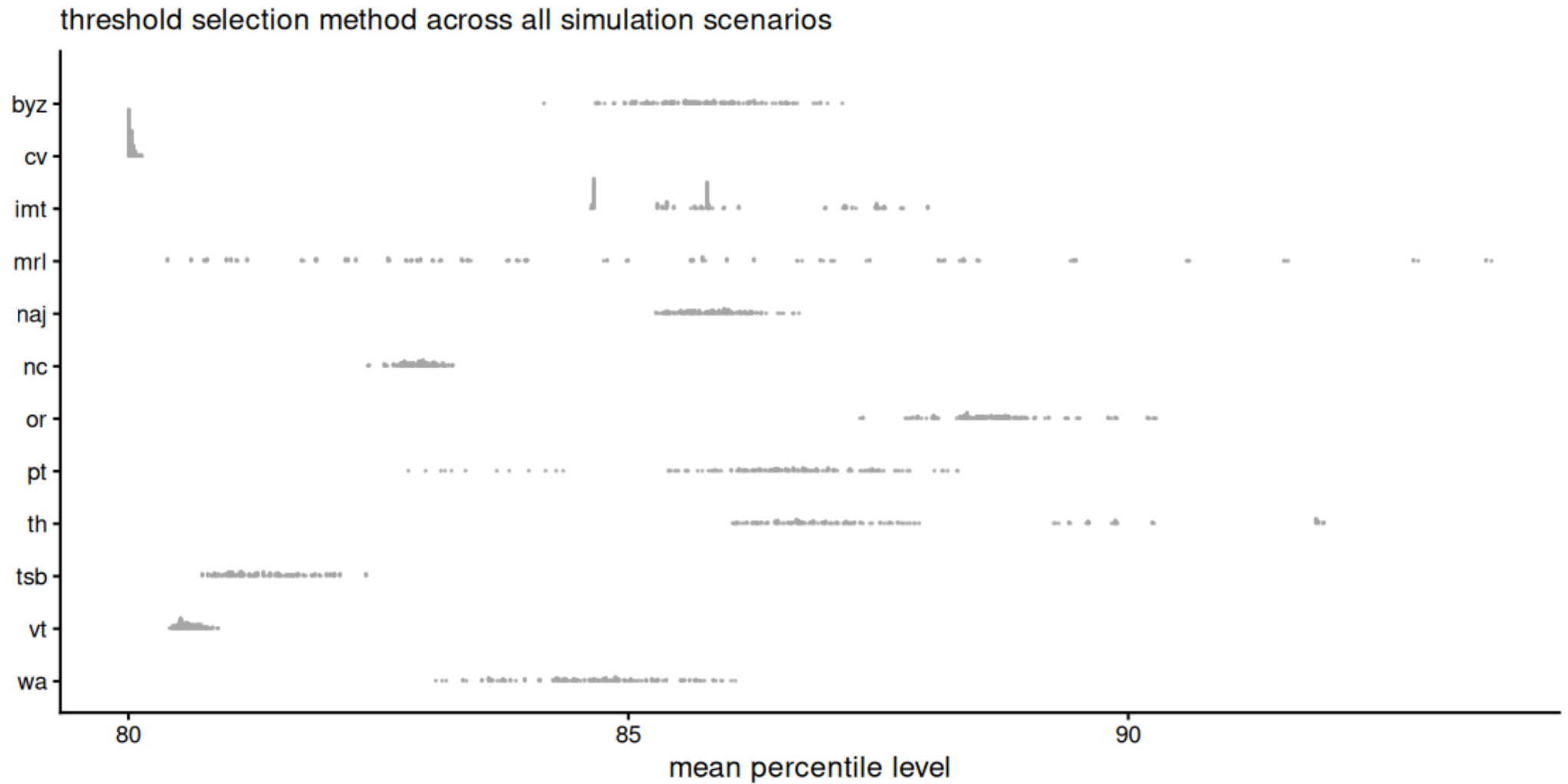
- amenable to different sampling schemes (censoring, non-identically distributed data, time-varying thresholds).
- computationally-intensive by design.

Simulation study: comparison of methods

We considered 13 different distributions from simulation studies in Choulakian and Stephens (2001) and Schneider, Krajina, and Krivobokova (2021).

- Consider 1000 replications of IID data, data with serial correlation in the tail, rounded observations, with different tail behaviour.
- Data of size $n \in \{1000, 2000\}$ with candidate thresholds at the sample $\{0.8, 0.81, \dots, 0.99\}$ quantiles, keeping a minimum of 20 exceedances.
- Evaluate bias, variance, RMSE of point estimator for 0.999 quantile.
- Compare to oracle (model with the closest quantile estimate among candidates).

Which quantile level on average?



Findings

- No universally better method.
- Many of the automated procedures return the lowest possible threshold.
- Using Forward stop method to account for multiple testing leads to thresholds that are much lower.

More comments

- The oracle method returns average threshold around the 87% and the 90% quantile.
- Mean residual life plots are uniformly scattered.
- Varty et al. (2021)'s metric diagnostic also leads to very small thresholds.
- Northrop, Attalides, and Jonathan (2017) and Thompson et al. (2009) lead to much less agreement and a greater variability of selected quantile levels for the thresholds. For the cross-validation approach, this is in line with findings of Murphy, Tawn, and Varty (2025).
- Wadsworth's sequential testing performs best with heavy tailed distributions, but otherwise is one of the worst in terms of ranking.

Conclusions and future work

Is the problem well-formulated? There is no “correct” threshold, so are we barking up the wrong tree?

Some alternatives:

- Weighting with different threshold choices to account for uncertainty ([Stein 2023](#)).
- Should we be fitting sub-asymptotic models to much more data?

Question period

- Thank you for your attention and thanks to the organizers.
- Preprint coming soon.
- Many numerical methods in **R** packages `mev` and `tea`.
- Slides at lbelzile.github.io/EVA2025-choosing-threshold

Semiparametric methods

The paper also compares 18 different semiparametric methods using Hill-type estimators for heavy-tailed data.

- Careful: many numerical implementations don't specify a minimum sample size!
- Primer: best methods include Caeiro and Gomes (2016), Gomes, Figueiredo, and Neves (2012), Wager (2014).

Don't use the following

- Methods based on minimization of the asymptotic mean squared error can break down catastrophically for particular data sets.
- Methods by Gomes et al. (2013) and Hall and Welsh (1985) behave erratically with small shape parameters: these procedures lead to strongly biased shape parameter estimates. This is due to them keeping more than 15% of the data for inference.
- Drees and Kaufmann (1998) bias-reduction method leads to large width of confidence intervals (unwanted variability), an
- Many other methods are extremely variable.

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