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Key Points:

- We propose nonasymptotic closed-form distribution for dependent maxima
- We introduce a new efficient generator of Markov chains with arbitrary marginals
- We contribute to develop more reliable data-rich-based analyses of extreme values

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On the Exact Distribution of Correlated Extremes in Hydrology

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Abstract The analysis of hydrological hazards usually relies on asymptotic results of extreme value theory, which commonly deals with block maxima or peaks over threshold (POT) data series. However, data quality and quantity of block maxima and POT hydrological records do not usually fulfill the basic requirements of extreme value theory, thus making its application questionable and results prone to high uncertainty and low reliability. An alternative approach to better exploit the available information of continuous time series and nonextreme records is to build the exact distribution of maxima (i.e., nonasymptotic extreme value distributions) from a sequence of low-threshold POT. Practical closed-form results for this approach do exist only for independent high-threshold POT series with Poisson occurrences. This study introduces new closed-form equations of the exact distribution of maxima taken from low-threshold POT with magnitudes characterized by an arbitrary marginal distribution and first-order Markovian dependence, and negative binomial occurrences. The proposed model encompasses and generalizes the independent-Poisson model and allows for analyses relying on significantly larger samples of low-threshold POT values exhibiting dependence, temporal clustering, and overdispersion. To check the analytical results, we also introduce a new generator (called Gen2Mp) of proper first-order Markov chains with arbitrary marginal distributions. An illustrative application to long-term rainfall and streamflow data series shows that our model for the distribution of extreme maxima under dependence takes a step forward in developing more reliable data-rich-based analyses of extreme values.

1. Introduction

The study of hydrological extremes is one of long history in research applied to design and management of water supply (e.g., Hazen, 1914) and flood protection works (e.g., Fuller, 1914). Almost half a century after the first pioneering empirical studies, Gumbel (1958) provided a general framework linking the theoretical properties of probabilities of extreme values (e.g., Fisher & Tippett, 1928) to the empirical basis of hydrological frequency curves. Since then, extreme value theory (EVT) applied to hydrological analyses has been a matter of primary concern in the literature (see, e.g., Papalexiou & Koutsoyiannis, 2013; Serinaldi & Kilsby, 2014 for detailed overview). EVT aims at modeling the extremal behavior of observed phenomena by asymptotic probability distributions, and observations to which such distributions are allegedly related should meet the following important conditions:

- 1. They should resemble the samples of independent and identically distributed (i.i.d.) random variables. Then, extreme events arise from a stationary distribution and are independent of one another.
- 2. Their number should be large. Defining how large their size should be depends on the characteristics of the parent distribution from which the extreme values are taken (e.g., the tail behavior) and the degree of precision we seek.

Most of these assumptions, commonly made in classical statistical analyses, are hardly ever realized in hydrological applications, especially when studying extremes. Specifically, the traditional analysis of hydrological extremes is based on statistical samples that are formed by selecting from the entire data series (e.g., at the daily scale) those values that can reasonably be considered as realizations of independent extremes, for example, annual maxima or peaks over a certain high threshold. Thus, many observations are discarded and the reduction of the already small size of common hydrological records significantly affects the reliability of

©2019. American Geophysical Union. All Rights Reserved. the estimates (Koutsoyiannis, 2004a, 2004b; Volpi et al., 2019). In addition, Koutsoyiannis (2004a) showed that the convergence to the asymptotic distributions can be extremely slow and may require a huge number of events. Thus, a typical number of extreme hydrological events does not guarantee convergence in applications.

Furthermore, the long-term behavior of the hydrological cycle and its driving forces provide the context to understand that correlations between hydrological samples not only occur, but they also can persist for a long time (see O'Connell et al., 2016, for a recent review). While Leadbetter (1974, 1983) demonstrated that distributions based on dependent events (with limited long-term persistence at extreme levels) share the same asymptotic properties of distributions based on independent trials, there is evidence that correlation has strong influence on the exact statistical properties of extreme values and it slows down the already slow rate of convergence (e.g., Bogachev & Bunde, 2012; Eichner et al., 2011; Serinaldi & Kilsby, 2016; Volpi et al., 2015). In essence, correlation inflates the variability of the expected values and the width of confidence intervals due to information redundancy, and a typical effect is reflected in the tendency of hydrological extremes to cluster in space and time (e.g., Serinaldi & Kilsby, 2018, and references therein). Moreover, focusing on extreme data values, such as annual maxima, hinders reliable retrieval of the dependence structure characterizing the underlying process because of sampling effects of data selection (Iliopoulou & Koutsoyiannis, 2019; Serinaldi et al., 2018). Then, correlation structures and variability of hydrological processes might easily be underestimated, further compromising the attempt to draw conclusions about trends spanning the period of records (see Serinaldi et al., 2018, for detailed discussion). In other words, the lately growing body of publications examining "nonstationarity" in hydrological extremes (see Salas et al., 2018, and references therein) may likely reflect time dependence of such extremes within a stationary setting, as observed patterns are usually compatible with stationary correlated random processes (Koutsoyiannis & Montanari, 2015; Luke et al., 2017; Serinaldi & Kilsby, 2018).

In classical statistical analyses of hydrological extremes, to form data samples, we commonly use two alternative strategies referred to as "block maxima" (BM) and "peaks over threshold" (POT) methods. The former is to choose the highest of all recorded values at each year (for a given time scale, e.g., daily rainfall) and form a sample with size equal to the number of years of the record. The POT method is to form a sample with all recorded values exceeding a certain threshold irrespective of the year they occurred, allowing to increase the available information by using more than one extreme value per year (Claps & Laio, 2003; Coles, 2001).

The fact that observed hydrological extremes tend to cluster in time increases the arguments toward the use of the POT sampling method, instead of block maxima approaches that tend to hide dependence (Iliopoulou & Koutsoyiannis, 2019). Such clustering reflects dependence (at least) in the neighboring excesses of a threshold, invalidating the basic assumption of independence made in classical POT analyses. Therefore, the standard approach in case studies is to fix a (somewhat subjective) high threshold and then filter the clusters of exceedances so as to obtain a set of observations that can be considered mutually independent. Such a declustering procedure involves using empirical rules to define clusters (e.g., setting a run length that represents a minimum timespan between consecutive clusters, meaning that a cluster ends when the separation between two consecutive threshold exceedances is greater than the fixed run length) and then selecting only the maximum excess within each cluster (Bernardara et al., 2014; Bommier, 2014; Coles, 2001; Ferro & Segers, 2003). Declustering results in significant loss of data that can potentially provide additional information about extreme values.

In this paper, we aim to overcome these problems by investigating the exact distribution of correlated extremes. Hence, we can set considerably lower thresholds with respect to the standard POT analyses and avoid declustering procedures whose effectiveness is called into question if we do not account for the process characteristics. The proposed approach provides new insight into probabilistic methods devised for extreme value analysis taking into account the clustering dynamics of extremes, and it is consistent with the general principle of allowing maximal use of information (Volpi et al., 2019).

In summary, hydrological applications have made wide recourse to asymptotes or limiting extreme value distributions, while exact distributions for real-world finite-size samples are barely used in stochastic hydrology because their evaluation requires the parent distribution to be known. However, the small size of common hydrological records (e.g., a few tens of years) and the impact of correlations on the information content of observed extremes cannot provide sufficient empirical evidence to estimate limiting

extreme value distributions with precision. Therefore, we believe that nonasymptotic analytical models for extremes arising from correlated processes should receive renewed research interest (Iliopoulou & Koutsoyiannis, 2019).

This paper is concerned with a theoretical approach to the exact distribution of high extremes based on the pioneering work by Todorovic and Zelenhasic (1970), who proposed a general stationary stochastic model to describe and predict behavior of the maximum term among a random number of random variables in an interval of time [0,t] assuming independence. As verified in several studies mentioned above, to make a realistic stochastic model of hydrological processes, we are forced to confront the fact that dependence should necessarily be taken into consideration. The dilemma is that dependence structures make for realistic models and also reduce the possibility for explicit probability calculations (i.e., analytical derivations of joint probability distributions are more complicated than under independence). The challenge of this paper is to propose a stochastic model of extremes with dependencies allowing for acceptable realism and also permitting sufficient mathematical tractability. In this context, short-range dependence structures, such as Pólya's and Markov's schemes, nicely make a trade-off between these two demands, when hydrological maxima satisfy Leadbetter's condition of the absence of long-range dependence (Koutsoyiannis, 2004a).

In the remainder of this paper, we first introduce a novel theoretical framework to model the exact distribution of correlated extremes in section 2. In section 3, we present a new generator, called Gen2Mp, of correlated processes with arbitrary marginal distributions and Markovian dependence and use it to validate the theoretical reasoning described in section 2. Then, section 4 deals with case studies in order to test the capability of our model to reproduce the statistical behavior of extremes of long-term rainfall and streamflow time series from the real world. Concluding remarks are reported in section 5.

2. Theoretical Framework

We use herein the POT approach to analyze the extreme maxima and assume the number of peaks (e.g., flood peak discharges or maximum rainfall depths) exceeding a certain threshold ξ and their magnitudes to be random variables. The threshold simplifies the study and helps focus the attention on the distribution tails, as they are important to know in engineering design (Papalexiou et al., 2013). In the following, we use upper case letters for random variables or distribution functions and lower case letters for values, parameters, or constants.

If we consider only those peaks Y_i in [0, t] exceeding ξ , then we can define the strictly positive random variable

$$Z_i = Y_i - \xi > 0 \tag{1}$$

for all i = 1, 2, ..., n, where *n* is the number of exceedances in [0, t]. Clearly, *n* is a nonincreasing function of ξ for a given *t*, but we assume herein that ξ is a fixed constant.

It is recalled from probability theory that given a fixed number *n* of i.i.d. random variables $\{Z_i\}$, the largest order statistic $X = \max\{Z_1, Z_2, ..., Z_n\}$ has a probability distribution $H_n(x)$ fully dependent on the joint distribution function of $\{Z_i\}$ that is

$$H_n(x) = \Pr\{Z_1 \le x, Z_2 \le x, ..., Z_n \le x\} = (F(x))^n$$
(2)

In hydrological applications, it may be assumed that the number *n* of values of $\{Z_i\}$ in [0,t] (e.g., the number of storms or floods per year), whose maximum is the variable of interest *X* (e.g., the maximum rainfall depth or flood discharge), is not constant but it is a realization of a random variable N (= 0,1,2,...). Therefore, we are interested in the maximum term *X* among a random number *N* of a sequence of random variables $\{Z_i\}$ in an interval of time [0, t].

In the following, we attempt to determine the one-dimensional distribution function of *X* that is defined as $H(x) = \Pr\{X \le x\}$. Since the magnitude of exceedances Z_i and their number *N* are supposed to be random variables, Todorovic (1970) derived the distribution of the extreme maximum of such a particular class of stochastic processes as

$$H(x) = \Pr\{N = 0\} + \sum_{k=1}^{\infty} \Pr\left\{\bigcap_{i=1}^{k} \{Z_i \le x\} \cap \{N = k\}\right\}$$
(3)

which represents the probability that all exceedances $Z_i > 0$ in [0, t] are less than or equal to x. If x = 0, then H(0) = Pr{N = 0} is the probability that there are no exceedances in [0, t].

Todorovic and Zelenhasic (1970) proposed the simplest form of the general model in equation (3) for use in hydrological statistics, which is now the benchmark against which we measure frequency analysis of extreme events (e.g., Koutsoyiannis & Papalexiou, 2017). Its basic assumptions are that $\{Z_i\}$ is a sequence of N independent random variables with common parent distribution $F(x) = \Pr\{Z_i \le x\}$ and N is a Poisson-distributed random variable independent of $\{Z_i\}$ with mean λ , that is, $\Pr\{N = k\} = (\lambda^k/k!)\exp(-\lambda)$.

Then, recalling that $\sum_{k=0}^{\infty} y^k / k! = \exp(y)$, equation (3) becomes

$$H(x) = \sum_{k=0}^{\infty} (F(x))^k \frac{\lambda^k}{k!} \exp(-\lambda) = \exp(-\lambda(1 - F(x)))$$
(4)

It can be shown that $H(x) \approx H_n(x)$ with satisfactory approximation (Koutsoyiannis, 2004a).

As stated above, the derivation of equation (4) includes strong assumptions, such as independence, and the purpose of this paper is to modify and test this equation under suitable dependence conditions.

First, we suppose that $\{Z_i\}$ is a sequence of *N* random variables with common parent distribution $F(x) = \Pr \{Z_i \le x\}$ and a particular Markovian dependence that give rise to the two-state Markov-dependent process (2Mp, see next section for further details). Specifically, we let the occurrences of the event $\{Z_i \le x\}$ evolve according to a Markov chain with two states, whose probabilities are

$$\begin{cases}
p_0 = \Pr\{Z_i \le x\} \\
p_1 = \Pr\{Z_i > x\} = 1 - p_0
\end{cases}$$
(5)

and the transition probabilities (see also Lombardo et al., 2017, appendix C) are

$$\begin{cases} \pi_{00} = \Pr\{Z_i \le x | Z_{i-1} \le x\} = p_0 + \rho_1 (1 - p_0) \\ \pi_{01} = \Pr\{Z_i \le x | Z_{i-1} > x\} = p_0 (1 - \rho_1) \\ \pi_{10} = \Pr\{Z_i > x | Z_{i-1} \le x\} = 1 - \pi_{00} \\ \pi_{11} = \Pr\{Z_i > x | Z_{i-1} > x\} = 1 - \pi_{01} \end{cases}$$

$$(6)$$

where ρ_1 is the lag-one autocorrelation coefficient of the Markov chain.

It follows that for the process $\{Z_i\}$, the probability of the state $\{Z_n \le x\}$ at a given time *n* depends solely on the state $\{Z_{n-1} \le x\}$ at the previous time step n-1. Then, for a fixed number of exceedances N = n, the Markov property yields

$$\Pr\{Z_n \leq x | Z_{n-1} \leq x, \dots, Z_1 \leq x\} = \Pr\{Z_n \leq x | Z_{n-1} \leq x\}$$

$$\tag{7}$$

Applying the chain rule of probability theory to the distribution function of the maximum term X, $H_n(x) = \Pr \{Z_1 \le x, Z_2 \le x, ..., Z_n \le x\}$, we obtain

$$H_n(x) = \Pr\{Z_n \le x | Z_{n-1} \le x\} \cdots \Pr\{Z_2 \le x | Z_1 \le x\} \Pr\{Z_1 \le x\}$$

$$\tag{8}$$

From the above it follows that $H_n(x)$ can be determined in terms of the conditional probabilities $Pr\{Z_i \le x | Z_{i-1} \le x\}$ and the parent univariate distribution function $F(x) = Pr\{Z_i \le x\}$. As the random variables $\{Z_i\}$ are identically distributed, they correspond to a stationary stochastic process, and then the function $Pr\{Z_i \le x | Z_{i-1} \le x\}$ is invariant to a shift of the origin. In this case, $H_n(x)$ is determined in terms of the second-order (bivariate) distribution $H_2(x) = Pr\{Z_1 \le x, Z_2 \le x\} = Pr\{Z_2 \le x | Z_1 \le x\}F(x)$ and the first-order (univariate) parent distribution F(x). Indeed, from equation (8) we obtain



$$H_n(x) = F(x) \left(\frac{H_2(x)}{F(x)}\right)^{n-1} = \frac{(F(x))^2}{H_2(x)} \left(\frac{H_2(x)}{F(x)}\right)^n$$
(9)

It can be easily shown that equation (9) reduces to equation (2) in case of independence, that is, $H_2(x) = (F(x))^2$.

Second, we assume that exceedances $\{Z_i\}$ have positively correlated occurrences causing a larger variance than if they were independent, that is, the occurrences are overdispersed with respect to a Poisson distribution, for which the mean is equal to the variance. Therefore, we assume that the random number of occurrences N in a specific interval of time [0, t] follows the negative binomial distribution (e.g., Calenda et al., 1977; Eastoe & Tawn, 2010), which allows adjusting the variance independently of the mean. The negative binomial distribution (known as the limiting form of the Pólya distribution, cf. Feller, 1968, p. 143) is a compound probability distribution that results from assuming that the random variable N is distributed according to a Poisson distribution whose mean λ_j varies randomly following a gamma distribution with shape parameter r > 0 and scale parameter $\alpha > 0$, so that its density is

$$g(\lambda_j) = \frac{\lambda_j^{r-1}}{\Gamma(r)\alpha^r} \exp\left(-\frac{\lambda_j}{\alpha}\right)$$
(10)

Then, the probability distribution function of *N* conditional on $\Lambda = \lambda_i$ is

$$\Pr\{N = k | \Lambda = \lambda_j\} = \frac{\lambda_j^k}{k!} \exp(-\lambda_j)$$
(11)

We can derive the unconditional distribution of *N* by marginalizing over the distribution of Λ , that is, by integrating out the unknown parameter λ_i as

$$\Pr\{N=k\} = \int_0^\infty \Pr\{N=k|\Lambda=\lambda_j\}g(\lambda_j) \,d\lambda_j \tag{12}$$

Substituting equations (10) and (11) into equation (12), we have

$$\Pr\{N=k\} = \frac{1}{k!\Gamma(r)\alpha^r} \int_0^\infty \lambda_j^{r+k-1} \exp\left(-\lambda_j\left(\frac{\alpha+1}{\alpha}\right)\right) d\lambda_j$$
(13)

Recalling that the gamma function is defined as $\Gamma(z) = \int_0^\infty x^{z-1} \exp(-x) dx$, then multiplying and dividing equation (13) by $(\alpha/(\alpha+1))^{r+k}$ and integrating by substitution, we obtain after algebraic manipulations

$$\Pr\{N=k\} = \left(\frac{\alpha}{\alpha+1}\right)^k \frac{\Gamma(r+k)}{k!\Gamma(r)} \left(\frac{1}{\alpha+1}\right)^r$$
(14)

To summarize, we specialize the general model in equation (3) for the following conditions:

- 1. {*Z_i*} is a sequence of *N* correlated random variables with 2Mp dependence and common parent distribution $F(x) = \Pr{Z_i \le x}$.
- 2. *N* is a negative binomial random variable independent of $\{Z_i\}$ with mean $\mu = r\alpha$ and variance $\sigma^2 = r\alpha(\alpha+1)$.

Under the above assumptions, from equation (3) we can derive the conditional distribution function of the maximum X as

$$H(x|\lambda_j) = \Pr\{N = 0|\Lambda = \lambda_j\} + \sum_{k=1}^{\infty} \Pr\{\bigcap_{i=1}^{k} \{Z_i \le x\}\} \Pr\{N = k|\Lambda = \lambda_j\}$$
(15)

where for $\{Z_i\}$ of 2Mp



$$\Pr\left\{\bigcap_{i=1}^{k} \{Z_i \le x\}\right\} = \frac{(F(x))^2}{H_2(x)} \left(\frac{H_2(x)}{F(x)}\right)^k$$
(16)

Substituting equations (11) and (16) in equation (15), we obtain

$$H(x|\lambda_j) = \exp(-\lambda_j) + \frac{(F(x))^2}{H_2(x)} \sum_{k=1}^{\infty} \left(\frac{H_2(x)}{F(x)}\right)^k \frac{\lambda_j^k}{k!} \exp(-\lambda_j)$$
(17)

Then, adding and subtracting the term $((F(x))^2/H_2(x))\exp(-\lambda_i)$ yields

$$H(x|\lambda_j) = \exp(-\lambda_j) - \frac{(F(x))^2}{H_2(x)} \exp(-\lambda_j) + \frac{(F(x))^2}{H_2(x)} \sum_{k=0}^{\infty} \left(\frac{H_2(x)}{F(x)}\right)^k \frac{\lambda_j^k}{k!} \exp(-\lambda_j)$$
(18)

and thus

$$H(x|\lambda_j) = \exp(-\lambda_j) - \frac{(F(x))^2}{H_2(x)} \exp(-\lambda_j) + \frac{(F(x))^2}{H_2(x)} \exp\left(-\lambda_j \left(1 - \frac{H_2(x)}{F(x)}\right)\right)$$
(19)

which is the conditional distribution function of the maximum term *X* among a Poisson-distributed random number *N* with gamma-distributed mean $\Lambda = \lambda_j$ of 2Mp random variables $\{Z_i\}$ in an interval of time [0, t]. It can be shown that equation (4) is easily recovered assuming independence, that is, $H_2(x) = \Pr\{Z_1 \le x, Z_2 \le x\} = (F(x))^2$ and $\Lambda = \lambda$ is a fixed constant.

The unconditional distribution of X is derived by substituting equations (14) and (16) into equation (3) as follows

$$H(x) = \left(\frac{1}{\alpha+1}\right)^r + \frac{(F(x))^2}{H_2(x)} \left(\frac{1}{\alpha+1}\right)^r \sum_{k=1}^{\infty} \left(\frac{H_2(x)}{F(x)}\right)^k \left(\frac{\alpha}{\alpha+1}\right)^k \frac{\Gamma(r+k)}{k!\Gamma(r)}$$
(20)

Then, adding and subtracting the term $((F(x))^2/H_2(x))/(\alpha+1)^r$ and denoting by $(r)_k = \Gamma(r+k)/\Gamma(r)$ the Pochhammer's symbol (Abramowitz & Stegun, 1972, p. 256) yields

$$H(x) = \left(\frac{1}{\alpha+1}\right)^r \left(1 - \frac{(F(x))^2}{H_2(x)} + \frac{(F(x))^2}{H_2(x)} \sum_{k=0}^{\infty} \frac{(r)_k}{k!} \left(\frac{\alpha H_2(x)}{(\alpha+1)F(x)}\right)^k\right)$$
(21)

Since $\alpha H_2(x)/((\alpha+1)F(x)) \in [0,1)$ and r > 0 is a real number, then this series is known as a binomial series (Graham et al., 1994, p. 162), and setting $y = \alpha H_2(x)/((\alpha+1)F(x))$, it converges to $(1-y)^{-r} = \sum_{k=0}^{\infty} \frac{(r)_k}{k!} (y)^k$; thus,

$$H(x) = (\alpha + 1)^{-r} \left(1 - \frac{(F(x))^2}{H_2(x)} + \frac{(F(x))^2}{H_2(x)} \left(1 - \frac{\alpha H_2(x)}{(\alpha + 1)F(x)} \right)^{-r} \right)$$
(22)

which is the unconditional distribution of the extreme maximum *X*. The parameters of the model in equation (22) are α and *r* along with those of the models chosen for both the parent distribution, *F*(*x*), and the bivariate distribution, *H*₂(*x*) (see section 4 for further details).

In the case of independence, where $H_2(x) = (F(x))^2$, equation (22) reduces to

$$H(x) = (1 + \alpha (1 - F(x)))^{-r}$$
(23)

As shown in later examples and case studies, equation (22) yields probabilities of nonexceedance that are systematically larger than those under independence, that is, $H_{dep}(x) > H_{indep}(x)$.

3. Gen2Mp: An Algorithm to Simulate the Two-State Markov-Dependent Process (2Mp) with Arbitrary Marginal Distribution

To check the performance of our stochastic model for correlated extremes, we need to simulate a random process $\{Z_i\}$ with any marginal distribution and Markovian dependence. Nevertheless, we must better clarify what the "Markovian dependence" refers to here. As stated in the previous section, we assume that a Markov chain with two states (which may represent, for example, flood or no flood and dry or wet year) governs the excursions above/below any level (threshold) x of the process $\{Z_i\}$ (see, e.g., Fernández & Salas, 1999). We refer to this process as 2Mp (Volpi et al., 2015). For such a process, the Markov property is valid because the probability of the state $\{Z_n \le x\}$ at a given time n depends solely on the state $\{Z_{n-1} \le x\}$ at the previous time step n - 1, that is, $\Pr\{Z_n \le x | Z_{n-1} \le x, ..., Z_1 \le x\} = \Pr\{Z_n \le x | Z_{n-1} \le x\}$.

One can be tempted to use the classical AR(1) (first-order autoregressive) model to simulate the 2Mp. However, this is not appropriate in general, as we show in the following by a numerical experiment that provides insights into an effective simulation strategy. Let us define the random variable S_j in such a way that for j = 1, 2,..., it is

$$\Pr\{S_j = j\} = \Pr\{Z_j \le x, Z_{j-1} \le x, \dots, Z_1 \le x\}$$

$$(24)$$

Then, by definition of conditional probability, we may write, for example, for j = 3

$$\Pr\{Z_3 \le x | Z_2 \le x, Z_1 \le x\} = \frac{\Pr\{Z_3 \le x, Z_2 \le x, Z_1 \le x\}}{\Pr\{Z_2 \le x, Z_1 \le x\}} = \frac{\Pr\{S_3 = 3\}}{\Pr\{S_2 = 2\}}$$
(25)

In our case the Markov property yields

$$\Pr\{Z_3 \le x | Z_2 \le x, Z_1 \le x\} = \Pr\{Z_3 \le x | Z_2 \le x\} = \frac{\Pr\{S_2 = 2\}}{\Pr\{S_1 = 1\}}$$
(26)

where $\Pr\{S_2 = 2\} = \Pr\{Z_2 \le x, Z_1 \le x\} = \Pr\{Z_3 \le x, Z_2 \le x\}$ because $\{Z_i\}$ is stationary. From equations (25) and (26), it is easily understood that we seek a modelling framework for which the ratio $\operatorname{rt}_j(x) = \Pr\{S_{j+1} = j+1\}/\Pr\{S_j = j\}$ should be constant for every *j*, depending solely on the value of the threshold *x*. In order to show that this is generally not valid for AR(1) processes, we compute such a ratio from a sequence of 100,000 random numbers generated by a standard Gaussian AR(1) model with lag-one correlation equal to 0.85. In particular, we calculate four ratios (*j* = 1,..., 4) for various threshold values x_k (k = 1,..., 100) selected randomly over the entire range of the standard Gaussian distribution. Then, as the ratio values depend on the threshold, for each x_k we "standardize" the results by taking the absolute difference between each ratio $\operatorname{rt}_j(x_k)$ and its mean $\mu_{\mathrm{rt}}(x_k)$ computed over j = 1,..., 4, that is, $\mu_{\mathrm{rt}}(x_k) = (1/4) \sum_{i=1}^{i} \operatorname{rt}_j(x_k)$, then dividing all by $\mu_{\mathrm{rt}}(x_k)$; hence, we obtain the relative difference $e_j(x_k) = |(\operatorname{rt}_j(x_k) - \mu_{\mathrm{rt}}(x_k))|/\mu_{\mathrm{rt}}^{i}(\overline{x}_k)|$.

We seek a model with a particular Markovian dependence so that $e_j(x) = 0$ for all *j* and *x*. In Figure 1, we show the boxplots depicting the variability of (percent) $e_j(x_k)$ over all threshold values x_k with j = 1,..., 4. In the left panel, we display the results for the AR(1) model described above. In contrast it can be noted that $e_j(x_k)$ values are not only significantly different from zero (especially if compared with results shown in the right panel of Figure 1, based on simulation algorithm described below), but their variability also changes strongly with the index *j*. Then, we conclude that AR(1) models are not appropriate for our purposes. As shown later, despite sharing similar dependence structures (see Figure 2), Gen2Mp outperforms AR(1) in terms of $e_j(x) = 0$.

3.1. Description of the Gen2Mp Simulation Algorithm

We introduce herein a new generator, which enables the Monte Carlo materialization of a 2Mp with any arbitrary marginal distribution. It is worth stressing that the theoretical considerations discussed above result in a conceptually simple simulation algorithm, whose scheme consists of an iteration procedure with the following steps:

1. We start by generating two sequences $\{a_i\}_{i=1}^n$ and $\{b_i\}_{i=1}^n$ of *n* independent random numbers with the same arbitrary distribution but conditional on being higher $(\{a_i\}_{i=1}^n)$ or lower $(\{b_i\}_{i=1}^n)$ than the median.



Figure 1. Box plots of four (j = 1,..., 4) relative differences $e_j(x_k) = |(rt_j(x_k) - \mu_{rt}(x_k))/\mu_{rt}(x_k)|$ for various threshold values x_k (k = 1,..., 100) selected at random from the parent (standard Gaussian) distribution, where $rt_j(x) = Pr\{S_{j+1} = j+1\}/Pr\{S_j = j\}$ and $\mu_{rt}(x_k) = (1/4) \sum_{j=1}^{\infty} rt_j(x_k)$. The red line inside each box is the median, and the box edges are the 25th and 75th percentiles of the samples. The left panel depicts results for AR(1) model, while right panel shows boxplots of synthetic data from Gen2Mp algorithm.

- 2. Then, we generate the series $\{c_i\}_{i=1}^n$ sampled from i.i.d. Bernoulli random variables taking values 1 and 0 with probability *p* and (1 p), respectively.
- 3. The events $\{c_i = 1\}$ in the Bernoulli series determine the alternation between the two states of our target process, that is, higher (state 1) and lower (state 2) than the median. In other words, the series $\{c_i\}_{i=1}^n$ determines the "holding times" before our process switches (jumps) from a state to the other one, because we assume that the state remains the same up to the "time" when there comes a state change $\{c_i = 1\}$. We can now simulate the state-of-generation sequence $\{d_i\}_{i=1}^n$ taking values 1 when the state of our process is higher than the median (i.e., $\{a_i\}_{i=1}^n$) and 2 if otherwise (i.e., $\{b_i\}_{i=1}^n$).
- 4. Consequently, the sequence $\{d_i\}_{i=1}^n$ is a sample of a Markov chain $\{D_i\}$ with state space $\{1, 2\}$. Since the holding times of each state are completely random, the state probabilities are $\Pr\{D_i = 1\} = \Pr\{D_i = 2\} = 0.5$. On the other hand, as the jumps arrive randomly according to the Bernoulli process, the transition probabilities are $\Pr\{D_i = 1 | D_{i-1} = 2\} = \Pr\{D_i = 2 | D_{i-1} = 1\} = p$ and $\Pr\{D_i = 1 | D_{i-1} = 2\} = \Pr\{D_i = 2 | D_{i-1} = 1\} = p$ and $\Pr\{D_i = 1 | D_{i-1} = 1\} = \Pr\{D_i = 2 | D_{i-1} = 2\} = 1 p$. Therefore, the dependence structure of $\{d_i\}_{i=1}^n$ is completely specified in terms of the lag-one autocorrelation coefficient $\rho_1 = 1 2p$ (see, e.g., Lombardo et al., 2017).
- 5. We can now obtain the target correlated sequence $\{z_i\}_{i=1}^n$ as follows:



Figure 2. Comparison of the empirical autocorrelation functions (EACFs) resulting from time series generated by Gen2Mp $\{z_i\}_{i=1}^n$ and the Markov chain $\{d_i\}_{i=1}^n$ with parameter p = 0.06, and by AR(1) model with lag-one correlation equal to 0.85.

$$z_i = \begin{cases} a_i & \text{if } d_i = 1\\ b_i & \text{otherwise} \end{cases}$$
(27)

6. As the resulting sequence {z_i}ⁿ_{i=1} generally does not satisfy the properties of the process we are interested in, we must subdivide each of the cases "> median" and "< median" into two subcases. Specifically, we generate the i.i.d. sequences {a'_i}ⁿ_{i=1}, {b'_i}ⁿ_{i=1} and {a''_i}ⁿ_{i=1}, {b''_i}ⁿ_{i=1} conditional on being, respectively, "> 75th percentile", "(median, 75th percentile)", "(25th percentile, median)" and "< 25th percentile". Then we generate other two Bernoulli series {c'_i}ⁿ_{i=1} and {c''_i}ⁿ_{i=1} with same parameter as above and consequently derive the corresponding state-of-generation sequences {d'_i}ⁿ_{i=1} (taking values 1 when the state of our process is higher than the 75th percentile and 2 if it belongs to the interval (median, 75th percentile)) and {d''_i}ⁿ_{i=1} (taking values 1 when the state belongs to the interval (25th percentile, median) and 2 if it is lower than the 25th percentile). We can now obtain the target correlated sequence {z_i}ⁿ_{i=1} as follows:

$$z_i = \begin{cases}
 a'_i & \text{if } d_i = 1 \text{ and } d'_i = 1 \\
 b'_i & \text{if } d_i = 1 \text{ and } d'_i = 2 \\
 a''_i & \text{if } d_i = 2 \text{ and } d''_i = 1 \\
 b''_i & \text{if } d_i = 2 \text{ and } d''_i = 2
 \end{cases}
 (27a)$$

7. We continue to subdivide until the relative difference $e_j(x_k)$ converges to zero for any *j*. In any subdivision step, we follow the same procedure as that described above with a fixed parameter *p*, until a convergence threshold is achieved (here a mean absolute error equal to 0.002 for $e_j(x_k)$ is used in the numerical examples below, which is obtained after 9 subdivision steps for p = 0.06).

3.2. Numerical Simulations

We show some Monte Carlo experiments assuming the standard Gaussian probability model as parent distribution, but it can be changed to any distribution function. We generate a correlated series of 100,000 standard Gaussian random numbers using Gen2Mp with parameter p = 0.06. Such a parameter completely determines the dependence structure of the 2Mp process. For 0 the process is positively correlated, while it reduces to white noise for <math>p = 0.5. For 0.5 we get an anticorrelated series. The particular value of <math>p = 0.06 is chosen in order to have the dependence structure of the generated series similar to that of the AR(1) model with lag-one correlation equal to 0.85 (see Figure 2). Such a value of p has been determined numerically exploiting the fact that the dependence structure of the generated series is closely related (showing slight downward bias) to that of the Markov chain $\{D_i\}$ defined above, whose lag-one autocorrelation is $\rho_1 = 1 - 2p$ (see Figure 2). Then, to a first approximation, we start assuming $\rho_1 = 0.85$, and progressively increase it until the dependence structures of the 2Mp and AR(1) match.

Then, even though Gen2Mp and the classical AR(1) algorithms generate time series exhibiting analogous dependence structures, the former significantly outperforms the latter in terms of $e_j(x) = 0$, as shown in Figure 1 (right panel). Furthermore, we generate an independent series of 100,000 standard Gaussian random numbers as a benchmark using classical generators (e.g., Press et al., 2007). As it can be noticed from the probability-probability (PP) and quantile-quantile (QQ) plots in Figure 3, the marginal distribution of the final dependent series (corresponding to a 2Mp) is the same as that of the benchmark series. In summary, the important achievement is that Gen2Mp does not alter the parent distribution, but it only induces time dependence in a Markov chain sense.

Focusing on the frequency analysis of maxima, we investigate the distribution of the maximum term *X* among a random number *N* of a sequence of standard Gaussian random variables $\{Z_i\}$. Specifically, we assume that *N* follows a negative binomial distribution in equation (14), while the variables $\{Z_i\}$ form a 2Mp stochastic process. Based on such hypotheses, in the previous section we derived the corresponding theoretical probability distribution function $H(x) = \Pr\{X \le x\}$ given by equation (22). To check this numerically, we generate the random numbers $\{n_k\}_{k=1}^m$ (where m = 450) from the negative binomial distribution with parameters r = 4 and $\alpha = 25$, then we form the target sample $\{x_k\}_{k=1}^m$ by taking the maximum of *m*



Figure 3. Probability-probability plot (left) and quantile-quantile plot (right) comparing the marginal distribution of a benchmark series (i.i.d. standard Gaussian random numbers) to that of the correlated series generated using Gen2Mp.

nonoverlapping sequences of n_k consecutive random numbers $\{z_i\}_{i=1}^{n_k}$. We allow two different dependence structures for $\{z_i\}_{i=1}^{n_k}$. In the first case we assume that $\{z_i\}_{i=1}^{n_k}$ are sampled from i.i.d. random variables; while in the second case $\{z_i\}_{i=1}^{n_k}$ are sampled from a 2Mp stochastic process with parameter p = 0.06, which is simulated by Gen2Mp.

Results in the form of PP plots are depicted in Figure 4. In the left panel, we show the independent case, and it can be noticed how the empirical distribution of $\{x_k\}_{k=1}^m$ is closely matched by equation (23), that is, the PP plot (blue line) follows a straight line configuration oriented from (0,0) to (1,1). In other words, when $\{Z_i\}$ are i.i.d,. equation (23) proves to be a good model for the theoretical distribution of *X*.

In the right panel of Figure 4, we show the dependent case where the joint probability $H_2(x) = \Pr\{Z_n \le x, Z_{n-1} \le x\}$ in equation (22) is determined numerically. Clearly, if we apply equation (23) to the correlated sample $\{x_k\}_{k=1}^m$, then the corresponding plot (blue line) shows a marked departure from the 45° line (i.e., the line of equality). By contrast, the theoretical distribution that we propose in equation (22) reasonably models the empirical distribution of correlated maxima $\{x_k\}_{k=1}^m$ in all respects (see black line). Therefore,



Figure 4. Probability-probability plots of the maximum term *X* among a (negative binomial) random number *N* of a sequence of i.i.d. (left panel) and 2Mp (right panel) standard Gaussian random variables $\{Z_i\}$.

when the $\{Z_i\}$ belong to 2Mp, equation (22) (black line) largely outperforms equation (23) (blue line) in modelling the extreme maxima.

4. Applications to Rainfall and Streamflow Data

In order to provide some insights into the capability of the proposed methodology to reproduce the statistical pattern of observed hydrological extremes, the data sets used in the applications comprise long-term daily rainfall and streamflow time series with no missing values or as few as possible, to fulfil the requirements of POT analyses. In more detail, we use three daily precipitation time series recorded by rain gages located at Groningen (northeastern Netherlands), Middelburg (southwestern Netherlands), and Bologna (northern Italy) respectively ranging from 1847 to 2017 (171 years, no missing values), from 1855 to 2017 (163 years, no missing values), and from 1813 to 2018 (206 years, only three missing values). Raw data, retrieved through the Royal Netherlands Meteorological Institute Climate Explorer web site, are available at https://climexp. knmi.nl/data/bpeca147.dat (accessed on 24 November 2019) for Groningen station, at https://climexp.knmi. nl/data/bpeca2474.dat (accessed on 24 November 2019) for Middelburg station, and at https://climexp. knmi.nl/data/pgdcnITE00100550.dat (accessed on 24 November 2019) for Bologna station in the period 1813-2007 (see Klein Tank et al., 2002; Menne et al., 2012). For the most recent period, 2008-2018, daily data for Bologna station are provided by the Dext3r public repository (http://www.smr.arpa.emr.it/dext3r/) (accessed 24 November 2019) of the Regional Agency for Environmental Protection and Energy (Arpae) of Emilia Romagna, Italy (retrieved and processed by Koutsoyiannis for the book: Stochastics of Hydroclimatic Extremes, in preparation for 2020).

Furthermore, we analyze one daily streamflow time series of the Po River recorded at Pontelagoscuro, northern Italy (see Montanari, 2012, for further details). The data series, spanning from 1920 to 2017 (98 years, no missing values), is made publicly available by Professor Alberto Montanari at https://distart119.ing.unibo.it/ albertonew/sites/default/files/uploadedfiles/po-pontelagoscuro.txt (accessed on 24 November 2019) for the period 1920–2009, while the remainder (2010–2017) has been retrieved through the Dext3r repository.

Since it has been shown that seasonality affects the distribution of hydrological extremes (Allamano et al., 2011), our analyses are performed on a seasonal basis; we distinguish four seasons, each consisting of three months such that the autumn comprises September, October, and November. Winter, spring, and summer are defined similarly. We prefer not to use deseasonalization procedures to avoid possible artifacts that may affect the results. Furthermore, as daily rainfall and streamflow processes exhibit very different marginal distributional properties, all recorded values exceeding a certain threshold are transformed to normality by normal quantile transformation for the sake of comparison (Krzysztofowicz, 1997). In practice, observed exceedances $\{z_i\}_{i=1}^n$ are transformed to $\psi_i = \Phi^{-1}(F_n(z_i))$, where Φ^{-1} is the quantile function of the standard Gaussian distribution and F_n is the Weibull plotting position of the ordered sample. In addition, all data sets used in this study have been preprocessed by removing leap days, because the 29 February was already removed from all leap years of the 1920–2009 Po river discharge data set.

We now investigate the frequency analysis of observed hydrological maxima. For each season of any data set, we use for example the value of the threshold corresponding to the 5th percentile (excluding zeros for rainfall data sets for simplicity, but we checked that results do not vary considerably if we include zeros), whose exceedances $\{z_i\}$ are normalized to $\{\psi_i\}$ for each sample. As stated in section 1, we are interested in the statistical behavior of the maximum term X among a random number of equally distributed random variables (i.e., belonging to a certain season) in an interval of time (we assume 1 year). Then, first, we form the POT samples for each year of the record, consisting of m (i.e., number of years) sequences of threshold excesses $\{\psi_i\}_{i=1}^{n_k}$ each of size n_k (for k = 1,..., m); second, we form the sample of annual extremes $\{x_k\}_{k=1}^m$ by taking the maximum of each POT series. In other words, $\{x_k\}_{k=1}^m$ is a sample of annual maxima of size m (i.e., the number of years of the given data set) taken from annual POT series of size n_k (i.e., the number of exceedances in the kth year for the considered season). It follows that the sample size used in classical BM analysis is m, while that used in our approach is $\sum_{k=1}^m n_k$. As detailed below, all parameter values (see, e.g., Tables 1 and 2) are estimated from the POT series by maximum likelihood method.

We compare the empirical distribution of *X* to the theoretical probability distribution function H(x) = Pr{ $X \le x$ } given by equation (4) (i.e., the classical method) assuming Poisson occurrences of independent

Tabl	e 1

Parameters Values for All Normalized Case Studies Detailed in the Text						
Station	Parameter/ season	Winter	Spring	Summer	Autumn	
Groningen	λ	50.04	41.56	45.04	50.05	
-	r	76.24	73.15	150.54	164.94	
	α	0.66	0.57	0.30	0.30	
	τ	0.08	0.04	0.02	0.10	
	ρ	0.10	0.05	0.04	0.13	
Middelburg	λ	48.41	40.00	38.42	47.16	
	r	35.71	40.22	35.47	61.68	
	α	1.36	0.99	1.08	0.76	
	τ	0.09	0.04	0.02	0.09	
	ρ	0.12	0.06	0.02	0.14	
Bologna daily	λ	20.92	25.39	16.59	24.67	
	r	7.20	22.57	20.98	21.14	
	α	2.91	1.13	0.79	1.17	
	τ	0.03	0.02	-0.05	-0.01	
	ρ	0.05	0.02	-0.06	0.01	
Bologna hourly	λ	127.59	128.87	54.09	129.74	
	r	5.27	14.14	4.55	12.32	
	α	24.22	9.12	11.90	10.53	
	τ	0.43	0.30	0.17	0.33	
	ρ	0.54	0.38	0.20	0.41	
Pontelagoscuro	λ	85.48	87.40	87.39	86.41	
	r	67.02	136.59	81.95	245.15	
	α	1.28	0.64	1.07	0.35	
	τ	0.82	0.81	0.84	0.84	
	ρ	0.92	0.92	0.94	0.93	

Note. λ for Poisson (P) occurrences (equation (4)); *r* and α for negative binomial occurrences (equation (22)); τ for Clayton (C) and Gumbel (G) copulas (equation (28) and (29)); ρ for Gaussian copula.

exceedances and by equation (22) (i.e., the proposed method) assuming negative binomial occurrences of 2Mp exceedances. Parameters of Poisson and negative binomial distributions are derived through a process of maximum likelihood estimation from the annual counts $\{n_k\}_{k=1}^m$ for each season of each data set. To a first approximation, we assume statistical independence of $\{n_k\}_{k=1}^m$ by checking that, for each data set, the empirical autocorrelations between the numbers of exceedances of subsequent years are negligible (not shown). Furthermore, we assume that the joint probability of exceedances $H_2(x) = \Pr\{Z_1 \le x, Z_2 \le x\}$ in equation (22) can be written in terms of the univariate marginal distribution F(x) (which is the standard normal in case of normal quantile transformation) and a bivariate copula that describes the dependence structure between the variables (Salvadori et al., 2007). Several bivariate families of copulas have been presented in the literature, allowing the selection of different dependence frameworks (Favre et al., 2004). For the sake of simplicity, we choose the following three types of copulas that have been in common use:

 The Gaussian copula (Salvadori et al., 2007, pp. 254-256), which implies the elliptical shape of isolines of the pairwise joint distribution *H*₂(*x*) that in our case is given by a bivariate normal distribution *N*₂

 $(\mathbf{0}, \mathbf{\Sigma})$ with zero mean and covariance matrix $\sum = \begin{pmatrix} 1 & \rho \\ \rho & 1 \end{pmatrix}$, where the parameter ρ is the average (over *m* years) lag-one autocorrelation

coefficient of the annual POT series { ψ_i }.

 The Clayton copula (Salvadori et al., 2007, pp. 237-240), which exhibits upper tail independence and lower tail dependence (Salvadori et al., 2007, pp. 170-175), and in our case yields

$$H_2 x = \max 2F x^{-\beta} - 1^{-\frac{1}{\beta}} 0 \tag{28}$$

where the parameter β can be written in terms of the Kendall's tau correlation coefficient as $\beta = 2\tau/(1 - \tau)$, which is the average (over *m* years) of lag-one Kendall's tau autocorrelation coefficient of the annual POT series { ψ_i }.

3. The Gumbel-Hougaard copula (Salvadori et al., 2007, pp. 236–237), which exhibits upper tail dependence and lower tail independence, and in our case yields

$$H_2(x) = \exp\left(-\left(2(-\ln(F(x)))^{\beta}\right)^{\frac{1}{\beta}}\right)$$
(29)

Parameters Values for All Models Used in the QQ Plots of Figure 10 Parameter/ Q50 Model threshold Q₇₅ Q_5 Q₂₅ Generalized -0.10-0.03-0.05-0.03γ Pareto σ 1220.16 1044.03 1065.80 998.06 ξ 653.00 998.00 1410.00 2133.00 Poisson λ 87.40 68.97 45.89 22.99 Negative Binomial 5.89 1.74 0.71 r 136.59 α 0.64 11.71 26.45 32.22 Clayton and τ 0.82 0.76 0.63 0.48 Gumbel copulas 0.61 Gaussian copula 0.91 0.86 0.75 ρ GEV -0.11-0.11-0.08-0.07Х Ө 1463.94 1463.94 1399.01 1273.31 и 3309.91 3309.91 3369.76 3739.46

where the parameter β is again written in terms of the Kendall's tau correlation coefficient as $\beta = 1/(1-\tau)$.

All parameter values for all seasons and data sets are reported in Table 1.

In Figures 5–7 we may observe that for all daily rainfall data sets the magnitudes of extreme events taken from excesses of a low threshold (the 5th percentile of the nonzero sample) can be considered independent and identically distributed, and this is consistent with the results shown in the literature using different approaches (see, e.g., Marani & Ignaccolo, 2015; Zorzetto et al., 2016; De Michele & Avanzi, 2018). In addition, we may notice that the classical model of POT analyses assuming Poisson occurrences (see equation (4)) seems to be appropriate to study rainfall extremes. Analogous considerations obviously apply to higher thresholds (not shown). Our model of correlated extremes in equation (22) is capable of capturing such a behavior with precision.

Table 2





Figure 5. Probability-probability plots of Groningen data set of daily rainfall. The empirical distributions of maximum terms $\{x_k\}_{k=1}^m$ among annual exceedances of the 5th percentile threshold for winter (top left), spring (top right), summer (bottom left), and autumn (bottom right) seasons are compared to the corresponding theoretical distributions assuming both Poisson (P) occurrences (with parameter λ) of independent exceedances (equation (4)), and negative binomial occurrences (with parameters *r* and α) of correlated exceedances (equation (22)) with pairwise joint distribution described by the Gaussian (N), Clayton (C, equation (28)), and Gumbel (G, equation (29)) copulas, with parameters ρ and τ as detailed in the text. All parameter values are reported in Table 1.

After showing the results with daily rainfall, we also analyze rainfall records at finer time resolution (hourly scale) whose correlation can be stronger than that pertaining to daily data. To this end, we use hourly rainfall data of "Bologna idrografico" station for the period 1990–2013 provided by the Dext3r repository (23 years full coverage, while the entire 2008 is missing). We checked that such hourly rainfall data agrregated at the daily scale are consistent with the daily data recorded in the same period by Bologna station above (not shown).



Figure 6. Same as Figure 5 for Middelburg data set of daily rainfall.



Figure 7. Same as Figure 5 for Bologna data set of daily rainfall.

Comparing Figures 7 and 8, it is noted that extremes of hourly rainfall data are more affected by correlation than daily data (see, e.g., winter and autumn seasons, respectively top left and bottom right panels). This is also the case if we consider the same period of record (1990–2013) for both data sets (not shown). Then, we may conclude that low thresholds can be used for classical POT analyses (assuming independence) of rainfall time series at the daily scale (or above), while further investigations of different data sets are required to describe the impact of dependence on the extremal behavior of the rainfall process at finer time scales. Besides, other interesting future analyses could investigate the extremes of areal rainfall, as for example weather radar data will become more reliable and will accumulate in time providing samples with lenghts adequate enough to enable reliable investigation of the probability distribution of areal rainfall (Lombardo, Napolitano & Russo 2006; Lombardo, Napolitano, Russo, et al., 2006; Lombardo et al., 2009).



Figure 8. Same as Figure 5 for Bologna data set of hourly rainfall.



Figure 9. Same as Figure 5 for the Po River data set of daily discharge.

By contrast, results change significantly when analyzing extremes of streamflow time series. In fact, we present a case study that shows how models assuming independence among magnitudes of extreme events prove to be inadequate to study the probability distribution of discharge maxima.

In Figure 9, we show the PP plots of the distribution of extreme maxima taken from annual exceedances of the 5th percentile thresholds for the four seasons of the Po River discharge data set, recorded at Pontelagoscuro station. Contrary to the rainfall case studies, the classical model assuming independent magnitudes with Poisson (P) occurrences shows marked departures from the 45° line. The theoretical distribution is usually much lower than its empirical counterpart, meaning that, under the popular assumption of independent extremes, the theoretical probability of an extreme event of given magnitude being exceeded is significantly higher than the corresponding observed frequency of exceedance. Figure 9 shows that our 2Mp model of correlated extremes outperforms the widely used independent model. In particular, the distribution of maxima that has a Gumbel copula seems to be more consistent with observed extreme values, denoting dependence in the upper tail of the bivariate distribution H_2 (x) = Pr{ $Z_1 \le x, Z_2 \le x$ } (Schmidt, 2005). In summary, daily streamflow extremes may exhibit noteworthy departures from independence that are consistent with a stochastic process characterized by a 2Mp behavior and upper tail depedence.

The above results are also evident if we compare theoretical and empirical distributions of streamflow maxima by plotting their quantiles against each other. We use real values for this example (i.e., we do not apply the normal quantile transformation to the data series); therefore, empirical quantiles equal the observed annual maxima. Theoretical quantiles referring to equations (4) and (22) (the latter specializes for Gaussian, Clayton, and Gumbel copulas) are computed by numerically solving for the root of the equation H(x) - p = 0 for a given probability value, p (i.e., the Weibull plotting position of observed annual maxima), assuming the classical generalized Pareto (GPD) with zero lower bound as parent distribution of threshold excesses:

$$F(x) = \begin{cases} 1 - \left(1 + \gamma \frac{x}{\sigma}\right)^{-\gamma} & \text{for } \gamma \neq 0\\ 1 - \exp\left(-\frac{x}{\sigma}\right) & \text{otherwise} \end{cases}$$
(30)

where γ is the shape parameter and σ is the scale parameter, which we estimate through the maximum likelihood method applied to the entire POT series of each season.



Figure 10. Quantile-quantile plots of Po river discharge (m^3/s) for spring season. The observed maximum terms among annual peaks over the 5th percentile (top left), 25th percentile (top right), 50th percentile (bottom left), and 75th percentile (bottom right) thresholds are compared to the corresponding theoretical quantiles. In all cases, we assume the GPD as parent distribution of daily streamflow (with shape $\gamma \in \mathbb{R}$, scale $\sigma > 0$ and threshold $\xi > 0$ parameters) and compute quantiles specializing equation (22) for Poisson (P) occurrences (with parameter λ , equation (4)) of independent exceedances, and for negative binomial occurrences (with parameters r and α) of correlated exceedances with pairwise joint distribution described by the Gaussian (N), Clayton (C), and Gumbel (G) copulas, with parameters ρ and τ as detailed in the text. We also plot theoretical quantiles from GEV distribution (with shape $\chi \in \mathbb{R}$, scale $\theta > 0$ and location $\mu > 0$ parameters) fitted to the observed annual maxima. All parameter values are reported in Table 2.

In Figure 10, QQ plots of Po river discharge for the spring season are shown when varying the threshold ξ (from the 5th, Q_5 , to the 75th, Q_{75} , percentiles) to form POT series. It can be noticed that for low thresholds there is a shift in variance between theoretical (i.e., derived from equation (22) with Gumbel copula) and empirical quantiles, namely, the variance of theoretical annual maxima underestimates its empirical counterpart. This can be due to the fitting performance of the marginal GPD, which does not reproduce well the tail behavior of observed data (not shown). Figure 10 shows that increasing the threshold value helps focus the attention on the distribution tail to better capture the behavior of maxima. This is also the case if we compare streamflow quantiles resulting from our model with those estimated through "classical" generalized extreme value (GEV) distribution fitted to the observed annual maxima. All parameter values are reported in Table 2. We note that three GEV parameters are estimated on m = 98 data points, while the 5 parameters of our model in equation 22 (α , r, τ or ρ , and the two parameters of the GPD with zero lower bound) are estimated on $\sum_{k=1}^{m} n_k$ data, which are 8565, 6759, 4497, and 2253 for Q_5 , Q_{25} , Q_{50} , Q_{75} , respectively.

As threshold increases evidence of persistence is progressively reduced as expected, but we also note in Figure 10 that the theoretical quantiles derived from the classical independent Poisson method always show a shift in mean with respect to observed maxima (i.e., under independence, theoretical streamflow quantiles systematically and significantly overestimate observed streamflow maxima).

To summarize, our model provides a closed-form expression of the exact distribution for dependent hydrological maxima, which is capable of capturing the behavior of observed extremes of long-term hydrological records. In particular, while rainfall extremes do not seem to be significantly affected by correlation at the daily scale so that the classical Poisson model can be appropriate for use in POT analyses of daily rainfall time series, the influence of correlation is prominent in the streamflow process at the daily scale and it is important to preserve in simulation and analysis of extremes.

5. Conclusions

The study of hydrological extremes faces the chronic lack of sufficient data to perform reliable analyses. This is partly related to the inherent nature of extreme values, which are rare by definition, and partly related to the relative shortness of systematic records from hydrometeorological gauge networks. The limited

availability of data poses serious problems for an effective and reliable use of asymptotic results provided by EVT.

Alternative methods focusing on the exact distribution of extreme maxima extracted from POT sequences of random size over fixed time windows have been proposed in the past. However, closed-form analytical results were developed only for independent data with Poisson occurrences. Even though these assumptions may be sufficiently reliable for high-threshold POT values, this type of data still generates relatively small sample size. In order to better exploit the available information, it can be convenient to consider lower thresholds. However, the effect of lower thresholds is twofold: on the one side the sample size increases, but on the other side the hypotheses of independent magnitudes and Poisson occurrences of POT values are no longer reliable.

In this study, we have introduced closed-form analytical formulae for the exact distribution of maxima from POT sequences that generalize the classical independent model, overcoming its limits and enabling the study of maxima taken from dependent low-threshold POT values with arbitrary marginal distribution, first-order Markov dependence structure, and negative binomial occurrences, and tested real data against this hypothesis. Even though the framework can be further generalized by introducing arbitrary dependence structures and models for POT occurrences, first-order Markov chains and negative binomial distributions provide a good compromise between flexibility and the possibility to obtain simple ready-to-use formulae. In this respect, it should be noted that our model of correlated extremes can cover a sufficient range of cases. We have shown that the modulation of the lag-one autocorrelation coefficient of the annual sequences of POT values (i.e., the Markov chain parameter) gives a set of extremal distributions that include the empirical distribution of maxima for rainfall data series, and for highly correlated low-threshold discharge POT series. On the other hand, the negative binomial model is a widely used and theoretically well-established model for occurrences exhibiting clustering and overdispersion, which are common characteristics of POT events resulting from persistent processes, such as river discharge.

The relationship between our model and its classical independent version (i.e., equations (22) and (4)) along with results of the case studies show that distribution of extreme maxima under dependence yields probabilities of exceedance that are systematically lower than those under independence and are also consistent with traditional approaches (GEV), based on extreme value theory, applied to long annual maxima series.

Finally, we stress that our model of the exact distribution of correlated extremes requires knowledge or fitting of a bivariate distribution (and therefore its univariate marginal distribution). In particular, while the extremal behavior of the rainfall process does not seem to be significantly affected by dependence at the daily scale so that the classical Poisson model can be appropriate for use in POT analyses of daily rainfall time series, the influence of correlation is prominent in the streamflow process at the daily scale and it appears also in the rainfall process at the hourly scale. Then, it is important to account for such dependence in the extreme value analyses, which are crucial to hydrological design and risk management because critical values can be less extreme and more frequent than expected under the classical independent models. Comparing the Gaussian, Clayton, and Gumbel bivariate copulas, describing different dependence structures, and the standard Gaussian and Generalized Pareto marginal distributions, we found that the distribution of maxima that has a Gumbel copula seems to be more consistent with streamflow extreme values, denoting dependence in the upper tail of the bivariate distribution. However, these aspects require further investigation from both theoretical and empirical standpoints, and will be the subject of future research. In the spirit of the recent literature on the topic, we believe that the present study will contribute to develop more reliable data-rich-based analyses of extreme values.

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