

Extreme value comparison of different climate model simulations and observations

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Abstract. We explore similarities and differences in extreme behavior between climate simulations and an observation product. We fit a Bayesian hierarchical threshold exceedance model to replicates of CanCM4 climate simulations. Three simulation classes are analyzed and compared: decadal, historical, and pre-industrial control. We extend the comparison to an observation product. To assess the extremes of the series considered we fit a generalized Pareto model to the exceedances over a threshold. This model is applied to data over California and the contiguous United States. Marginal comparisons are made visually with posterior parameter intervals and numerically using the Bhattacharyya distance between probability densities. We find that in some domains, the simulations are in agreement among themselves and with the observations, but in others they are quite different. We also perform a bivariate extreme value analysis using simple Pareto processes.

1 Introduction

The main focus of this paper is to compare the extreme values of an observation product with those of CanCM4 climate simulations. We address at least two questions: 1) Does using the same climate model in different ways produce similar extreme behavior? and 2) How well do the simulations agree with the observations? Answers to these questions would inform us whether the climate model is a reasonable representation of observed extremes.

The Fourth Generation Coupled Global Climate Model (CanCM4) from the Canadian Centre for Climate Modeling and Analysis (CCCma) is made up of an atmospheric component, CanAM4 (von Salzen et al., 2013), and an ocean component, CanOM4. The two components are coupled daily to produce climate predictions of a variety of variables on a roughly 2.5° degree grid over the globe (see Merryfield et al. (2013)). Two variables will be analyzed: precipitation (labeled `pr`, in meters) and maximum temperature (labeled `tasmax`, in Kelvin). We further restrict our attention to analyzing two seasons—summer and winter—and two regions—California and the U.S.

Three experimental classes that are of particular interest are decadal, historical, and pre-industrial control runs. The decadal simulations provide climate estimates for ten years into the future, after conditioning on the state of the ocean at the starting time of the simulation. We consider two decades in this analysis: 1962–1971 and 1990–1999, which are conditioned on ocean states in 1961 and 1989, respectively. Historical simulations are obtained for the years 1961–2005 and are noted for including events that affect the climate such as volcanoes. The pre-industrial control, or simply control, simulations begin at climate conditions comparable to those preceding the industrial revolution and are run over a thousand years. The purpose of the control runs is to provide some measure of internal variability of the climate system.

Decadal and historical simulations are run at $R = 10$ different initial conditions. To obtain $R = 10$ “replicates” for the control simulations, we randomly select ten non-overlapping 10-year periods.

An observation product is obtained from Maurer et al. (2002). The observations are based on daily measurements from weather stations throughout the United States and are interpolated onto a fine grid (about $1/8^\circ$ degree spacing). To make the observations comparable to the climate simulations, we take weighted sums or averages of the climate simulations and just sums or averages of the observations. See section 2 for details, along with other changes made to the data in preparation for analysis.

The classic approach to analyzing extreme values is to model block maxima (e.g. annual maxima). It can be shown that under certain conditions the block maxima of independent random variables have a distribution which belongs to the generalized extreme value (GEV) family of distributions (Coles, 2001). Such an approach naturally requires omitting a large portion of the data. This can be remedied by using a threshold exceedance model which involves selecting some large threshold and fitting the exceedances to the generalized Pareto distribution (GPD). See Coles (2001) for an excellent introduction to these and other approaches in extreme value analysis.

We take a Bayesian approach to modeling threshold exceedances (section 3.1). Under this framework, taking into account the replicated experiments is naturally addressed with a hierarchical model.

We attempt to answer the questions posed earlier by comparing posterior intervals for statistical model parameters and other quantities such as return level (section 3.3) and Bhattacharyya distance (section 3.4). Additionally, the bivariate analysis (section 4) allows us to measure the strength of tail dependence between simulations and observations.

2 Data processing

2.1 Aggregation

We will analyze precipitation and temperature over both California and the United States. In each case, we take the climate grid cell locations and create non-overlapping cells, or rectangles, such that each location is roughly in the center of the cell (see Figure 1). Then we count the number of locations from the observation product that are contained within each cell. The number of locations within the cells are used to weight the climate simulations (the right plot in Figure 1 shows which climate simulation locations have non-zero weight). For precipitation, we take a weighted sum and for temperature a weighted average. No weighting is used for the observations. Instead, a straight sum or average of all locations within our region of interest (either California or U.S.) is used. This method places the simulations and the observations on the same scale and yields daily time-series.

2.2 De-trending

Climate data are often non-stationary series characterized by complicated trends and cycles. As such, these present problems when studying extremes. Since we are interested in the

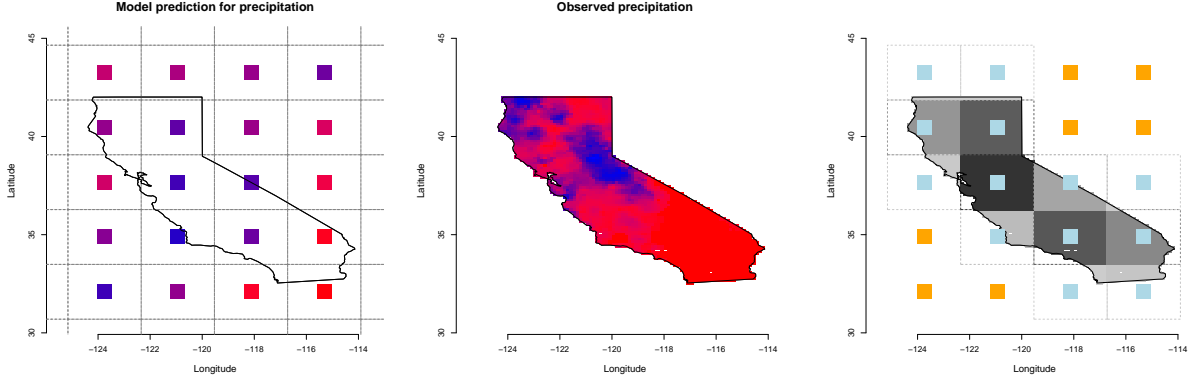


Figure 1: Left: CanCM4 simulation grid cells. Center: Observation locations. Right: method for computing weighted sum or average for CanCM4 to make values comparable with observations; the lighter gray points mean less weight is applied to the climate simulations and the darker gray means more weight. The data shown are from a single day in January.

behavior of the extremes, each time-series is “de-trended” prior to parameter estimation. This is accomplished through the use of dynamic linear models (DLMs). We will review some basic concepts for DLMs, see Prado and West (2010) chapter 4 for more details.

A normal DLM is specified by the quadruple $\{\mathbf{F}_t, v_t, \mathbf{G}_t, \mathbf{W}_t\}$ which determine how a univariate time series y_1, \dots, y_T is modeled over time. We assume

$$\begin{aligned} y_t &= \mathbf{F}_t^\top \boldsymbol{\theta}_t + \nu_t, & \nu_t &\sim N(0, v_t) \\ \boldsymbol{\theta}_t &= \mathbf{G}_t \boldsymbol{\theta}_{t-1} + \mathbf{w}_t & \mathbf{w}_t &\sim N(\mathbf{0}, \mathbf{W}_t) \end{aligned} \quad (1)$$

where $\boldsymbol{\theta}_t$ is the length- p state vector, \mathbf{F}_t is a length p vector of known constants are regressors, ν_t is observation noise, \mathbf{G}_t is the known $p \times p$ state evolution matrix, and \mathbf{w}_t is the state evolution noise. Note that ν_s and \mathbf{w}_t are independent and mutually independent.

An advantage to model (1) is its capability in yielding a smooth and flexible mean across time. After conditioning on the data up to time T , we extrapolate back over time to obtain the posterior distributions $p(\boldsymbol{\theta}_t | D_T)$ for all $t < T$, which have mean \mathbf{a}_t . Using these distributions, and given \mathbf{F}_t , the mean of y_t is simply $\mathbf{F}_t^\top \mathbf{a}_t$ (we refer the reader to Prado and West (2010) for the algorithmic details).

We must omit further details in the interest of space. Our DLM is finalized in the following way. We construct \mathbf{F}_t and \mathbf{G}_t such that the evolution of $\boldsymbol{\theta}_t$ has annual and semi-annual periods, i.e. the first and second harmonics. Higher harmonics did not seem to make significant contributions in modeling the time-series. A discount factor of $\delta = 0.9999$ was chosen, signifying low systematic variance. We assume the prior for v_t is an inverse gamma having sensible shape and scale parameters.

In the end, we are left with the residuals. See Figure 2. The blue line in the left plot is the mean of y_t , $\mathbf{F}_t^\top \mathbf{a}_t$, given the whole time series. The interior of the vertical green lines mark the summer months. The right plot is the result of subtracting the observation y_t with the mean from the DLM, which produces a roughly stationary sequence. Thus, in our extreme value analysis we fit our model to the residuals, or anomalies.

For each time-series to be analyzed, we fit a DLM having the characteristics described above to obtain the anomalies. When working within a specific season, either winter (De-

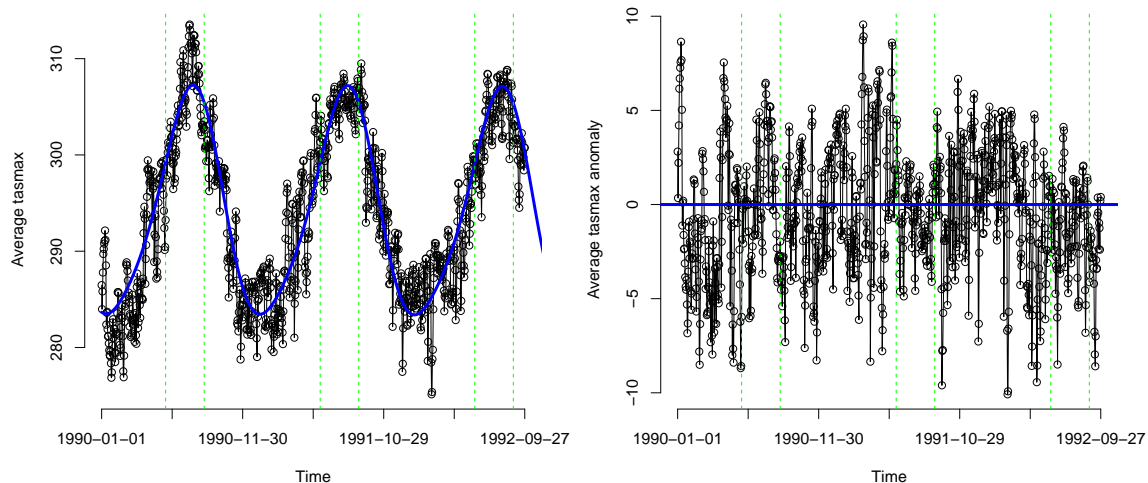


Figure 2: One of the DLMS used to calculate the anomalies. Shown is one of the decadal replicates of average `tasmax` in California for about the first two and one-half years of the time-series. The green dashed lines mark the beginning and the end of the summer months.

ember, January, February) or summer (June, July, August), we concatenate across years to form a single time series of seasonal anomalies. So, for example in winter, 28 February is followed immediately by 1 December.

3 Marginal analysis

In this section we describe our method for analyzing each of our four data sources (the three climate simulation types and the observations). For each source we have four factors having two levels each. This provides us with $4 \times 4^2 = 64$ total data sets to which we apply the models and methods of this section.

3.1 Threshold exceedance model

3.1.1 Univariate

Under some mild assumptions, for random variable X and for large enough u , the distribution of $X - u$ (the exceedance), conditional on $X > u$ is approximately

$$P(X - u \leq y | X > u) \approx H(y) = 1 - \left(1 + \frac{\xi y}{\sigma}\right)^{-1/\xi} \quad (2)$$

defined on $\{y : y > 0 \text{ and } (1 + \xi y/\sigma) > 0\}$. $H(y)$ is the distribution function for a generalized Pareto random variable with shape parameter $\xi \in \mathbb{R}$ and scale $\sigma > 0$.

Let X_1, \dots, X_n be a sequence of i.i.d. random variables and u be a high threshold. Define

$Y_i = X_i - u$ for $X_i > u$ be the k exceedances. The likelihood of (ξ, σ) is derived from (2) as

$$L(y_1, \dots, y_k; \sigma, \xi) = \sigma^{-k} \sum_{i=1}^k \left(1 + \frac{\xi y_i}{\sigma} \right)_+^{-1/\xi-1} \quad (3)$$

where $z_+ = \max(z, 0)$. This provides the basis for an extreme value analysis. For example, after declustering, the cluster maxima (which are roughly independent) may be fit using likelihood (3).

3.1.2 Hierarchical model

Suppose we have R replicates or computer simulations, each with n_i observations, for $i = 1, \dots, R$. Let X_{ij} denote the j th observation in replicate i . We assume

$$X_{ij} \sim F_i, \quad i = 1, \dots, R, \quad j = 1, \dots, n_i$$

and all X_{ij} are mutually conditionally independent. For a fixed u and each i , define the following sets:

$$A_i = \{j : x_{ij} \leq u\}, \quad A_i^c = \{j : x_{ij} > u\}$$

where $|A_i| = n_i - k_i$ and $|A_i^c| = k_i$ with k_i being the number of exceedances in replicate i . We define our exceedances as

$$y_{ij} = (x_{ij} - u) \cdot \mathbf{1}_{(j \in A_i^c)}$$

so that all observations not exceeding u are marked as 0. Let $\mathbf{y}_i = (y_{i,1}, \dots, y_{i,n_i})^\top$ and $\mathbf{y} = (\mathbf{y}_1^\top, \dots, \mathbf{y}_R^\top)^\top$.

The likelihood is given by

$$\begin{aligned} L(\mathbf{y}; \boldsymbol{\sigma}, \boldsymbol{\xi}, \boldsymbol{\zeta}) &= \prod_{i=1}^R f_{Y_i}(\mathbf{y}_i | \sigma_i, \xi_i, \zeta_i) \\ &\approx \prod_{i=1}^R \left[(1 - \zeta_i)^{n_i - k_i} \zeta_i^{k_i} \prod_{j \in A_i^c} \frac{1}{\sigma_i} \left(1 + \xi_i \frac{y_{ij}}{\sigma_i} \right)_+^{-1/\xi_i-1} \right] \end{aligned} \quad (4)$$

Note that the parameters describing the tail of F_i (i.e. ξ_i, σ_i) depend only on those observations which exceed u . The parameter $\zeta_i = P(X_{ij} > u)$, which is necessary for calculating return levels (section 3.3), is based only on the number of exceedances. This justifies the use of cluster maxima for \mathbf{y}_i .

We complete the hierarchical model formulation by specifying the following priors:

$$\begin{aligned} \xi_i | \xi, \tau^2 &\sim \text{Normal}(\xi, \tau^2) \\ \sigma_i | \alpha, \beta &\sim \text{Gamma}(\alpha, \beta) \\ \zeta_i | \zeta, \eta &\sim \text{Beta}(\zeta\eta, (1 - \zeta)\eta) \end{aligned} \quad (5)$$

$$\begin{aligned} \xi &\sim \text{Normal}(m, s^2) & \tau^2 &\sim \text{InvGamma}(a_\tau, b_\tau) \\ \alpha &\sim \text{Gamma}(a_\alpha, b_\alpha) & \beta &\sim \text{Gamma}(a_\beta, b_\beta) \\ \zeta &\sim \text{Beta}(a_\zeta, b_\zeta) & \eta &\sim \text{Gamma}(a_\eta, b_\eta) \end{aligned}$$

By combining (4) and (5) we obtain the full posterior distribution. Samples are obtained via MCMC. The inverse gamma is parametrized to have mean $b/(a - 1)$ and the gammas have mean a/b .

3.2 Extremal Index

The threshold exceedance model described in section 3.1 relies on an assumption of independence which is unrealistic for a time-series. When there is dependence between the random variables, the extremes are related according to the so-called extremal index (Leadbetter, 1983), denoted by $\theta \in [0, 1]$, which arises in the following way, as summarized in Ferro and Segers (2003). For $\{X_n\}_{n \geq 1}$ a strictly stationary sequence of random variables with marginal distribution F , the sequence has extremal index θ if for each $\tau > 0$ there is a sequence $\{u_n\}_{n \geq 1}$ such that,

$$\lim_{n \rightarrow \infty} n(1 - F(u_n)) \rightarrow \tau \text{ and}$$

$$\lim_{n \rightarrow \infty} P(\max(X_1, \dots, X_n) \leq u_n) \rightarrow \exp(-\theta\tau).$$

The extremal index describes the behavior of exceedances in the limit and can be loosely interpreted as

$$\theta = (\text{limiting mean cluster size})^{-1}.$$

As an example, suppose $\theta = 0.5$, then we would expect exceedances of a large threshold to occur in pairs; for $\theta = 0.33$, in groups of 3.

Ferro and Segers (2003) show that the extremal index arises in the limiting distribution of the times between exceedances of a threshold. If T_θ is the random variable for interexceedance times in the limit, then T_θ is distributed according to the mixture

$$(1 - \theta)\epsilon_0 + \theta\mu_\theta \tag{6}$$

where ϵ_0 is the degenerate probability distribution at 0 and μ_θ is an exponential distribution with mean θ^{-1} . This means that the role of θ is two-fold: it is both the proportion of non-zero interexceedance times and the inverse mean of non-zero interexceedance times. This poses a challenge when estimating θ since it is impossible to observe an interexceedance time of zero in practice.

We next describe the hierarchical model used to estimate θ . This is distinct from the threshold exceedance model and is used only in getting a single estimate for θ , which is used to decluster the exceedances and to calculate return levels.

3.2.1 Estimation

Suppose we have observations X_1, \dots, X_n . For a threshold u , the N exceedances $Y_i = X_i - u$ given $X_i > u$ occur at times $1 \leq j_1 < \dots < j_N \leq n$. The observed interexceedance times are given by $T_i = j_{i+1} - j_i$ for $i = 1, \dots, N - 1$. Ferro and Segers (2003) provide the following

log-likelihood

$$l(\theta, p; \mathbf{T}) = m_1 \log(1 - \theta p^\theta) + (N - 1 - m_1) \{\log(\theta) + \log(1 - p^\theta)\} \\ + \theta \log(p) \sum_{i=1}^{N-1} (T_i - 1) \quad (7)$$

where p is the probability of not exceeding the threshold. We require that this likelihood be used in a hierarchical model.

Suppose we have R replicates from a climate model with values from replicate i denoted $X_{i,1}, \dots, X_{i,n}$. If we assume these simulations are independent from each other, then we expect there to be R unique extremal indices $\theta_1, \dots, \theta_R$. However, since these all come from the same climate model, we may wish to assume that the θ_i come from a common distribution,

$$\theta_i \stackrel{iid}{\sim} \text{Beta}(\theta\nu, (1 - \theta)\nu),$$

having mean $\theta\nu/(\theta\nu + (1 - \theta)\nu) = \theta$. Under model (7), we place a similar prior on the p_i ,

$$p_i \stackrel{iid}{\sim} \text{Beta}(p\tau, (1 - p)\tau).$$

The model is completed by choosing priors for θ , ν , p , and τ —the latter two parameters being required only for model (7). We assume

$$\begin{aligned} \theta &\sim \text{Beta}(a_\theta, b_\theta) \\ \nu &\sim \text{Gamma}(a_\nu, b_\nu) \\ p &\sim \text{Beta}(a_p, b_p) \\ \tau &\sim \text{Gamma}(a_\tau, b_\tau) \end{aligned}$$

with the hyperparameters chosen to be

$$\begin{array}{ll} \theta: & a_\theta = 1 \quad b_\theta = 1/2 \\ \nu: & a_\nu = 1 \quad b_\nu = 1/10 \\ p: & a_p = 100\hat{F} \quad b_p = 100(1 - \hat{F}) \\ \tau: & a_\tau = 1 \quad b_\tau = 1/10 \end{array}$$

where $\hat{F} = \sum_{i=1}^R \sum_{j=1}^n \mathbf{1}(X_{i,j} \leq u)$. Our parametrization for the gamma random variables are such that $X \sim \text{Gamma}(\alpha, \beta)$ has mean α/β . The prior values for θ attempt to mitigate some of the issues surrounding model (7)

By assuming conditional independence between the simulations, we can construct the following log-likelihood

$$L = \sum_{i=1}^R l(\theta_i, p_i; \mathbf{T}^{(i)}) \quad (8)$$

where $\mathbf{T}^{(i)}$ is the vector of interexceedance times for replicate i having length N_i . In the univariate setting for the observation product, only model (7) is needed.

Süveges (2007) proposed on an alternative likelihood for estimating the extremal index which dealt with some of the issues noted in Ferro and Segers (2003). This likelihood was extended in Süveges and Davison (2010). Though there are advantages to the alternative likelihood, we prefer to use that given in (7). In a separate simulation study, both likelihoods performed very similarly, with some preference to model (7), within the hierarchical setting.

3.2.2 Declustering

Declustering is done as given in Ferro and Segers (2003). Each replicate is declustered separately. Let $\hat{\theta}_i$ be the posterior mean of the extremal index of each replicate. Calculate $C_i = \lfloor \hat{\theta}_i N_i \rfloor + 1$, the estimated number of independent clusters. Let T_{C_i} be the C_i th largest interexceedance time in $\mathbf{T}^{(i)}$. In the case of ties, decrement C_i by one until T_{C_i+1} is strictly greater than T_{C_i} . Clusters are formed by grouping the exceedances that are separated in time by no more than T_{C_i} . In other words, two exceedances are in the same cluster if their interexceedance time is less than or equal to T_{C_i} .

The C_i clusters that will be formed using the above scheme are assumed to be independent. For each cluster we compute the cluster maximum, this being the ultimate quantity used in our inference.

3.3 Return levels

A most useful quantity in an extreme value analysis is the return level. Generally, for a distribution G , the return level x_m is the solution to

$$G(x_m) = 1 - \frac{1}{m} \quad (9)$$

and has the convenient interpretation as the quantity that is exceeded on average once every m observations.

When working with the generalized Pareto model (2), it can be shown that the m -observation return level is

$$x_m = u + \frac{\sigma}{\xi} \left[(m\zeta\theta)^\xi - 1 \right] \quad (10)$$

where the terms ζ and θ account for the probability of exceeding u and being within a cluster, respectively. We can obtain a distribution for x_m using MCMC samples for (ξ, σ, ζ) . Posterior samples for θ are obtained separately from (ξ, σ, ζ) , and so we choose to use the posterior mean for θ when computing return levels. The intention here is avoid possible complications due to the fact we do not have samples of the joint vector $(\xi, \sigma, \zeta, \theta)$.

3.4 Bhattacharyya distance

Since our focus is on comparing the climate model simulations with the observations, we must to assess differences between a variety of posterior distributions. A naive approach may be to simply determine whether posterior intervals overlap. Though we make use of visuals in our comparison, we desire a more quantitative approach. Here, we make use of Bhattacharyya distance.

Bhattacharyya (1943) proposed a means for measuring the degree of similarity between two probability distributions. For two continuous random variables on support \mathcal{X} with densities p and q , the Bhattacharyya coefficient is defined as

$$BC(p, q) = \int_{\mathcal{X}} \sqrt{p(x)q(x)} dx \quad (11)$$

and the Bhattacharyya distance is

$$D_B(p, q) = -\log BC(p, q). \quad (12)$$

We use kernel density estimation to calculate p and q along a grid of the support and then approximate the integral in (11). If the support is different for the two random variables (as will typically be the case when comparing random variables whose parameters determine the support such as the generalized Pareto), we will integrate over the intersection of the supports.

Our approach is to compute distances from the replicates to their mean and determine whether the observations could have reasonably come from the climate model. Taking the shape parameter as an example, from the hierarchical model in 3.1.2 we have posterior samples for ξ_1^c, \dots, ξ_R^c for, say, some decadal simulations. We also have posterior samples for the mean ξ^c . Using the kernel density estimation mentioned earlier, we obtain R Monte Carlo estimates $D_B(\xi_i^c, \xi^c)$, for $i = 1, \dots, R$. From the univariate model 3.1.1 we have the shape parameter ξ^o for the observation product. Finally, we calculate $D_B(\xi^o, \xi^c)$ and ask whether this quantity falls within the range of $D_B(\xi_i^c, \xi^c)$. When this occurs, we say ξ^o is “similar” to the ξ_i^c since the observation differs from the mean climate model in a similar way as the replicates differ from the mean.

4 Bivariate analysis

The marginal analysis described in section 3 may reveal comparable extreme behavior among some of the simulations and observations, but it is insufficient to describe any tail dependence. We perform several bivariate analyses using Pareto processes to model the joint tail behavior.

4.1 Simple Pareto processes

The primary result we use is Theorem 3.2 from Ferreira and de Haan (2014). Let $C(S)$ be the space of continuous real functions on S , equipped with the supremum norm, where S is a compact subset of \mathbb{R}^d . Let X be from $C(S)$. Then the conditions of their Theorem 3.1 imply

$$\lim_{t \rightarrow \infty} P \left(T_t X \in A \mid \sup_{s \in S} T_t X(s) > 1 \right) = P(W \in A)$$

with $A \in \mathcal{B}(C_1^+(S))$, $P(\partial A) = 0$, W some simple Pareto process, and

$$T_t X = \left(1 + \xi \frac{X - u_t}{\sigma_t} \right)_+^{1/\xi}.$$

We assume that t is large enough that the theorem kicks in (implying $u = u_t$ and $\sigma = \sigma_t$). Being interested in the bivariate case, we have data at only two fixed locations $s_1, s_2 \in S$. The particular values of s_1 and s_2 are irrelevant since we are comparing climate simulations to observations which have no quantitative meaning as far as their position in S is concerned; we only require that the labels for the observations be appropriately distinguished.

We further assume that the parameters ξ and σ are indexed by $s \in S$, so that our transformation is

$$T_t X(s) = \left(1 + \xi(s) \frac{X(s) - u_t(s)}{\sigma_t(s)} \right)_+^{1/\xi(s)}.$$

The first stage of our analysis involves estimating $\xi(s)$ and $\sigma(s)$ marginally, which is accomplished by selecting a high threshold $u(s)$ and fitting the generalized Pareto distribution to the excesses $X(s) - u(s)$. The posterior means for $\xi(s)$ and $\sigma(s)$ are then used for the transformation from $X(s)$ to $W(s)$.

Every observation of $X(s)$, say $X_1(s), \dots, X_{n(s)}(s)$ is transformed with

$$W_i(s) = T_t X_i(s) = \left(1 + \hat{\xi}(s) \frac{X_i(s) - u_t(s)}{\hat{\sigma}(s)} \right)_+^{1/\hat{\xi}(s)}, \quad i = 1, \dots, n(s) \quad (13)$$

forming the vector $\mathbf{W}(s) = (W_1(s), \dots, W_{n(s)}(s))^\top$. After performing this transformation, two components are combined to form a joint vector $(W(s_1), W(s_2))$ having realizations $\mathbf{W}_{12} = (\mathbf{W}(s_1), \mathbf{W}(s_2))$, an $n(s) \times 2$ matrix. Note that when we perform the bivariate analysis, we guarantee that $n(s) = n(s_1) = n(s_2)$.

By Theorem 3.2, \mathbf{W}_{12} has rows that are realizations of a simple Pareto process. By the constructive definition of a simple Pareto process, we can write \mathbf{W}_{12} as

$$\mathbf{W}_{12} = \begin{pmatrix} Y_1 V_1(s_1) & Y_1 V_1(s_2) \\ Y_2 V_2(s_1) & Y_2 V_2(s_2) \\ \vdots & \vdots \\ Y_n V_n(s_1) & Y_n V_n(s_2) \end{pmatrix}$$

where Y_i is a standard Pareto, $V_i(s_j) \geq 0$ ($j = 1, 2$), and $V_i(s_1) \vee V_i(s_2) = 1$, for $i = 1, \dots, n = n(s)$. This is easily obtained by

$$Y_i = W_i(s_1) \vee W_i(s_2), \quad \text{and} \quad V_i(s_j) = W_i(s_j) / Y_i \quad (j = 1, 2), \quad \text{for } i = 1, \dots, n.$$

The points $(V_i(s_1), V_i(s_2))$ fall along the curve of the non-negative unit sphere with supremum norm $\{(v_1, v_2) : \|(v_1, v_2)\|_\infty = 1, v_1 \geq 0, v_2 \geq 0\}$ which is thus one dimensional. An alternative representation is to specify $(V_i(s_1), V_i(s_2))$ in terms of a scaled angle

$$\phi_i = \frac{2}{\pi} \arctan \left(\frac{V_i(s_2)}{V_i(s_1)} \right) \in [0, 1].$$

We scale ϕ_i to be in $[0, 1]$ so we can model the angle using a mixture of beta distributions and (possibly) point masses at zero and one. The theorem holds when $\sup_{s \in S} T_t X(s) > 1$ which is equivalent to $Y_i > 1$. Therefore, we need only those ϕ_i for which $Y_i > 1$, ignoring the rest. Beyond this point we assume that ϕ_i have been relabeled to include only the appropriate angles which are indexed $i = 1, \dots, k \leq n$.

Under this framework, our experience is that the distribution of ϕ_i is not easily modeled with standard distributions, including a mixture of a few beta distributions. We use Bernstein polynomial Dirichlet process prior to model the density of ϕ_i . This has the advantage of capturing a large variety of shapes on the unit interval.

ϕ_i 's that are in $[0, 0.005]$ are treated as zero and those in $[0.995, 1]$ are treated as one. The remainder are used in the BDP model, which has difficult modeling point masses.

4.2 Asymptotic dependence

The distribution function for $W_i \equiv W(s_i)$ is

$$F_{W_i}(w) = 1 - \frac{E(V_i)}{w}$$

where $V_i \equiv V(s_i)$ and $w > 1$. Using this fact, we can standardize W_i to be uniform and compute χ in this way

$$\begin{aligned} \chi &= \lim_{u \rightarrow 1} P(F_{W_1}(W_1) > u | F_{W_2}(W_2) > u) \\ &= E\left(\frac{V_1}{E(V_1)} \wedge \frac{V_2}{E(V_2)}\right) \end{aligned}$$

The major downside to the simple Pareto process is that we do not allow for asymptotic independence ($\chi = 0$) unless $P(V_1 \wedge V_2$

5 Results

For each of the four data sources (i.e. the three climate simulation classes and the observation produce), there are four factors with two levels each. The factors, with their levels, are:

1. Variable — precipitation or maximum temperature
2. Season — winter or summer
3. Decade — 1962–1971 or 1990–1999
4. Region — California or U.S.A.

There are then 16 combinations of the factors to be made. For each combination, the hierarchical model described in section 3.1.2 is fit to the decadal, historical, and control runs; the univariate model in section 3.1.1 is fit to the observation product since this data source does not have replicates.

Thresholds are chosen to be the 0.95 quantile for the climate simulations and 0.85 for the observations. These quantiles were found to be quite reasonable for this data while still giving us enough data with which to fit the models. The values of the thresholds themselves are not too important since different thresholds may produce similar return levels.

Figures 3 through 10 show posterior parameters and other quantities of interest. For the hierarchical model, we show the results of the *mean* process. For example, in Figure 3 the parameter shown is the posterior for ξ , the mean of ξ_1, \dots, ξ_R . This is in opposition to inference on an unknown replicate which would require sampling, among other things, a new shape parameter ξ^* . Therefore, the intervals are more narrow than if we looked at the posterior predictive distribution for a new replicate, but the parameters will be comparable to those from the univariate model with the observations and give us a sense of how the climate simulation performs on average.

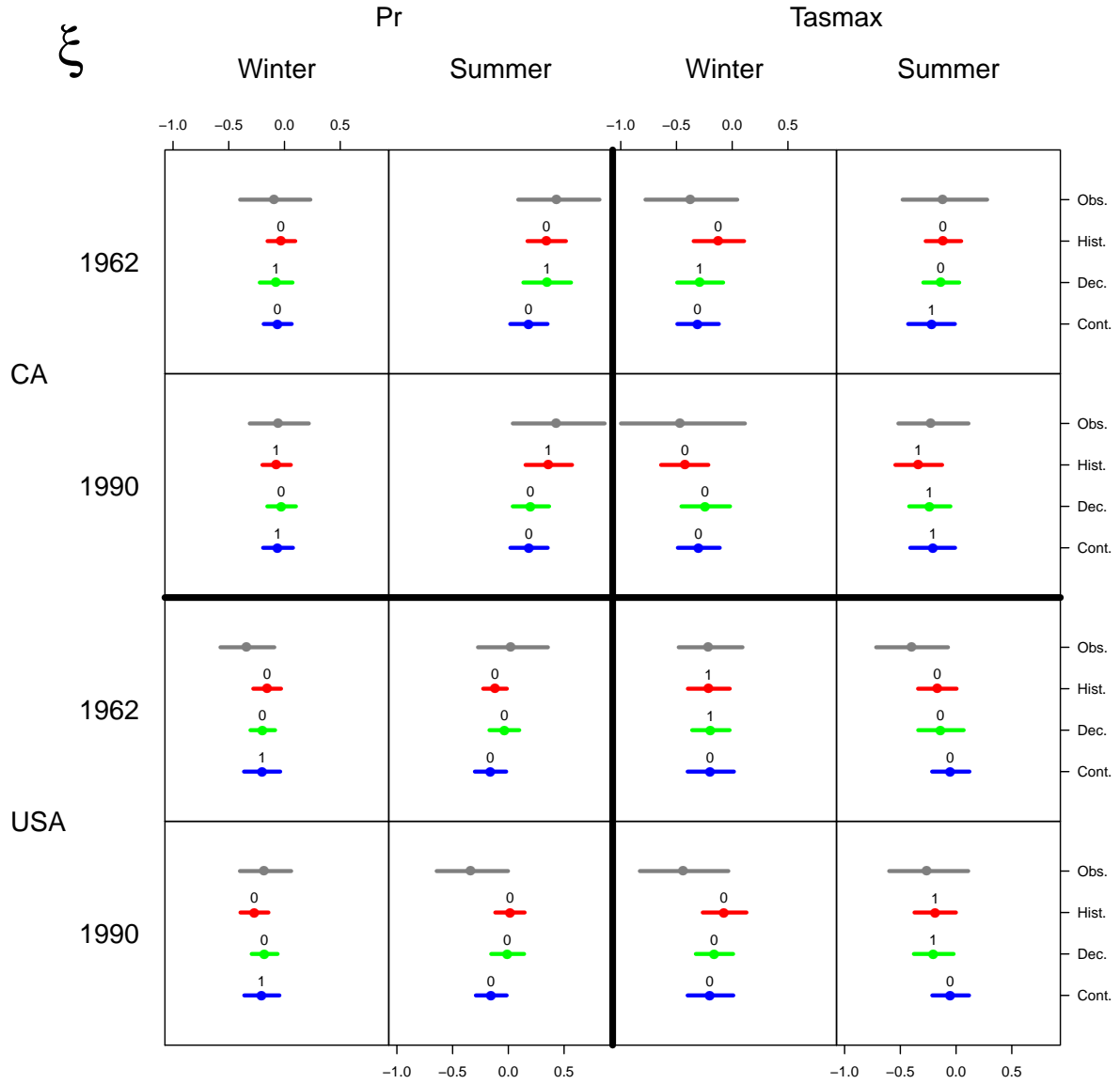


Figure 3: Posterior shape parameter, ξ , under each domain and each of the four data types. The points are the means and the lines mark the 95% h.p.d. intervals. Note: The x -axes are the same for every plot. The y -axes (for this and all subsequent figures) denote only the data type and thus hold no quantitative meaning.

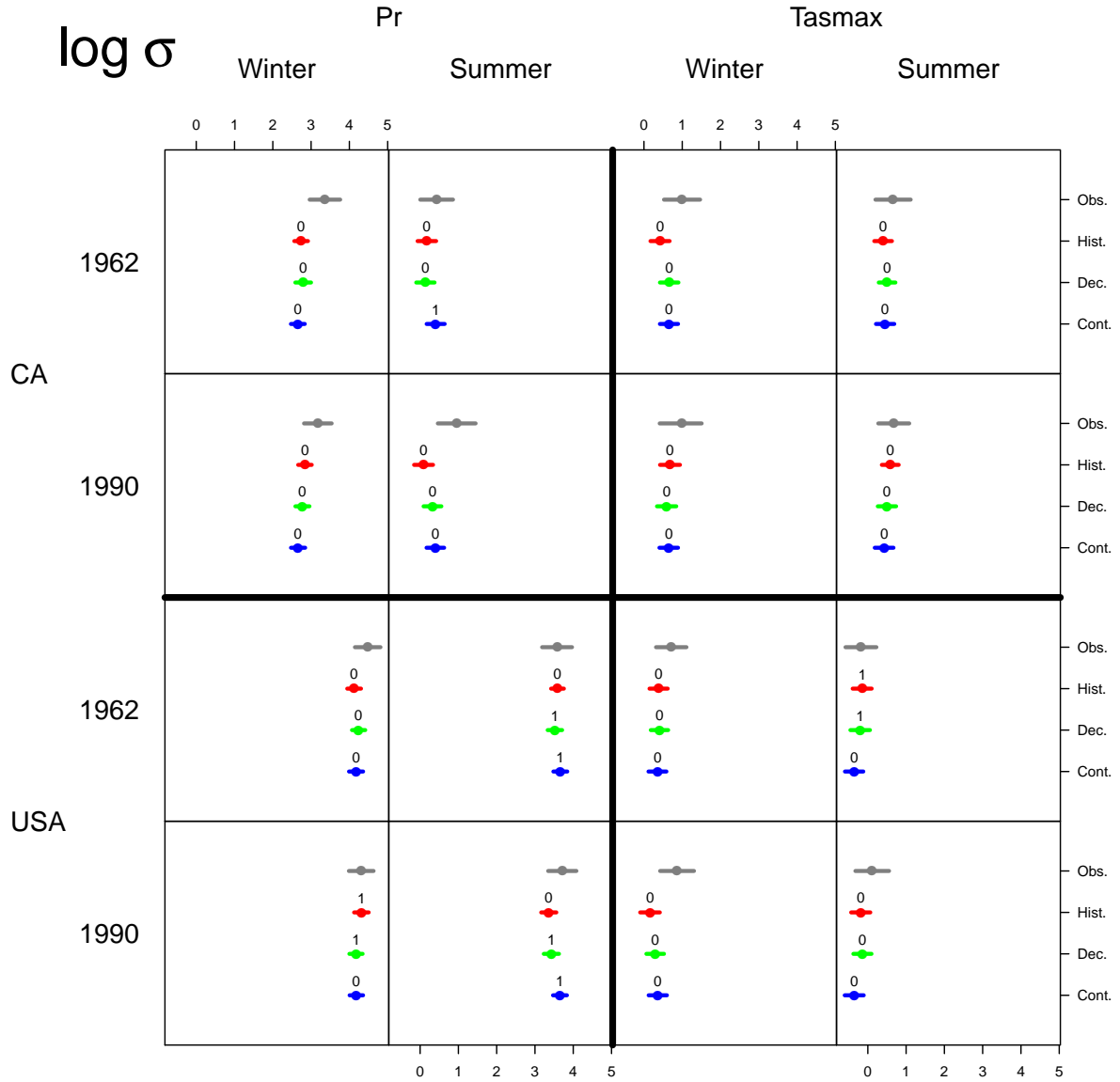


Figure 4: Natural logarithm of the posterior scale. For the CanCM4 simulations, the parameter shown is $\log(\alpha/\beta)$ (the mean scale) because σ_i follows a Gamma distribution with mean α/β . No change of variables is necessary for the observations. Note: The x -axes are the same for every plot.

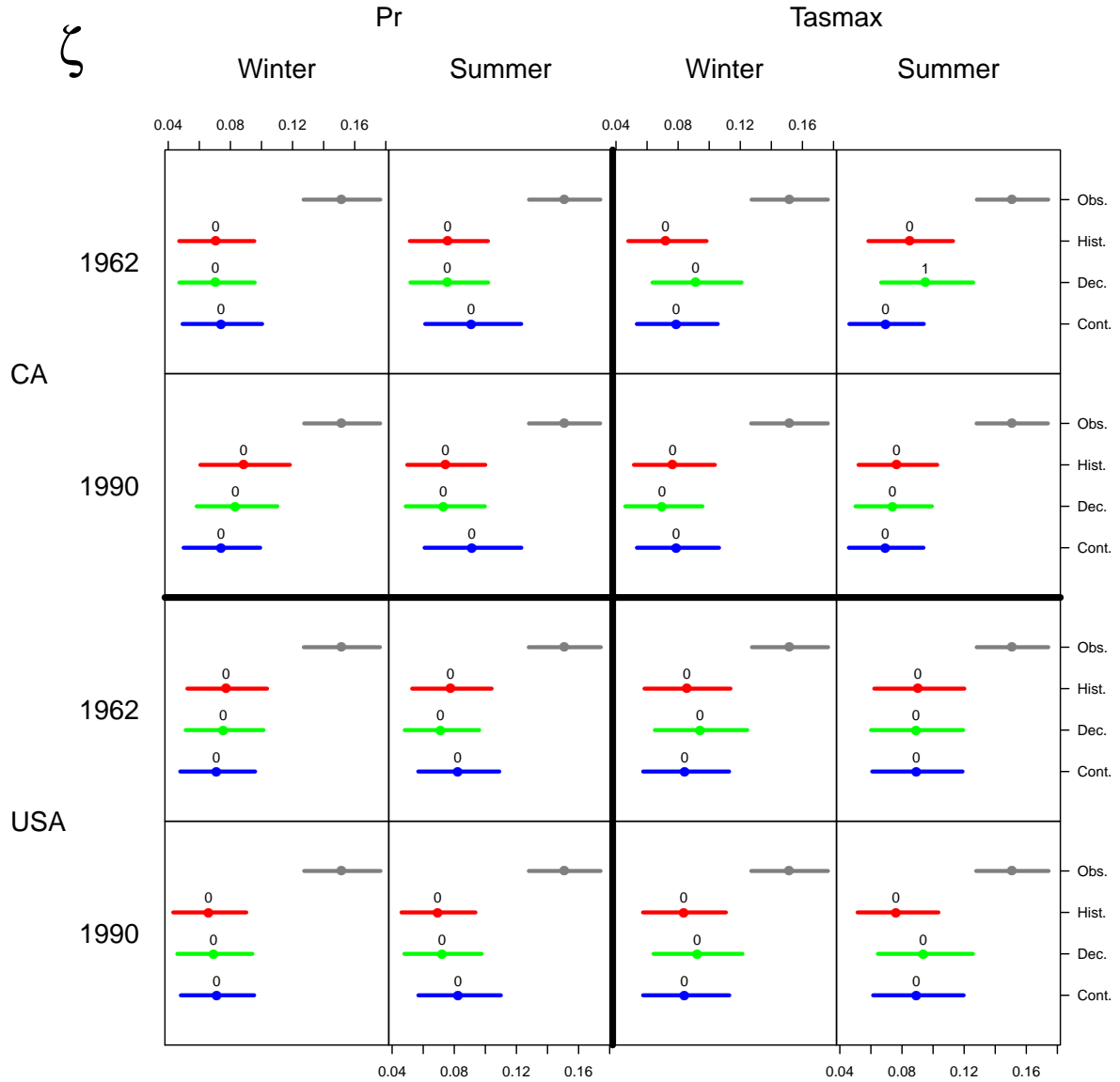


Figure 5: The probability of exceeding the threshold. These parameters are closely tied to the threshold, a user-specified quantity. Since we chose thresholds as the 0.95 quantile for the climate simulations and the 0.85 quantile for the observations, we do not expect there to be much overlap between these posteriors.

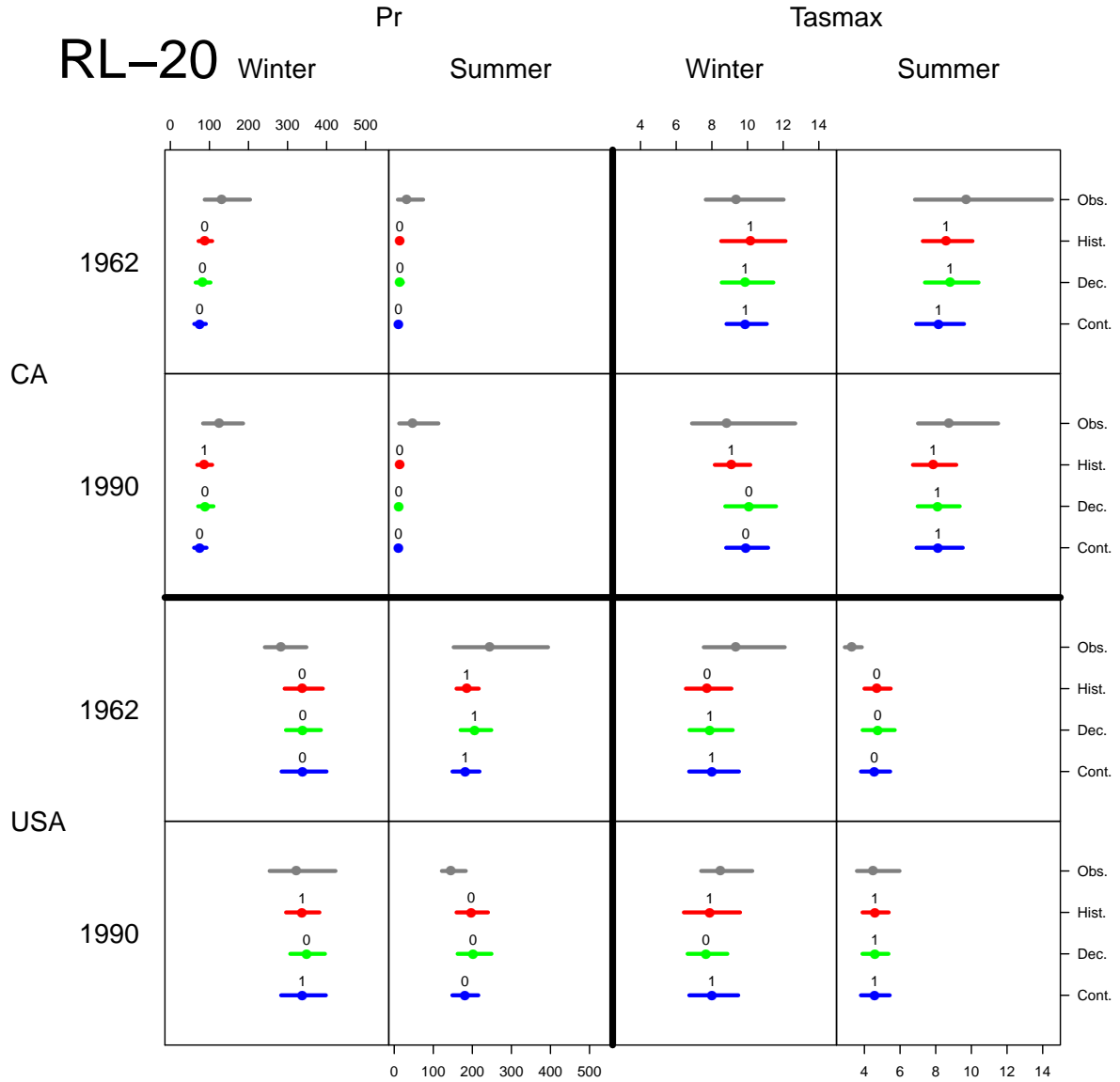


Figure 6: 20-year return levels. Note: The left two columns have the same x -axes, which are different than those in the right two columns, which have the same.

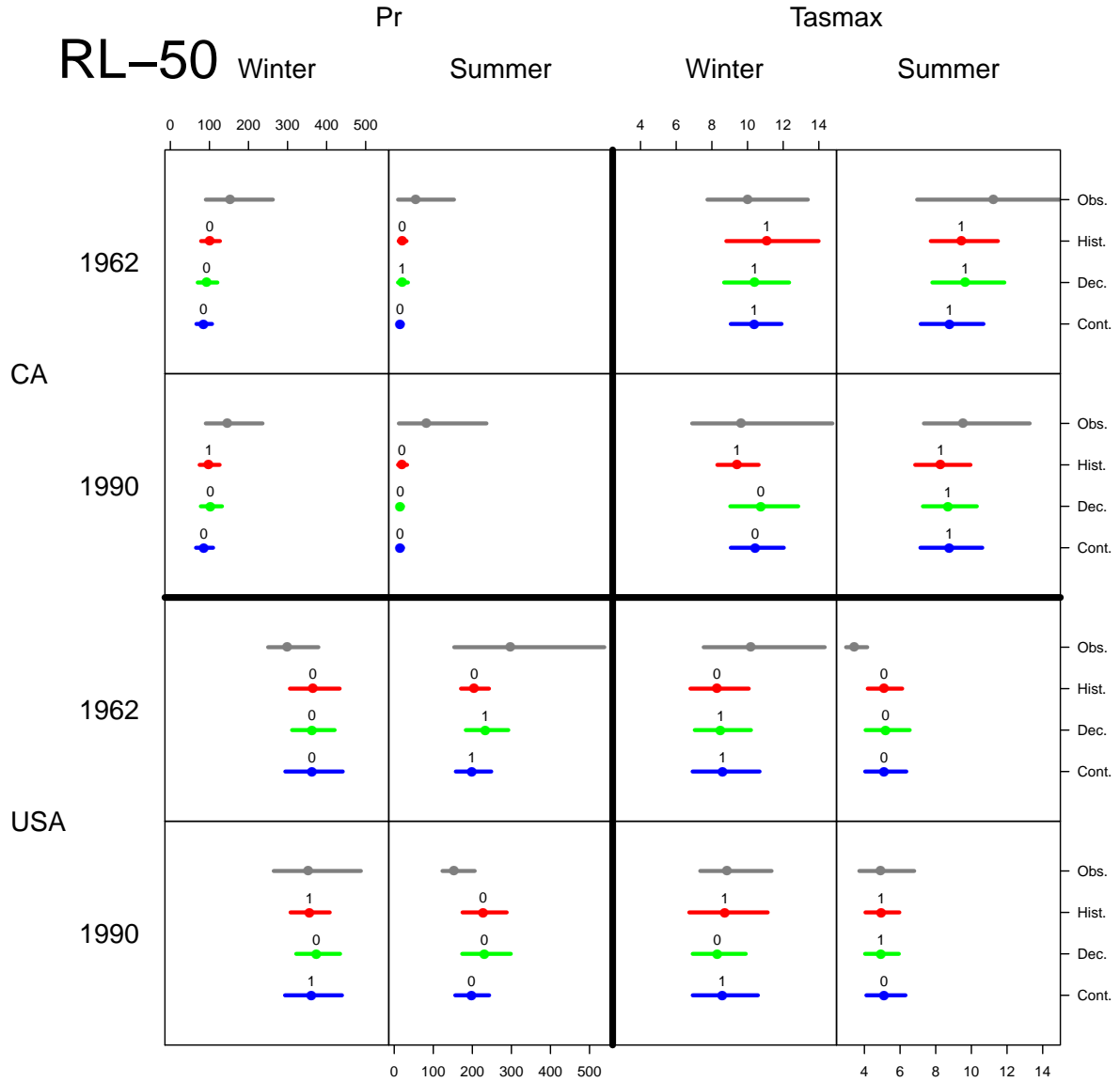


Figure 7: 50-year return levels. The x -axes are the same as those in Figure 6.

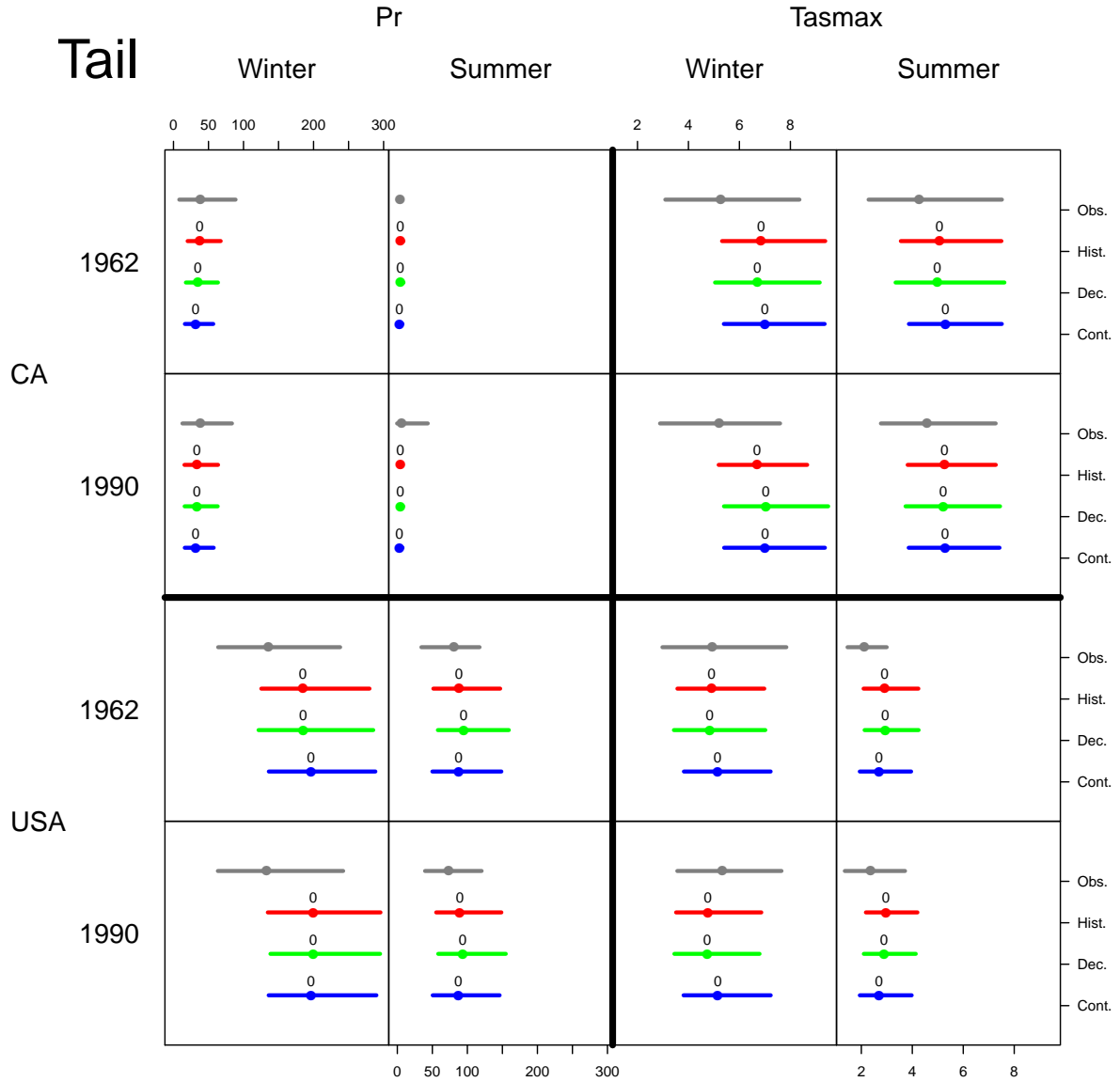


Figure 8: Mean and 95% h.p.d. for the upper tail (i.e. the generalized Pareto) of the ensemble average. As in Figures 6 and 7, the left two columns have the same x -axes and the right two columns have the same x -axes.

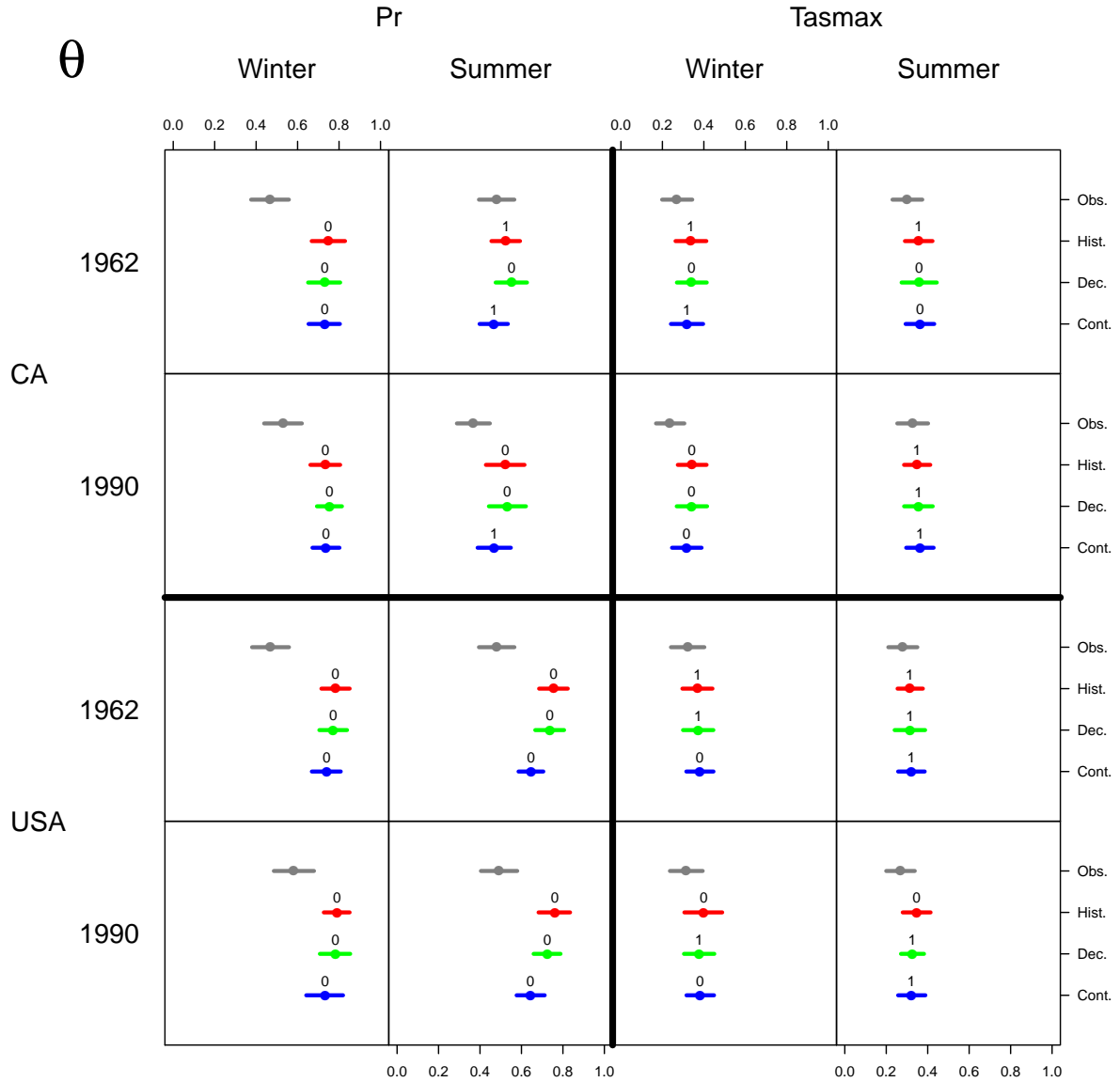


Figure 9: The mean extremal index. Like the parameters shown in Figures 3 and 4, the hierarchical mean is shown for the CanCM4 simulations.

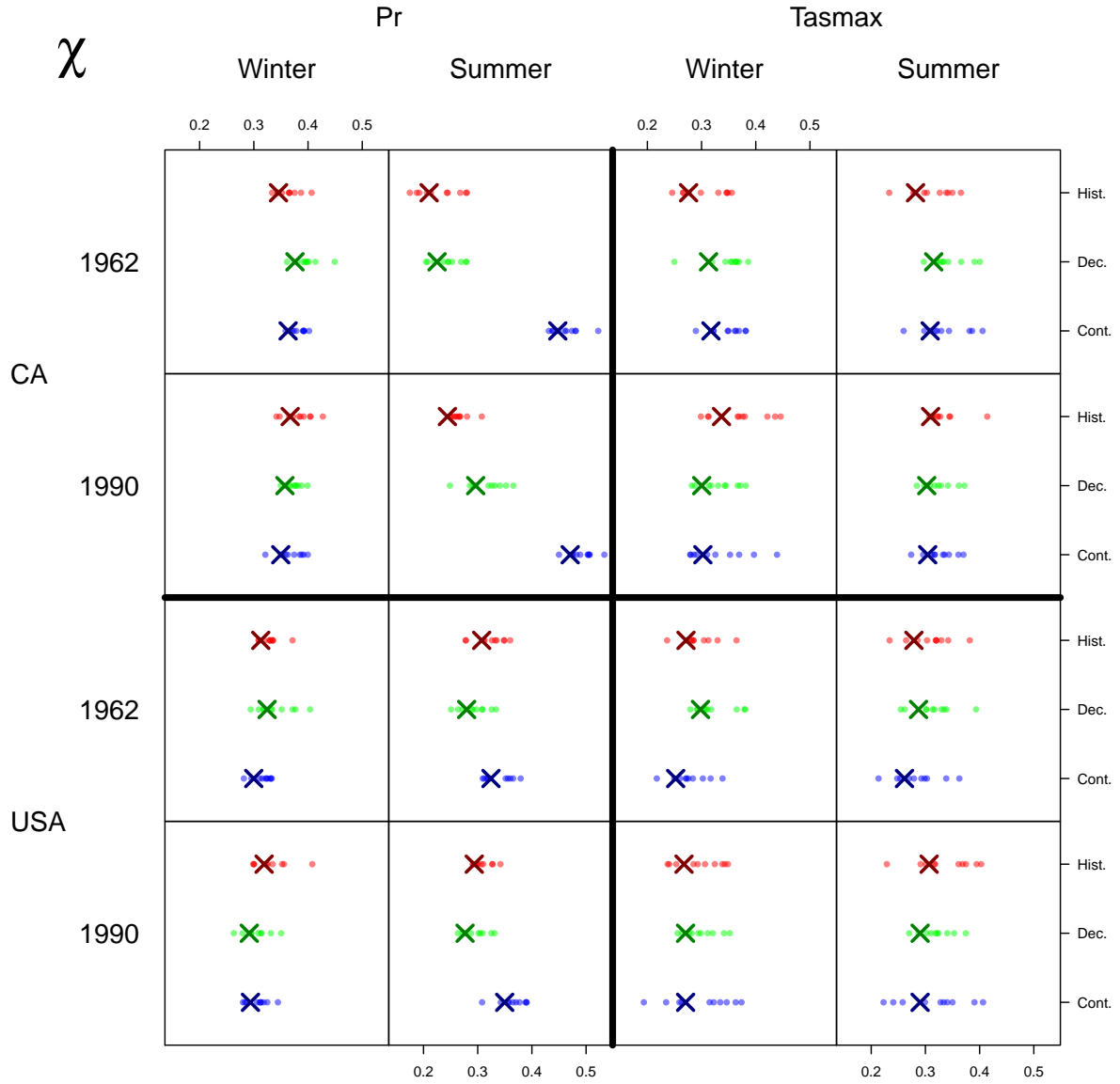


Figure 10: Measure of asymptotic dependence, calculated using results from the simple Pareto process. The dots mark the ξ for each replicate against the observations. The X marks the value of ξ when all replicates are taken together.

The posterior shape parameters in Figure 3 show overlapping bounds in many cases, but in some combinations of factors we can see some departure from the observations. The numbers shown above the lines are indicators for whether the posterior from the observations is similar to the posterior of the replicates—in the sense of Bhattacharyya distance described in section 3.4—for a particular simulation class.

Figure 4 shows the logarithm of the mean scale parameter, $\log(\sigma)$ for the observations and $\log(\alpha/\beta)$ for the simulations, and the posteriors for ζ , the probability of exceeding the threshold, are given in Figure 5.

Figures 6 and 7 give the 20- and 50-year return levels, respectively. In these figures, we have the same x -axes for the columns in total precipitation and for the columns under average maximum temperature. Some intervals are difficult to see given the scale, but we can still inspect how the return levels from the observations differ from those of the climate simulations. There is evidence of similarity between the data sources for maximum temperature (except for the 1962–1971 decade in the United States). The simulations struggle to find agreement with the observations when considering precipitation, only having two domains where the observations are similar to at least two of the climate sources (1990 USA Winter and 1962 USA Summer).

Figure 8 shows posterior predictive samples (based on the mean parameters for the hierarchical model) drawn from the generalized Pareto distribution, conditioned on the random variables being greater than the given threshold. With respect to the 95% posterior intervals, there is significant overlap of the simulations with the observations, but this is not the case with Bhattacharyya distance. This is because nearly all replicates were very close to their mean, having a distance of roughly 0.02, while the observations were different enough to have a distance of about 0.2 from the mean process. Despite all this, we saw in Figures 6 and 7 that there is still some similarity between the important return level quantities, which account for ζ and θ .

The posterior mean and 95% highest posterior density intervals for the extremal index are shown in Figure 10. The climate simulations seem to be consistent with the observations for the temperature, but not so for precipitation where they tend to overestimate θ . The Bhattacharyya distance was not computed for these parameters.

6 Conclusions

We have proposed a hierarchical threshold model to handle replicates of climate simulations. The model was applied to a variety of factor combinations and compared to a univariate threshold model for observations. A handful of similarities and differences were found to exist between simulations and the observation product.

We have accounted for trends in the time-series by subtracting out the mean of dynamic linear models, thus yielding anomalies. This poses issues of interpretability and practicality. Our extreme value analysis are in terms of the anomalies and this may not be terribly useful when making comparisons. Should we be interested in the precipitation or temperature extremes in real terms, we must add back the (possibly unknown) mean. However, with a sufficient model on the mean, it would not be unreasonable to use our analysis for projecting extremes in the future.

Some improvement on our use of the Bhattacharyya distance can be made. In this paper we decided that the observations were “similar” enough to the climate simulations if the Bhattacharyya distance fell within the distances from the replicates to their mean. A better approach may be to compute a bootstrap sample of the distances and then calculate the proportion of times this exceeds the distance from the observations.

The next step is to perform a bivariate analysis, with a vision toward the multivariate setting. This bivariate analysis is complicated due to there being replicates in the climate simulations as well as a large variety of factor combinations at which to make the comparisons.

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