

Cross-validatory extreme value threshold selection and uncertainty with application to ocean storm severity

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Abstract

Designs conditions for marine structures are typically informed by threshold-based extreme value analyses of oceanographic variables, in which excesses of a high threshold are modelled by a generalized Pareto (GP) distribution. Too low a threshold leads to bias from model mis-specification; raising the threshold increases the variance of estimators: a bias-variance trade-off. Many existing threshold selection methods do not address this trade-off directly, but rather aim to select the lowest threshold above which the GP model is judged to hold approximately. In this paper Bayesian cross-validation is used to address the trade-off by comparing thresholds based on predictive ability at extreme levels. Extremal inferences can be sensitive to the choice of a single threshold. We use Bayesian model-averaging to combine inferences from many thresholds, thereby reducing sensitivity to the choice of a single threshold. The methodology is applied to significant wave height datasets from the northern North Sea and the Gulf of Mexico.

Keywords. Cross-validation; Extreme value theory; Generalized Pareto distribution; Predictive inference; Threshold

1 Introduction

Ocean and coastal structures, including breakwaters, ships and oil and gas producing facilities are designed to withstand extreme environmental conditions. Marine engineering design codes stipulate that estimated failure probabilities of offshore structures, associated with one or more return periods, should not exceed specified values. To characterize the environmental loading on an offshore structure, return values for winds, waves and ocean currents corresponding to a return period of typically 100 years, but sometimes to 1,000 and 10,000 years are required. The severity of waves in a storm is quantified using significant wave height. Extreme value analyses of measured and hindcast samples of significant wave height are undertaken to derive environmental design criteria, typically by fitting a GP distribution to excesses of a high threshold. The selection of appropriate threshold(s) is important because inferences can be sensitive to threshold.

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1.1 Storm peak significant wave height datasets

The focus of this paper is the analysis of two sequences of hindcasts of storm peak significant wave height, shown in Figure 1. Significant wave height (H_s) is a measure of sea surface roughness. It is defined as four times the standard deviation of the surface elevation of a *sea state*, the ocean surface observed for a certain period of time (3 hours for our datasets). Cardone et al. (2014) gives the largest H_s value ever generated by a hindcast model as 18.33m and the largest value ever measured in the ocean as 20.63m. Hindcasts are samples from physical models of the ocean environment, calibrated to observations of pressure, wind and wave fields.

For each of the datasets raw data have been declustered, using a procedure described in Ewans and Jonathan (2008), to isolate cluster maxima (storm peaks) that can reasonably be treated as being mutually independent. We also assume that storm peaks are sampled from a common distribution. Even in this simplest of situations practitioners have difficulty in selecting appropriate thresholds. The first dataset, from an unnamed location in the northern North Sea, contains 628 storm peaks from October 1964 to March 1995, but restricted to the period October to March within each year. The other dataset contains 315 storm peaks from September 1900 to September 2005. It is one of the datasets from a spatial grid of time series in the Gulf of Mexico analysed in Northrop and Jonathan (2011): the location is at the centre of this spatial grid.

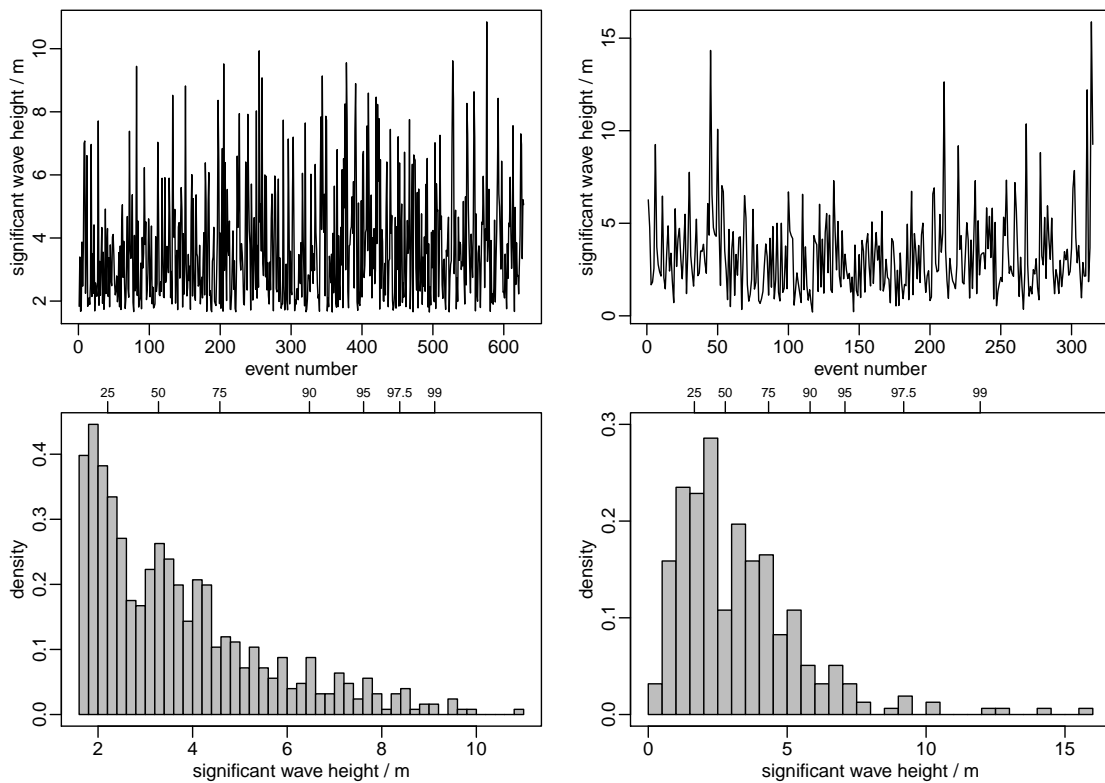


Figure 1: Storm peak significant wave height hindcast datasets. Left: North Sea data (628 observations). Right: Gulf of Mexico data (315 observations). Top: time series plots. Bottom: histograms. The upper axis scales give the sample quantile levels.

In the northern North Sea the main fetches are the Norwegian Sea to the North, the Atlantic Ocean to the west, and the North Sea to the south. Extreme sea states from the directions of Scandinavia to the east and the British Isles to the southwest are not possible, owing to the shielding effects of these land masses. At the location under consideration, the most extreme sea states are associated with storms from the Atlantic Ocean (Ewans and Jonathan, 2008). With up to several tens of storms impacting the North Sea each winter, the number of events for analysis is typically larger than for locations in regions such as the Gulf

of Mexico, where hurricanes produce the most severe sea states. Most hurricanes originate in the Atlantic Ocean between June and November and propagate west to northwest into the Gulf producing the largest sea states with dominant directions from the southeast to east directions. It is expected that there is greater potential for very stormy sea conditions in the Gulf of Mexico than in the northern North Sea and therefore that the extremal behaviour is different in these two locations.

1.2 Extreme value threshold selection

Extreme value theory provides asymptotic justification for a particular family of models for excesses of a high threshold. Let X_1, X_2, \dots, X_n be a sequence of independent and identically distributed random variables, with common distribution function H , and u_n be a threshold, increasing with n . Pickands (1975) showed that if there is a non-degenerate limiting distribution for appropriately linearly rescaled excesses of u_n then this limit is a Generalized Pareto (GP) distribution. In practice, a suitably high threshold u is chosen empirically. Given that there is an exceedance of u , the excess $Y = X - u$ is modelled by a $\text{GP}(\sigma_u, \xi)$ distribution, with positive threshold-dependent scale parameter σ_u , shape parameter ξ and distribution function

$$G(y) = \begin{cases} 1 - (1 + \xi y / \sigma_u)_+^{-1/\xi}, & \xi \neq 0, \\ 1 - \exp(-y / \sigma_u), & \xi = 0, \end{cases} \quad (1)$$

where $y > 0$, $x_+ = \max(x, 0)$. The $\xi = 0$ case is defined in the limit as $\xi \rightarrow 0$. When $\xi < 0$ the distribution of X has a finite upper endpoint of $u - \sigma_u / \xi$; otherwise, X is unbounded above. The frequency with which the threshold is exceeded also matters. Under the current assumptions the number of exceedances u has a binomial(n, p_u) distribution, where $p_u = P(X > u)$, giving a $\text{BGP}(p_u, \sigma_u, \xi)$ model (Coles, 2001, chapter 4)

Many threshold diagnostic procedures have been proposed: Scarrott and MacDonald (2012) provides a review. Broad categories of methods include: assessing stability of model parameter estimates with threshold (Drees et al., 2000; Wadsworth, 2015); goodness-of-fit tests (Davison and Smith, 1990; Dupuis, 1998); approaches that minimize the asymptotic mean-squared error of estimators of ξ or of extreme quantiles, under particular assumptions about the form of the upper tail of H (Hall and Welsh, 1985; Hall, 1990; Ferreira et al., 2003; Beirlant et al., 2004); specifying a model for (some or all) data below the threshold (Wong and Li, 2010; MacDonald et al., 2011; Wadsworth and Tawn, 2012). In the latter category, the threshold above which the GP model is assumed to hold is treated as a model parameter and *threshold uncertainty* is incorporated by averaging inferences over a posterior distribution of model parameters. In contrast, in a *single threshold approach* threshold level is viewed as a tuning parameter, whose value is selected prior to the main analysis and is treated as fixed and known when subsequent inferences are made.

Single threshold selection involves a bias-variance trade-off (Scarrott and MacDonald, 2012): the lower the threshold the greater the estimation bias due to model misspecification; the higher the threshold the greater the estimation uncertainty. Many existing approaches do not address the trade-off directly, but rather examine sensitivity of inferences to threshold and/or aim to select the lowest threshold above which the GP model appears to hold approximately. We seek to deal with the bias-variance trade-off based on the main purpose of the modelling, i.e. prediction of future extremal behaviour. We make use of a data-driven method commonly used for such purposes: cross-validation (Stone, 1974). We consider only the simplest of modelling situations, i.e. where observations are treated as independent and identically distributed. However, selecting the threshold level is a fundamental issue for all threshold-based extreme value analyses and we anticipate that our general approach can have much wider applicability.

We illustrate some of the issues involved in threshold selection by applying to the significant wave height datasets two approaches that assess parameter stability. In the top row of Figure

2 maximum likelihood (ML) estimates $\widehat{\xi}$ of ξ are plotted against threshold. The aim is to choose the lowest threshold above which $\widehat{\xi}$ is approximately constant in threshold, taking into account sampling variability summarized by the confidence intervals. It is not possible to make a definitive choice and different viewers may choose rather different thresholds. In both of these plots *our* eyes are drawn to the approximate stability of the estimates at around the 70% sample quantile. One could argue for lower thresholds, to incur some bias in return for reduced variance, but it is not possible to assess this objectively from these plots. In practice, it is common not to consider thresholds below the apparent mode of the data because the GP distribution has its mode at the origin. For example, based on the histogram of the Gulf of Mexico data in Figure 1 one might judge a threshold below the 25% sample quantile to be unrealistic. However, we *will* consider such thresholds because it is interesting to see to what extent the bias expected is offset by a gain in precision.

The inherent subjectivity of this approach, and other issues such as the strong dependence between estimates of ξ based on different thresholds, motivated more formal assessments of parameter stability (Wadsworth and Tawn, 2012; Northrop and Coleman, 2014; Wadsworth, 2015). The plots in the bottom row of Figure 2 are based on Northrop and Coleman (2014). A subasymptotic piecewise constant model (Wadsworth and Tawn, 2012) is used in which the value of ξ may change at each of a set of thresholds, here set at the 0%, 5%, . . . , 95% sample quantiles. For a given threshold the null hypothesis that the shape parameter is constant from this threshold upwards is tested. In the plots p -values from this test are plotted against threshold. Although these plots address many of the inadequacies of the parameter estimate stability plots subjectivity remains because one must decide how to make use of the p -values. One could prespecify a size, e.g. 0.05, for the tests or take a more informal approach by looking for a sharp increase in p -value. For the North Sea data the former would suggest a very low threshold and the latter a threshold in the region of the 70% sample quantile. For the Gulf of Mexico data respective thresholds close to the 10% and 55% sample quantiles are indicated.

An argument against selecting a single threshold is that this ignores uncertainty concerning the choice of this threshold. As mentioned above, one way to account for this uncertainty is to embed a threshold parameter within a model. We use an approach based on Bayesian Model Averaging (BMA), on which Hoeting et al. (1999) provide a review. Sabourin et al. (2013) have recently used a similar approach to combine inferences from different multivariate extreme value models. We treat different thresholds as providing competing models for the data. Predictions of extremal behaviour are averaged over these models, with individual models weighted in proportion to the extent to which they are supported by the data. There is empirical and theoretical evidence (Hoeting et al., 1999) that averaging inferences in this way results in better average predictive ability than provided by any single model.

For the most part we work in a Bayesian framework because prediction is handled naturally and regularity conditions required for making inferences using ML (Smith, 1985) and probability-weighted moments (PWM) (Hosking and Wallis, 1987), namely $\xi > -1/2$ and $\xi < 1/2$ respectively, can be relaxed. This requires a prior distribution to be specified for the parameters of the BGP model. Initially, we consider three different ‘reference’ prior distributions, in the general sense of priors constructed using formal rules (Kass and Wasserman, 1996). Such priors can be useful when information provided by the data is much stronger than prior information from other sources. This is more likely to be the case for a low threshold than for a high threshold. We use simulation to assess the utility of these priors for our purpose, that is, making predictive statements about future extreme observations and use the results to formulate an improved prior. For high thresholds, when the data are likely to provide only weak information, it may be important to incorporate at least some basic prior information in order to avoid making physically unrealistic inferences.

In section 2 we use Bayesian cross-validation to estimate a measure of threshold performance for use in the selection of a single threshold. In section 3 we consider how to use this

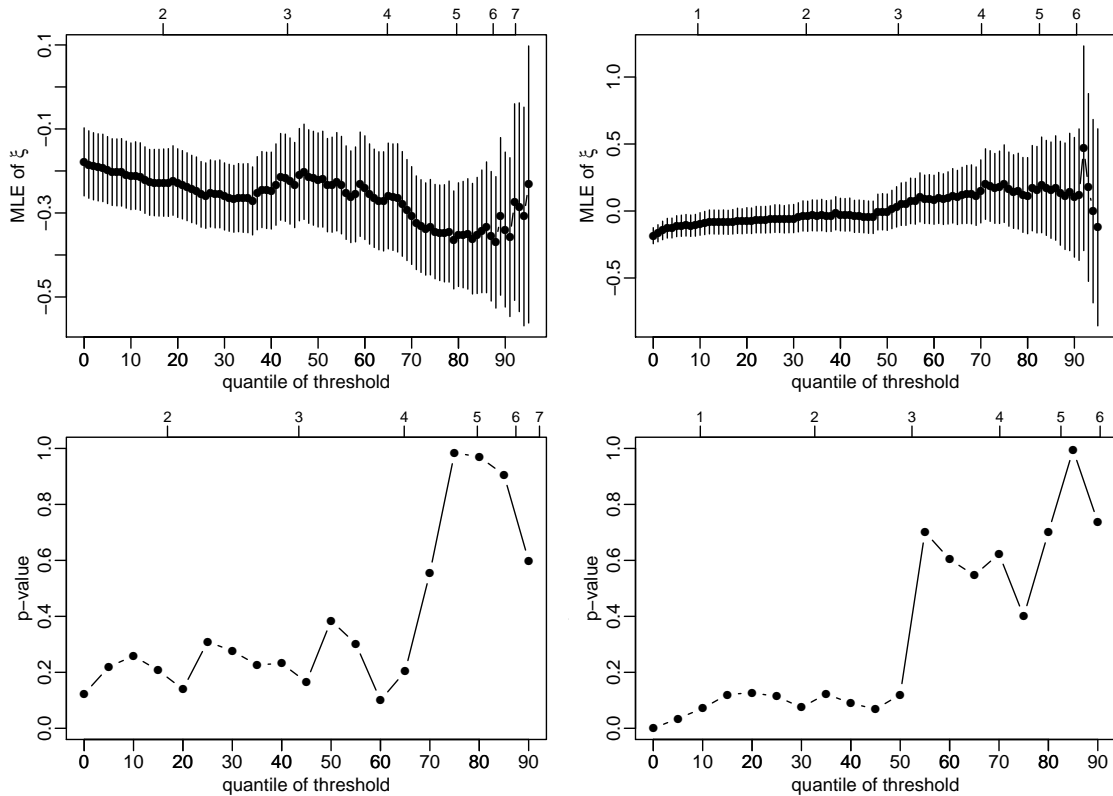


Figure 2: Threshold diagnostic plots for the storm peak significant wave height hindcast datasets. Left: North Sea data. Right: Gulf of Mexico data. Top: parameter stability plots for MLEs of ξ , with 95% pointwise profile likelihood-based confidence intervals. Bottom: p -values associated with a test of constant shape parameter against the lowest threshold considered. The upper axis scales give the level of the threshold in metres.

measure to combine inferences over many thresholds. Sections 2.1 and 2.2 describe the cross-validation procedure and its role in selecting a single threshold. In section 2.3 we discuss two related formulations of the objective of an extreme value analysis and, in section 2.4, we use one of these formulations in a simulation study to inform the choice of prior distribution for GP parameters. Another simulation study, in section 3.1, compares the strategies of choosing a single ‘best’ threshold and averaging of many thresholds. In sections 2.5 and 3.2 we use our methodology to make inferences about extreme significant wave heights in the North Sea and in the Gulf of Mexico. In section 3.3 we illustrate how one could incorporate prior information about the GP shape parameter ξ to avoid physically unrealistic inferences.

2 Single threshold selection

We use a Bayesian implementation of leave-one-out cross-validation to compare the predictive ability of BGP inferences based on different thresholds. We take a predictive approach, averaging inferences over the posterior distribution of parameters, to reflect differing parameter uncertainties across thresholds: uncertainty in GP parameters will tend to increase as the threshold is raised. A point estimate of GP model parameters can give a zero likelihood for a validation observation: this occurs if $\hat{\xi} < 0$ and this observation is greater than the estimated upper endpoint $u - \hat{\sigma}_u / \hat{\xi}$. In this event an estimative approach would effectively rule out the threshold u . Accounting for parameter uncertainty alleviates this problem by giving weight to parameter values other than a particular point estimate.

A naive implementation of leave-one-out cross-validation is computationally intensive. To avoid excessive computation we use importance sampling to estimate cross-validation predictive densities based on Bayesian inferences from the entire dataset. One could use a similar

strategy in a frequentist approximation to predictive inference based on large sample theory or bootstrapping (Young and Smith, 2005, chapter 10). However, large sample results may provide poor approximations for high thresholds (small number of excesses) and the GP observed information is known to have poor finite-sample properties (Süveges and Davison, 2010). Bootstrapping, of ML or PWM estimates, increases computation time further and is subject to the regularity conditions mentioned in the introduction.

2.1 Assessing threshold performance using cross-validation

Suppose that $\mathbf{x} = (x_1, \dots, x_n)$ is a random sample of raw (unthresholded) data from H . Without loss of generality we assume that $x_1 < \dots < x_n$. Consider a *training threshold* u . A BGP(p_u, σ_u, ξ) model is used at threshold u , where $p_u = P(X > u)$ and (σ_u, ξ) are the parameters of the GP model for excesses of u . Let $\theta = (p_u, \sigma_u, \xi)$ and $\pi(\theta)$ be a prior density for θ . Let \mathbf{x}^s denote a subset of \mathbf{x} , possibly equal to \mathbf{x} . The posterior density $\pi_u(\theta | \mathbf{x}^s) \propto L(\theta; \mathbf{x}^s, u)\pi(\theta)$, where

$$L(\theta; \mathbf{x}^s, u) = \prod_{i: x_i \in \mathbf{x}^s} f_u(x_i | \theta),$$

$$f_u(x_i | \theta) = (1 - p_u)^{I(x_i \leq u)} \{p_u g(x_i - u; \sigma_u, \xi)\}^{I(x_i > u)},$$

$I(x) = 1$ if x is true and $I(x) = 0$ otherwise, and

$$g(x; \sigma_u, \xi) = \sigma_u^{-1} \left(1 + \frac{\xi x}{\sigma_u}\right)_+^{-(1+1/\xi)},$$

is the density of a GP(σ_u, ξ) distribution.

We quantify the ability of BGP inferences based on threshold u to predict (out-of-sample) at extreme levels. For this purpose we introduce a *validation threshold* $v \geq u$. If $1 + \xi(v - u)/\sigma_u > 0$ then a BGP(p_u, σ_u, ξ) model at threshold u implies a BGP(p_v, σ_v, ξ) model at threshold v , where $\sigma_v = \sigma_u + \xi(v - u)$ and $p_v = P(X > v) = (1 + \xi(v - u)/\sigma_u)^{-1/\xi} p_u$. Otherwise, $p_v = 0$ and excesses of v are impossible. For a particular value of v we wish to compare the predictive ability of the implied BGP(p_v, σ_v, ξ) model across a range of values of u .

We employ a leave-one-out cross-validation scheme in which $\mathbf{x}_{(r)} = \{x_i, i \neq r\}$ forms the training data and x_r the validation data. The *cross-validation predictive densities* at validation threshold v , based on a training threshold u , are given by

$$f_v(x_r | \mathbf{x}_{(r)}, u) = \int f_v(x_r | \theta, \mathbf{x}_{(r)}) \pi_u(\theta | \mathbf{x}_{(r)}) d\theta, \quad r = 1, \dots, n. \quad (2)$$

Suppose that the $\{x_i\}$ are conditionally independent given θ . If $p_v > 0$ then

$$f_v(x_r | \theta, \mathbf{x}_{(r)}) = f_v(x_r | \theta) = (1 - p_v)^{I(x_r \leq v)} \{p_v g(x_r - v; \sigma_v, \xi)\}^{I(x_r > v)}. \quad (3)$$

If $p_v = 0$ then $f_v(x_r | \theta, \mathbf{x}_{(r)}) = I(x_r \leq v)$. Suppose that we have a sample $\theta_j^{(r)}, j = 1, \dots, m$ from the posterior $\pi_u(\theta | \mathbf{x}_{(r)})$. Then a Monte Carlo estimator of $f_v(x_r | \mathbf{x}_{(r)}, u)$ based on (2) is given by

$$\widehat{f}_v(x_r | \mathbf{x}_{(r)}, u) = \frac{1}{m} \sum_{j=1}^m f_v(x_r | \theta_j^{(r)}, \mathbf{x}_{(r)}). \quad (4)$$

Evaluation of estimator (4), for $r = 1, \dots, n$, is computationally intensive because it involves generating samples from n different posterior distributions. To reduce computation time we use an importance sampling estimator (Gelfand, 1996; Gelfand and Dey, 1994) that enables

estimation of $\widehat{f}_v(x_r | \mathbf{x}_{(r)}, u)$, for $r = 1, \dots, n-1$, using a single sample only. We rewrite (2) as

$$f_v(x_r | \mathbf{x}_{(r)}, u) = \int f_v(x_r | \theta, \mathbf{x}_{(r)}) q_r(\theta) h(\theta) d\theta, \quad r = 1, \dots, n, \quad (5)$$

where $q_r(\theta) = \pi_u(\theta | \mathbf{x}_{(r)})/h(\theta)$ and $h(\theta)$ is a density whose support must include that of $\pi_u(\theta | \mathbf{x}_{(r)})$. In the current context a common choice is $\pi_u(\theta | \mathbf{x})$ (Gelfand and Dey, 1994, page 511). However, the support of $\pi_u(\theta | \mathbf{x})$: $\xi > -\sigma_u/(x_n - u)$, does not contain that of $\pi_u(\theta | \mathbf{x}_{(n)})$: $\xi > -\sigma_u/(x_{n-1} - u)$, since $x_n > x_{n-1}$. Therefore we use $h(\theta) = \pi_u(\theta | \mathbf{x})$ only for $r \neq n$.

Suppose that we have a sample $\theta_j, j = 1, \dots, m$ from the posterior $\pi_u(\theta | \mathbf{x})$. For $r = 1, \dots, n-1$ we use the importance sampling ratio estimator

$$\widehat{f}_v(x_r | \mathbf{x}_{(r)}, u) = \frac{\sum_{j=1}^m f_v(x_r | \theta_j) q_r(\theta_j)}{\sum_{j=1}^m q_r(\theta_j)}, \quad (6)$$

$$= \frac{\sum_{j=1}^m f_v(x_r | \theta_j)/f_u(x_r | \theta_j)}{\sum_{j=1}^m 1/f_u(x_r | \theta_j)}, \quad (7)$$

where $q_r(\theta) = \pi_u(\theta | \mathbf{x}_{(r)})/\pi_u(\theta | \mathbf{x}) \propto 1/f_u(x_r | \theta)$. If we also have a sample $\theta_j^{(n)}, j = 1, \dots, m$ from the posterior $\pi_u(\theta | \mathbf{x}_{(n)})$ then $\widehat{f}_v(x_n | \mathbf{x}_{(n)}, u) = (1/m) \sum_{j=1}^m f_v(x_n | \theta_j^{(n)})$.

We use

$$\widehat{T}_v(u) = \sum_{r=1}^n \log \widehat{f}_v(x_r | \mathbf{x}_{(r)}, u) \quad (8)$$

as a measure of predictive performance at validation threshold v when using training threshold u .

2.2 Comparing training thresholds

Consider k training thresholds $u_1 < \dots < u_k$, resulting in estimates $\widehat{T}_v(u_1), \dots, \widehat{T}_v(u_k)$. Up to an additive constant, $\widehat{T}_v(u)$ provides an estimate of the negated Kullback-Leibler divergence between the BGP model at validation threshold v and the true density (see, for example, Silverman (1986, page 53)). Thus, $u^* = \arg \max_u \widehat{T}_v(u)$ has the property that, of the thresholds considered, it has the smallest estimated Kullback-Leibler divergence.

Some inputs are required: u_1, \dots, u_k, v and $\pi(\theta)$. Choosing a set u_1, \dots, u_k of plausible thresholds that span the range over which the bias-variance trade-off is occurring is the starting point for many threshold selection methods. An initial graphical diagnostic, such as a parameter stability plot, can assist this choice. In particular, u_k should not be so high that little information is provided about GP parameters. There is no definitive rule for limiting u_k but Jonathan and Ewans (2013) suggest that there should be no fewer than 50 threshold excesses. Applying this rule would restrict u_k to be no higher than the 84% and 92% sample quantiles for the Gulf of Mexico and North Sea datasets respectively, but later we will use u_k that break the rule and examine the consequences.

We need $v \geq u_k$, but the larger v is the fewer excesses of v there are and the smaller the information from data thresholded at v . Consider two validation thresholds: $v_1 = u_k$ and $v_2 > u_k$. If we use v_2 we lose validation information: if $v_1 < x_r \leq v_2$ then in (3) x_r is censored rather than entering into the GP part of the predictive density; and gain nothing: the prediction of $x_r > v_2$ is unaffected by the choice of v_1 or v_2 because $p_{v_1} g(x - v_1; \sigma_{v_1}; \xi) = p_{v_2} g(x - v_2; \sigma_{v_2}; \xi)$. Therefore, we should use $v = u_k$.

In section 2.4 we compare predictive properties of three ‘reference’ priors for GP parameters. Such priors are intended for use when substantial prior information is not available and it is anticipated that information provided by the data will dominate the posterior distribution (O’Hagan, 2006). For high thresholds this may not be the case, with the possible

consequence that a diffuse posterior places non-negligible posterior probability on unrealistically large values of ξ and produces unrealistic extrapolation to long future time horizons. In this event one needs to incorporate more information, perhaps by using a more considered prior distribution, and/or limit the time horizon of interest. We return to this issue in section 3.3.

2.3 Prediction of extreme observations

In an extreme value analysis the main focus is often the estimation of extreme quantiles called *return levels*. Let M_N denote the largest value observed over a time horizon of N years. The N -year return level $z(N)$ is defined as the value exceeded by an annual maximum M_1 with probability $1/N$. In off-shore engineering design criteria are usually expressed in terms of return levels, for values of N such as 100, 1000, 10,000. A related approach defines the quantity of interest as the random variable M_N , rather than particular quantiles of M_1 . Under a BGP(p_u, σ_u, ξ) model, for $z > u$,

$$F(z; \theta) = P(X \leq z) = 1 - p_u \left\{ 1 + \xi \left(\frac{z - u}{\sigma_u} \right) \right\}^{-1/\xi}.$$

Then $z(N) = z(N; \theta)$ satisfies $F(z(N); \theta)^{n_y} = 1 - 1/N$, where n_y is the mean number of observations per year. Similarly, for $z > u$, $P(M_N \leq z) = F(z; \theta)^{n_y N}$. For large N ($N = 100$ is sufficient), $z(N)$ is approximately equal to the 37% quantile of the distribution of M_N (Cox et al., 2002). In an estimative approach, based on a point estimate of θ , the value of $z(N)$ is below the median of M_N . A common interpretation of $z(N)$ is the level exceeded on average once every N years. However, for large N (again $N = 100$ is sufficient) and under an assumption of independence at extreme levels, $z(N)$ is exceeded 0, 1, 2, 3, 4 times with respective approximate probabilities of 37%, 37%, 18%, 6% and 1.5%. It may be more instructive to examine directly the distribution of M_N , rather than very extreme quantiles of the annual maximum M_1 .

The relationship between these two approaches is less clear under a predictive approach, in which posterior uncertainty about θ is incorporated into the calculations. The N -year (*posterior*) *predictive return level* $z_P(N)$ is the solution of

$$P(M_1 \leq z_P(N) \mid \mathbf{x}) = \int F(z_P(N); \theta)^{n_y} \pi(\theta \mid \mathbf{x}) \, d\theta = 1 - 1/N,$$

and the predictive distribution function of M_N is given by

$$P(M_N \leq z \mid \mathbf{x}) = \int F(z; \theta)^{n_y N} \pi(\theta \mid \mathbf{x}) \, d\theta. \quad (9)$$

As noted by Smith (2003, section 1.3), accounting for parameter uncertainty tends to lead to larger estimated probabilities of extreme events, that is, $z_P(N)$ tends to be greater than an estimate $\hat{z}(N)$ based on, for example, the MLE $\hat{\theta}$. The strong non-linearity of $F(z; \theta)^{n_y}$ for large z , and the fact that it is bound above by 1, mean that averages of $F(z; \theta)^{n_y}$ over areas of the parameter space relating to the extreme upper tail of M_1 tend to be smaller than point values near the centre of such areas. This phenomenon is less critical when working with the distribution of M_N because now central quantiles of M_N also have relevance, not just particular extreme tail probabilities. Numerical results in section 2.5 (Figure 7) show that $z_P(N)$ can be rather greater than the median of the predictive distribution of M_N , particularly when posterior uncertainty about θ is large.

For a given value of N , we estimate $P(M_N \leq z \mid \mathbf{x})$ using the sample $\theta_j, j = 1, \dots, m$ from the posterior density $\pi(\theta \mid \mathbf{x})$ to give

$$\hat{P}(M_N \leq z \mid \mathbf{x}) = \frac{1}{m} \sum_{j=1}^m F(z; \theta_j)^{n_y N}. \quad (10)$$

The solution $\widehat{z}_P(N)$ of $\widehat{P}(M_1 \leq \widehat{z}_P(N) \mid \mathbf{x}) = 1 - 1/N$ provides an estimate of $z_P(N)$.

2.4 Simulation study 1: priors for GP parameters

We compare approaches for predicting future extreme observations: a predictive approach using different prior distributions and an estimative approach using the MLE. We use Jeffreys' prior $p_u \sim \text{beta}(1/2, 1/2)$ for p_u , so that $p_u \mid \mathbf{x} \sim \text{beta}(n_u + 1/2, n - n_u + 1/2)$, where n_u is the number of threshold excesses. Initially we consider three prior distributions for GP parameters: a Jeffreys' prior

$$\pi_J(\sigma_u, \xi) \propto \sigma_u^{-1} (1 + \xi)^{-1} (1 + 2\xi)^{-1/2}, \quad \sigma_u > 0, \xi > -1/2; \quad (11)$$

a maximal data information (MDI) prior

$$\pi_M(\sigma_u, \xi) \propto \sigma_u^{-1} e^{-(\xi+1)} \quad \sigma_u > 0, \xi \geq -1, \quad (12)$$

truncated from $\xi \in \mathbb{R}$ to $\xi \geq -1$; and a flat prior

$$\pi_F(\sigma_u, \xi) \propto \sigma_u^{-1}, \quad \sigma_u > 0, \xi \in \mathbb{R}. \quad (13)$$

Motivated by findings presented later in this section we generalize (12) to an MDI(a) prior:

$$\pi_A(\sigma, \xi_u; a) \propto \sigma_u^{-1} a e^{-a(\xi+1)} \quad \sigma_u > 0, \xi \geq -1, a > 0. \quad (14)$$

These priors are improper. Let n_u be the number of threshold excesses. Castellanos and Cabras (2007) show that the Jeffreys' prior yields a proper posterior for $n_u \geq 1$ and Northrop and Attalides (2014) show that under the flat prior a sufficient condition for posterior propriety is $n_u \geq 3$. Northrop and Attalides (2014) also show that for any sample size, if, and only if, ξ is bounded below *a priori*, the MDI prior, and the generalized MDI prior, yield a proper posterior. At the particular bound of -1 used in (12) the GP distribution reduces to a uniform distribution on $(0, \sigma)$ and corresponds to a change in the behaviour of the GP density: for $\xi < -1$, this density increases without limit as it approaches its mode at the upper end point $-\sigma_u/\xi$, behaviour not expected in extreme value analyses. Figure 3 compares the Jeffreys', MDI and generalized MDI prior (for $a = 0.6$) as functions of ξ . The Jeffreys' prior (11) is

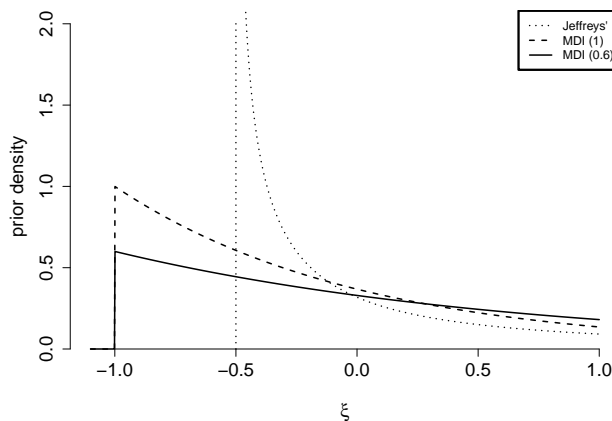


Figure 3: Jeffreys', truncated MDI and generalized MDI priors as functions of ξ .

unbounded as $\xi \downarrow -1/2$. If there are small numbers of threshold excesses this can result in a bimodal posterior distribution, with one mode at $\xi = -1/2$. In this simulation study we also find that the Jeffreys' prior results in poorer predictive performance than the truncated MDI and flat priors.

Let Z_{new} be a future N -year maximum, sampled from a distribution with distribution function $F(z; \theta)^{n_u N}$. If the predictive distribution function (9) is the same as that of Z_{new} then

$P(M_N \leq Z_{\text{new}} \mid \mathbf{x})$ has a $U(0,1)$ distribution. In practice this can only hold approximately: the closeness of the approximation under repeated sampling provides a basis for comparing different prior distributions. Performance of an estimative approach based on the MLE $\hat{\theta}$ can be assessed using $F(Z_{\text{new}}; \hat{\theta})^{n_y N}$. For a given prior distribution and given values of N, n_y and n , the simulation scheme is:

1. simulate a dataset \mathbf{x}_{sim} of n independent observations from a $\text{BGP}(p_u, \sigma_u, \xi)$ model and then a sample $\theta_j, j = 1, \dots, m$ from the posterior $\pi(\theta \mid \mathbf{x}_{\text{sim}})$;
2. simulate an observation z_{new} from the distribution of M_N , that is, $\max(X_1, \dots, X_{N_u})$, where $N_u \sim \text{bin}(n_y N, p_u)$ and $X_i \stackrel{\text{i.i.d.}}{\sim} \text{GP}(\sigma_u, \xi), i = 1, \dots, N_u$;
3. use (10) to evaluate $\hat{P}(M_N \leq z_{\text{new}} \mid \mathbf{x})$.

Steps 1. to 3. are repeated 10,000 times, providing a putative sample of size 10,000 from a $U(0, 1)$ distribution. In the estimative approach step 3 is replaced by evaluation of $F(z_{\text{new}}; \hat{\theta})^{n_y N}$. Here, and throughout this paper, we produce samples of size m from the posterior distribution $\pi(\theta \mid \mathbf{x})$ using the generalized ratio-of-uniforms method of Wakefield et al. (1991), following their suggested strategy of relocating the mode of $\pi(\theta \mid \mathbf{x})$ to the origin and setting a tuning parameter r to $1/2$. In the simulation studies we use $m = 1,000$ and when analyzing real data we use $m = 10,000$.

We assess the closeness of the $U(0,1)$ approximation graphically (Geweke and Amisano, 2010), comparing the proportion of simulated values in each $U(0,1)$ decile to the null value of 0.1. To aid the assessment of departures from this value we superimpose approximate pointwise 95% tolerance intervals based on number of points within each decile having a $\text{bin}(10,000, 0.1)$ distribution, i.e. $0.1 \pm 1.96 (0.1 \times 0.9/10,000)^{1/2} = 0.1 \pm 0.006$. We use $p_u \in \{0.1, 0.5\}$, $\sigma_u = 1$ and values of ξ suggested approximately in section 2.5 by the analyses of the Gulf of Mexico data ($\xi \approx 0.1$) and the North Sea data ($\xi \approx -0.2$).

The plots in Figures 4 ($\xi = 0.1, p_u = 0.5$) and 5 ($\xi = -0.2, p_u = 0.1$) are based on simulated datasets of length $n = 500$ and $n_y = 10$, i.e. 50 years of data with a mean of 10 observations per year, for $N = 100, 1000, 10000$ and 100000 years. Note that the plots on the bottom right have much wider y -axis scales than the other plots. It is evident that the estimative approach based on the MLE produces too few values in deciles 2 to 9 and too many in deciles 1 and 10. When the true BGP distribution of M_N (from which z_{new} is simulated in step 2) is wider than that inferred from data (in step 1 and using (10)) we expect a surplus of values in the first and last deciles. The estimative approach fails to take account of parameter uncertainty, producing distributions that tend to be too concentrated and resulting in underprediction of large values of z_{new} and overprediction of small values of z_{new} .

The predictive approaches perform much better. Although departures from desired performance are relatively small, and vary with N in some cases, some general patterns appear. In Figure 4 the flat prior tends to overpredict large values and small values. The MDI prior tends to result in underprediction of large values. The Jeffreys prior underpredicts large values, to a greater extent than the MDI prior, and also tends to underpredict small values. All these tendencies are slightly more pronounced for $\xi = 0.1, p_u = 0.1$ (not shown).

Figure 5 gives similar findings, although the $N = 100$ case behaves a little differently to the larger values of N . The Jeffreys' prior is replaced by a control plot based on values sampled from a $U(0,1)$ distribution. For $\xi = -0.2$ and with small numbers of threshold excesses the Jeffreys' prior occasionally produces a posterior that is also unbounded as $\xi \downarrow -1/2$, making sampling from the posterior difficult.

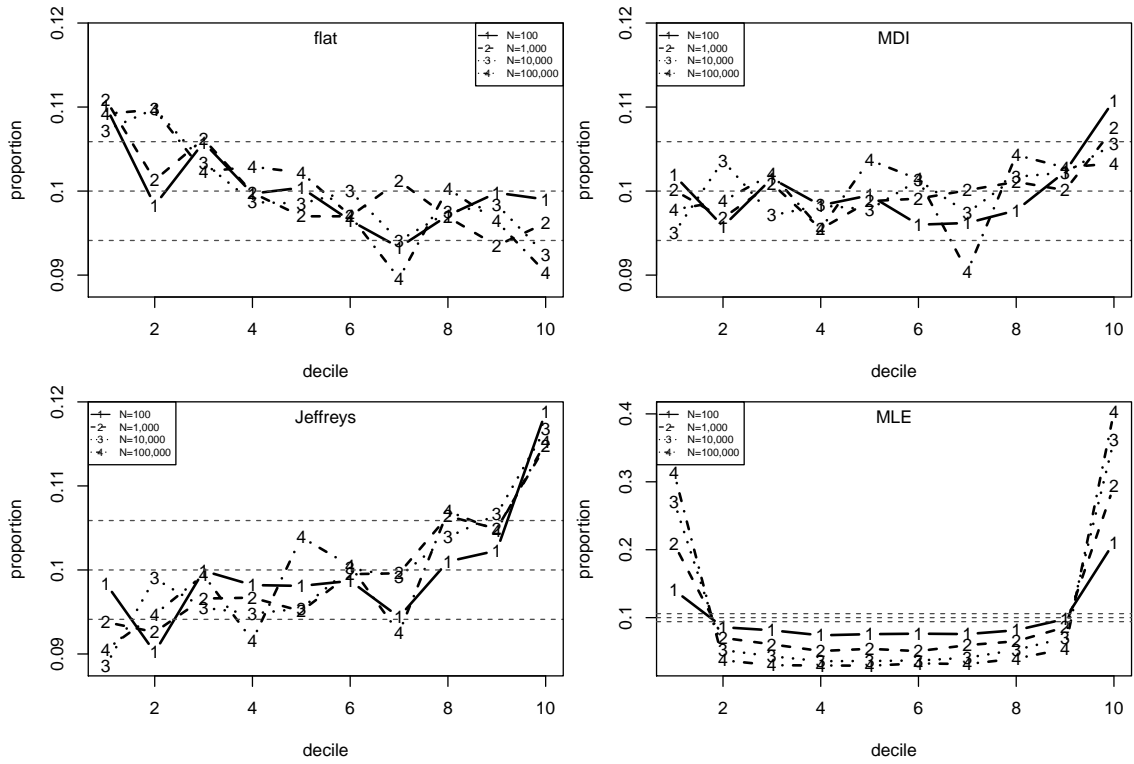


Figure 4: Proportions of simulated values of $\hat{P}(M_N \leq z_{\text{new}} | \mathbf{x})$ falling in U(0,1) deciles for the case $\xi = 0.1$ and $p_u = 0.5$. The prior is labelled on the plots. Separate lines are drawn for $N = 100, 1000, 10000$ and 100000 . 95% tolerance limits are superimposed.

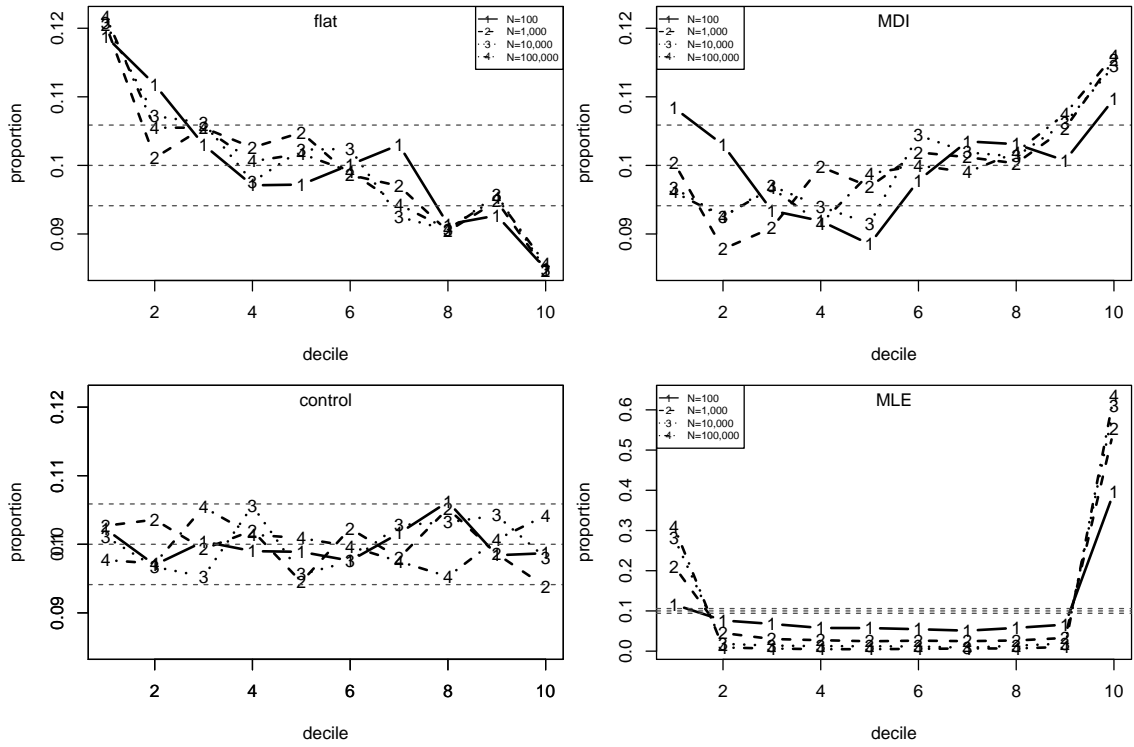


Figure 5: Proportions of simulated values of $\hat{P}(M_N \leq z_{\text{new}} | \mathbf{x})$ falling in U(0,1) deciles for the case $\xi = -0.2$ and $p_u = 0.1$. The Jeffreys' prior is replaced by a control plot based on random U(0,1) samples. Separate lines are drawn for $N = 100, 1000, 10000$ and 100000 . 95% tolerance limits are superimposed.

These results suggest that, in terms of predicting M_N for large N , the MDI prior performs better than the flat prior and the Jeffreys' prior. However, a prior for ξ that is in some sense intermediate between the flat prior and the MDI prior could possess better properties. To explore this we consider the prior (14) for $0 < a \leq 1$. Letting $a \rightarrow 0$ produces flat prior for ξ on the interval $[-1, \infty)$. In order to explore quickly a range of values for a we reuse the posterior samples based on the priors $\pi_F(\sigma, \xi)$ and $\pi_M(\sigma, \xi)$. We use the importance sampling ratio estimator (6) to estimate $P(M_N \leq Z_{\text{new}} | \mathbf{x})$ twice, once using $\pi_F(\theta | \mathbf{x})$ as the importance sampling density $h(\theta)$ and once using $\pi_M(\theta | \mathbf{x})$. We calculate an overall estimate of $P(M_N \leq Z_{\text{new}} | \mathbf{x})$ using a weighted mean of the two estimates, with weights equal to the reciprocal of the estimated variances of the estimators (Davison, 2003, page 603).

Figure 6 shows plots based on the MDI(0.6) prior. This value of a has been selected based on plots for $a \in \{0.1, 0.2, \dots, 0.9\}$. We make no claim that this is optimal, just that it is a reasonable compromise between the flat and MDI priors, providing relatively good predictive properties for the cases we have considered.

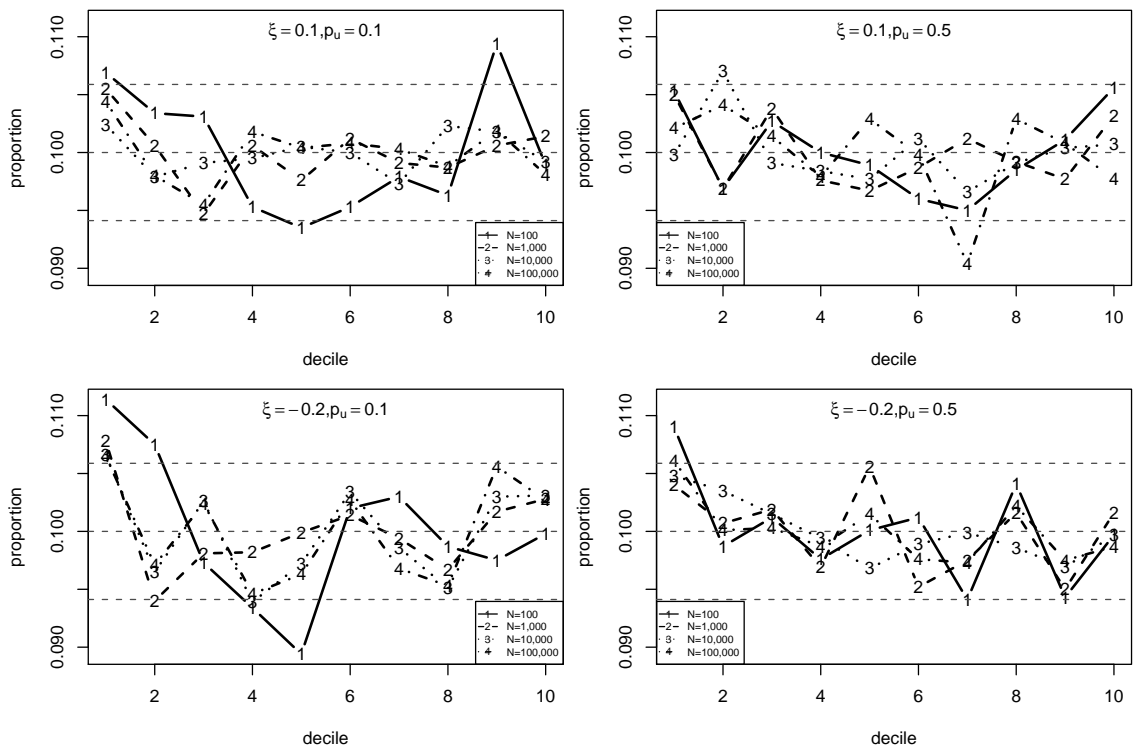


Figure 6: Proportions of simulated values of $\hat{P}(M_N \leq z_{\text{new}} | \mathbf{x})$ falling in $U(0,1)$ deciles for different combinations of ξ and p_u under the MDI(0.6) prior. Separate lines are drawn for $N = 100, 1000, 10000$ and 100000 . 95% tolerance limits are superimposed.

2.5 Significant wave height data: single thresholds

We analyse the North Sea and Gulf of Mexico storm peak significant wave heights using the MDI(0.6) prior suggested by the simulation study in section 2.4. We use the methodology proposed in section 2.1 to quantify the performance of different training thresholds. We use *training thresholds* set at the $(0, 5, \dots, u_k)\%$ sample quantiles, for different u_k . We define the estimated *threshold weight* associated with training threshold u_i , assessed at *validation threshold* $v (= u_k)$, by

$$w_i(v) = \exp\{\hat{T}_v(u_i)\} / \sum_{j=1}^k \exp\{\hat{T}_v(u_j)\}, \quad (15)$$

where $\hat{T}_v(u)$ is defined in (8). The ratio $w_2(v)/w_1(v)$, an estimate of a *pseudo-Bayes factor* (Geisser and Eddy, 1979), is a measure of the relative performance of threshold u_2 compared

to threshold u_1 . In section 3 these weights will be used to combine inferences from different training thresholds.

The top row of Figure 7 shows plots of the estimated training weights against training threshold based for different u_k . For the North Sea data training thresholds in the region of the sample 25-35% quantiles (for which the MLE of $\xi \approx -0.2$) have relatively large threshold weight and there is little sensitivity to u_k . For the Gulf of Mexico data training thresholds in the region of the 60-70% sample quantiles (for which the MLE of $\xi \approx 0.1$) are suggested, and the threshold at which the largest weight is attained is more sensitive to u_k . As expected from the histogram of the Gulf of Mexico data in Figure 1 training thresholds below the 25% sample quantile have low threshold weight. Note that for the Gulf of Mexico data the 90% and 95% thresholds have far fewer excesses (32 and 16) than the suggested 50.

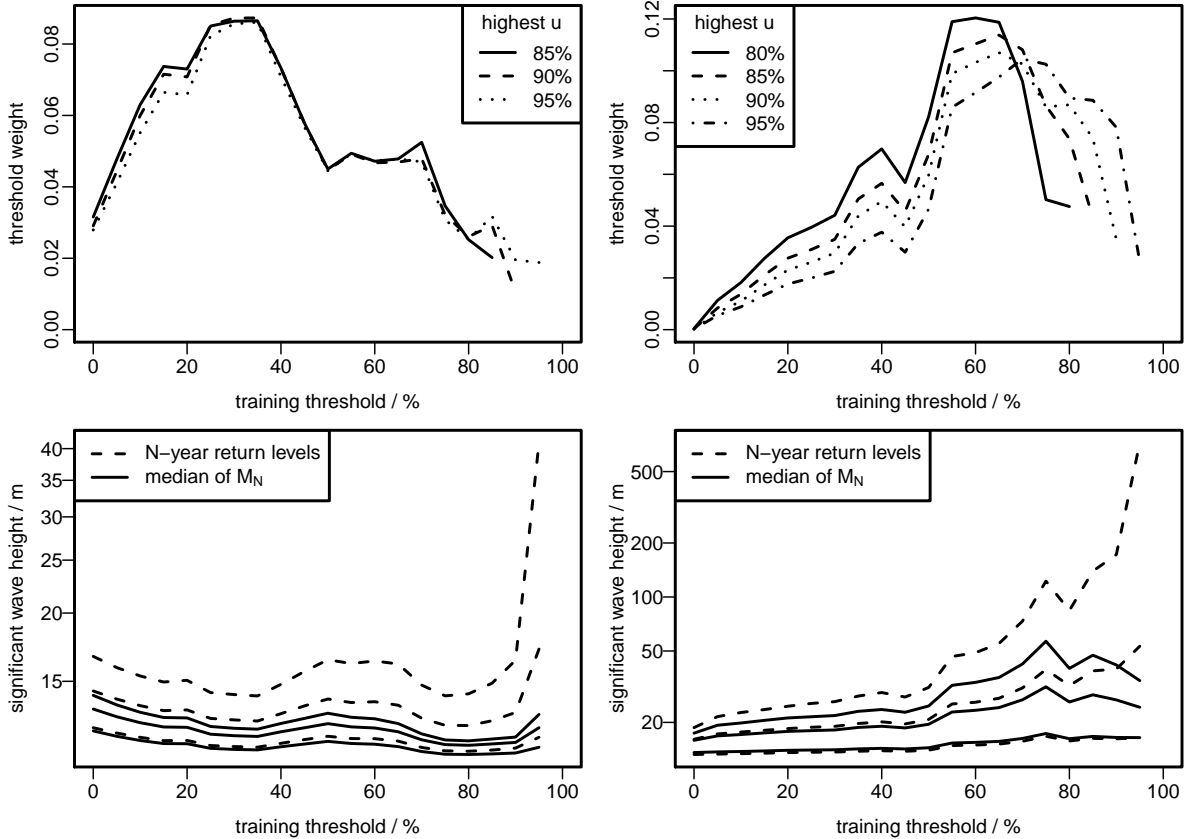


Figure 7: Analyses of significant wave height data by training threshold u . Top row: estimated threshold weights by the highest training threshold considered. Bottom row: N -year predictive return levels and medians of the predictive distribution of M_N for $N = 100, 1000$ and 10000 . Left: North Sea. Right: Gulf of Mexico.

The bottom row of Figure 7 shows that the N -year predictive return levels and the medians of the predictive distribution of N -year maxima M_N are close for $N = 100$, where little or no extrapolation is required, but for $N = 1000$ and $N = 10000$ the former is much greater than the latter. For the North Sea data the results appear sensible and broadly consistent with estimates from elsewhere. From the 55% training threshold upwards, which includes thresholds that have high estimated training weights, estimates of the median of M_{1000} and M_{10000} from the Gulf of Mexico data are implausibly large, e.g. 31.6m and 56.7m for the 75% threshold. The corresponding estimates of the predictive return levels are even less credible. The problem is that high posterior probability of large positive values of ξ , caused by high posterior uncertainty about ξ , translates into large predictive estimates of extreme quantiles. Figure 8 gives examples of the posterior samples of σ_u and ξ underlying the plots in Figure 7. The marginal posterior distributions of ξ are positively skewed, particularly so for the 95% training thresholds, mainly because for fixed σ_u , ξ is bounded below by $\sigma_u/(x_n - u)$. The

higher the threshold the larger the posterior uncertainty and the greater the skewness towards values of ξ that correspond to a heavy-tailed distribution. For the Gulf of Mexico data at the 95% threshold $\hat{P}(\xi > 1/2 \mid \mathbf{x}) \approx 0.20$ and $\hat{P}(\xi > 1 \mid \mathbf{x}) \approx 0.05$. This issue is not peculiar to a Bayesian analysis: frequentist confidence intervals for ξ and for extreme quantiles are also unrealistically wide.

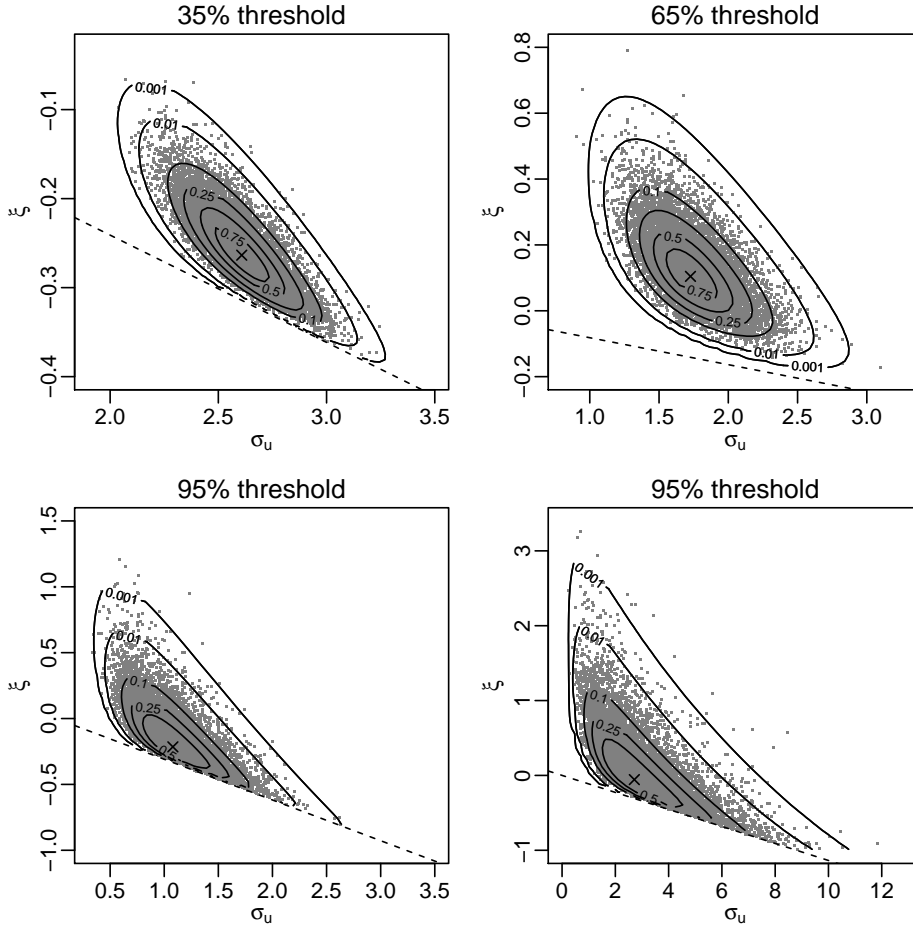


Figure 8: Samples from $\pi(\sigma_u, \xi \mid \mathbf{x})$, with density contours scaled to unity at the posterior mode (\times). Dashed lines show the support of the posterior distribution: $\xi > \sigma_u / (x_n - u)$. Top: ‘best’ training threshold (for u_k set at the 85% sample quantile). Bottom: 95% training threshold. Left: North Sea. Right: Gulf of Mexico.

Physical considerations suggest that there is a finite upper limit to storm peak H_s (Jonathan and Ewans, 2013). However, if there is positive posterior probability on $\xi \geq 0$ then the implied distribution of H_s is unbounded above and on extrapolation to a sufficiently long time horizon, N_l say, unrealistically large values will be implied. This may not be a problem if N_l is greater than the time horizon of practical interest, that is, the information in the data is sufficient to allow extrapolation over this time horizon. If this is not the case then one should incorporate supplementary data (perhaps by pooling data over space as in Northrop and Jonathan (2011)) or prior information. Some practitioners assume that $\xi < 0$ *a priori*, in order to ensure a finite upper limit, but such a strategy may sacrifice performance at time horizons of importance and produce unrealistically small estimates for the magnitudes of rare events. In section 3.3 we consider how one might specify a prior for ξ that allows the data to dominate when it contains sufficient information to do so and avoids unrealistic inferences otherwise.

3 Accounting for uncertainty in threshold

We use Bayesian model-averaging (Hoeting et al., 1999; Gelfand and Dey, 1994) to combine inferences based on different thresholds. Consider a set of k training thresholds u_1, \dots, u_k and a particular validation threshold v . We view the k BGP models associated with these thresholds as competing models. There is evidence that one tends to get better predictive performance by interpolating smoothly between all models entertained as plausible *a priori*, than by choosing a single model (Hoeting et al., 1999, section 7). Suppose that we specify prior probabilities $P(u_i), i = 1, \dots, k$ for these models. In the absence of more specific prior information, and in common with Wadsworth and Tawn (2012), we use a discrete uniform prior $P(u_i) = 1/k, i = 1, \dots, k$. We suppose that the thresholds occur at quantiles that are equally spaced on the probability scale. We prefer this to equal spacing on the data scale because it seems more natural than an equal spacing on the data scale and retains its property of equal spacing under data transformation.

Let $\theta_i = (p_i, \sigma_i, \xi_i)$ be the BGP parameter vector under model u_i , under which the prior is $\pi(\theta_i | u_i)$. By Bayes' theorem, the *posterior threshold weights* are given by

$$P_v(u_i | \mathbf{x}) = \frac{f_v(\mathbf{x} | u_i) P(u_i)}{\sum_{j=1}^k f_v(\mathbf{x} | u_j) P(u_j)},$$

where $f_v(\mathbf{x} | u_i) = \int f_v(\mathbf{x} | \theta_i, u_i) \pi(\theta_i | u_i) d\theta_i$ is the prior predictive density of \mathbf{x} based on validation threshold v under model u_i . However, $f_v(\mathbf{x} | u_i)$ is difficult to estimate and is improper if $\pi(\theta_i | u_i)$ is improper. Following Geisser and Eddy (1979) we use $\prod_{r=1}^n f_v(x_r | \mathbf{x}_{(r)}, u_i) = \exp\{\widehat{T}_v(u_i)\}$ as a surrogate for $f_v(\mathbf{x} | u_i)$ to give

$$\widehat{P}_v(u_i | \mathbf{x}) = \frac{\exp\{\widehat{T}_v(u_i)\} P(u_i)}{\sum_{j=1}^k \exp\{\widehat{T}_v(u_j)\} P(u_j)}. \quad (16)$$

Let $\theta_{ij}, j = 1, \dots, m$ be a sample from $\pi(\theta_i | \mathbf{x})$, the posterior distribution of the GP parameters based on threshold u_i . We calculate a threshold-averaged estimate of the predictive distribution function of M_N using

$$\widehat{P}_v(M_N \leq z | \mathbf{x}) = \sum_{i=1}^k \widehat{P}(M_N \leq z | \mathbf{x}, u_i) \widehat{P}_v(u_i | \mathbf{x}), \quad (17)$$

where, by analogy with (10), $\widehat{P}(M_N \leq z | \mathbf{x}, u_i) = (1/m) \sum_{j=1}^m F(z; \theta_{ij})^{n_y N}$. The solution $\widehat{z}_{PM}(N)$ of

$$\widehat{P}_v(M_1 \leq \widehat{z}_{PM}(N) | \mathbf{x}) = 1 - 1/N \quad (18)$$

provides a threshold-averaged estimate of the N -year predictive return level, based on validation threshold v .

3.1 Simulation study 2: single and multiple thresholds

We compare inferences from a single threshold to those from averaging over many thresholds, based on random samples simulated from three distributions, chosen to represent qualitatively different behaviours. With knowledge of the simulation model we should be able to choose a suitable single threshold, at least approximately. In practice this would not be the case and so it is interesting to see how well the strategies of choosing the 'best' threshold u^* (section 2), and of averaging inferences over different thresholds (section 3), compare to this choice and how the estimated weights $\widehat{P}_v(u_i | \mathbf{x})$ in (16) vary over u_i .

The three distributions are now described. A (unit) exponential distribution has the property that a GP(1,0) model holds above any threshold. Therefore, choosing the lowest available

threshold is optimal. For a (standard) normal distribution the GP model does not hold for any finite threshold, the quality of a GP approximation improving slowly as the threshold increases. In the limit $\xi = 0$, but at finite levels the effective shape parameter is negative (Wadsworth and Tawn, 2012) and one expects a relatively high threshold to be indicated. A uniform-GP hybrid has a constant density up to its 75% quantile and a GP density (here with $\xi = 0.1$) for excesses of the 75% quantile. Thus, a GP distribution holds only above the 75% threshold.

In each case we simulate 1000 samples each of size 500, representing 50 years of data with an average of 10 observations per year. We set training thresholds at the 50%, 55%, \dots , 90% sample quantiles, so that there are 50 excesses of the (90%) validation threshold. For each sample, and for values of N between 100 and 10,000, we solve $\widehat{P}_v(M_N \leq z \mid \mathbf{x}) = 1/2$ for z (see (10)) to give estimates of the median of M_N . We show results for three single thresholds: the threshold one might choose based on knowledge of the simulation model; the ‘best’ threshold u^* (see section 2); and another (clearly sub-optimal) threshold chosen to facilitate further comparisons. We compare these estimates, and a threshold-averaged estimate based on (17) to the true median of M_N , $H^{-1}((1/2)^{10N})$, where H is the distribution function of the underlying simulation model.

The results for the exponential distribution are summarized in Figure 9. As expected, all strategies have negligible bias. The threshold-averaged estimates match closely the behaviour of the optimal strategy (the 50% threshold). The best single threshold results in slightly greater variability, offering less protection than threshold-averaging against estimates that are far from the truth. In the normal case (Figure 10) the expected underestimation is evident for large N : this is substantial for a 50% threshold but small for a 90% threshold. The CV-based strategies have greater bias than those based on a 90% threshold, because inferences from lower thresholds contribute, but have much smaller variability. Similar findings are evident in Figure 11 for the uniform-GP hybrid distribution: contributions from thresholds lower than the 75% quantile produce negative bias but threshold-averaging achieves lower variability than the optimal 75% threshold.

In all these examples the CV-based strategies seem preferable to a poor choice of a single threshold, and, in a simple visual comparison of bias and variability, are not dominated clearly by a (practically unobtainable) optimal threshold. Using threshold-averaging to account for threshold uncertainty is conceptually attractive but, the exponential example aside, compared to the ‘best’ threshold strategy it’s reduction in variability is at the expense of slightly greater bias. A more definitive comparison would depend on problem-dependent losses associated with over- and under-estimation.

Figure 12 summarizes how the posterior threshold weights vary with training threshold. For a few datasets the 90% training threshold receives highest weight. This occurs when inferences about ξ using a 90% threshold differ from those using each lower threshold. This effect is diminished if the number of excesses in the validation set is increased. In the exponential and hybrid cases the average weights behave as expected: decreasing in u in the exponential case, and peaking at approximately the 70% quantile (i.e. lower than the 75% quantile) in the uniform-GP case. In the exponential example the best available threshold (the 50% quantile) receives the highest weight with relatively high probability. In the hybrid example the 70% quantile receives the highest weight most often. The 75% quantile is the lowest threshold at which the GP model for threshold excesses is correct. The 70% quantile performs better than the 75% quantile by trading some model mis-specification bias for increased precision resulting from larger numbers of threshold excesses. In the normal case there is no clear-cut optimal threshold. This is reflected in the relative flatness of the graphs, with the average weights peaking at approximately the 70-80% quantile and the 50% threshold being the ‘best’ slightly more often than higher thresholds. Given the slow convergence in this case it may be that much higher thresholds should be explored, requiring much larger simulated sample sizes, such as those used by Wadsworth and Tawn (2012).

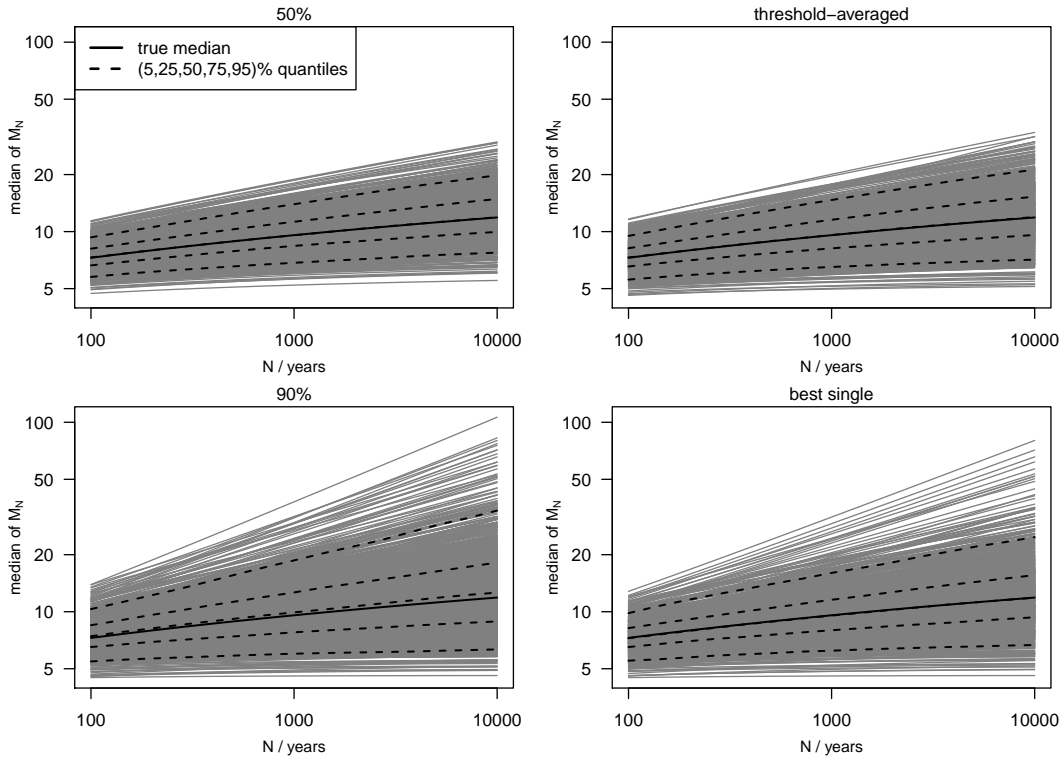


Figure 9: Exponential example. Predictive median of M_N by N : individual datasets (grey), N -specific (5,25,50,75,95)% sample quantiles (dashed) and true median (solid). Threshold strategies: median (top left); 90% quantile (bottom left); threshold-averaged (top right); ‘best’ threshold (bottom right).

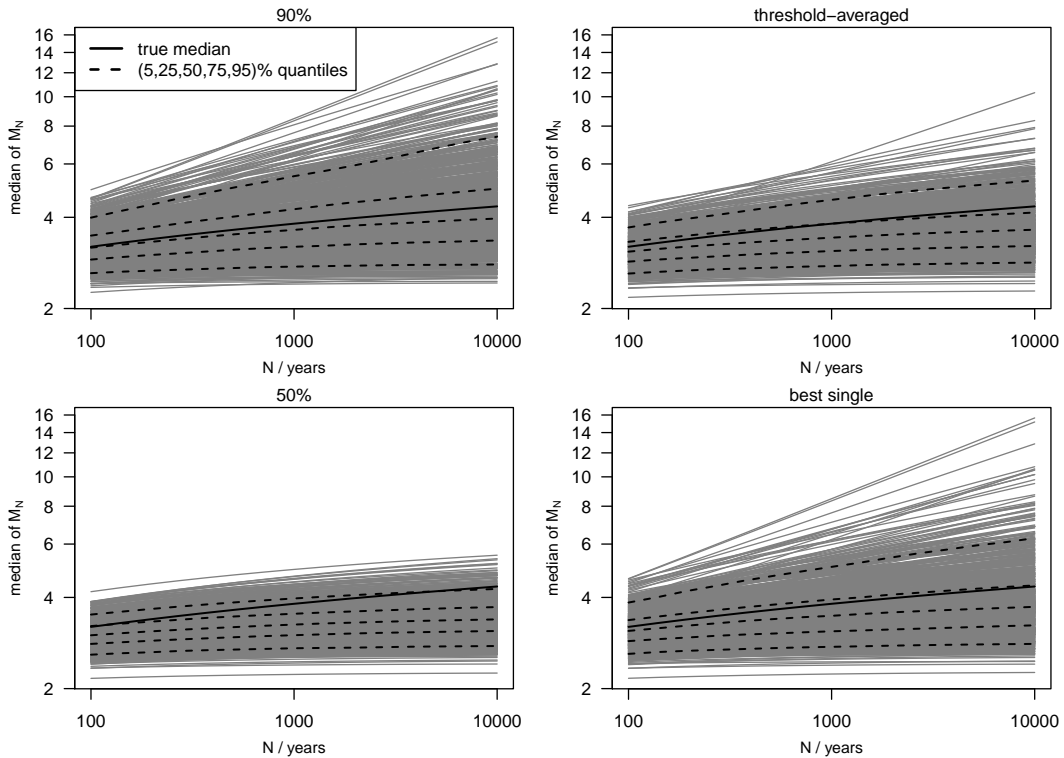


Figure 10: Normal example. Predictive median of M_N by N : individual datasets (grey), N -specific (5,25,50,75,95)% sample quantiles (dashed) and true median (solid). Threshold strategies: 90% quantile (top left); median (bottom left); threshold-averaged (top right); ‘best’ threshold (bottom right).

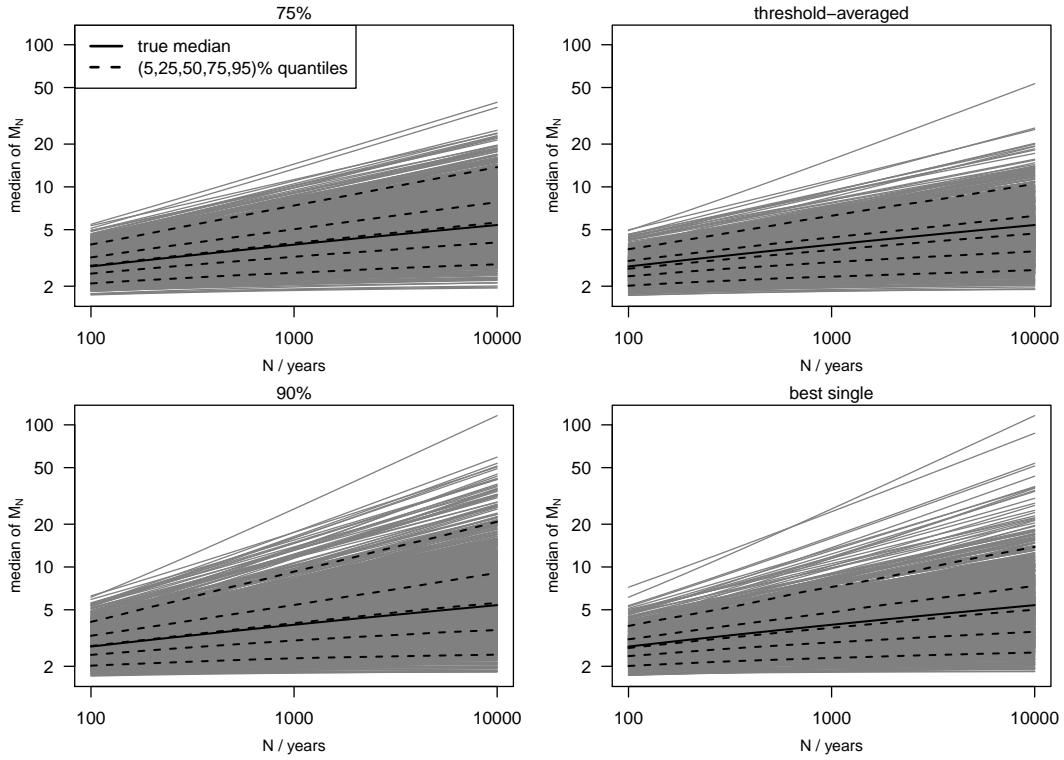


Figure 11: Uniform-GP hybrid example. Predictive median of M_N by N : individual datasets (grey), N -specific (5,25,50,75,95)% sample quantiles (dashed) and true median (solid). Threshold strategies: 75% quantile (top left); 90% quantile (bottom left); threshold-averaged (top right); ‘best’ threshold (bottom right).

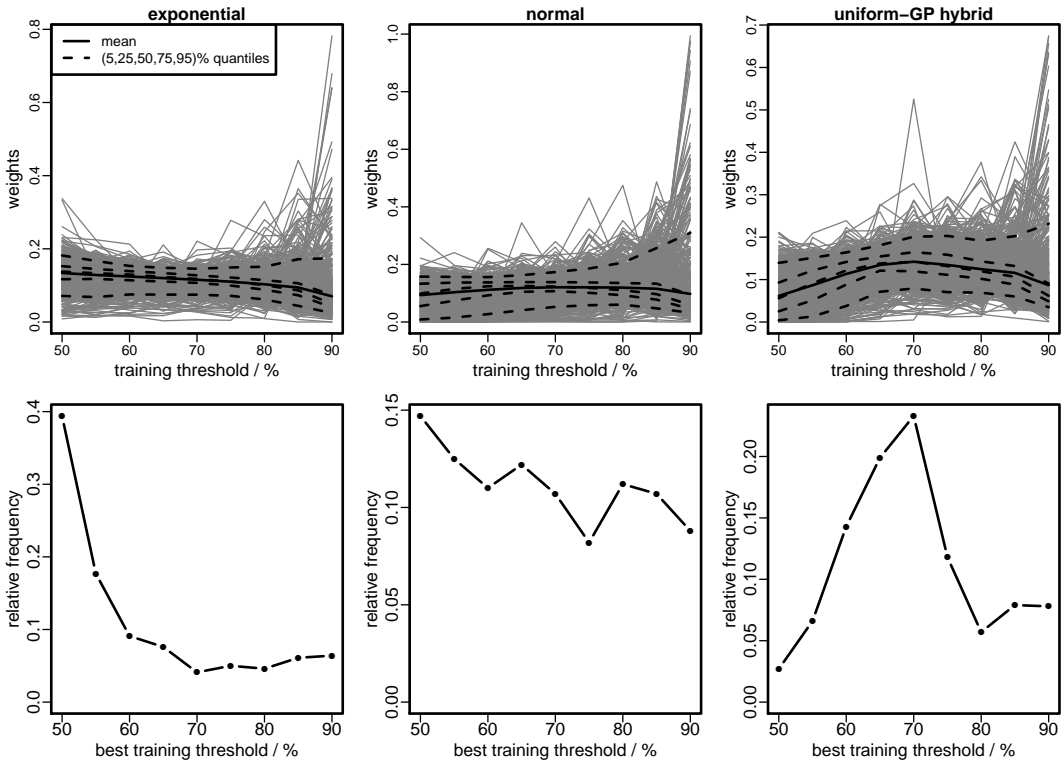


Figure 12: Threshold weights by training threshold. Top: individual datasets (grey) with threshold-specific sample means (solid black) and (5,25,50,75,95)% sample quantiles (dashed). Bottom: relative frequency with which each threshold has the largest weight. Left: exponential. Middle: normal. Right: uniform-GP hybrid.

3.2 Significant wave height data: threshold uncertainty

We return to the significant wave height datasets, using the methodology of section 3 to average extreme value inferences obtained from different thresholds. Figure 13 shows the estimated threshold-specific predictive distribution functions of M_{100} and M_{1000} . Also plotted are estimates from the weighted average (17) over thresholds, for different choices of the highest threshold u_k . For the North Sea data there is so little sensitivity to u_k that the black curves are indistinguishable. For the Gulf of Mexico data there is greater sensitivity to u_k , although based on the discussion in section 2.2 setting u_k at the 95% sample quantile is probably inadvisable with only 315 observations. However, for both choices of u_k , averaging inferences over thresholds has provided some protection against the high probability of unrealistically large values of H_s estimated under some individual thresholds.

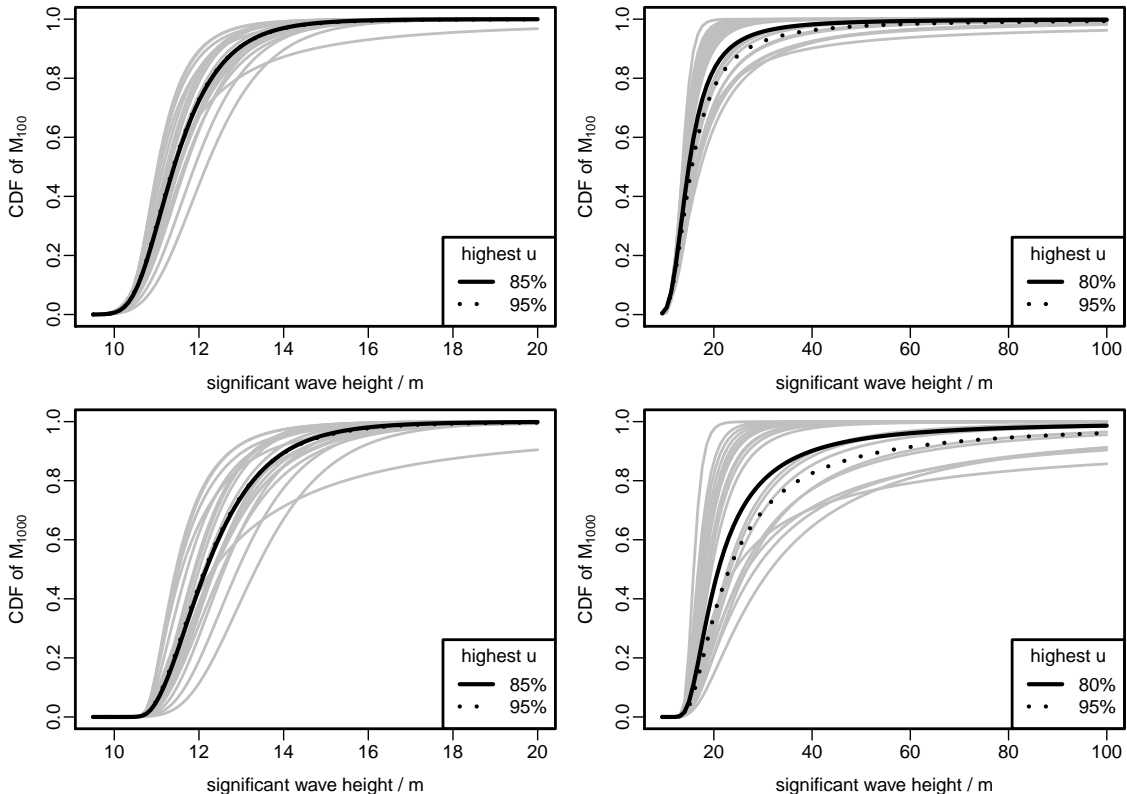


Figure 13: Threshold-specific (grey lines) and threshold-averaged (black lines) predictive distribution functions of M_{100} (top) and M_{1000} (bottom). Left: North Sea data. Right: Gulf of Mexico data.

3.3 A weakly-informative prior for ξ

We have used prior distributions for model parameters that are constructed without reference to the particular problem in hand. This strategy is inadvisable when the data contain insufficient information to dominate such priors, because inferences are then influenced strongly by a generically-chosen prior. When analysing the Gulf of Mexico data in section 2.5 we found that for the highest thresholds there was insufficient information in the data to avoid unrealistic extreme value extrapolations at long time horizons. If small sample sizes cannot be avoided and long time horizons are important then unrealistic inferences can be avoided by providing application-specific prior information. This prior could be elicited from an expert, as in Coles and Tawn (1996), or specified to reflect general experience of the quantity under study, such as the beta-type prior for ξ on $-1/2 \leq \xi \leq 1/2$ used by Martins and Stedinger (2001) for river flows and rainfall totals.

Here, we illustrate the effects on the Gulf of Mexico analysis of providing weak information about ξ , with the aim of preventing unrealistic inferences rather than specifying strong prior information. The particular prior we use is

$$\pi_C(\sigma_u, \xi; A) \propto \sigma_u^{-1} (1 + \xi^2/A^2)^{-1} \quad \sigma_u > 0, \xi \geq -1, A > 0, \quad (19)$$

so that the marginal prior for ξ is a Cauchy distribution, truncated to $\xi \geq -1$. The prior mode is zero, reflecting the expectation that ξ somewhat close to zero. The use of a Cauchy density is motivated by Gelman (2006), who uses a half-Cauchy prior for a group-level standard deviation in a hierarchical model. The scale parameter A is set, guided by expert information from oceanographers (see below), to downweight *a priori* unrealistically large values of ξ . The tails of a Cauchy density have a gentle slope, so information from the data can be influential if it conflicts with this prior.

Let m_N be the median of M_N . Oceanographers with expert knowledge of the hurricane-induced storms in the Gulf of Mexico have provided approximate values of $\widehat{m}_{100} = 15\text{m}$ and 1.5 for the ratio m_{10000}/m_{100} . Estimating m_1 directly from the data gives $\widehat{m}_1 = 4.55\text{m}$. With a view to downweighting substantially *a priori* only very unrealistically large values of ξ we set $\widehat{m}_{10000} = 3\widehat{m}_{100} = 45\text{m}$, that is, we double the ratio provided by the experts. Based on theory concerning the limiting behaviour of the maximum of i.i.d. random variables (Coles, 2001, chapter 3) we suppose that, for $N \geq 1$, M_N has a generalized extreme value GEV(μ_N, σ_N, ξ) distribution. Then $\mu_N = \mu_1 + \sigma_1[(N/\ln 2)^\xi - 1]/\xi$ and the quantity $R = (m_{10000} - m_1)/(m_{100} - m_1)$ is a function only of ξ . If $\xi = 0.229$ then $R = (\widehat{m}_{10000} - \widehat{m}_1)/(\widehat{m}_{100} - \widehat{m}_1)$, suggesting a prior for which $P(\xi > 0.229)$ is small. We use $A = 0.154$, for which $P(\xi > 0.229) = 0.2$ and $P(\xi > 1/2) = 0.1$. Note that under the MDI(0.6) prior $P(\xi > 1/2) = 0.41$.

Figure 14 contains graphical summaries of posterior samples under the Cauchy and MDI(0.6) priors for 65% and 95% training thresholds. For the 65% threshold, comparing the marginal posterior densities for ξ (top left) shows that the change of prior has little effect, because at this threshold there is sufficient information in the data to dominate these priors. Similarly, the plot of the joint posterior (bottom left) is very similar to the corresponding plot in the top right of Figure 8. For the 95% threshold the posterior for ξ under the Cauchy prior is much less diffuse than under the MDI(0.6) prior and far less posterior density is associated with large values of ξ . Compare also the bottom right plots in Figures 8 and 14. Lack of information the data means that the posterior based on the Cauchy prior is only slightly less diffuse than the prior itself.

The plots on the left of Figure 15 show, by comparison with the plots on the right of Figure 7, the effect of the change of prior on the threshold weights and the threshold-specific predictive extreme value inferences. The change from the MDI(0.6) prior to the Cauchy prior has changed little the threshold weights. The main difference is that the highest thresholds benefit more greatly from the increased prior information than the lower thresholds, resulting in a slight increase in their estimated threshold weights. For the highest thresholds the Cauchy prior has prevented the very unrealistic estimates that were obtained under the MDI(0.6) prior. This can also be seen by comparing the plots on the right of Figures 13 and 15: the grey curves corresponding to high thresholds have shifted to the left, that is, towards giving higher density to smaller values of M_N , with a similar knock-on effect on the threshold-averaged black curves.

4 Discussion

We have proposed new methodology for extreme value threshold selection based on a GP model for threshold excesses. It can be used either to inform the choice of a ‘best’ single threshold or to reduce sensitivity to a particular choice of threshold by averaging extremal inferences from several thresholds, weighting thresholds with better cross-validatory predictive

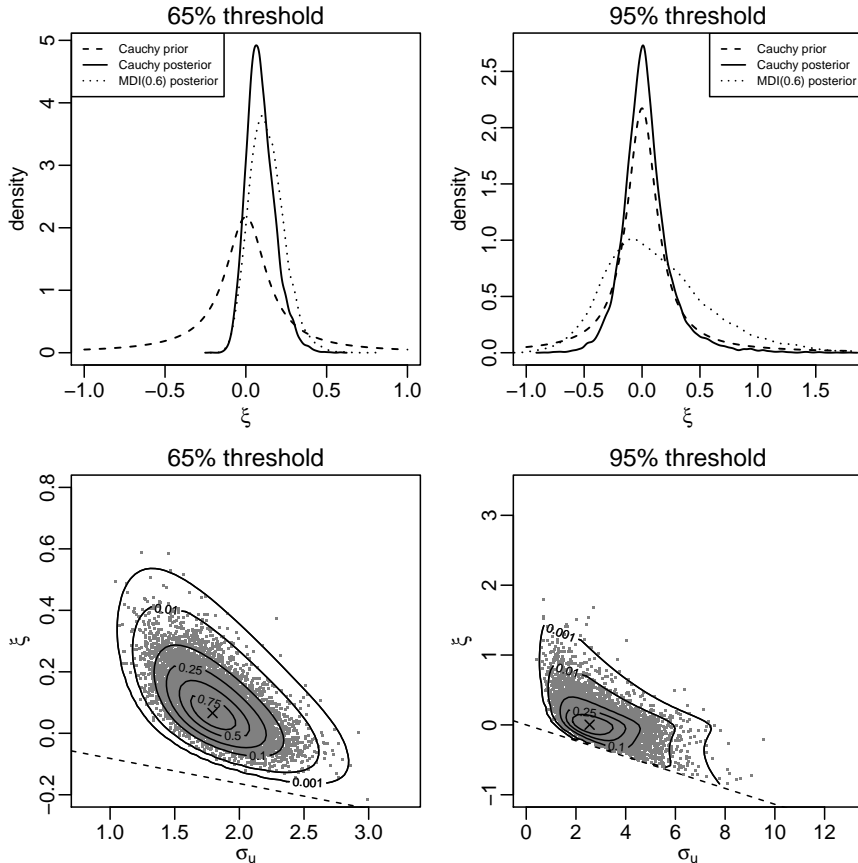


Figure 14: Posterior samples for the Gulf of Mexico data. Top: kernel estimates of the posterior density for ξ under the Cauchy prior (solid) and MDI(0.6) prior (dotted), with the Cauchy prior density (dashed). Bottom: samples from $\pi(\sigma_u, \xi | \mathbf{x})$ under the Cauchy prior, with density contours scaled to unity at the mode (\times). Dashed lines show the support of the posterior distribution. Left: 65% threshold. Right: 95% threshold.

performance more heavily than those with poorer performance. The simulation study in section 3.1 shows that the estimated threshold weights behave as expected in cases where the GP model holds exactly above some threshold and illustrates the potential benefit of averaging different estimated tail behaviours to perform extreme value extrapolation.

The methodology has been applied to significant wave height datasets from the northern North Sea and the Gulf of Mexico. For the latter dataset the highest thresholds result in physically unrealistic extrapolation to long future time horizons. Averaging inferences over different thresholds avoids basing inferences solely on one of these thresholds, but we also explored how the incorporation of basic prior information can be used to address this problem. Stronger prior information about GP model parameters, or indeed prior information about the threshold level itself, could also be used.

In common with many existing methods we need a set of thresholds u_1, \dots, u_k or an interval (u_{min}, u_{max}) for threshold, where u_k (or u_{max}) does not exceed the highest threshold from which it is judged that meaningful inferences can be made. This is also true of other methods, such as those that assess formally stability of parameter estimates and those for which the form of a standard extreme value model is extended to model data below the threshold. In the latter case results can be sensitive to the form of the model specified below the threshold and to u_{min} .

The fact that our methodology is based on inferences from standard unmodified extreme value models makes it relatively amenable to generalization. In significant wave height examples considered in this paper it is standard to extract event maxima from raw data, thereby producing observations that are treated as approximately independent. Otherwise, data may exhibit short-term temporal dependence at extreme levels, leading to clusters of extremes.

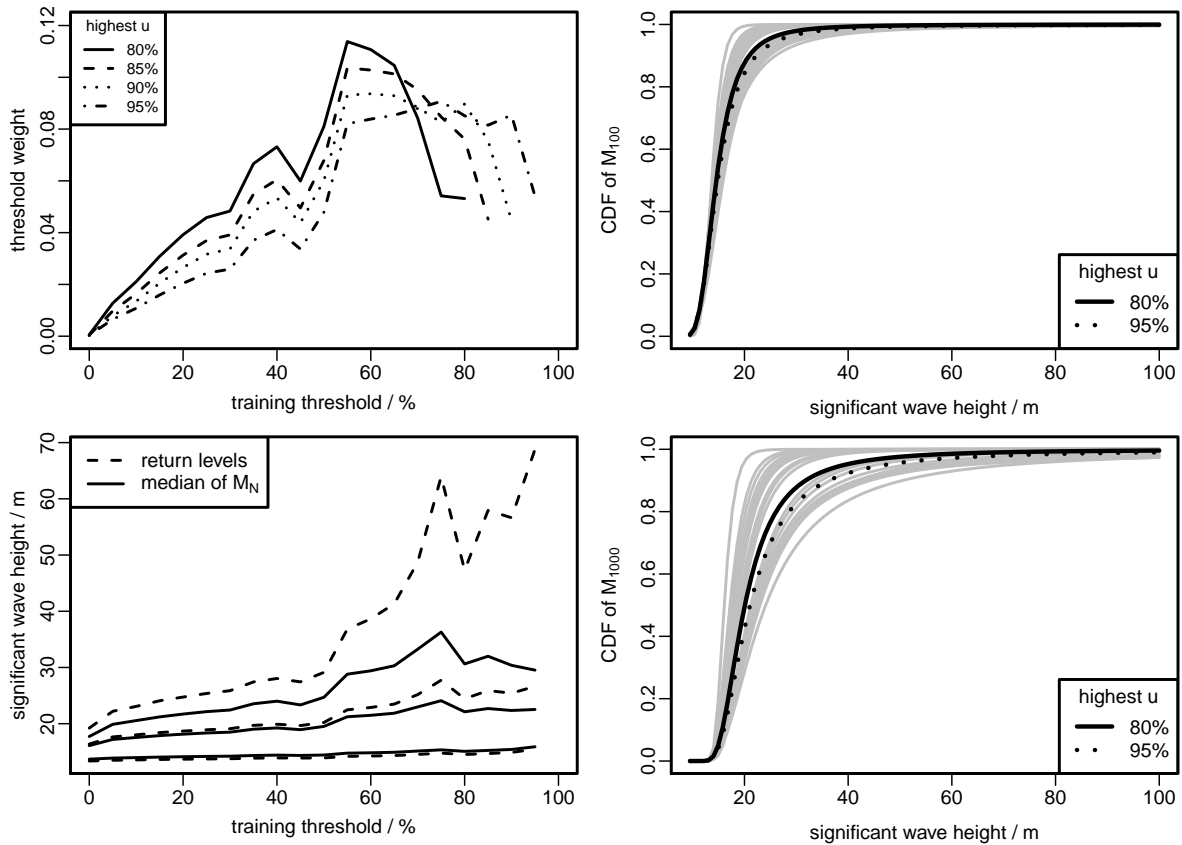


Figure 15: Extreme value inferences for the Gulf of Mexico data using a Cauchy prior for ξ . Top left: estimated threshold weights by the highest training threshold. Bottom left: N -year predictive return levels and medians of the predictive distribution of M_N for $N = 100, 1000$ and 10000 . Right: threshold-specific (grey lines) and threshold-averaged (black lines) predictive distribution functions of M_{100} (top) and M_{1000} (bottom).

In on-going work we are extending our general approach to this situation and to deal with other important issues: the presence of covariate effects; the choice of measurement scale; and inference for multivariate extremes.

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