Exact unconditional sampling from max-stable random vectors

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The 'mev' package was originally introduced to implement the exact unconditional sampling algorithms in Dombry et al. (2016). Thetwo algorithms therein allow one to simulate simple max-stable random vectors. The implementation will work efficiently for moderate dimensions and do not currently take approximate schemes for large spatial domains.

1 Functions and use

There are two main functions, rmev and rmevspec. rmev samples from simple max-stable processes, meaning it will return an $n \times d$ matrix of samples, where each of the column has a sample from a unit Frechet distribution. In constrast, rmevspec returns sample on the unit simplex from the spectral (or angular) measure. One could use this to test estimation based on spectral densities, or to construct samples from Pareto processes.

The syntax is

```
library(mev)
#Sample of size 1000 from a 5-dimensional logistic model
x <- rmev(n=1000, d=5, param=0.5, model="log")
#Marginal parameters are all standard Frechet, meaning GEV(1,1,1)
apply(x, 2, function(col){ismev::gev.fit(col, show=FALSE)$mle})

## [,1] [,2] [,3] [,4] [,5]
## [1,] 0.9727 1.0145 1.0005 1.001 0.9946
## [2,] 0.9526 1.0179 0.9799 0.986 0.9753
## [3,] 0.9916 0.9817 0.9605 0.981 0.9497

#Sample from the corresponding spectral density
w <- rmevspec(n=1000, d=5, param=0.5, model="log")
#All rows sum to 1 by construction
head(rowSums(w))

## [1] 1 1 1 1 1 1</pre>
```

```
#The marginal mean is 1/d round(colMeans(w),2)
## [1] 0.2 0.2 0.2 0.2 0.2 0.2
```

2 Description of the models implemented

The different models implemented are described in Dombry et al. (2016), but some other models can be found and are described here.

1. logistic distribution (log) The logistic model of Gumbel (1960) (the Gumbel Archimedean copula)

$$P(X \le x) = \exp \left[-\left(\sum_{i=1}^{n} \left(\frac{1}{x_i}\right)^{\alpha}\right)^{\frac{1}{\alpha}} \right]$$

for $\alpha > 1$. By default, rmev will transform an argument in (0,1) without warning, to conform with the implementation. The spectral measure density is

$$h_{\boldsymbol{W}}(\boldsymbol{w}) = \frac{1}{d} \frac{\Gamma(d-\alpha)}{\Gamma(1-\alpha)} \alpha^{d-1} \left(\prod_{j=1}^{d} w_j \right)^{-(\alpha+1)} \left(\sum_{j=1}^{d} w_j^{-\alpha} \right)^{1/\alpha-d}, \quad \boldsymbol{w} \in \mathbb{S}_d$$

2. **asymmetric logistic distribution** (alog) This model was proposed by Tawn (1990). It shares the same parametrization as the evd package, merely replacing the algorithm for the generation of logistic variates. Let \mathbb{B}_d be the collection of all nonempty subsets of $\{1, ..., d\}$. The distribution function of the d-variate asymmetric logistic distribution is

$$P(X \le x) = \exp \left[-\sum_{b \in \mathbb{B}_d} \left(\sum_{i \in b} \left(\frac{\theta_{i,b}}{x_i} \right)^{\alpha_b} \right)^{\frac{1}{\alpha_b}}, \right]$$

while the spectral density is

$$h_{\boldsymbol{W}}(\boldsymbol{w}) = \frac{1}{d} \frac{\prod_{j=1}^{d-1} (j\alpha_b - 1)}{\prod_{i \in b} w_i} \left(\prod_{i \in b} \frac{\theta_{i,b}}{w_i} \right)^{\alpha_b} \left(\sum_{i \in b} \left(\frac{\theta_{i,b}}{w_i} \right)^{\alpha_b} \right)^{1/\alpha_b - d}$$

The parameters $\theta_{i,b}$ must be provided in a list and represent the asymmetry parameter. The sampling algorithm, from Stephenson (2003) gives some insight on the construction mechanism as a max-mixture of logistic distributions. Consider sampling \mathbf{Z}_b from a logistic distribution of dimension |b| (or Fréchet variates if |b|=1) with parameter α_b (possibly recycled). Each marginal value corresponds to the maximum of the weighted corresponding entry. That is, $X_i = \max_{b \in \mathbb{B}} \theta_{i,b} Z_{i,b}$ for all $i=1,\ldots,d$. The max-mixture is valid provided that $\sum_{b \in \mathbb{B}_d} \theta_{i,b} = 1$ for $i=1,\ldots,d$. As such, empirical estimates of the spectral measure will almost surely place mass on the inside of the simplex rather than on subfaces.

3. **negative logistic distribution** (neglog) The distribution function of the min-stable distribution due to Galambos (1975) is

$$P(X \le x) = \exp \left[-\sum_{b \in \mathbb{B}} (-1)^{|b|} \left(\sum_{i \in b} x_i^{-\alpha} \right)^{\frac{1}{\alpha}} \right]$$

for $\alpha \le 0$ and corresponds to the model in Dombry et al. (2016). The density is (Kotz & Nadarajah, 2000)

$$h_{\mathbf{W}}(\mathbf{w}) = \frac{1}{d} \sum_{\substack{b \in B \\ c \in h}} (-1)^{|b|} \left(\prod_{j=1}^{d-1} (1 - j\alpha) \right) \left(\prod_{i \in c} w_i^{-1} \right)^{-(\alpha+1)} \left(\sum_{i \in c} w_i^{-\alpha} \right)^{1/\alpha - d}$$

4. **asymmetric negative logistic distribution** (aneglog) The asymmetric negative logistic model is alluded to in Joe (1990) as a generalization of the Galambos model. It is constructed in the same way as the asymmetric logistic distribution; see Theorem 1 in Stephenson (2003). Let $\alpha_b \leq 0$ for all $b \in \mathbb{B}_d$ and $\theta_{i,b} \geq 0$ with $\sum_{b \in \mathbb{B}_d} \theta_{i,b} = 1$; the distribution function is then

$$\mathsf{P}\left(\boldsymbol{X} \leq \boldsymbol{x}\right) = \exp\left[-\sum_{b \in \mathbb{B}} \sum_{c \in b} (-1)^{|c|} \left(\sum_{i \in c} \left(\frac{\theta_{i,b}}{x_i}\right)^{\alpha_b}\right)^{\frac{1}{\alpha_b}}\right].$$

The density is (Kotz & Nadarajah, 2000)

$$h_{\boldsymbol{W}}(\boldsymbol{w}) = \frac{1}{d} \sum_{\substack{b \in B \\ c \subset b}} (-1)^{|b|} \left(\prod_{j=1}^{d-1} (1-j\alpha) \right) \left(\prod_{i \in c} w_j \right)^{-(\alpha+1)} \left(\prod_{j \in b} \theta_{j,b} \right)^{\alpha_b} \left(\sum_{i \in c} \left(\frac{\theta_{i,b}}{w_i} \right)^{\alpha_b} \right)^{1/\alpha_b - d}$$

In particular, it does not correspond to the "negative logistic distribution" given in e.g. §4.2 of Coles & Tawn (1991) or §3.5.3 of Kotz & Nadarajah (2000). The latter is not a valid distribution function in dimension $d \ge 3$ as the constraints therein on the parameters $\theta_{i,b}$ are necessary, but not sufficient.

Joe (1990) mentions generalizations of the distribution as given above but the constraints were not enforced elsewhere in the literature. The proof that the distribution is valid follows from Theorem 1 of Stephenson (2003) as it is a max-mixture. Note that the parametrization of the asymmetric negative logistic distribution does not match the bivariate implementation of rbvevd.

5. **multilogistic distribution** (bilog) This multivariate extension of the logistic, proposed by Boldi (2009), places mass on the interior of the simplex. Let $W \in \mathbb{S}_d$ be the solution of

$$\frac{W_j}{W_d} = \frac{C_j U_j^{-\alpha_j}}{C_d U_d^{-\alpha_d}}, \quad j = 1, \dots, d$$

where $C_j = \Gamma(d - \alpha_j)/\Gamma(1 - \alpha_j)$ for j = 1, ..., d and $\boldsymbol{U} \in \mathbb{S}_d$ follows a d-mixture of Dirichlet with the jth component being $\mathcal{D}(1 - \delta_j \alpha_j)$, so that the mixture has density function

$$h_{\mathbf{U}}(\mathbf{u}) = \frac{1}{d} \sum_{j=1}^{d} \frac{\Gamma(d-\alpha_j)}{\Gamma(1-\alpha_j)} u_j^{-\alpha_j}$$

for $0 < \alpha_i < 1$, j = 1, ..., d. The spectral density is given by

$$h_{\boldsymbol{W}}(\boldsymbol{w}) = \frac{1}{d} \left(\sum_{i=1}^{d} \alpha_{i} u_{i} \right)^{-1} \left(\prod_{i=1}^{d} \alpha_{i} u_{d} \right) \left(\sum_{j=1}^{d} \frac{\Gamma(d-\alpha_{j})}{\Gamma(1-\alpha_{j})} u_{j}^{-\alpha_{j}} \right) \prod_{i=1}^{d} w_{j}^{-1}$$

for $\alpha_i \in (0,1)$

6. Coles and Tawn Dirichlet distribution (ct) The Dirichlet model of Coles & Tawn (1991)

$$h_{\boldsymbol{W}}(\boldsymbol{w}) = \frac{1}{d} \frac{\Gamma\left(1 + \sum_{j=1}^{d} \alpha_{j}\right)}{\prod_{j=1}^{d} \alpha_{j} w_{j}} \left(\sum_{j=1}^{d} \alpha_{j} w_{j}\right)^{-(d+1)} \prod_{j=1}^{d} \alpha_{j} \prod_{j=1}^{d} \left(\frac{\alpha_{j} w_{j}}{\sum_{k=1}^{d} \alpha_{k} w_{k}}\right)^{\alpha_{j}-1}$$

for $\alpha_i > 0$.

7. **scaled extremal Dirichlet** (dir) The angular density of the scaled extremal Dirichlet model with parameters $\rho > 0$ and $\alpha \in \mathbb{R}^d_+$ is given, for all $\mathbf{w} \in \mathbb{S}_d$, by

$$h_{\boldsymbol{W}}(\boldsymbol{w}) = \frac{\Gamma(\bar{\alpha} + \rho)}{d\rho^{d-1} \prod_{i=1}^{d} \Gamma(\alpha_{i})} \left\langle \{\boldsymbol{c}(\boldsymbol{\alpha}, \rho)\}^{1/\rho}, \boldsymbol{w}^{1/\rho} \right\rangle^{-\rho - \bar{\alpha}} \prod_{i=1}^{d} \{c(\alpha_{i}, \rho)\}^{\alpha_{i}/\rho} w_{i}^{\alpha_{i}/\rho - 1}.$$

where $c(\alpha, \rho)$ is the *d*-vector with entries $\Gamma(\alpha_i + \rho)/\Gamma(\alpha_i)$ for i = 1, ..., d.

8. **scaled negative extremal Dirichlet** (negdir) The angular density of the scaled negative extremal Dirichlet model with parameters $0 < \rho < \min(\alpha_1, ..., \alpha_d)$ and α is given, for all $w \in \mathbb{R}^d_+$, by

$$h_{\boldsymbol{W}}(\boldsymbol{w}) = \frac{\Gamma(\bar{\alpha} - \rho)}{d\rho^{d-1} \prod_{i=1}^{d} \Gamma(\alpha_{i})} \left\langle \{\boldsymbol{c}(\boldsymbol{\alpha}, -\rho)\}^{-1/\rho}, \boldsymbol{w}^{-1/\rho} \right\rangle^{\rho - \bar{\alpha}} \prod_{i=1}^{d} \{\boldsymbol{c}(\alpha_{i}, -\rho)\}^{-\alpha_{i}/\rho} w_{i}^{-\alpha_{i}/\rho - 1}.$$

9. **Hüsler-Reiss** (hr), due to Hüsler & Reiss (1989). It is a special case of the Brown–Resnick process. While Engelke et al. (2015) state that Hüsler–Reiss variates can be sampled following the same scheme, the spatial analog is conditioned on a particular site (s_0), which complicates the comparisons with the other methods.

Let $I_{-j} = \{1, \ldots, d\} \setminus \{j\}$ and $\lambda_{ij}^2 \ge 0$ be entries of a strictly conditionally negative definite matrice Λ , for which $\lambda_{ij}^2 = \lambda_{ji}^2$. Then, following Nikoloulopoulos et al. (2009) (Remark 2.5) and Huser & Davison (2013), we can write the distribution function as

$$\mathsf{P}\left(\boldsymbol{X} \leq \boldsymbol{x}\right) = \exp\left[-\sum_{j=1}^{d} \frac{1}{x_{j}} \Phi_{d-1,\boldsymbol{\Sigma}_{-j}}\left(\lambda_{ij} - \frac{1}{2\lambda_{ij}} \log\left(\frac{x_{j}}{x_{i}}\right), i \in I_{-j}\right)\right].$$

where the partial correlation matrix Σ_{-i} has elements

$$\varrho_{i,k;j} = \frac{\lambda_{ij}^2 + \lambda_{kj}^2 - \lambda_{ik}^2}{2\lambda_{ij}\lambda_{kj}}$$

and $\lambda_{ii} = 0$ for all $i \in I_{-j}$ so that the diagonal entries $\varrho_{i,i;j} = 1.^1$

Engelke et al. (2015) uses the covariance matrix with entries are $\varsigma = 2(\lambda_{ij}^2 + \lambda_{kj}^2 - \lambda_{ik}^2)$, so the resulting expression is evaluated at $2\lambda_{.j}^2 - \log\left(\frac{x_j}{x_{-j}}\right)$ instead. We recover the same expression by standardizing, since this amounts to division by the standard deviations $2\lambda_{.j}$

The evd package implementation has a bivariate implementation of the Hüsler–Reiss distribution with dependence parameter r, with $r_{ik} = 1/\lambda_{ik}$ or $2/r = \sqrt{2\gamma(h)}$ for $h = \|s_i - s_i\|$ for the Brown–Resnick model. In this setting, it is particularly easy since the only requirement is nonnegativity of the parameter. For inference in dimension d > 2, one needs to impose the constraint $\Lambda = \{\lambda_{ij}^2\}_{i,j=1}^d \in \mathcal{D}$ (cf. Engelke et al. (2015), p.3), where

$$\mathcal{D} = \left\{ \mathbf{A} \in [0, \infty)^{d \times d} : \mathbf{x}^{\top} \mathbf{A} \mathbf{x} < 0, \ \forall \ \mathbf{x} \in \mathbb{R}^d \setminus \{ \mathbf{0} \} \right\}$$

$$\text{with } \sum_{i=1}^d x_i = 0, a_{ij} = a_{ji}, a_{ii} = 0 \ \forall \ i, j \in \{1, \dots, d\} \right\}$$

denotes the set of symmetric conditionally negative definite matrices with zero diagonal entries. An avenue to automatically satisfy these requirements is to optimize over a symmetric positive definite matrix parameter $\Sigma = \mathbf{L}^{\top} \mathbf{L}$, where \mathbf{L} is an upper triangular matrix whose diagonal element are on the log-scale to ensure uniqueness of the Cholesky factorization; see Pinheiro & Bates (1996). By taking

$$\Lambda(\Sigma) = \begin{pmatrix} 0 & \operatorname{diag}(\Sigma)^{\top} \\ \operatorname{diag}(\Sigma) & 1 \operatorname{diag}(\Sigma)^{\top} + \operatorname{diag}(\Sigma) \mathbf{1}^{\top} - 2\Sigma \end{pmatrix}$$

one can perform unconstrained optimization for the non-zero elements of L which are in one-to-one correspondence with those of Λ .

It easily follows that generating Z from a d-1 dimensional log-Gaussian distribution with covariance $Cov(Z_i,Z_k)=2(\lambda_{ij}^2+\lambda_{kj}^2-\lambda_{ik}^2)$ for $i,k\in I_{-j}$ with mean vector $-2\lambda_{\bullet j}^2$ gives the finite dimensional analog of the Brown–Resnick process in the mixture representation of Dombry et al. (2016).

The rmev function checks conditional negative definiteness of the matrix. The easiest way to do so negative definiteness of Λ with real entries is to form $\tilde{\Lambda} = \mathbf{P}\Lambda\mathbf{P}^{\top}$, where \mathbf{P} is an $d \times d$ matrix with ones on the diagonal, -1 on the (i,i+1) entries for $i=1,\ldots d-1$ and zeros elsewhere. If the matrix $\Lambda \in \mathcal{D}$, then the eigenvalues of the leading $(d-1) \times (d-1)$ submatrix of $\tilde{\Lambda}$ will all be negative.

10. **Brown–Resnick** (br) The Brown–Resnick process is the extension of the Hüsler–Reiss distribution, and is a max-stable process associated with the log-Gaussian distribution.

It is often in the spatial setting conditioned on a location (typically the origin). Users can provide a variogram function or the resulting covariance function. See Engelke et al. (2015) or Dombry et al. (2016) for more information.

11. Extremal Student (extstud) of Nikoloulopoulos et al. (2009), eq. 2.8, with unit Fréchet margins is

$$\mathsf{P}\left(\boldsymbol{X} \leq \boldsymbol{x}\right) = \exp\left[-\sum_{j=1}^{d} \frac{1}{x_{j}} T_{d-1,\nu+1,\mathbf{R}_{-j}} \left(\sqrt{\frac{\nu+1}{1-\rho_{ij}^{2}}} \left[\left(\frac{x_{i}}{x_{j}}\right)^{1/\nu} - \rho_{ij}\right], i \in I_{-j}\right)\right],$$

where T_{d-1} is the distribution function of the d-1 dimensional Student-t distribution and the partial correlation matrix \mathbf{R}_{-i} has diagonal entry

$$r_{i,i;j} = 1,$$
 $r_{i,k;j} = \frac{\rho_{ik} - \rho_{ij}\rho_{kj}}{\sqrt{1 - \rho_{ij}^2}\sqrt{1 - \rho_{kj}^2}}$

for $i \neq k, i, k \in I_{-i}$.

The user must provide a valid correlation matrix (the function checks for diagonal elements), which can be obtained from a variogram.

12. **Dirichlet mixture** (dirmix) proposed by Boldi & Davison (2007), see Dombry et al. (2016) for details on the mixture. The spectral density of the model is

$$h_{W}(\boldsymbol{w}) = \sum_{k=1}^{m} \pi_{k} \frac{\Gamma(\alpha_{1k} + \dots + \alpha_{dk})}{\prod_{i=1}^{d} \Gamma(\alpha_{ik})} \left(1 - \sum_{i=1}^{d-1} w_{i}\right)^{\alpha_{dk} - 1} \prod_{i=1}^{d-1} w_{i}^{\alpha_{ik} - 1}$$

The argument param is thus a $d \times m$ matrix of coefficients, while the argument for the m-vector weights gives the relative contribution of each Dirichlet mixture component.

13. **Smith model** (smith), from the unpublished report of Smith (1990). It corresponds to a moving maximum process on a domain X. The de Haan representation of the process is

$$Z(x) = \max_{i \in \mathbb{N}} \zeta_i h(x - \eta_i), \qquad \eta_i \in \mathbb{X}$$

where $\{\zeta_i, \eta_i\}_{i \in \mathbb{N}}$ is a Poisson point process on $\mathbb{R}_+ \times \mathbb{X}$ with intensity measure $\zeta^{-2} d\zeta d\eta$ and h is the density of the multivariate Gaussian distribution. Other h could be used in principle, but are not implemented.

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