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Testing One Hypothesis Multiple Times: The Multidimensional Case

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ABSTRACT

The identification of new rare signals in data, the detection of a sudden change in a trend, and the selection of competing models are among the most challenging problems in statistical practice. These challenges can be tackled using a test of hypothesis where a nuisance parameter is present only under the alternative, and a computationally efficient solution can be obtained by the “testing one hypothesis multiple times” (TOHM) method. In the one-dimensional setting, a fine discretization of the space of the non identifiable parameter is specified, and a global p -value is obtained by approximating the distribution of the supremum of the resulting stochastic process. In this article, we propose a computationally efficient inferential tool to perform TOHM in the multidimensional setting. Here, the approximations of interest typically involve the expected Euler characteristics (EC) of the excursion set of the underlying random field. We introduce a simple algorithm to compute the EC in multiple dimensions and for arbitrarily large significance levels. This leads to an highly generalizable computational tool to perform hypothesis testing under nonstandard regularity conditions. Supplementary materials for this article are available online.

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1. Introduction

In applied sciences, searches for new signals in data often reduce to a problem of detecting an unexpected mode, a variation in a trend, or a sudden change in the association among the variables considered. From a statistical perspective, all these scenarios can be characterized by a structural change in the underlying model.

The search for dark matter is one of many examples that falls under this framework. Dark matter is the substance postulated in the 1930s by Jan Oort and Fritz Zwicky (Oort 1932; Zwicky 1933, 1937) to account for missing mass in the universe. Understanding its nature and proving experimentally its existence are a hot topic in both particle physics and astronomy. One of the main physics collaborations focusing on the discovery of dark matter is the Fermi Large Area Telescope (LAT) collaboration. The experiments conducted at Fermi LAT provide measurements of photons emission over large regions of the sky with the goal of finding evidence for emission due to dark matter over emission due to the cosmic background.

To illustrate one of the many statistical challenges which arise in this context, we consider a simplified example. Here, the locations of photons emitted by the cosmic background are assumed to be uniformly distributed over the search region, whereas photon locations from the dark matter source are assumed to be distributed as a bivariate Gaussian. Specifically, let $(x_1, y_1), \dots, (x_n, y_n)$ be the coordinates at which n photons are observed, and assume that the pairs (x_i, y_i) are iid realizations

of a random vector (X, Y) with density

$$h(x, y | \theta_1, \theta_2) = (1 - \eta) \frac{1}{\lambda(\Theta)} + \eta \frac{1}{k_{\theta_1 \theta_2}} \times \exp \left\{ -\frac{1}{2\nu^2} \left[(x - \theta_1)^2 + (y - \theta_2)^2 \right] \right\}, \quad (1)$$

where $\eta \in [0, 1]$ is the relative intensity of the dark matter emission centered at $\theta = (\theta_1, \theta_2)$, Θ is the search region with area $\lambda(\Theta)$, and $k_{\theta_1 \theta_2}$ is a normalizing constant. To reduce the computational cost of the simulations proposed in Section 3, we assume that ν is a known constant and we fix it to 0.5. While the model in (1) represents a simplified scenario, it is straightforward to extend it to more realistic applications for dark matter searches (e.g., Anderson et al. 2016).

To assess if the signal of a dark matter source is present or not we test

$$H_0 : \eta = 0 \quad \text{versus} \quad H_1 : \eta > 0. \quad (2)$$

Notice that under H_0 , θ does not appear in (1) and is unidentifiable. Thus, classical inferential procedures (e.g., Wilks 1938; Chernoff 1954) do not apply to the test in (2). The most common approach to address this issue is to conduct a Monte Carlo simulation of the distribution of the tests statistic considered. Alternatively, it is possible to test for the presence of a signal at each possible location over a fine grid and then correcting for the multiplicity of tests of hypothesis conducted (see Conrad 2015, for an extensive discussion on statistical methods used in the

search of dark matter and their limitations). More recent works in physics literature discuss solutions based on random fields theory (Pilla, Loader, and Taylor 2005; Vitells and Gross 2011).

Unfortunately, when dealing with stringent significance requirements, as is typically the case in the most crucial (astro)physics discoveries (Lyons 2013), deriving the distribution of the test statistics via a Monte Carlo simulation or resampling methods (e.g., Efron and Tibshirani 1993) may be computationally prohibitive. This is further aggravated when dealing with complex models for which even a single Monte Carlo replicate can be computationally expensive. Conversely, a multiple hypothesis testing approach may be of limited use because it may be overly conservative (e.g., Bonferroni 1935, 1936), may inflate the probability of a Type I error (e.g., Benjamini and Hochberg 1995), or require independence among the tests being conducted (e.g., Hochberg 1988). Finally, although solutions based on random fields appear promising, in their current formulation they require substantial mathematical derivations of the main quantities involved (e.g., Pilla and Loader 2005; Pilla, Loader, and Taylor 2005) or may be challenging to implement computationally in more than two dimensions. In this article, we discuss a simple solution, namely “testing one hypothesis multiple times” (TOHM), which generalizes the methods proposed by Gross and Vitells (2010) and Vitells and Gross (2011), and provides a novel computational strategy. The solution proposed in this article allows us to approximate small p -values while avoiding the need for case-by-case mathematical computations and reduces drastically the number of Monte Carlo or bootstrap samples required. As additional advantage, the accuracy of the approximation proposed increases under stringent significance requirements; thus, it is particularly well suited for astrophysical and other physics searches which typically impose stringent detection thresholds.

1.1. TOHM at a Glance

In general terms, the structural change that we aim to test for can be specified via a nuisance parameter, denoted by θ , which characterizes the alternative model but becomes meaningless under the null hypothesis. For instance, in the dark matter search in (1), the parameter θ characterizing the location of the signal has no meaning when the signal intensity, η , is zero. Thus, the problem is reduced to a test of hypothesis in presence of non-identifiability. In this setting, the null hypothesis can be tested versus a sequence of *sub-alternative hypotheses*, $H_1(\theta)$, one for each possible value of θ over a fine grid. The observed *sub-test statistics* are then combined into a *global test statistic* from which the global p -value is obtained. Hence, the name: *Testing One Hypothesis Multiple times*. When θ is one-dimensional, this leads to a stochastic process indexed by θ , and a global p -value is obtained by approximating the tail probability of the supremum of this process (e.g., Davies 1977, 1987). In Algeri and van Dyk (2019), the global p -value is efficiently computed by defining a simple expansion for the expectation of the number of upcrossings of the underlying process to bound the tail probability of its supremum. The advantage of this expansion is that its leading term can be computed using a Monte Carlo

simulation that is much smaller than the one required by a full simulation of the null distribution of the global test statistic. In addition to its computational advantages, Algeri and van Dyk (2019) generalizes the approximation/bound of Davies (1977, 1987) and Gross and Vitells (2010) for the likelihood ratio test (LRT), to the supremum of a wider class of stochastic processes. Like Davies (1977, 1987), however, Algeri and van Dyk (2019) is limited to the case of θ being one-dimensional.

1.2. TOHM and Multiple Hypothesis Testing

In principle, the problem of detecting a structural change in data can be formulated as a multiple hypothesis testing problem, where an ensemble of local p -values, one for each possible value of θ over a fine grid, is produced. The main goal is to identify an adequate correction for the smallest of these p -values to guarantee the desired family-wise probability of Type I error or rate of false discoveries. In TOHM, on the other hand, an overall correction for the probability of Type I error is generated intrinsically by exploring the topology of the stochastic process of interest to obtain the global p -value.

1.3. TOHM in Multiple Dimensions: Framework and Challenges

To perform TOHM in multiple dimensions, we rely on fundamental results pertaining to the distribution of the suprema of random fields (Worsley 1994; Taylor and Adler 2003; Adler and Taylor 2007; Taylor and Worsley 2008). Specifically, we consider a random field indexed by the non-identifiable multidimensional parameter, θ , and we use the *mean Euler characteristic (EC) of the excursion set* of the random field (to be introduced Section 2) to approximate the global p -value. As discussed in Section 2.3.1, this approximation relies on the so called EC heuristic and thus we verify it numerically in our applied examples. Furthermore, closed-form expressions for the expected EC typically depend on complicated functionals, such as the so-called Lipschitz–Killing curvatures (see Section 2), whose analytical form is often hard to derive explicitly. Finally, numerical methods may be computationally challenging in multiple dimensions or when the threshold at which the excursion occurs is particularly high. Hence, there is a need for novel computational tools to adequately estimate these quantities.

1.4. Main Contributions of this Article

To overcome these difficulties, we develop a novel algorithm, based on graph theory, to efficiently compute the EC in multiple dimensions. The resulting outputs can then be used in a system of linear equations whose solution provides an estimate of the Lipschitz–Killing curvatures. The method proposed can efficiently perform bump-hunting in two or more dimensions and tackle other problems where structural changes can be characterized by a multidimensional parameter (e.g., the dark matter example above and Examples 2 and 3 in Section 2). Additionally, from a theoretical perspective, the ability to test when a multidimensional parameter is present only under the alternative further generalizes classical inferential procedures,

such as the LRT, beyond the standard regularity conditions including non nested models comparisons (Algeri, Conrad, and van Dyk 2016; Algeri and van Dyk 2019) as shown in our Example 2. Finally, the R package TOHM (Algeri 2019b) aims to facilitate the implementation of TOHM in practical applications using R programming (R Core Team 2019).

The remainder of the article is organized as follows. In Section 2, we introduce the theoretical framework of TOHM in multiple dimensions. In Section 3, we present both a suite of simulation studies that validates the results of Section 2, and three applications of TOHM to real data in the context of bump-hunting, non nested models comparison, and break-point regression. A general discussion appears in Section 4. Proofs, regularity conditions, and additional results are collected in Appendices 4 and 4. Supplementary materials of this article are available online.

2. TOHM in Multiple Dimensions

2.1. Motivating Examples

Here, we extend the results of Algeri and van Dyk (2019) to the case where the data distribution under H_1 is characterized by a multidimensional parameter, θ , that is not identifiable under H_0 . In addition to the dark matter search example introduced in Section 1, hereafter referred to as Example 1, we consider the following two examples.

Example 2 (Non nested model comparison). As discussed in Algeri, Conrad, and van Dyk (2016) and Algeri and van Dyk (2019), to choose between two non nested models $f(y, \gamma)$ and $g(y, \theta)$, we consider the comprehensive model

$$h(y, \eta, \gamma, \theta) = (1 - \eta)f(y, \gamma) + \eta g(y, \theta) \quad (3)$$

with $y \in \mathbb{R}^q$, $\eta \in [0, 1]$, $\gamma \in \Gamma \subseteq \mathbb{R}^p$, $\theta \in \Theta$ and $\Theta \subset \mathbb{R}^D$. We test both (2), and

$$H_0 : \eta = 1 \quad \text{versus} \quad H_1 : \eta < 1. \quad (4)$$

Specifically, suppose we aim to distinguish between a gamma and a log-normal distribution. Equation (3) becomes

$$(1 - \eta) \frac{e^{-y/\tau} y^{\gamma-1}}{k_{\tau\gamma}} + \eta \frac{\exp\left\{-\frac{\ln y - \mu}{2\sigma^2}\right\}}{y k_{\mu\sigma}}, \quad (5)$$

where $\eta \in [0, 1]$, $\gamma > 0$, $\tau > 0$, $k_{\tau\gamma}$ and $k_{\mu\sigma}$ are normalizing constants. In this case, the parameter which is present only under the alternative is $\theta = (\mu, \sigma)$ when testing (2) and $\theta = (\gamma, \tau)$ when testing (4). The informative scenarios arising from (2) and (4) are the following:

- if H_0 in (2) is rejected and H_0 in (4) is not, the log-normal model is selected,
- if H_0 in (4) is rejected and H_0 in (2) is not, the gamma model is selected.

In all other cases (2) and (4) are insufficient or inappropriate to select between the models being compared.

Example 3 (Break-point regression with a change of trend). We consider a logistic-regression model where the presence of a break-point θ may introduce a polynomial relationship between the logit of the probability of success and the explanatory variable x , that is

$$\log\left(\frac{\pi_i}{1 - \pi_i}\right) = \phi_1 + \phi_2 x_i + \xi(x_i - \theta)^\alpha \mathbb{1}_{\{x_i \geq \theta\}} \quad \text{for } i = 1, \dots, n, \quad (6)$$

where $x_i \in \mathbb{R}$ for all $i = 1, \dots, n$ and are considered as fixed, $\mathbb{1}_{\{\cdot\}}$ is the indicator function, $\theta = (\theta, \alpha)$, $\pi_i = P(Y_i = 1)$, $Y_i \sim \text{Binomial}(m_i, \pi_i)$, and m_i is the number of observations available for each value x_i . In this case, the test of hypothesis is

$$H_0 : \xi = 0 \quad \text{versus} \quad H_1 : \xi \neq 0. \quad (7)$$

The goal of Section 2.2 is to establish a general framework to perform tests of hypothesis like those in (2), (4), or (7).

2.2. Theoretical Framework

To formalize the general setting, consider a random sample $y = (y_1, \dots, y_n)$, with independent components distributed as the random variables or random vectors Y_i . Let $\theta \in \Theta \subset \mathbb{R}^D$, with $D \geq 1$, and suppose that for all $\theta \in \Theta$, it is possible to specify a sub-test statistic, $W_n(\theta)$, which is a function of the Y_i , and whose asymptotic or exact distribution under H_0 is known to be the same as some statistic $W(\theta)$, with known distribution. Similarly, letting θ vary, we can consider a D -dimensional random field indexed by θ , namely $\{W_n(\theta)\} = \{W_n(\theta), \theta \in \Theta\}$, whose exact or asymptotic distribution under H_0 is known to be the same as a random field $\{W(\theta)\} = \{W(\theta), \theta \in \Theta\}$. We define the *global test statistics* to be $\sup_{\theta \in \Theta} \{W_n(\theta)\}$ which, by the continuous mapping theorem, follows the same distribution as $\sup_{\theta \in \Theta} \{W(\theta)\}$ (exactly or asymptotically). To perform tests of hypothesis such as those in (2), (4), or (7) we consider the global p -value

$$P\left(\sup_{\theta \in \Theta} \{W(\theta)\} > c\right), \quad c \in \mathbb{R}, \quad (8)$$

where c is the observed value of $\sup_{\theta \in \Theta} \{W_n(\theta)\}$.

In Examples 1 and 2, we choose $W_n(\theta)$ to be the LRT statistic, that is, $W_n(\theta)$ is the difference of the log-likelihood under H_1 and H_0 multiplied by a factor of two and evaluated at the maximum likelihood estimates (MLEs) of the unknown parameters. The models involved in both Examples 1 and 2 are special cases of (3). Hence, the asymptotic distribution of $\{W_n(\theta)\}$ and its components can be derived on the basis of existing results in literature for mixture models. Specifically, for each value of θ fixed, (3) is characterized by $\gamma \in \Gamma \subseteq \mathbb{R}^p$ and the one-dimensional parameter η , which is tested on the boundary of its parameter space. Self and Liang (1987, Theorem 3, Case 5) showed that, under suitable regularity conditions, the asymptotic distribution of the LRT is given by $Z^2 \mathbb{1}_{\{Z \geq 0\}}$, which corresponds to a $\tilde{\chi}_{01}^2$ random variable (as defined in Shapiro 1985; Lin and Lindsay 1997; Takemura and Kuriki 1997) and distributed as a 50:50 mixture of χ_1^2 and zero.

The asymptotic joint distribution of $\{W_n(\theta)\}$ can be specified following the approach of Ghosh and Sen (1985), who derived the asymptotic distribution of the LRT for finite mixture models

of the form (3) with $\mathbf{y} \in \mathbb{R}$ and $\Theta \subset \mathbb{R}$. However, since in our setting both \mathbf{y} and θ are allowed to be multidimensional, in Appendix A we restate the regularity conditions of Ghosh and Sen (1985) accordingly. These assumptions allow us to establish the following result.

Proposition 2.1. Consider the mixture model in (3), for which we test either (2) or (4). If assumptions A0–A5 in Appendix A hold, under H_0 , the LRT random field $\{W_n(\theta)\}$ converges to

$$\{W(\theta)\} = \{Z(\theta)\}^2 \mathbb{1}_{\{Z(\theta) \geq 0\}} \quad \text{as } n \rightarrow \infty, \quad (9)$$

where $\{Z(\theta)\}$ is a Gaussian random field with mean zero, unit variance, and covariance function depending on θ .

Equation (9) implies that $\{W(\theta)\}$ is distributed as a $\bar{\chi}_{01}^2$ random field (as defined in Taylor and Worsley 2013, Remark 2), that is, a “patchwork” of a χ_1^2 random field and a random field which is zero everywhere, with components marginally distributed as $\bar{\chi}_{01}^2$ random variables. In the supplementary materials, we assess the validity of assumptions A0–A5 for Examples 1 and 2 which guarantee the applicability of Proposition 2.1 (see Appendix A for the proof).

In Example 3, we let $W_n(\theta)$ to be the signed-root-LRT, that is

$$W_n(\theta) = \text{sign}(\hat{\xi}) \sqrt{\text{LRT}_n(\theta)}, \quad (10)$$

where $\hat{\xi}$ is the MLE of ξ , and $\text{LRT}_n(\theta)$ is the LRT statistic evaluated at θ . In Moran (1970) (see also Davies 1977), the signed-root-LRT test statistic is shown to be equivalent to the normalized score function in (10), and asymptotically normally distributed with mean-zero and unit variance. Therefore, it is sufficient to show the asymptotic normality of the normalized score random field indexed by $\theta = (\theta, \alpha)$ to guaranteed that, for large samples, $\{W_n(\theta)\}$ is also a Gaussian random field. Proposition 2.2 establishes this result for Example 3 (see Appendix A for the proof).

Proposition 2.2. Consider the model in (6) under which we test (7), and assume that the classical Cramer’s conditions which guarantee normality and consistency of the MLE (Cramer 1946, p. 500) hold. Under H_0 , the random field $\{W_n(\theta)\}$, with components defined as in (10), converges to

$$\{W(\theta)\} = \{Z(\theta)\} \quad \text{as } m_i \rightarrow \infty \text{ for all } i = 1, \dots, n, \quad (11)$$

where $\{Z(\theta)\}$ is a Gaussian random field with mean zero, unit variance, and covariance function depending on θ .

As discussed in Section 3.2, several challenges arise from a theoretical perspective in Example 3 due to the lack of smoothness of $\{Z(\theta)\}$ in (11) and necessary to approximate (8) (see Section 2.3). Numerically, we find that the approximation obtained for the global p -value is less accurate than in Examples 1 and 2 and leads to an upper bound for (8).

Although we focus on test statistics based on the LRT, our method can in principle be applied to any test statistic whose asymptotic or exact distribution is known. For instance, one may consider, among others, the normalized score, Lagrange multipliers, or Wald test statistics. While these choices circumvent the computational burden of the optimization involved in the LRT,

they require the calculation of the Fisher information matrix and the covariance function of the resulting processes. This may introduce a substantial level of computational complexity when the integrals involved can only be computed numerically. Additionally, even when the sample size is moderately large, the asymptotic distribution may not be achieved (see, e.g., Algeri, Conrad, and van Dyk 2016, where a realistic dark matter search is conducted using only 200 events). These concerns are carefully investigated via a suite of numerical studies in Algeri et al. (2016).

2.3. Approximating Global p -Values

In the one-dimensional setting, (8) is equivalent to the probability of observing at least one upcrossing of $\{W(\theta)\}$ above c . Specifically, we say that the process $\{W(\theta)\}$ has an *upcrossing* of a threshold $c \in \mathbb{R}$ at $\theta_0 \in \Theta \subseteq \mathbb{R}$ if, for some $\epsilon > 0$, $\{W(\theta)\} \leq c$ in the interval $(\theta_0 - \epsilon, \theta_0)$ and $\{W(\theta)\} \geq c$ in the interval $[\theta_0, \theta_0 + \epsilon)$ (e.g., Adler 2000). This definition, however, is unhelpful in the multidimensional setting. Therefore, our first aim is to identify a generalization of the number of upcrossings in the context of random fields. This can be done by means of a heuristic argument known as *EC heuristic* (Adler 1981, 2000); this is described in Section 2.3.1.

2.3.1. The Euler Characteristic Heuristic

Hasofer (1978) noted that the relationship between (8) and the probability of an upcrossing can be extended to multiple dimensions by considering the number of local maxima¹ of $\{W(\theta)\}$ that exceed c , namely M_c , hence

$$P\left(\sup_{\theta \in \Theta} \{W(\theta)\} > c\right) = P(M_c \geq 1) \leq E(M_c), \quad (12)$$

where the inequality follows from Markov’s inequality. Unfortunately, analytical expressions for $E(M_c)$ are known only asymptotically in c , and thus cannot be exploited to derive multidimensional counterpart of Algeri and van Dyk (2019), which rely on evaluating $E(M_{c_0})$ at an arbitrarily small c_0 . A quantity that is more amenable and for which analytical expressions are known exactly, is the *expected EC of the excursion set of $\{W(\theta)\}$ above c* . A clear description of the EC requires a few concepts from geometry that we now summarize (see Adler 2000).

Definition 2.3. The excursion set of $\{W(\theta)\}$ above c is the set of points

$$\mathcal{A}_c = \{\theta \in \Theta : W(\theta) \geq c\}. \quad (13)$$

Definition 2.4. The EC, $\phi(A)$, of a compact set $A \subset \mathbb{R}^D$ is the additive, integer-valued functional of A uniquely determined by the following properties:

$$\phi(A) = \begin{cases} 1 & \text{if } A \text{ is homeomorphic to a} \\ & D\text{-dimensional sphere;} \\ 0 & \text{otherwise} \end{cases} \quad (14)$$

that is, there exists a continuous mapping between A and the D -dimensional sphere, whose inverse is also continuous, and

$$\phi(A \cup B) = \phi(A) + \phi(B) - \phi(A \cap B).$$

¹We are interested in scenarios where local maxima become rarer and rarer as $c \rightarrow +\infty$. Hence, we are implicitly assuming that no ridges above c occur.

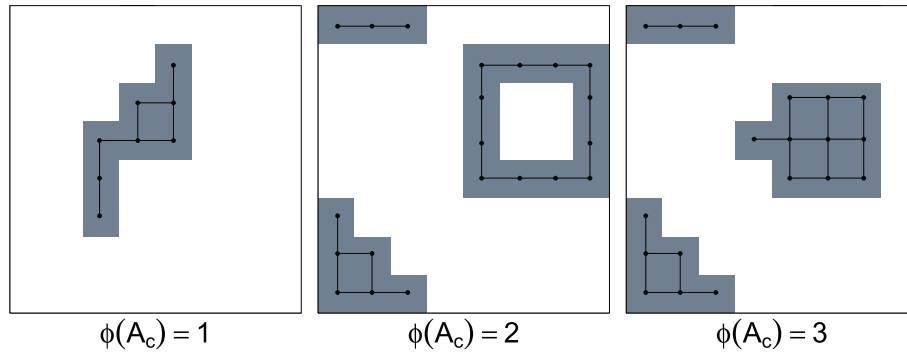


Figure 1. The shaded regions illustrate three possible excursion sets \mathcal{A}_c . The Euler characteristic (EC) of \mathcal{A}_c in the left, central, and right panels are 1, 2, and 3, respectively. The EC can be obtained by counting the number of connected components less the number of holes of \mathcal{A}_c . Alternatively, considering a quadrilateral mesh of the image (black points and black edges), the same EC is given by the number of points less the number of edges plus the number of faces (squares).

Intuitively, in two dimensions the EC of \mathcal{A}_c is the number of its connected components less its number of “holes,” see Figure 1.

The heuristic argument of Adler (1981, chap. 6) (see also Adler (2000)), aims to approximate $P(\sup_{\theta \in \Theta} \{W(\theta)\} > c)$ using the expected EC. Specifically, Adler (1981) noticed that the maxima of $\{W(\theta)\}$ above large values of c can be approximated by elliptic paraboloids, which correspond to simple connected components of \mathcal{A}_c (in two dimensions, e.g., “simple” refers to components which are connected but do not contain any hole). Additionally, as $c \rightarrow \infty$, the holes within the components of \mathcal{A}_c disappear and EC approaches the number of simple connected components around each local maxima exceeding c . It follows that the left hand side of the inequality in (12) can be approximated by the expected EC of \mathcal{A}_c , namely, $E[\phi(\mathcal{A}_c)]$, and thus we write

$$P\left(\sup_{\theta \in \Theta} \{W(\theta)\} > c\right) \approx E[\phi(\mathcal{A}_c)] \quad \text{as } c \rightarrow \infty, \quad (15)$$

where the \approx sign indicates that the difference between the right and the left hand side approaches zero as $c \rightarrow \infty$ (see Adler and Taylor 2007, chap. 14). It is important to point out that, unlike $E[M_c]$, in principle $E[\phi(\mathcal{A}_c)]$ can be negative. This means that $E[\phi(\mathcal{A}_c)]$ does not bound the left hand side of (15) from above.

Unfortunately, since the approximation in (15) is based on a purely heuristic argument, its validity is not clear. Furthermore, evaluating its accuracy is a particularly challenging task, and the error of the approximation is known only in a limited number of cases (e.g., Taylor, Takemura, and Adler 2005; Taylor and Worsley 2008). Since existing results cannot be easily extended to the applications considered in this article, we rely on a numerical assessment of the accuracy of (15) for Examples 1–3 by means of Monte Carlo simulations (see Section 3.2 and Figure 6).

2.3.2. The Expected Euler Characteristic

Another difficulty arising from the EC heuristic is the computation of $E[\phi(\mathcal{A}_c)]$. Worsley (1994, 1995) and Adler (2000) among others, gave analytical expressions of $E[\phi(\mathcal{A}_c)]$ for isotropic random fields. The seminal work of Taylor and Adler (2003), Adler and Taylor (2007), and Taylor and Worsley (2008) generalizes these approaches to non isotropic random fields of arbitrarily large dimension. They provided a convenient expansion of $E[\phi(\mathcal{A}_c)]$ for Gaussian-related (e.g., χ^2 , t , F , $\bar{\chi}^2$, etc.) random fields. Specifically, if $\{W(\theta)\}$ is a real-valued random field that

can be written as a function of iid mean zero, unit variance and suitably regular² Gaussian random fields, its expected EC can be written as

$$E[\phi(\mathcal{A}_c)] = \sum_{d=0}^D \mathcal{L}_d(\Theta) \rho_d(c), \quad (16)$$

where the $\rho_d(c)$ are functionals known as EC densities, and only depend on the (identical) marginal distribution of each component of $\{W(\theta)\}$ (see Appendix B). For example, $\rho_0(c) = P(W(\theta) > c)$. Closed-form expressions of $\rho_d(c)$ are available in literature for Gaussian, χ^2 , F , and other Gaussian-related random fields (Taylor and Adler 2003; Adler and Taylor 2007; Taylor and Worsley 2008). The functionals $\mathcal{L}_d(\Theta)$ are known as the Lipschitz–Killing curvatures of Θ . Intuitively, they measure the intrinsic volume of Θ , that is, they account for its volume, surface area, and boundaries. Their analytical forms typically rely on the covariance structure and partial derivatives of $\{W(\theta)\}$.

Unfortunately, obtaining closed-form expressions for $\mathcal{L}_d(\Theta)$ is challenging for non isotropic fields (Adler and Taylor 2007). Even in the isotropic case this may require tedious calculations and knowledge of the distribution of the derivatives of $\{W(\theta)\}$. In Sections 2.4 and 2.5, we introduce a simple approach to estimate the $\mathcal{L}_d(\Theta)$ in (16), and consequently, to compute the approximation of the global p -value in (15).

2.4. Methodological Setup

In this section, we extend the results of Algeri and van Dyk (2019) with the goal of efficiently computing the right hand side of (16).

This can be done following the approach of Vitells and Gross (2011) in two dimensions, and further formalized in Adler et al. (2017) in a multidimensional setting. Specifically, we consider a sequence of constants $c_1 \neq c_2 \neq \dots \neq c_D$, with $c_k \in \mathbb{R}$ for $k = 1, \dots, D$. Notice that, under suitable smoothness conditions (Taylor and Adler 2003, see also Section 3.2), (16) holds for any value c . Hence, we can specify the following system of linear

²A Gaussian random field $\{Z(\theta)\}$ is said to be “suitably regular” if it has almost surely continuous partial derivatives up to the second order, and if the two-tensor field induced by $\{Z(\theta)\}$ satisfies the additional mild conditions specified in Definition 3.2 of Taylor and Adler (2003).

equations

$$\begin{cases} E[\phi(\mathcal{A}_{c_1})] - \mathcal{L}_0(\Theta)\rho_0(c_1) &= \sum_{d=1}^D \mathcal{L}_d(\Theta)\rho_d(c_1) \\ E[\phi(\mathcal{A}_{c_2})] - \mathcal{L}_0(\Theta)\rho_0(c_2) &= \sum_{d=1}^D \mathcal{L}_d(\Theta)\rho_d(c_2) \\ \vdots & \\ E[\phi(\mathcal{A}_{c_D})] - \mathcal{L}_0(\Theta)\rho_0(c_D) &= \sum_{d=1}^D \mathcal{L}_d(\Theta)\rho_d(c_D), \end{cases} \quad (17)$$

where the \mathcal{A}_{c_k} are the excursion sets of $\{W(\theta)\}$ above the constants c_k and $E[\phi(\mathcal{A}_{c_k})]$ are the expected EC of \mathcal{A}_{c_k} .

In (17), the Lipschitz–Killing curvature for $d = 0$, $\mathcal{L}_0(\Theta)$, is known and corresponds to the EC of Θ (Taylor and Worsley 2008) (e.g., $\mathcal{L}_0(\Theta)$ is 0, 1, 1 or 2 if Θ is a circle, a disc, a square, or a cube, respectively). Thus, $\mathcal{L}_0(\Theta)$ need not to be estimated. Whereas, given the linear independence of the EC densities $\rho_d(\cdot)$ evaluated at different c_k , the solutions $\mathcal{L}_d^*(\Theta)$, $d = 1, \dots, D$, of (17) provide expressions for the Lipschitz–Killing curvatures for $d > 0$. Hence, we can rewrite (16) as

$$E[\phi(\mathcal{A}_c)] = \mathcal{L}_0(\Theta)P(W(\theta) > c) + \sum_{d=1}^D \mathcal{L}_d^*(\Theta)\rho_d(c), \quad (18)$$

Finally, the approximation to the global p -value in (15) can be restated, on the basis of (18), as

$$P\left(\sup_{\theta \in \Theta} \{W(\theta)\} > c\right) \approx \mathcal{L}_0(\Theta)P(W(\theta) > c) + \sum_{j=1}^D \mathcal{L}_j^*(\Theta)\rho_j(c) \quad (19)$$

for large values of c . Notice that when the $E[\phi(\mathcal{A}_{c_k})]$ are known, (18) is an exact equivalence and thus the accuracy of the approximation in (19) is the same as (15).

In practice the $E[\phi(\mathcal{A}_{c_k})]$ are unknown and estimated via a Monte Carlo simulation (details are given in Section 2.5). In Section 3, we discuss choices of the constants c_k to reduce the computational time while preserving the accuracy of the Monte Carlo estimates of $E[\phi(\mathcal{A}_{c_k})]$.

2.5. Computing the Mean Euler Characteristic via Graphs

To compute the Lipschitz–Killing curvatures $\mathcal{L}_d^*(\Theta)$ involved in the approximation of the global p -value in (19), we estimate the quantities $E[\phi(\mathcal{A}_{c_k})]$, for c_1, \dots, c_D via a Monte Carlo simulation. This requires the evaluation of $\phi(\mathcal{A}_{c_k})$ for a sequence of realizations of $\{W(\theta)\}$. In this section, we propose a convenient algorithm to achieve this goal.

To simplify notation, we assume that Θ is the cross product of the parameter spaces of components θ . Specifically, $\Theta = \Theta_1 \times \dots \times \Theta_D$, where Θ_d is the parameter space of component d of θ ; the same reasoning easily applies when $\Theta \subset \Theta_1 \times \dots \times \Theta_D$ (e.g., Example 1 described in Section 3.1). In practice, we can only evaluate $\{W(\theta)\}$ on a finite set of values for θ . We do so by placing a grid of R_d points on Θ_d , for $d = 1, \dots, D$ and evaluating $\{W(\theta)\}$ at $\theta_r = (\theta_{r1}, \dots, \theta_{rD})$ for $r = 1, \dots, R$, with $R = R_1 \times \dots \times R_D$, so that the evaluation points are the cross products of the component-wise grids. Finally, we let $\tilde{\Theta}_d$ be the ordered set of evaluation points of component d of θ and let Θ_\times be the full set of evaluation points of θ over the cross product of $\tilde{\Theta}_1, \dots, \tilde{\Theta}_D$, that is, $\Theta_\times = \{\theta_r, r = 1, \dots, R\}$. For each constant c_k in (17), we define the excursion sets of $\{W(\theta_r)\}$

above c_k to be the set of evaluation points $\tilde{\mathcal{A}}_{c_k} = \{\theta_r \in \Theta_\times : W(\theta_r) \geq c_k\}$, hence $\tilde{\mathcal{A}}_{c_k} \subseteq \Theta_\times$ provides a discretization of \mathcal{A}_{c_k} . To compute $\phi(\mathcal{A}_{c_k})$ numerically, we consider a quadrilateral mesh of $\tilde{\mathcal{A}}_{c_k}$ (Taylor and Worsley 2008), that is, the set of vertices composed of the points in $\tilde{\mathcal{A}}_{c_k}$ and the edges that connect them to form a partition of $\tilde{\mathcal{A}}_{c_k}$ into D -dimensional hyperrectangles, and denoted by \mathcal{M}_k . Specifically, we consider the set of edges, E_k^1 , such that two vertices θ_r and θ_s in $\tilde{\mathcal{A}}_{c_k}$ are joined by an edge if and only if

$$\mathfrak{d}_\varphi(\theta_r, \theta_s) = \sqrt{\sum_{d=1}^D (\varphi_d(r) - \varphi_d(s))^2} = 1, \quad (20)$$

where, $\varphi_d(r)$ is the index of component d of θ_r within its (ordered) grid of evaluation points $\tilde{\Theta}_d$ and $\mathfrak{d}_\varphi(\theta_r, \theta_s)$ is the Euclidean distance between the D indexes of the D components of θ_r and θ_s within the component-wise grids $\tilde{\Theta}_1, \dots, \tilde{\Theta}_D$. In \mathcal{M}_k , the lengths of the edges in E_k^1 are the Euclidean distances between θ_r and θ_s , that is, $\mathfrak{d}(\theta_r, \theta_s) = \sqrt{\sum_{d=1}^D (\theta_{rd} - \theta_{sd})^2}$. In quadrilateral meshes involving only unit hypercubes $\mathfrak{d}(\theta_r, \theta_s) = \mathfrak{d}_\varphi(\theta_r, \theta_s)$.

An underlying assumption of our approach is that the resolution of Θ_\times is sufficiently high to guarantee that \mathcal{A}_{c_k} is well approximated by \mathcal{M}_k . In Example 1 for instance, choosing a grid of $R = 2500$ points (Figure 2, central panel), leads to a good approximation of \mathcal{A}_1 . Specifically, we consider as benchmark for the true excursion set \mathcal{A}_1 a computation of the random field over a grid of resolution $R = 14,641$ (left panel of Figure 5). Conversely, selecting a grid of size $R = 961$ leads to a poor approximation of \mathcal{A}_1 since several among the connected components disappear (right panel of Figure 5). Hence, when the size R of the grid is not dictated by the experiment under study, the choice of R should be supported by a sensitivity analysis based on a small simulation of the random field under H_0 (e.g., Figure 2, see also Algeri and van Dyk (2019)).

The EC is calculated by alternatively adding and subtracting the number of d -dimensional hyperrectangles for $d = 0, \dots, D$ in \mathcal{M}_k (e.g., Gruber 2007, p. 268). In two dimensions for instance, the EC is obtained by counting the number of vertices, subtracting the number of edges and adding the number of rectangles (Worsley 1995; Taylor and Worsley 2008), for example, Figure 1.

To ease computations in higher dimensions, one way to count the number of hyperrectangles of arbitrarily large dimension d is summarized in Algorithm 1 and described below. The goal of Algorithm 1 is to construct graphs where the number of d -dimensional complete subgraphs (or cliques, in the second paragraph following) is equal to the number of d -dimensional hyperrectangles in \mathcal{M}_k . This can be done as follows.

For each constant c_k in (17), and for each dimension $d = 1, \dots, D$, consider an undirected unweighted graph, $\mathcal{G}_k^d = (\tilde{\mathcal{A}}_{c_k}, E_k^d)$, with vertices $\tilde{\mathcal{A}}_{c_k}$ and edges E_k^d such that two vertices θ_r and θ_s are joined by an edge if and only if

$$1 \leq \mathfrak{d}_\varphi(\theta_r, \theta_s) \leq \sqrt{d}, \quad (21)$$

where \sqrt{d} corresponds to the length of the longest diagonal of a d -dimensional unit hypercube.

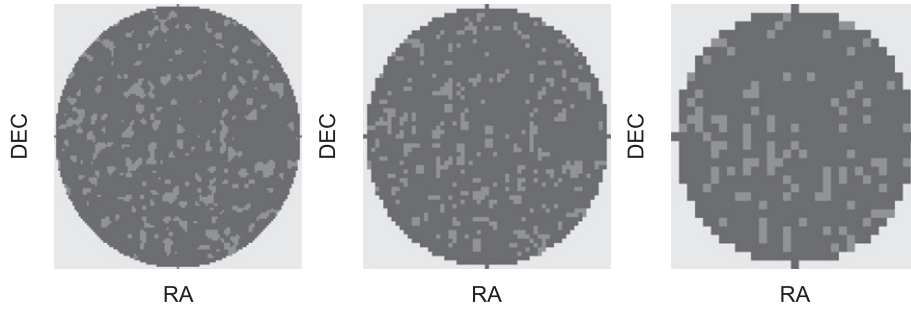


Figure 2. Approximated excursion sets of $W_n(\theta)$ in Example 1 with respect to $c_k = 1$ with $R = 14,641$, $R = 2500$, $R = 961$. The grid points in Θ_\times are chosen at distance 0.5, 1, and 2 in the left, central and right panels, respectively. As R decreases, the excursion set \mathcal{A}_k is poorly approximated by \mathcal{M}_k .

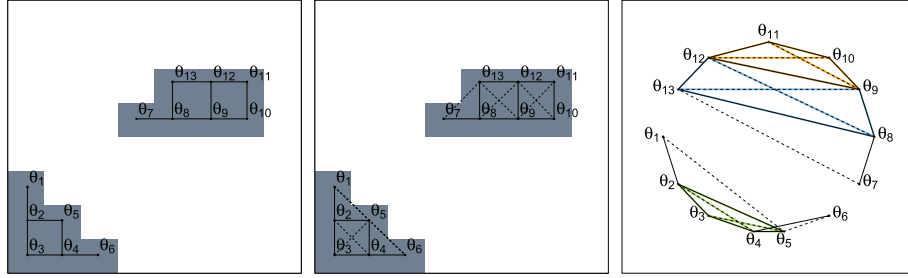


Figure 3. Left panel: Quadrilateral mesh \mathcal{M}'_k of the excursion set \mathcal{A}_k (gray area), with set of vertices $\tilde{\mathcal{A}}_{c_k}$ (black dots) and edges E_k^1 allocated according to (20) (black solid segments) of unit length. Central panel: Quadrilateral mesh \mathcal{M}'_k and diagonals of length $\sqrt{2}$ (black dashed segments). Right panel: graph $\mathcal{G}_k^2 = (\tilde{\mathcal{A}}_{c_k}, E_k^2)$ in which the three four-dimensional cliques in \mathcal{C}_k^2 are highlighted in orange, blue, and green. As expected, each clique in \mathcal{G}_k^2 corresponds to a square in \mathcal{M}'_k .

Algorithm 1 Computing $\phi(\mathcal{A}_{c_k})$ via graphs

Input 1: Constant c_k .

Step 1: For all pairs (θ_r, θ_s) in $\tilde{\mathcal{A}}_{c_k}$ calculate the distance $d_\varphi(\theta_r, \theta_s)$ in (20);

Step 2: construct the undirected graph $\mathcal{G}_k^D = (\tilde{\mathcal{A}}_{c_k}, E_k^D)$ where the edges E_k^D are allocated according to (21), with $d = D$;

Step 3: set $j = 1$;

Step 4: while $j < D$:

- (i) set $d = D - j$;
- (ii) obtain \mathcal{G}_k^d from \mathcal{G}_k^{d+1} by removing edges in E_k^{d+1} for which (21) does not hold;
- (iii) count $|\mathcal{C}_k^d|$ in \mathcal{G}_k^d via Eppstein, Löffler, and Strash (2010);
- (iv) $j = j + 1$;

Step 5: calculate $\phi(\mathcal{A}_{c_k})$ via (22).

Output: Value of $\phi(\mathcal{A}_{c_k})$.

A graph $\mathcal{G} = (V, E)$ has a clique of dimension Q if there exists a subset of Q vertices in V such that every pair of distinct vertices of the subset are connected by an edge. We denote the set of all 2^d -dimensional cliques in \mathcal{G}_k^d by \mathcal{C}_k^d . The distance between points in $\tilde{\mathcal{A}}_{c_k}$ does not affect the enumeration of the hyperrectangles in \mathcal{M}_k . Specifically, since the allocation of the edges E_k^1 only depends on the indexes $\varphi_d(r)$ of the θ_{rd} within $\tilde{\Theta}_d$, for $d = 1, \dots, D$, the number of d -dimensional hyperrectangles in \mathcal{M}_k is equal to the number of d -dimensional unit hypercubes

in a “unit” mesh, denoted by \mathcal{M}'_k , with vertices $\tilde{\mathcal{A}}_{c_k}$ and edges E_k^1 of unit length.

It follows that the 2^d vertices of each clique in \mathcal{C}_k^d is a subset of points in $\tilde{\mathcal{A}}_{c_k}$ which are at least one unit, and at most \sqrt{d} , apart one another. By construction, this implies that each clique in \mathcal{C}_k^d corresponds to a unit d -dimensional hypercube in \mathcal{M}'_k , which in turn corresponds to a d -dimensional hyperrectangle in \mathcal{M}_k . For illustrative purposes, in Figure 3 we give an example in two dimensions, where for simplicity the points θ_r are equally spaced over unit intervals in each $\tilde{\Theta}_d$, $d = 1, 2$, and thus $\mathcal{M}_k = \mathcal{M}'_k$.

Therefore, in general terms, we can compute $\phi(\mathcal{A}_{c_k})$ as

$$\phi(\mathcal{A}_{c_k}) = \sum_{d=0}^D (-1)^d |\mathcal{C}_k^d| \quad (22)$$

$$= |\tilde{\mathcal{A}}_{c_k}| - |E_k| + \sum_{d=2}^D (-1)^d |\mathcal{C}_k^d|, \quad (23)$$

where $|\cdot|$ is the cardinality of the set considered. Equation (23) follows from (22) since by construction $\mathcal{G}_k^0 = \tilde{\mathcal{A}}_{c_k}$, \mathcal{G}_k^1 is the unweighted graph with the same vertices and edges of \mathcal{M}_k and \mathcal{M}'_k ; thus $|\mathcal{C}_k^0| = |\tilde{\mathcal{A}}_{c_k}| = \sum_{r=1}^R \mathbb{1}_{\{w(\theta_r) > c_k\}}$ and $|\mathcal{C}_k^1| = |E_k^1| = \sum_{r=1}^R \sum_{s=1}^R \mathbb{1}_{\{d_\varphi(\theta_r, \theta_s)=1\}}$.

Naively, computing $|\mathcal{C}_k^d|$ by sequentially considering each subset of $\tilde{\mathcal{A}}_{c_k}$ of size 2^d requires a complexity $O(|\tilde{\mathcal{A}}_{c_k}|^{2^d} 4^D)$ to evaluate (22), a massive computation load unless D is quite

³ Notice that the main difference between the mesh \mathcal{M}_k (or \mathcal{M}'_k) and the graph \mathcal{G}_k^D is that the former depends on the position of its vertices in Θ and their distance; whereas the latter only accounts for their connectivity.

small. The advantage of converting the hyperrectangles enumeration problem into a clique-finding problem is that several efficient algorithms exist to address this challenge in near-optimal time (e.g., Bron and Kerbosch 1973; Johnston 1976; Eppstein, Löffler, and Strash 2010). In our implementations in Section 3, we use the algorithm proposed by Eppstein, Löffler, and Strash (2010), and implemented in the R function `cliques` in the `igraph` R package (Csardi and Nepusz 2006). Specifically, Eppstein, Löffler, and Strash (2010) propose a variation of the Bron-Kerbosch algorithm (Bron and Kerbosch 1973) for sparse graphs where the running time is of $O(h|\tilde{\mathcal{A}}_k|^{\frac{h}{3}})$, with $h = 2^D - 1$. This is particularly convenient in our context where the constants c_k can be chosen arbitrarily to reduce both the size of the graph and its sparsity. Hence, in Algorithm 1 we recommend a top-down approach where \mathcal{G}_k^D is constructed first, and the constants c_k can be adequately adjusted between Step 2 and Step 3 to increase sparsity in \mathcal{G}_k^D . The graphs \mathcal{G}_k^d , for $d = 0, \dots, D-1$, are obtained subsequently by removing edges for which (21) is not satisfied as d decreases. An additional advantage of this approach is that \mathcal{G}_k^D provides a simple two-dimensional representation of the D -dimensional excursion sets \mathcal{A}_{c_k} .

Finally, since in practice the $E[\phi(\mathcal{A}_{c_k})]$ are unknown, we can estimate them via Monte Carlo simulation. Specifically, for arbitrary choices of c_k , Monte Carlo estimates of $E[\phi(\mathcal{A}_{c_k})]$, namely $E[\widehat{\phi}(\mathcal{A}_{c_k})]$, can be obtained via Algorithm 1 with a small set of Monte Carlo replicates of $\{W(\theta_r)\}$ and averaging over the values $\phi(\mathcal{A}_{c_k})$ obtained at each replicate. The reader is referred to Section 3 for a discussion on the accuracy of $E[\widehat{\phi}(\mathcal{A}_{c_k})]$. Consequently, we can estimate the right-hand sides of (19) with

$$\mathcal{L}_0(\Theta)P(W(\theta_1) > c) + \sum_{j=1}^D \widehat{\mathcal{L}}_d^*(\Theta) \rho_d(c), \quad (24)$$

where $\widehat{\mathcal{L}}_d^*(\Theta)$ are the solution of the system of equation in (17) with $E[\phi(\mathcal{A}_{c_k})]$ in the left hand sides of each equation replaced by their Monte Carlo estimates $E[\widehat{\phi}(\mathcal{A}_{c_k})]$.

3. Numerical Results

3.1. Case Studies: Description

In this section, we apply TOHM to the three examples introduced in Sections 1 and 2, that is, a dark matter signal search, a non-nested model comparison and a logistic regression with a break point and change of trend.

In Example 1, we consider a realistic simulation of the Fermi LAT obtained with the *gtobssim* package⁴ in the FermiTools Python interface (Razzano 2013). This analysis represents a simplified example in the context of searches for γ -ray lines in galaxy clusters (Weniger 2012; Ackermann et al. 2015; Anderson et al. 2016). Our goals are (i) to assess the presence of a photon emission due to a dark matter source in addition to cosmic background photons, and (ii) to identify the location at which maximum evidence in favor of the suspected source is achieved. The cosmic background is uniformly distributed over the search region Θ which in this case corresponds to a

disc in the sky of 30° radius and centered at (195 RA, 28 DEC), where RA and DEC are the coordinates in the sky, and thus in (1) $x \in [165, 195]$, $y \in [28 - \sqrt{30^2 - (x - 195)^2}, 28 + \sqrt{30^2 - (x - 195)^2}]$. In our simulation, the dark matter source emission is located at $(\theta_1, \theta_2) = (174.952, 37.986)$ and realistic representations of the systematic errors, as well as the calibration of the detector, are included. This set up led to 139,821 background events and 144 dark matter events; these data are available in the supplementary materials.

In Example 2, we apply TOHM to the *Compressive strength and strain of maize seeds dataset* available in the R package `goft` (Gonzalez-Estrada and Villesanor-Alva 2017). The dataset records the compression strength in Newtons of 90 seeds and the goal is to choose between a gamma and a log-normal distribution for the data. To ease our computations, in (5) we let $y \in (0, 1000]$ and $(\mu, \sigma^2) = [1, 10] \times [0.2, 5]$.

Finally, in Example 3 we consider the *Down Syndrome dataset* available in the R package `segmented` (Muggeo 2008). The dataset records whether babies born to 354,880 women are affected by Down Syndrome. Our goal is to use TOHM to assess the presence of a break point when regressing the logit of the probability π_i that a woman of age x_i delivers a baby with down syndrome, where $x_i \in [17, 47]$, and we let $\theta \in [20, 44]$. In contrast to the analysis in Algeri and van Dyk (2019) we allow a change of trend after the break point. Specifically, we allow for a quadratic trend, a change of the linear trend or a break due to a change of the intercept, that is, $\alpha \in \{0, 1, 2\}$. The data for Examples 1–3 are plotted in Figure 4.

3.2. Goodness of the Approximations

Our first task is to assess the accuracy of the approximation of $P(\sup_{\theta \in \Theta} \{W(\theta)\} > c)$ in (24), as $c \rightarrow \infty$.

In Examples 1 and 2, $\{W(\theta)\}$ is a function of a zero mean and unit variance Gaussian random field $\{Z(\theta)\}$, whereas in Example 3, $\{W_n(\theta)\}$ is asymptotically distributed as $\{Z(\theta)\}$ (see Propositions 2.1 and 2.2). As discussed in footnote 2, among the conditions of Taylor and Adler (2003, p. 347) which guarantee the validity of (16), $\{Z(\theta)\}$ must have almost surely continuous partial derivatives up to the second order, and the two-tensor field induced by $\{Z(\theta)\}$ must be not degenerate over Θ . These assumptions guarantee smoothness of $\{W(\theta)\}$ but unfortunately, they are often difficult to verify directly as they require knowledge of both the joint distribution of the random fields and their derivatives. In this article, we limit our assessment of the smoothness of $\{W(\theta)\}$ to a small Monte Carlo simulation, and in Figure 5 we report the results of one of the Monte Carlo replicates obtained for each of the examples considered.

Both Examples 1 and 2 exhibit smooth random fields (left and central panel) under H_0 . This is not surprising since the covariance function of the underlying Gaussian process $\{Z(\theta)\}$ can be specified as a function of the respective Fisher information matrix (Ghosh and Sen 1985, p. 16), and which can be shown to be twice differentiable (see page 4 of the supplementary materials). Conversely, the signed-root LRT random field in Example 3 appears to be particularly irregular. This is due to the fact that $\{W_n(\theta)\}$ in (10) is continuous, but not continuously differentiable with respect to the parameter θ , which indexes the location of the break-point. Keeping α fixed, the EC heuristic is

⁴<http://fermi.gsfc.nasa.gov/ssc/data/analysis/software>

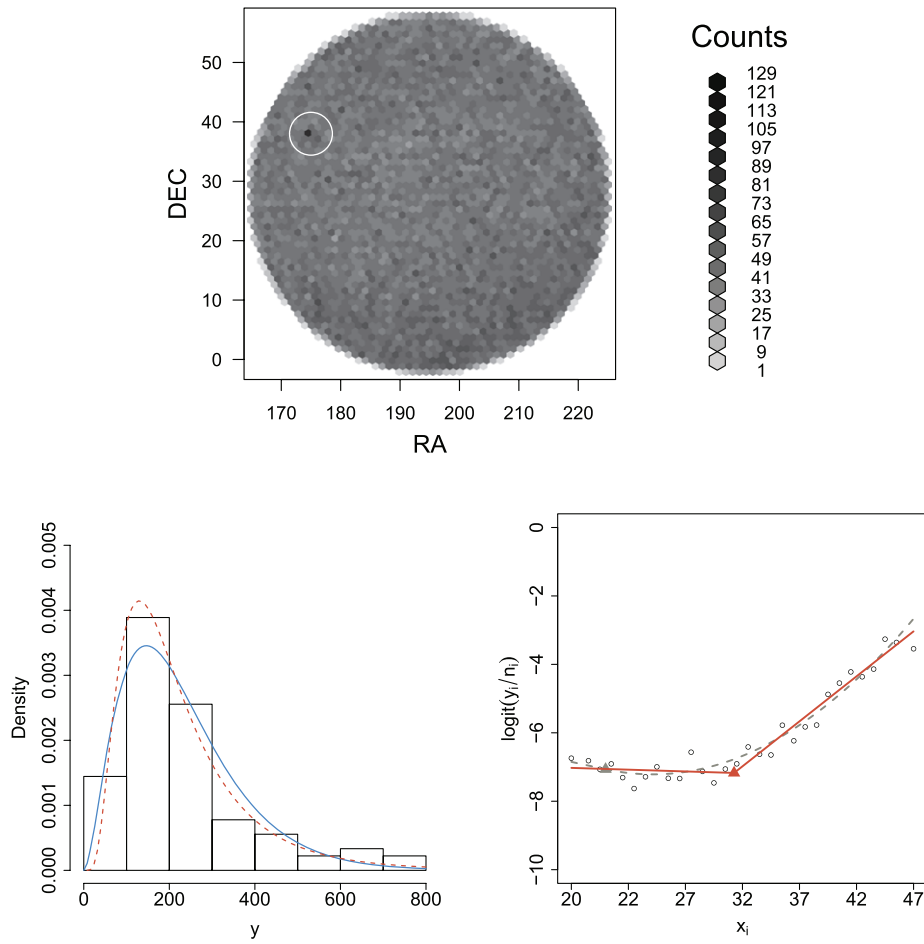


Figure 4. Top panel: Two-dimensional histogram of the Fermi-LAT realistic data simulation for Example 1. The white circle indicates the location at which the LRT-process achieves is maximum, that is, $\theta = (175, 38)$ with estimated intensity $\hat{\eta} = 0.001$. Bottom left panel: Histogram of maize seeds strength in Example 2. The null model in (2) (blue solid curve) is fitted as a gamma distribution with $(\hat{\tau}, \hat{\gamma}) = (2.762, 83.007)$. The null model in (4) (red dashed line) is fitted as a log-normal distribution with $(\hat{\mu}, \hat{\sigma}) = (5.243, 0.614)$. Bottom right panel: Down syndrome data, the model in (6) selected by THOM is a break point logistic regression with linear trend (red solid lines), that is, $\alpha = 1$ and $\theta = 31.265$ (red triangle). For comparison, a break point logistic regression with change of trend from linear to quadratic (gray dashed line) is also fitted while fixing $\alpha = 2$. In this case the breakpoint occurs at $\theta = 20$ (gray triangle).

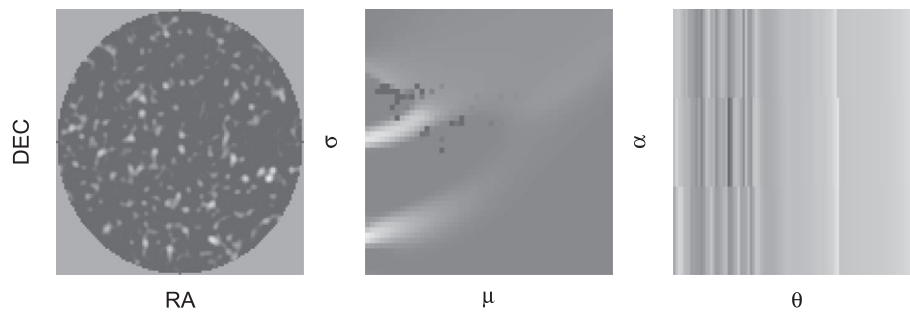


Figure 5. Simulated random fields under H_0 for Examples 1–3 (left, central, right panels, respectively).

not affected by this lack of regularity if the number of jumps in the derivative with respect to θ is finite (see Davies 1987); however, when considering $\theta = (\theta, \alpha)$, the number of jumps easily diverges if the parameter space of α is continuous. In Example 3, this effect is mitigated by choosing $\alpha \in \{0, 1, 2\}$, but this introduces a different source of non-regularity since $\{W_n(\theta)\}$ is no longer continuous with respect to α . Therefore, it is particularly interesting to assess if (24) can still be used as a reliable approximation for $P(\sup_{\theta \in \Theta} \{W(\theta)\} > c)$ despite this lack of regularity.

In Figure 6, we show as red dashed lines the Monte Carlo estimates of $P(\sup_{\theta \in \Theta} \{W(\theta)\} > c)$ obtained by simulating data under the null model for each of example; the Monte Carlo errors are given by the pink areas. For Examples 1 and 3, we simulated 130,000 datasets, each of size 10,000. For Example 2, a sample size of 100,000 was needed to guarantee the marginal $\bar{\chi}_{01}^2$ asymptotic distribution of the LRT statistics. This, along with the three-dimensional constrained optimization needed to compute the LRT for each θ at each replicate, drastically reduced the computational speed. Therefore, for this example we only

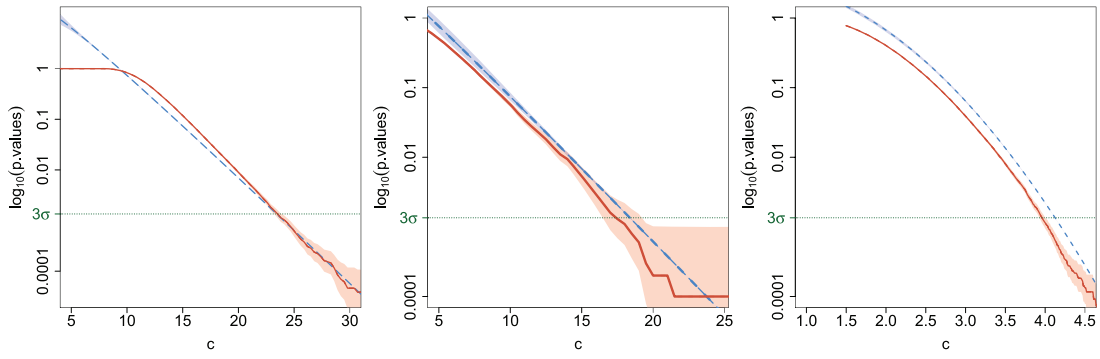


Figure 6. Estimated approximations in (24) (blue dashed line), Monte Carlo estimates of $P(\sup_{\theta \in \Theta} \{W(\theta)\} > c)$ (red solid line) in \log_{10} -scale, and Monte Carlo errors (pink areas) for increasing values of the threshold c , for Example 1 (left panel), Example 2 (central panel), and Example 3 (right panel). Monte Carlo errors associated with $E[\phi(\mathcal{A}_{c_k})]$ in (24) are plotted as gray areas.

Table 1. Computational times needed to compute the plots in Figure 6.

Example	n	R	Blue dashed lines			Red solid lines		
			# MC rep.	# cores used	Clock time	# MC rep.	# cores used	Clock time
Example 1	10,000	2821	100	1	~42 min	130,000	10	~91 hr
Example 2	100,000	2500	100	1	~154 min	10,000	10	~27 hr
Example 3	100,000	150	100	1	~1 min	130,000	10	~2 hr

NOTE: The of approximated p -value in (24) plotted in Figure 6 as blue dashed lines are conducted with a smaller number of Monte Carlo replicates (fourth column) and the one required by a full Monte Carlo simulation of $P(\sup_{\theta \in \Theta} \{W(\theta)\} > c)$ (seventh column) and plotted in Figure 6 as red solid lines. Consequently, the computational time is substantially reduced (sixth and ninth columns).

considered a simulation of 10,000 Monte Carlo replicates. The Monte Carlo estimates of $P(\sup_{\theta \in \Theta} \{W(\theta)\} > c)$ are compared with the approximation in (24) plotted as blue dashed lines as c increases (x -axis). Specifically, (24) has been computed via a set of 100 Monte Carlo replicates (each of size 10,000 for Examples 1 and 3 and 100,000 for Example 2), to estimate the quantities $E[\phi(\mathcal{A}_{c_k})]$ in (17), and the Lipschitz–Killing curvatures $\mathcal{L}_d^*(\Theta)$ in (18). The computational time needed to implement the approximation in (24) and the one required by a full simulation of $P(\sup_{\theta \in \Theta} \{W(\theta)\} > c)$ are reported in Table 1. In all the examples considered, (24) reduces the computational time from hours to minutes.

For Example 1 (left panel of Figure 6), we consider a grid of size $R = 2821$ over the 30° radius circular search region centered at (195 RA, 28 DEC). Since in this case Θ is given by a disc, its EC is one and thus $\mathcal{L}_0(\Theta) = 1$. To estimate $\mathcal{L}_1(\Theta)$ and $\mathcal{L}_2(\Theta)$ we consider $c_1 = 1$ and $c_2 = 8$, which lead to $\widehat{\mathcal{L}}_1^*(\Theta) = -244.053$ and $\widehat{\mathcal{L}}_2^*(\Theta) = 644.244$. As shown in the left panel of Figure 6, the approximation becomes particularly accurate as c approaches 24; this corresponds to the threshold required for a 3σ detection (see (25) for a definition of “ σ -significance”).

For Example 2 in the central panel of Figure 6, we define a grid of size $R = 2500$ over the square $[1, 10] \times [0.2, 5]$. Again $\mathcal{L}_0(\Theta) = 1$, and we chose $c_1 = 2$ and $c_2 = 3$. The resulting estimates for the Lipschitz–Killing curvatures are $\widehat{\mathcal{L}}_1^*(\Theta) = 30.11037$ and $\widehat{\mathcal{L}}_2^*(\Theta) = 30.52665$. In this case, despite the small simulation size, a good approximation of $P(\sup_{\theta \in \Theta} \{W(\theta)\} > c)$ is quickly achieved as c approaches 10.

Finally for Example 3, the parameter space Θ corresponds to $[-12, 12] \times \{0, 1, 2\}$, and we let $R = 150$ as we only allow values of α equal to 0, 1, and 2. Selecting $c_1 = 0.5$ and $c_2 = 1$, we

obtain $\widehat{\mathcal{L}}_1^*(\Theta) = 16.724$ and $\widehat{\mathcal{L}}_2^*(\Theta) = 23.291$. Despite the lack of smoothness in the underlying process, (24) leads to an upper bound for the global p -value, see the right panel of Figure 6. We conjecture that the fact that we obtain an upper bound on the p -value rather than a close approximation is due to the several jumps in the derivative of $\{W(\theta)\}$ with respect to θ . This is despite the fact that we only consider three possible values of α . Thus, the resulting excursion set only involves simple connected components and no holes. Therefore, in this setting, $E[\phi(\mathcal{A}_{c_k})]$ corresponds to the expected number of local maxima and, as shown in (12), provides a bound on the global p -value, $P(\sup_{\theta \in \Theta} \{W(\theta)\} > c)$, from above.

3.2.1. Guidelines for Setting c_k

Our goal is to reduce the computational time needed to compute the right hand side of (19) while guaranteeing that the difference between the right and left hand sides of (19) approaches zero. Since (18) holds for any choice of c_k , $k = 1, \dots, D$, this can be done by selecting the thresholds c_k sufficiently small so that the excursion sets \mathcal{A}_{c_k} are composed by a reasonably high number of connected components. This reduces the size of the Monte Carlo simulation needed to accurately estimate the quantities $E[\phi(\mathcal{A}_{c_k})]$. Hence, the c_k should be chosen to be small enough that \mathcal{A}_{c_k} is non empty with high probability. Additionally, since both the size and the sparsity of the graph \mathcal{G}_k^D affect the running time of Algorithm 1, c_k should be selected accordingly. These points can be assessed with a sensitivity analysis. Specifically, for a given c_k , \mathcal{G}_k^D allows a two-dimensional visualization of the D -dimensional mesh \mathcal{M}_k , and thus after step 2 in Algorithm 1, c_k can be increased to increase sparsity and decrease the size of \mathcal{G}_k^D before proceeding with Steps 3–5.

Table 2. TOHM p -values computed via (24) and σ -significance.

Example	Selected θ	p -value (significance)	Monte Carlo error (significance interval)
Example 1	(θ_1, θ_2) (175, 38)	1.092×10^{-26} (10.629 σ)	9.272×10^{-28} [10.621 σ , 10.637 σ]
Example 2	(μ, σ) (5.041, 0.5)	0.034 (1.801 σ)	0.012 [1.663 σ , 1.988 σ]
$H_0 : \eta = 0$ versus $H_0 : \eta > 0$	(γ, τ) (6.510, 70)	0.596 (0.00 σ)	0.093
Example 2	(θ, α) (31.265, 1)	5.663×10^{-30} (11.313 σ)	7.881×10^{-31} [11.301 σ , 11.326 σ]
$H_0 : \eta = 1$ versus $H_0 : \eta < 1$			
Example 3			

NOTE: The last column refers to the Monte Carlo errors associated with the estimates of $E[\phi(\mathcal{A}_{c_k})]$ that are used in (17) to obtain (24) as described in the Section 2.5.

In principle, the choice of c_1, \dots, c_D should also take into account the possibility that the ECs computed at different thresholds c_k , $\phi(\mathcal{A}_{c_k})$, may be positively correlated leading to inflation of the variance of the estimators of $E[\phi(\mathcal{A}_{c_k})]$. However, since we are interested in the limit as $c \rightarrow \infty$ and the Monte Carlo error associated with (24) become extremely small as c increases, such correlation may be of little concern. This may be true even when, as in Figure 6, the quantities $\phi(\mathcal{A}_{c_k})$ have been computed on the same set of Monte Carlo simulations for each c_k considered.

3.3. Data Analysis

We calculated the TOHM p -value in (24) for the case studies introduced in Section 3.1. The results are summarized in Table 2; the codes used to generate these analyses are collected in the file `Analyses_Tab1.R` downloadable among the Supplementary Materials. In addition to the p -values, we report the respective σ -significance, a quantity typically used in physics to quantify the statistical evidence in support of new discoveries, that is

$$\#\sigma = \Phi^{-1}(1 - p\text{-value}), \quad (25)$$

where Φ is the standard normal cumulative function.

In Example 1, we performed $R = 2821$ tests over our circular search region centered at (195 RA, 28 DEC). In our realistic simulation, the true dark matter emission was located at (174.952 RA, 37.986 DEC) and the LRT-process used in TOHM achieves its maximum at $\theta = (175 \text{ RA}, 38 \text{ DEC})$ with about 10σ significance. Notice that our original dataset includes 51,098 background events and only 39 dark matter events; hence the procedure appears to be particularly powerful even in presence of a low signal-to-noise ratio. The location at which the maximum LRT statistics has been observed is plotted as a white circle in the upper panel of Figure 4.

In Example 2, we set $R = 2500$ when testing (2) and the gamma model is rejected at a 0.05 significance level by the THOM p -value. Whereas, when testing (4), the log-normal model cannot be rejected; the resulting p -value is 0.596. Thus, the log-normal model is selected for the maize seeds strength data, and the LRT-process achieves its maximum at $\mu = 5.004$ and $\sigma = 0.633$. The log-normal fitted model is plotted in the bottom left panel of Figure 4 as a red solid line.

Finally in Example 3, testing (7) $R = 150$ times, (24) provides strong evidence ($\sim 11\sigma$) in favor of a linear trend ($\alpha = 1$)

with a break point at $\theta = 31.265$. Hence, we expect the risk of giving birth to a child with down syndrome to increase when the mother is 31 years old or older. The model selected is displayed as a red solid line in the bottom right panel of Figure 4, with the break-point indicated by a red triangle. For the sake of comparison, we also plot the fitted model when allowing a quadratic trend ($\alpha = 2$) with a break point chosen at $\hat{\theta} = 20$.

4. Discussion

In this article, we propose a novel computational method to perform TOHM in the multidimensional setting. The resulting inferential tool generalizes classical inferential methods, such as the LRT, beyond standard regularity conditions including non-identifiability of multidimensional parameters and non-nestedness of the models under comparison. From a more practical perspective, the procedure proposed provides a computationally efficient solution to the bump hunting problem in multiple dimensions, and implicitly introduces a Type I error correction for dependent tests. It also simplifies the estimation of the so called Lipschitz–Killing curvatures involved in the computation of the TOHM p -value on the basis of Taylor and Worsley (2008).

Despite its simplicity and efficiency in computation, the main limitation of TOHM is that it requires the specification of a parametric form for the alternative model. In the context of signal identification, for instance, this implies that the researcher can specify the density function of the events associated to the signal (e.g., a Gaussian bump). In situations where this cannot be done, one possibility is to refer to nonparametric inferential methods (e.g., Chen, Genovese, and Wasserman 2016; Mukhopadhyay 2017; Algeri 2019a). More work is needed to extend TOHM to discrete regions Θ and provide a formal justification of its validity in non-regular setting, such as the one in Example 3.

It is important to note that, in the context of multiple hypothesis testing and large-scale inference, TOHM allows us to reduce the dimensionality of the tests being conducted from R to one by exploring the topology of the random field associated with the test statistics of interest. From this perspective, TOHM may offer a path forward to solve the long-standing problem of identifying an unknown number of signals, in one or multiple dimensions.

Appendix A: Regularity Conditions and Proofs

A.1. Regularity Conditions A0–A5

Ghosh and Sen (1985) show that when testing (2) for (3) with $\mathbf{y} \in \mathbb{R}$ and $\Theta \subset \mathbb{R}$, the LRT converges to a $\bar{\chi}_{01}^2$ process under suitable assumptions. In our examples, the nuisance parameter θ is allowed to be multidimensional, that is, $\Theta \subset \mathbb{R}^D$, $D \geq 1$, and the data can be multivariate, that is, $\mathbf{y} \in \mathbb{R}^q$, $q \geq 1$; therefore, we restate the regularity conditions of Ghosh and Sen (1985) accordingly below.

(A0) The mixture model in (3) is strongly identifiable, that is

$$\begin{aligned} \text{if } \eta \in (0, 1) \text{ and } h(\mathbf{y}, \eta, \boldsymbol{\gamma}, \boldsymbol{\theta}) &= h(\mathbf{y}, \eta', \boldsymbol{\gamma}', \boldsymbol{\theta}') \\ \Rightarrow (\eta, \boldsymbol{\gamma}, \boldsymbol{\theta}) &= (\eta', \boldsymbol{\gamma}', \boldsymbol{\theta}'). \end{aligned}$$

(A1) For each θ fixed, let $S(\theta) = \nabla \log h(\mathbf{y}, \eta, \boldsymbol{\gamma}, \boldsymbol{\theta})$ be the score vector and $S_j(\theta)$ its element j . Denote them with $S^0(\theta)$ and $S_j^0(\theta)$ when evaluated at the true values of $(\eta, \boldsymbol{\gamma})$ under H_0 , that is, $(0, \boldsymbol{\gamma}_0)$. Similarly, for each θ fixed, let $H(\theta)$ be the Hessian matrix of $\log h(\mathbf{y}, \eta, \boldsymbol{\gamma}, \boldsymbol{\theta})$ and $H_{jk}(\theta)$ its element in position (j, k) . Denote them with $H^0(\theta)$ and $H_{jk}^0(\theta)$ when evaluated at $(\eta, \boldsymbol{\gamma}) = (0, \boldsymbol{\gamma}_0)$. We require the following.

- (i) Γ is an open interval in \mathbb{R}^p and Θ is a compact subset of \mathbb{R}^D .
- (ii) $h(\mathbf{y}, \eta, \boldsymbol{\gamma}, \boldsymbol{\theta})$ is continuous in $(\eta, \boldsymbol{\gamma}, \boldsymbol{\theta})$ and twice continuously differentiable with respect to $(\eta, \boldsymbol{\gamma})$.
- (iii) $E[S_j^0(\theta)] = 0$ for all $j = 1, \dots, p+1$ and $\theta \in \Theta$.
- (iv) $E[H_{jk}^0(\theta)] = -E[S_j^0(\theta)S_k^0(\theta)]$ for all $j, k = 1, \dots, p+1$ and $\theta \in \Theta$.
- (v) For $j, k = 1, \dots, p+1$,

$$\lim_{\delta \rightarrow 0} E \left[\sup_{\|(\eta, \boldsymbol{\gamma}, \boldsymbol{\theta}) - (0, \boldsymbol{\gamma}_0, \boldsymbol{\theta})\| < \delta} |H_{jk}(\boldsymbol{\theta}) - H_{jk}^0(\boldsymbol{\theta})| \right] = 0,$$

where all the expectations above and those to follow are taken with respect to $f(\mathbf{y}, \boldsymbol{\gamma}_0)$.

(A2) There exists a compact neighborhood \mathcal{N} of $(\eta, \boldsymbol{\gamma})$ such that $E[\psi(\mathbf{y}, \boldsymbol{\theta})] < 0$, where

$$\psi(\mathbf{y}, \boldsymbol{\theta}) = \sup_{(\eta, \boldsymbol{\gamma}) \in [0, 1] \times \mathcal{N}^c} \log \frac{h(\mathbf{y}, \eta, \boldsymbol{\gamma}, \boldsymbol{\theta})}{f(\mathbf{y}, \boldsymbol{\gamma})}.$$

Further $\psi(\mathbf{y}, \boldsymbol{\theta})$ is continuous on Θ and there exist a function $w(\mathbf{y})$ such that $|\psi(\mathbf{y}, \boldsymbol{\theta})| \leq w(\mathbf{y})$ and $E[w(\mathbf{y})] \leq \infty$ for all $\theta \in \Theta$.

(A3) For each $(\eta, \boldsymbol{\gamma}) \neq (0, \boldsymbol{\gamma}_0)$ there exists an open ball with center at $(\eta, \boldsymbol{\gamma})$ and radius δ_0 , namely $B(\delta_0)$, such that

$$|\psi_B(\mathbf{y}, \boldsymbol{\theta})| \leq w(\mathbf{y}) \quad \text{for all } \theta \in \Theta,$$

where

$$\begin{aligned} \psi_B(\mathbf{y}, \boldsymbol{\theta}) &= \sup_{(\eta', \boldsymbol{\gamma}') \in B(\delta_0) \cap [0, 1] \times \Gamma} \log \frac{h(\mathbf{y}, \eta', \boldsymbol{\gamma}', \boldsymbol{\theta})}{f(\mathbf{y}, \boldsymbol{\gamma}_0)} \quad \text{and} \\ &\times E[w(\mathbf{y})] \leq \infty. \end{aligned}$$

(A4) $I(\theta) = E[-H^0(\theta)]$ is continuous in θ and positive definite uniformly over Θ .

(A5) $E \left| \frac{g(\mathbf{y}, \boldsymbol{\theta})}{f(\mathbf{y}, \boldsymbol{\gamma}_0)} - \frac{g(\mathbf{y}, \boldsymbol{\theta}^\dagger)}{f(\mathbf{y}, \boldsymbol{\gamma}_0)} \right|^\xi \leq K \|\theta - \theta^\dagger\|^{1+\lambda}$ for some $\xi, \lambda > 0$, and $K \geq 0$.

Proof of Proposition 2.1. Under conditions A0 and A1, it follows from Ghosh and Sen (1985, eqs. (2.7) and (2.8)) that the random field $\{W_n(\theta)\}$ can be written as

$$\begin{aligned} \{W_n(\theta)\} &= \{R_n(\theta)\} \quad \text{over the set } \Theta_{0n} = \{\theta : T_n(\theta) < 0\} \\ \{W_n(\theta)\} &= \{T_n(\theta)\} + \{R_n(\theta)\} \quad \text{over the set } \Theta_{1n} = \{\theta : T_n(\theta) \geq 0\}, \end{aligned} \quad (\text{A.1})$$

where $\{R_n(\theta)\}$ is a reminder term such that, if A2 and A3 hold, $\{R_n(\theta)\} = o_p(1)$, uniformly in θ . Whereas, $\{T_n(\theta)\}$ is a random field such that, if A4 and A5 hold and under H_0 , it converges weakly to a Gaussian random field $\{Z(\theta)\}$ with mean zero, unit variance and covariance function depending on θ . \square

Proof of Proposition 2.2. Let ϕ_{10} and ϕ_{20} be the true values of ϕ_1 and ϕ_2 in (6) when H_0 in (7) is true. Denote with $U_n^*(\theta)$ the normalized score function of (6) for $\theta = (\theta, \alpha)$ fixed. Under H_0 , $U_n^*(\theta)$ can be specified by

$$U_n^*(\theta|H_0) = \sum_{i=1}^n \frac{Z_i}{\sigma} \sqrt{m_i \pi_{0i} (1 - \pi_{0i})} (x_i - \theta)^\alpha \mathbb{1}_{\{x_i \geq \theta\}}, \quad (\text{A.2})$$

where $\sigma = \sum_{i=1}^n m_i \pi_{0i} (1 - \pi_{0i}) (x_i - \theta)^{2\alpha} \mathbb{1}_{\{x_i \geq \theta\}}$, $Z_i = \frac{Y_i - m_i \pi_{0i}}{\sqrt{m_i \pi_{0i} (1 - \pi_{0i})}}$, $Y_i \sim \text{Binomial}(m_i, \pi_i)$, $\pi_i = [1 + \exp\{-\phi_1 + \phi_2 x_i + \xi(x_i - \theta)^\alpha \mathbb{1}_{\{x_i \geq \theta\}}\}]^{-1}$ and such that, under H_0 , $\pi_i = \pi_{0i} = [1 + \exp\{-\phi_{10} + \phi_{20} x_i\}]^{-1}$. Under the Cramer's classical conditions (Cramer 1946, p. 500), $U_n^*(\theta|H_0)$ is asymptotically normally distributed with mean zero and variance one. Since ϕ_{10} and ϕ_{20} are unknown, we can estimate them by the respective MLEs $\hat{\phi}_{10}$ and $\hat{\phi}_{20}$. The latter are \sqrt{n} -consistent estimators of ϕ_{10} and ϕ_{20} , therefore, when substituting $\hat{\pi}_{0i} = [1 + \exp\{-\hat{\phi}_{10} + \hat{\phi}_{20} x_i\}]^{-1}$ to π_{0i} in (A.2), the Gaussian asymptotic distribution of $U_n^*(\theta|H_0)$ is preserved. By De Moivre–Laplace theorem, under H_0 , each Z_i follows a standard normal distribution as $m_i \rightarrow \infty$. Therefore, for each θ fixed, $U_n^*(\theta|H_0)$ is a linear function of the Z_i ; hence the joint distribution of $\{U_n^*(\theta|H_0)\}$ is asymptotically Gaussian as $m_i \rightarrow \infty$ for all $i = 1, \dots, n$. Finally, by virtue of the equivalence between $U_n^*(\theta)$ and $W_n(\theta)$ (Moran 1970; Davies 1977), $\{W_n(\theta)\}$ is also asymptotically distributed as a Gaussian random field under H_0 . \square

Appendix B: EC Densities for Gaussian, χ^2 , and $\bar{\chi}_{01}^2$ Random Fields

B.1. Gaussian Case

If $\{W(\theta)\}$ is such that $W(\theta) \sim N(0, 1)$ for all θ , the EC densities $\rho_d(c)$, $d = 0, \dots, 5$ are given by

$$\begin{aligned} \rho_0(c) &= 1 - \Phi(c), \quad \rho_1(c) = \frac{e^{-\frac{c^2}{2}}}{2\pi}, \quad \rho_2(c) = \frac{e^{-\frac{c^2}{2}}}{(2\pi)^{3/2}}, \\ \rho_3(c) &= \frac{(c^2 - 1)e^{-\frac{c^2}{2}}}{(2\pi)^2}, \quad \rho_4(c) = \frac{(c^3 - 3c)e^{-\frac{c^2}{2}}}{(2\pi)^{5/2}}, \\ \rho_5(c) &= \frac{(c^4 - 4c^2 + 3)e^{-\frac{c^2}{2}}}{(2\pi)^3}, \end{aligned}$$

where $\Phi(\cdot)$ is the cumulative function of a standard normal.

B.2. χ_s^2 Case

If $\{W(\theta)\}$ is such that $W(\theta) \sim \chi_s^2$ for all θ , the EC densities $\rho_d(c)$, $d = 0, \dots, 3$ are given by

$$\begin{aligned} \rho_0(c) &= 1 - F_\chi(c), \quad \rho_1(c) = \frac{c^{\frac{s-1}{2}}}{\Gamma(\frac{s}{2})} \sqrt{\frac{2}{\pi}} e^{-\frac{c}{2}}, \\ \rho_2(c) &= \left(\frac{c}{2}\right)^{\frac{s}{2}-1} \frac{e^{-\frac{c}{2}}}{2\pi} [c - (s-1) \mathbb{1}_{\{s \geq 2\}}], \end{aligned}$$

$$\rho_3(c) = \frac{c^{\frac{s-3}{2}} e^{-\frac{c}{2}}}{(2\pi)^{3/2} \Gamma(\frac{s}{2}) 2^{\frac{s-2}{2}}} \left[(s-1)(s-2) \mathbb{1}_{\{s \geq 3\}} - 2(s-1)c \mathbb{1}_{\{s \geq 2\}} + (c^2 - c) \mathbb{1}_{\{s \geq 1\}} \right],$$

where $F_\chi(\cdot)$ is the cumulative function of a χ_s^2 and $\mathbb{1}_{\{\cdot\}}$ is the indicator function.

B.3. $\bar{\chi}_{01}^2$ Case

From Taylor and Worsley (2013), it follows that the EC densities of a $\bar{\chi}_{01}^2$ random field are given by the sum of the EC densities of a χ_0^2 random field (i.e., a random field which is zero everywhere) and those of a χ_1^2 random field, each multiplied by the respective weight, that is, 0.5. Consequently, when $\Theta \subset \mathbb{R}^2$ as in Examples 1 and 2, (15) specifies as

$$E[\phi(\mathcal{A}_c)] = \frac{c^{\frac{1}{2}} e^{-\frac{c}{2}}}{(2\pi)^{\frac{3}{2}}} \mathcal{L}_2(\Theta) + \frac{e^{-\frac{c}{2}}}{2\pi} \mathcal{L}_1(\Theta) + \frac{P(\chi_1^2 > c)}{2} \mathcal{L}_0(\Theta), \quad (\text{B.1})$$

where the functions of c multiplying $\mathcal{L}_0(\Theta), \dots, \mathcal{L}_2(\Theta)$ are the EC densities of a two-dimensional χ_1^2 random field divided by 2. Because the EC densities of a two-dimensional χ_0^2 random field evaluated at $c > 0$ are all zero, they do not contribute in (B.1).

See Adler and Taylor (2007, p. 426) for higher order EC densities.

Supplementary Materials

The folder *TOHM_Codes_and_files* collects the data used in Example 1 and the codes used for the analyses in Figure 4 and Table 2. The interested reader is directed to the supplementary file *Codes_description.pdf* for a more detailed description of all the codes and files available. In the file *Supp_JCGS.pdf* we assess the validity of assumptions A0–A5 for Examples 1 and 2.

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