MODELING THE VARIABILITY OF RANKINGS

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For better or for worse, rankings of institutions, such as universities, schools and hospitals, play an important role today in conveying information about relative performance. They inform policy decisions and budgets, and are often reported in the media. While overall rankings can vary markedly over relatively short time periods, it is not unusual to find that the ranks of a small number of “highly performing” institutions remain fixed, even when the data on which the rankings are based are extensively revised, and even when a large number of new institutions are added to the competition. In the present paper, we endeavor to model this phenomenon. In particular, we interpret as a random variable the value of the attribute on which the ranking should ideally be based. More precisely, if \( p \) items are to be ranked then the true, but unobserved, attributes are taken to be values of \( p \) independent and identically distributed variates. However, each attribute value is observed only with noise, and via a sample of size roughly equal to \( n \), say. These noisy approximations to the true attributes are the quantities that are actually ranked. We show that, if the distribution of the true attributes is light-tailed (e.g., normal or exponential) then the number of institutions whose ranking is correct, even after recalculation using new data and even after many new institutions are added, is essentially fixed. Formally, \( p \) is taken to be of order \( n^C \) for any fixed \( C > 0 \), and the number of institutions whose ranking is reliable depends very little on \( p \). On the other hand, cases where the number of reliable rankings increases significantly when new institutions are added are those for which the distribution of the true attributes is relatively heavy-tailed, for example, with tails that decay like \( x^{-\alpha} \) for some \( \alpha > 0 \). These properties and others are explored analytically, under general conditions. A numerical study links the results to outcomes for real-data problems.

1. Introduction.

There are many contemporary settings in which ranking plays an important role. For example, universities, schools and hospitals are regularly ranked in a variety of contexts, the results of which typically generate interest and can often drive policy decisions. In many of these situations, a given ranking can carry a high degree of uncertainty, with this effect particularly pronounced in high-dimensional cases; that is, where there are very many populations or institutions to be ranked.
Despite this, one feature of many rankings reported over time is that the ordering at the extreme top or bottom remains relatively invariant. For example, in the THE-QS university rankings,\(^1\) Harvard University has ranked first for each of the years 2005–2008, while New York University’s rankings are 56, 43, 49 and 40. If we believe that the observed data used for ranking are measures of true underlying values, distorted by noise, then we can reinterpret this behavior as a tendency to obtain correct rankings at extremes, but not otherwise. It is this phenomenon that we explore in this paper, using both theoretical and numerical arguments.

Intuitively, this behavior has a natural explanation. Those scores at the extreme of a range are more likely to be sufficiently “spaced out” to overcome the problems of data noise, whereas less extreme scores are likely to be bunched more closely together. We introduce models that describe this behavior and explore their properties. Related to this, it turns out that one important consideration for correct ranking at the extremes is whether the possible scores used for ranking have infinite support but nevertheless have light tails. If this is the case and the tail of the distribution of the underlying scores is smooth, we can expect accurate ranking of the top portion of the institutions, even when dimension is very large. Moreover, even when the support is bounded, there remains potential for correct ranking at extremes, although now there is greater likelihood that the ranking will change if new institutions are added. Such results have a variety of practical implications; we briefly present two of these here, with more detail provided in the numerical section.

**Example 1 (University rankings).** Suppose we attempt to rank universities and other research institutions by counting how many papers their faculty members publish in *Nature*\(^2\) each year. This is a high-dimensional example due to the large number of institutions competing to be published. Figure 1 shows the ranking of the top 50 institutions on this measure. The institutions are aligned along the horizontal axis, with the each dot denoting the point estimate of the rank and the vertical line a corresponding estimated 90% prediction interval. The four plots show how the confidence intervals change as we increase the number of years, \(n\), of data used for the ranking.

The two main observations are that the prediction intervals are widest when a smaller number of years are considered, and that the prediction intervals for the highest ranked universities are the smallest. In fact, the intervals are small enough in the extremes to give us genuine confidence in that aspect of the ranking. Even when \(n = 1\), we can be reasonably sure that the top ranked institution (Harvard University) is in fact ranked correctly. When \(n = 15\), the top four universities are known with a high degree of certainty, and the next set of ten or so is fairly stable.

\(^1\)www.topuniversities.com.
\(^2\)www.nature.com/nature/index.html.
too. Thus, it is possible to have correctness in the upper extreme of this ranking, even when the lower ranks remain highly variable. In the present paper, we model this phenomenon by addressing the underlying stochastic properties of the institutions; the data provide only a noisy measure of this random process, and we assess the impact of the noise on the ranking.

**EXAMPLE 2 (Microarray data).** We take the colon microarray data first analysed by Alon et al. (1999). It consists of 62 observations in total, each of which indicates either a normal colon or a tumor. For each observation, there are also expression levels for $p = 2000$ genes. It is of interest to determine which genes are most closely related to the response, so that they can be investigated further. This of course amounts to a ranking and we are interested in stability at the extreme, since we seek only a small number of genes. Here, the genes are ranked based on the Mann–Whitney U test statistic, which is a nonparametric assessment of the difference between the two distributions.

Figure 2 plots the top 30 genes, ranked by the lower tail of an estimated 90% prediction interval, rather than the point estimate of the rank. In this situation, we cannot authoritatively conclude that any of the top genes are ranked exactly correctly, but the top four genes appear much more stable than the others. This stability is highly important; if the length of all prediction intervals were roughly the same as the average length (1400 genes), then there would be little hope of discovering useful genes from such datasets.
There is a literature on the bootstrap in connection with rankings. See Goldstein and Spiegelhalter (1996), who discuss bootstrap methods for constructing prediction intervals for rankings; Langford and Leyland (1996), who address bootstrap methods for ranking the performance of doctors; Cesário and Barreto (2003), Hui, Modarres and Zheng (2005) and Taconeli and Barreto (2005), who take up the problem of bootstrap methods for ranked set sampling; Mukherjee et al. (2003), who develop methods for gene ranking using bootstrapped $p$-values; and Xie, Singh and Zhang (2009) and Hall and Miller (2009), who focus on consistent bootstrap methods for assessing rankings. More generally, there is a vast literature on ranking problems in statistics, and we cite here only the more relevant items since 2000. Joe (2000, 2001) discusses ranking problems in connection with random utility models, and points to connections to multivariate extreme value theory. Murphy and Martin (2003) develop mixture-based models for rankings. Mease (2003) and Barker et al. (2005) treat methods for ranking football players. McHale and Scarf (2005) study the problem of ranking immunisation coverage in U.S. states. Brijs et al. (2006, 2007) introduce Bayesian models for the ranking of hazardous road sites, with the aim of better scheduling road safety policies. Chen, Stansy and Wolfe (2006) discuss ranking accuracy in ranked-set sampling methods, and Opgen-Rhein and Strimmer (2007) examine the accuracy of gene rankings in high-dimensional problems involving genomic data. Nordberg (2006) addresses the reliability of performance rankings. Corain and Salmaso (2007) and Quevedo, Bahamonde and Luaces (2007) discuss ways of constructing rankings.

Section 2 describes our model for the ranking problem, and discusses the main properties of this framework. The formal theoretical results which underpin the discussion in Section 2 are given in Section 3. Section 4 presents simulated and real-data numerical work, including details on the examples presented above. Technical proofs are deferred to Section 5.
2. Model. We consider a set of underlying parameters $\theta_1, \ldots, \theta_p$ corresponding to the objects to be ranked, hereafter referred to as items. The error in the estimation is controlled by the number of observed data points, $n$. In our analysis, we take $p = p(n)$ to diverge with $n$ as the latter increases. An obvious difficulty here is in establishing where the newly added items should fit into the ranking. A natural solution is to take the $\theta_j$’s to be randomly generated from some distribution function. In the setup below, we interpret the $\Theta_j$’s as values of means; see the end of this section for generalizations.

Let $\Theta_1, \ldots, \Theta_p$ denote independent and identically distributed random variables, and write

$$\Theta_1(1) \leq \cdots \leq \Theta_1(p)$$

for their ordered values. There exists a permutation $R = (R_1, \ldots, R_p)$ of $(1, \ldots, p)$ such that $\Theta_1(j) = \Theta_{R_j}$ for $1 \leq j \leq p$. If the common distribution of the $\Theta_j$’s is continuous, then the inequalities in (2.1) are all strict and the permutation is unique.

We typically do not observe the $\Theta_j$’s directly, only in terms of noisy approximations which can be modelled as follows. Let $Q_i = (Q_{i1}, \ldots, Q_{ip})$ denote independent and identically distributed random $p$-vectors with finite variance and zero mean, independent also of $\Theta = (\Theta_1, \ldots, \Theta_p)$. Suppose we observe

$$X_i = (X_{i1}, \ldots, X_{ip}) = Q_i + \Theta$$

for $1 \leq i \leq n$. The mean vector

$$\bar{X} = (\bar{X}_1, \ldots, \bar{X}_p) = \frac{1}{n} \sum_{i=1}^{n} X_i = \bar{Q} + \Theta$$

is an empirical approximation to $\Theta$. (Here, $\bar{Q} = n^{-1} \sum_i Q_i$ equals the mean of the $p$-vectors $Q_i$.) The components of $\bar{X}$ can also be ranked, as

$$\bar{X}_1(1) \leq \cdots \leq \bar{X}_1(p),$$

and there is a permutation $\hat{R}_1, \ldots, \hat{R}_p$ of $1, \ldots, p$ such that $\bar{X}_1(j) = X_{\hat{R}_j}$ for each $j$. If the common distribution of the $\Theta_j$’s is continuous then, regardless of the distribution of the components of $Q_i$, the inequalities in (2.4) are strict with probability 1.

The permutation $\hat{R} = (\hat{R}_1, \ldots, \hat{R}_p)$ serves as an approximation to $R$, and we wish to determine the accuracy of that approximation. In particular, for what values of $j_0 = j_0(n, p)$, and for what relationships between $n$ and $p$, is it true that

$$P(\hat{R}_j = R_j \text{ for } 1 \leq j \leq j_0) \to 1$$

as $n$ and $p$ diverge? That is, how deeply into the ranking can we go before the connection between the true ranking and its empirical form is seriously degraded by noise?

The answer to this question depends to some degree on the extent of dependence among the components of each $Q_i$. To elucidate this point, let us consider the case
where all the components of $Q_i$ are identical; this is an extreme case of strong dependence. Then the components of $\bar{Q}$ are also identical. Clearly, in this setting $\bar{R}_j = R_j$ for each $j$, and so (2.5) holds in a trivial and degenerate fashion. Other strongly dependent cases, although not as clear-cut as this one, can also be shown to be ones where $\bar{R}_j = R_j$ with high probability for many values of $j$.

The case which is most difficult, that is, where the strongest conditions are needed to ensure that (2.5) holds, occurs when the components of $Q_i$ are independent. To emphasize this point we give sufficient conditions for (2.5), and show that when the components of each $Q_i$ are independent, those conditions are also necessary. Our arguments can be modified to show that the conditions continue to be necessary under sufficiently weak dependence, for example if the components are $m$-dependent where $m = m(n)$ diverges sufficiently slowly as $n$ increases.

The assumptions under which (2.5) holds are determined mainly by the lower tail of the common distribution of the $\Theta_j$’s. If that distribution has an exponentially light left-hand tail, for example, if the tail is like that of a normal distribution, then a sufficient condition for (2.5) is that $j_0$ should increase at a strictly slower rate than $n^{1/4} \left( \log n \right)^c$, where the constant $c$, which can be either positive or negative, depends on the rate of decay of the exponential lower tail of the distribution of $\Theta$. For example, $c = 0$ if the distribution decays like $e^{-|x|}$ in the lower tail, and $c = -\frac{1}{4}$ if it is normal. As indicated in the previous paragraph, the condition $j_0 = o\left\{ n^{1/4} \left( \log n \right)^c \right\}$ is also necessary for (2.5) if the components of the $Q_i$’s are independent.

These results have several interesting aspects, including: (a) the exponent $\frac{1}{4}$ in the condition $j_0 = o\left\{ n^{1/4} \left( \log n \right)^c \right\}$ does not change among different types of distribution with exponential tails; (b) the exponent is quite small, implying that the empirical rankings $\hat{R}_j$ quite quickly become unreliable as predictors of the true rankings $R_j$; and (c) the critical condition $j_0 = o\left\{ n^{1/4} \left( \log n \right)^c \right\}$ does not depend on the value of $p$. (We assume that $p$ diverges at no faster than a polynomial rate in $n$, but we impose no upper bound on the degree of that polynomial.)

The condition on $j_0$ such that (2.5) holds changes in important ways if the lower tail of the distribution of the $\Theta_j$’s decays relatively slowly, for example, at the polynomial rate $x^{-\alpha}$ as $x \to \infty$. Examples of this type include Pareto, nonnormal stable and Student’s $t$ distributions, and more generally, distributions with regularly varying tails. Here a sufficient condition for (2.5) to hold is $j_0 = o\left\{ (n^{\alpha/2} p)^{1/(2\alpha+1)} \right\}$, and this assumption is necessary if the components of the $Q_i$’s are independent. In this setting, unlike the exponential case, the value of dimension, $p$, plays a major role in addition to the sample size, $n$, in determining the number of reliable rankings.

In practical terms, a major way in which this heavy-tailed case differs from the light-tailed setting considered earlier is that if a polynomially large number of new items are added to the competition in the heavy-tailed case, and all items are reranked, the results will change significantly and the number of correct rankings
will also alter substantially. By way of contrast, if a polynomially large number of new items are added in the light-tailed, or exponential, case then there will again be many changes to the rankings, but now there will be relatively few changes to the number of items that are correctly ranked.

The exponential case can be regarded as the limit, as $\alpha \to \infty$, of the polynomial case. More generally, note that as the left-hand tail of the common distribution of the $\Theta_j$’s becomes heavier, the value of $j_0$ can be larger before (2.5) fails. That is, if the distribution of the $\Theta_j$’s has a heavier left-hand tail then the empirical rankings $\hat{R}_j$ approximate the true rankings $R_j$ for a greater number of values of $j$, before they degenerate into noise.

The analysis above has focused on cases where the ranks of the $X_j$’s are estimated by ranking empirical means of noisy observations of those quantities; see (2.4). However, similar results are obtained if we rank other measures of location. Such a measure need only satisfy moderate deviation properties similar to (5.3) and (5.4) in the proof of Theorem 1. Thus, the results are applicable to a wide range of ranking contexts. For example, $L_q$ location estimators for general $q \geq 1$ enjoy moderate deviation properties under appropriate assumptions. Therefore, if we take the variables $Q_{ij}$ to have zero median, rather than zero mean, and continue to define $X_i$ by (2.2) but replace the ranking in (2.4) by a ranking of medians, then the results above and those in Section 3 continue to hold, modulo changes to the regularity conditions. Other suitable measures include the Mann–Whitney test used in the genomic example, quantiles and some correlation-based measures.

The model suggested by (2.2), where data on $\Theta$ arise in the form of $p$-vectors $X_1, \ldots, X_n$, is attractive in a number of high-dimensional settings, for example, genomics. There, the $j$th component $X_{ij}$ of $X_i$ would typically represent the expression level of the $j$th gene of the $i$th individual in a sample. However, in other cases the means $\bar{X}_1, \ldots, \bar{X}_p$ at (2.3), or medians or other location estimators, might be computed from quite different datasets, one for each component index $j$. Moreover, those datasets might be of different sizes, $n_1, \ldots, n_p$ say, and then the argument that they arise naturally in the form of vectors would be inappropriate. This can happen when data are used to rank items, for example schools where the ranking is based on individual student performance. The conclusions discussed earlier in this section, and the theoretical properties developed in Section 3 below, continue to apply in this case provided there is an “average” value, $n$ say, of the $n_j$’s which represents all of them, in the sense that

$$n = O\left( \min_{1 \leq j \leq p} n_j \right) \quad \text{and} \quad \max_{1 \leq j \leq p} n_j = O(n)$$

as $n$ diverges. Additionally, in such cases it is often realistic to make the assumption that the corresponding centred means (or medians, etc.) $\bar{Q}_j = n^{-1} \sum_i Q_{ij}$ are stochastically independent of one another, and so the particular results that are valid in this case are immediately available.
The distribution of the $\Theta_j$’s has been taken to be continuous. This is usually appropriate although there can be contexts in which the distribution is discrete. Note that assumption of discreteness of the $\Theta_j$’s is different from that of discreteness of the observations $X_{ij}$. In such cases, the analysis still holds, except that allowance must be made for ties (any reordering of tied $\Theta_j$’s is still “correct”), and the tail density assumptions should be characterized in integral form.

The model has been set up so that it focuses on the populations with lowest parameters $\Theta_j$. Obviously, similar arguments apply to the largest parameters too, so the results are applicable to both the most highly and lowly ranked populations.

3. Theoretical properties. For the most part, we shall assume one of two types of lower tail for the common distribution function, $F$, of the random variables $\Theta_j$: either it decreases exponentially fast, in which case we suppose that $F(-x) \asymp x^\beta \exp(-C_0 x^\alpha)$ as $x \to \infty$, where $\alpha > 0$ and $-\infty < \beta < \infty$; or it decreases polynomially fast, in which case $F(-x) \asymp x^{-\alpha}$ as $x \to \infty$, where $C_0, \alpha > 0$. [The notation $f(x) \asymp g(x)$, for positive functions $f$ and $g$, will be taken to mean that $f(x)/g(x)$ is bounded away from zero and infinity as $x \to \infty$.] The former case covers distributions such as the normal, exponential and Subbotin; the latter, distributions such as the Pareto, Student’s $t$ and nonnormal stable laws (e.g., the Cauchy).

It is convenient to impose the shape constraints on the densities, which we assume to exist in the lower tail, rather than on the distribution functions. Therefore, we assume that one of the following two conditions hold as $x \to \infty$:

\[(d/dx)F(-x) \asymp (d/dx)x^\beta \exp(-C_0 x^\alpha), \quad (3.1)\]
\[(d/dx)F(-x) \asymp (d/dx)x^{-\alpha}. \quad (3.2)\]

In both (3.1) and (3.2), $\alpha$ must be strictly positive, but $\beta$ in (3.1) can be any real number. The constant $C_0$ in (3.1) must be positive. We assume too that

\[\text{for fixed constants } C_1, \ldots, C_5 > 0, \text{ where } C_2 > 2(C_1 + 1) \text{ and } C_4 < C_5, \quad \text{(3.3)}\]

\[p = O(n^{C_1}) \text{ as } n \to \infty, \text{ and, for each } j \geq 1, E|Q_j|^C_2 \leq C_3, E(Q_j) = 0, \text{ and } E(Q_j^2) \in [C_4, C_5].\]

Recall from Section 1 that we wish to examine the probability that the true ranks $R_j$, and their estimators $\hat{R}_j$, are identical over the range $1 \leq j \leq j_0$. We consider both $j_0$ and $p$ to be functions of $n$, so that the main dependent variable can be considered to be $n$. With this interpretation, define

\[\nu_{\exp} = \nu_{\exp}(n) = n^{1/4} (\log n)^{(1/\alpha) - 1/2}, \quad (3.4)\]
\[\nu_{\pol} = \nu_{\pol}(n) = (n^{\alpha/2} p)^{1/(2\alpha + 1)},\]

where the subscripts denote “exponential” and “polynomial,” respectively, and refer to the respective cases represented by (3.1) and (3.2). In the theorem below, we impose the additional condition that, for some $\epsilon > 0$,

\[n = O(p^{4-\epsilon}). \quad (3.5)\]
This restricts our attention to problems that are genuinely high dimensional, in the sense that, with probability converging to 1, not all the rankings are correct. Cases where \( p \) diverges sufficiently slowly as a function of \( n \) are easier and will generally permit all ranks to be correctly determined with high probability. Assumption (3.5) is also very close, in both the exponential and polynomial cases, to the basic condition \( j_0 \leq p \), as can be seen via a little analysis starting from (3.6) and (3.7) in the respective cases; yet, at the same time, (3.5) is suitable to both cases, and so helps to unify our account of their properties. Note too that (3.5) implies that, in both the exponential and polynomial cases, \( v_{\exp} = O(p^{1-\delta}) \) and \( v_{\text{pol}} = O(p^{1-\delta}) \) for some \( \delta > 0 \).

**Theorem 1.** Assume (3.3), (3.5) and that either (a) (3.1) or (b) (3.2) holds. In case (a), if

\[
(3.6) \quad j_0 = o(v_{\exp})
\]

as \( n \to \infty \) then (2.5) holds. Conversely, when the components of the vectors \( Q_i \) are independent, (3.6) is necessary for (2.5). In case (b), if

\[
(3.7) \quad j_0 = o(v_{\text{pol}}),
\]

then (2.5) holds. Conversely, when the components of the vectors \( Q_i \) are independent, (3.7) is necessary for (2.5).

It can be deduced from Theorem 1 that when a new item (e.g., an institution) enters the competition that leads to the ranking, we are still able to rank the top \( j_0 \) institutions correctly. In this sense, the institutions that make up the cohort of size \( j_0 \) do not need to be fixed.

It is also of interest to consider cases where the common distribution, \( F \), of the \( \Theta_j \)'s is bounded to the left, for example, where \( F(x) \approx x^\alpha \) as \( x \downarrow 0 \). However, it can be shown that in this context, unless \( p \) is constrained to be a sufficiently low degree polynomial function of \( n \), very few of the estimated ranks \( \hat{R}_j \) will agree with the correct values \( R_j \).

To indicate why, we first recall the model introduced in Section 1, where the estimated ranks \( \hat{R}_j \) are derived by ordering the values of \( \bar{Q}_j + \Theta_j \). Here \( \bar{Q}_j = n^{-1} \sum_{1 \leq i \leq n} Q_{ij} \) is the average value of \( n \) independent and identically distributed random variables with zero mean. Therefore the means, \( \bar{Q}_j \), are of order \( n^{-1/2} \). By way of contrast, if we take \( \alpha = 1 \) in the formula \( F(x) \approx x^\alpha \) as \( x \downarrow 0 \), for example, if \( F \) is the uniform distribution on \([0, 1]\), then the spacings of the order statistics \( \Theta_{(1)} \leq \cdots \leq \Theta_{(p)} \) are approximately of size \( p^{-1} \). (More concisely, they are of size \( Z/p \) where \( Z \) has an exponential distribution; an independent version of \( Z \) is used for each spacing.) Therefore, if \( p \) is of larger order than \( n^{1/2} \) then the errors of the “estimators” \( \bar{Q}_j + \Theta_j \) of \( \Theta_j \), for \( 1 \leq j \leq p \), are an order of magnitude larger than the spacings among the \( \Theta_j \)'s. This can make it very difficult to estimate the ranks
of the $\Theta_j$’s from the ranks of values of $\bar{Q}_j + \Theta_j$. Indeed, it can be shown that, in the difficult case where the components of the $Q_i$’s are independent, and even for fixed $j_0$, if $\alpha = 1$ and $p$ is of larger order than $n^{1/2}$ then in contrast to (2.5),

$$P(\hat{R}_j = R_j \text{ for } 1 \leq j \leq j_0) \rightarrow 0.$$  

(3.8)

This explains why, when $F(x) \backsimeq x^\alpha$, it can be quite rare for the estimated ranks $\hat{R}_j$ to match their true values. Indeed, no matter what the value of $\alpha$ and no matter what the value of $j_0$, property (2.5) will typically fail to hold unless $p$ is no greater than a sufficiently small power of $n$, in particular unless $p = o(n^{\alpha/2})$, as the next result indicates. Thus, the differences between the cases of bounded and unbounded distributions are stark, as can be seen by contrasting Theorem 1 with the properties described below.

THEOREM 2. Assume that $(d/dx)F(x) \backsimeq x^{\alpha-1}$ as $x \downarrow 0$, where $\alpha > 0$, and that (3.3) holds. Part (a): instances where (2.5) holds and $p^2/n^{\alpha} \rightarrow 0$. Under the latter condition, (i) if $\alpha < \frac{1}{2}$ then (2.5) holds even for $j_0 = p$; (ii) if $\alpha = \frac{1}{2}$ then (2.5) holds provided that

$$(\log j_0)^2 (p^2/n^{\alpha}) \rightarrow 0;$$

and (iii) if $\alpha > \frac{1}{2}$ then (2.5) holds provided that

$$j_0 = o \left\{ (n^{\alpha/2} / p)^{1/(2\alpha - 1)} \right\}. $$

(3.9)

(3.10)

Part (b): converses to (a)(ii) and (a)(iii). If $p^2/n^{\alpha} \rightarrow 0$ and the components of the vectors $Q_i$ are independent then, if (2.5) holds, so too does (3.9) (if $\alpha = \frac{1}{2}$) or (3.10) (if $\alpha > \frac{1}{2}$). Part (c): instances where (3.8) holds. If $\alpha > 0$ and $p^2/n^{\alpha} \rightarrow \infty$, and if the components of the vectors $Q_i$ are independent, then (3.8) holds even for $j_0 = 1$.

The proof of Theorem 2 is similar to that of Theorem 1, and so is omitted. Theorem 1 is derived in Section 5. Both results continue to hold if the sample from which $\bar{X}_j$ is computed is of size $n_j$ for $1 \leq j \leq p$, rather than $n$, provided that (2.6) holds.

4. Numerical properties. This section discusses three real-data and three simulated examples linked to the theoretical properties in Section 3. The real-data examples make use of the bootstrap to create prediction intervals [Xie, Singh and Zhang (2009), Hall and Miller (2009)]. In each simulated example, the error is relatively light-tailed, and any discussion of tails refers to the distribution of the $\Theta_j$’s. In our real-data examples, the noise has been averaged and so it is also generally light-tailed. Thus, any heavy-tailed behavior present in the real-data examples is likely to be due to heavy tails of the distribution of the $\Theta_j$’s, rather than the noise.
EXAMPLE 1 (Continued). The originating institutions of *Nature* articles were obtained using the ISI Web of knowledge database\(^3\) for each of the years 1999 through 2008. A point ranking was obtained by taking the average number of articles published per year. Of course, there are implicit simplifying assumptions in doing this, most significantly concerning the independence of articles between years, and the stationarity of means time. These assumptions appear reasonable in context, and are consistent with most publication-based analyses.

When constructing prediction intervals the bootstrap resamples for each institution were drawn independently, conditional on the data. [See Hall and Miller (2009).] The number of observations in the resample can be varied to create different time windows, as illustrated in Figure 1. The most natural question from a ranking correctness viewpoint is determining the behavior at the right tail; there are many institutions with mean at or near the hard threshold of zero, so there is little hope for ranking correctness in the left tail. Furthermore, the right tail appears to be long. Harvard University has an average of 67.5 papers per year, followed by means of 34.6, 29.6 and 28.2 for Berkeley, Stanford and Cambridge, respectively.

A natural question to ask is what the tail shape for this example might be. Approaches to estimating the shape parameter of a distribution with regularly varying tails, such as the method of Hill (1975), are unstable for these data; the number of extreme data for which a linear fit is plausible is very small, implying that the decay rate is faster than polynomial. Indeed, the left panel of Figure 3 shows the QQ plot of the observed data against a random variable with distribution function \[ F(x) = 1 - \exp(-0.85x^{1/2}) \], which suggests that an exponential tail might be reasonable for the data. If this is the case then the number of institutions that we

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\(^3\)www.isiknowledge.com.
expect to be ranked correctly should depend, to first order, only on $n$, not on $p$, and
be of order up to $n^{1/4} (\log n)^{1/2}$. One way to explore this further is to take $j_0$ as
given, and to resample from the data, seeking, for example, the number of years, $n$, needed to obtain correct ranking of the first $j_0$ institutions at least 90\% of the time. A plot of $j_0$ against $n^{1/4} (\log n)^{1/2}$ should be roughly linear. The right-hand panel of Figure 3 plots results of this experiment and appears to support the hypothesis. The flatness between $j_0 = 3$ and $j_0 = 4$ indicates that these two institutions are quite difficult to separate from each other.

**EXAMPLE 2 (Continued).** The Mann–Whitney test statistic can be written as
\[
\max \left\{ \sum_{i,j} I(x_i < y_j), \sum_{i,j} I(x_i > y_j) \right\},
\]
where the $x_i$’s and $y_j$’s are the observed values of the two samples. Notice that this
statistic will have a hard lower threshold at $n_1 n_2 / 2$, where $n_1$ and $n_2$ are the sizes of
the two classes. Here, like the previous example, when the distributions differ only
in location the difference has to be quite large to be detectable. Figure 4 shows the
estimated density as well as the truncated normal density, which is the distribution
that the scores would have if none of the genes had systematically different means
for the two classes. This suggests that an assumption that the majority of genes is
unrelated to whether the tissue is tumorous is not valid here.

Bootstrapped versions of the dataset with different choices for $n$ were created
to indicate how many observations we need to obtain reasonable confidence in a
ranking. Table 1 shows the probability that the set of the top $j$ genes is identified
correctly out of the 2000 for various $j$ and $n$. Note that this is a slightly different
statistic from the one in (2.5), since we allow any permutation of the top $j$ genes
to be detected. The results suggest that we have nearly a 50\% chance of detecting

![Graph](image)

**FIG. 4.** Estimated sampling density genes under the Mann–Whitney test for colon data.
TABLE 1
Probability that set of top \( j \) genes is correct for colon data

| \( j \) | \( n \) | 62   | 100  | 150  | 200  | 250  |
|--------|--------|------|------|------|------|------|
| 1      |        | 0.251| 0.326| 0.437| 0.446| 0.490|
| 2      |        | 0.067| 0.109| 0.166| 0.218| 0.277|
| 4      |        | 0.022| 0.054| 0.094| 0.163| 0.193|
| 6      |        | 0.007| 0.018| 0.035| 0.040| 0.068|

The upper tail for this dataset again appears relatively light; the model \( F(x) = 1 - \exp(-0.19(x - 1)^2) \), for \( x > 1 \), produces a good fit to the upper tail.

Theorem 1 suggests that these probabilities should not depend on the choice of \( p \). We can obtain a sense of this by randomly sampling, without replacement, \( p = 500 \) or \( p = 1000 \) genes from the original \( p = 2000 \), for each simulation; and recalculating the values in Table 1. For \( j = 4 \) and \( n = 250 \), the respective probabilities were 0.183 and 0.170, quite close to the value 0.193 observed for \( p = 2000 \). While the equivalence appears good for \( j \geq 4 \), there are larger departures for \( j = 1 \) or 2, where the initial results for this particular realization tend to distort the calculation.

**EXAMPLE 3 (School rankings).** A third example of accuracy in the extremes of a ranking is based on student performance at 75 private schools in NSW, Australia. For each school the number of final year exams taken, and the number of these where a score of at least 90% was achieved, were recorded. The proportion of exams where 90% or more was scored can be used to rank the schools, and prediction intervals can be constructed by resampling from appropriate binomial distributions. The results in Figure 5 indicate the increased confidence we can have in the upper extreme, with the top school identified with reasonable certainty. In this example, the possible range of scores for ranking has finite support, being restricted to the interval \([0, 1]\); thus it is a context where Theorem 2 is applicable.

Hill’s (1975) estimator of \( \alpha \), when (3.2) holds, is relatively stable in this example and suggests that \( \alpha \approx 6 \). From (3.10), we can calculate that \((n^{\alpha/2} / p)^{1/(2\alpha-1)} \approx 4\), which is consistent with a small number of schools being correctly ranked. If the number were large, then we would expect a significant portion of the schools to be ranked with a high degree of accuracy. In the case of these data, however, the small value suggests that it might not be possible to obtain any correct ranks.

**EXAMPLE 4 (Simulation with exponential tails and infinite support).** Here, we simulate increasing \( n \) and \( p \) in the case of exponential tails. For a given \( n \), set
$$p = 0.0005n^2,$$ let the $\Theta_j$’s be drawn from a standard exponential distribution and the $Q_{ij}$’s be normal random variables with zero mean and standard deviation 3.5. Table 2 shows the results of 1000 simulations for various values of $n$, approximating (2.5) for different choices of $j_0$. Theorem 1 suggests that the results should converge to 1 if $j_0 = o(n^{1/4})$, and degrade, otherwise. This appears consistent with the results. The difficulty of the problem due to the quadratic growth of $p$ and the large error in $Q_{ij}$ is also evident; even when $j_0 = 1$ and $n$ is large, reliable prediction of the top rank is not assured.

**Example 5 (Simulation with polynomial tails and infinite support).** We use the same setup as in the previous example, except that the generating distribution for the $\Theta_j$’s is Pareto, $F(x) = 1 - x^{-\alpha}$ for $x \geq 1$, with $\alpha = 4$. Theorem 1 and (3.4)

**Table 2**

*Probability that the first $j_0$ rankings are correct in the case of exponential tails*

| $j_0$ | 500   | 1000  | 2000  | 5000  | 10,000 | 20,000 | 50,000 |
|-------|-------|-------|-------|-------|--------|--------|--------|
| 1     | 0.909 | 0.9365| 0.959 | 0.970 | 0.9745 | 0.9840 | 0.9910 |
| $n^{0.15}$ | 0.764 | 0.823 | 0.767 | 0.844 | 0.897  | 0.872  | 0.890  |
| $n^{0.20}$ | 0.591 | 0.700 | 0.655 | 0.683 | 0.667  | 0.664  | 0.743  |
| $n^{0.25}$ | 0.420 | 0.406 | 0.424 | 0.383 | 0.334  | 0.402  | 0.428  |
| $n^{0.30}$ | 0.183 | 0.188 | 0.180 | 0.116 | 0.101  | 0.079  | 0.069  |
| $n^{0.35}$ | 0.056 | 0.030 | 0.021 | 0.004 | 0.002  | 0.000  | 0.001  |
suggest that the rate $n^{4/18} p^{1/9} = n^{4/9}$ is critical for $j_0$, and this is consistent with the results in Table 3. This is an easier problem than that in the previous example, because of the polynomial decay of the tail. For instance, the top right-hand result in the table suggests that the top nine ranks can be correctly ascertained more than 90% of the time when $p > 50,000$, whereas the figure 0.890 in the last column of Table 2 suggests that, for the distribution represented there, only the top five ranks have this level of reliability.

**Example 6 (Simulation with polynomial tails with finite support).** Theorem 2 has many interesting consequences, but the present example focuses on case (iii), where $\alpha > \frac{1}{2}$. First, let the $\Theta_j$’s be uniformly distributed on $[0, 1]$, and consider a case where the entire ranking is correct. Using the notation of Section 3 and taking $\alpha = 1$, Theorem 2 implies that $p \asymp n^{1/4}$ defines the critical growth in dimension. For simulation, we took $p = 2n^k$ for various $k$, and scaled the (normally distributed) error for each $k$ such that the $n = 500$ case had probability approximately 0.5 of correctly identifying all ranks. Each simulation was repeated 10,000 times, with results summarized in Table 4. As predicted, growth rates in dimension
Table 5

| $n$  | $5 \times 10^3$ | $1 \times 10^4$ | $2 \times 10^4$ | $5 \times 10^4$ | $1 \times 10^5$ | $2 \times 10^5$ | $5 \times 10^5$ | $1 \times 10^6$ |
|------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|
| 0.05 | 0.500           | 0.539           | 0.553           | 0.583           | 0.603           | 0.609           | 0.628           | 0.641           |
| 0.07 | 0.502           | 0.532           | 0.506           | 0.546           | 0.558           | 0.580           | 0.555           | 0.591           |
| 1/11 | 0.497           | 0.486           | 0.489           | 0.516           | 0.489           | 0.463           | 0.513           | 0.496           |
| 0.11 | 0.497           | 0.481           | 0.471           | 0.432           | 0.461           | 0.447           | 0.447           | 0.452           | 0.421           |
| 0.13 | 0.506           | 0.492           | 0.461           | 0.481           | 0.445           | 0.427           | 0.387           | 0.385           |

sion slower than $n^{1/4}$ have probability of correct ranking tending to 1, while those faster than $n^{1/4}$ degrade.

Next, we examine the case $p = 5 \times 10^{-6}n^2$, where dimension grows at a quadratic rate; and $F(x) = x^\alpha$ on $[0, 1]$, with $\alpha = 6$, implying a reasonably severe tail. Theorem 2 suggests that if $j_0 = o(p^{1/22})$, or equivalently if $j_0 = o(n^{1/11})$, then (2.5) should hold. Table 5 shows the probability of ranking the smallest $j_0 = 10n^k$ scores correctly for various $k$ and $n$, with 10,000 simulations. Again the normal error is tuned so that the $n = 5000$ case has probability of close to $\frac{1}{2}$. The results suggest that $n^{1/11}$ indeed separates values of $k$ for which correct ranking is possible.

5. Technical arguments. We begin by giving a brief sketch of the proof of Theorem 1. Two steps in the proof are initially presented as lemmas, the first using moderate deviation properties to approximate sums related to the object of interest, and the second employing Taylor’s expansion applied to Rényi representations of order statistics to show that the gaps $\Theta_{(j+1)} - \Theta_{(j)}$ have a high probability of being of reasonable size. In the proof itself, we use Lemma 1 to bound the probability in (2.5) from below [see (5.19)] and then show that the last two terms in this expression converge to zero, implying that the probability converges to 1 if (3.6) holds. For the converse, assuming independence, we find an upper bound to the probability in (5.20) and show that if this probability tends to one then the sum $s(n)$, introduced at (5.21), must converge to zero, which in turn implies (3.6). Only the exponential tail case is presented in detail; comments at the end of the proof describe the main differences in the polynomial tail case.

Throughout, we let $\tilde{E}(j_0)$ denote the event that $\tilde{Q}_{R_j} + \Theta_{R_j} > \tilde{Q}_{R_{j_0}} + \Theta_{R_{j_0}}$ for $j_0 + 1 \leq j \leq p$, we define $E_j$ to be the event that $\Theta_{(j+1)} - \Theta_{(j)} \geq -(\tilde{Q}_{R_{j+1}} - \tilde{Q}_{R_j})$, and we take $\tilde{E}(j_0)$ and $\tilde{E}_j$ to be the respective complements. Also, we let $\zeta_j = \Theta_{(j+1)} - \Theta_{(j)}$ denote the $j$th gap, where $\Theta_{(0)} = -\infty$ for convenience.

In Lemma 1 below, we write $\mathcal{O}$ to denote the sigma-field generated by the $\Theta_j$’s, $N$ for a standard normal random variable independent of $\mathcal{O}$, $\delta_n$ for any given se-
quence of positive constants $\delta_n$ converging to zero, and $\Delta$ for a generic random variable satisfying $P(|\Delta| \leq \delta_n) = 1$.

**Lemma 1.** For any positive integer $j_0 < p$, let $J$ denote the set of positive, even integers less than or equal to $j_0$. Put

$$T_{1j} = \frac{\min(\xi_{j-1}, \xi_j)}{2(\text{var} \bar{Q}_{R_j})^{1/2}}, \quad T_{2j} = \frac{\xi_j}{\text{var}(\bar{Q}_{R_{j+1}} - \bar{Q}_{R_j})^{1/2}}.$$

Then

$$\sum_{j=1}^{j_0} P\left( \frac{1}{2} \min(\xi_{j-1}, \xi_j) > |\bar{Q}_{R_j}| \right) = 2\{1 + o(1)\} \sum_{j=1}^{j_0} P(|N| > T_{1j}) + o(1).$$

(5.1)

If in addition the components of the $Q_i$'s are independent then

$$E\left[ \exp\left\{ -\sum_{j \in J} P(\bar{\xi}_j | \mathcal{O}) \right\} \right] \leq \{1 + o(1)\} E\left[ \exp\left\{ -(1 + \Delta) \sum_{j \in J} P(N > T_{2j} | \mathcal{O}) \right\} \right].$$

(5.2)

**Proof.** Using the arguments of Rubin and Sethuraman (1965) and Amosova (1972), it can be shown that, if the constant $C_2$ in (3.3) satisfies $C_2 > B^2 + 2$ where $B > 0$, then as $n$ (and hence, also $p$) diverges,

$$P\{ |\bar{Q}_j| > x(\text{var} \bar{Q}_j)^{1/2} \} = \{1 + o(1)\} 2\{1 - \Phi(x)\},$$

(5.3)

$$P[ -\bar{Q}_{j_1} - \bar{Q}_{j_2} ] \geq x(\text{var} \bar{Q}_{j_1} - \bar{Q}_{j_2})^{1/2} = \{1 + o(1)\} \{1 - \Phi(x)\},$$

(5.4)

uniformly in $0 < x < B(\log p)^{1/2}$ and $j, j_1, j_2 \geq 1$ such that $j_1 \neq j_2$. Expression (5.4) requires the independence assumption. Therefore, since $C_2 > 2(C_1 + 1)$ in (3.3), we can take $B = (2 + \epsilon)^{1/2}$ for some $\epsilon > 0$, and then (5.3) and (5.4) hold uniformly in $0 < x < (2 + \epsilon) \log p)^{1/2}$. Thus, as $n \to \infty$, they hold uniformly in all $x > 0$, modulo an $o(p^{-1})$ term. We use (5.3) to derive (5.1), while (5.4) implies that

$$\sum_{j \in J} P(\bar{\xi}_j) = \{1 + o(1)\} \sum_{j \in J} P(N > T_{2j}) + o(1),$$

which leads to (5.2). \qed
Lemma 2. If (3.1), indicating the case of exponential tails, holds then there exist \(B_4, B_5 > 0\) such that, for any choice of constants \(c_1, c_2\) satisfying \(0 < c_1 < c_2 < (4 - \varepsilon)^{-1}\) with \(\varepsilon\) as in (3.5), and for all \(B_6 > 0\),

\[
\inf_{j \in [1, n^{c_1 + \varepsilon}]} P\{\zeta_j Z_{j+1}^{-1} (\log n)^{1-(1/\alpha)} \geq B_4 n^{-c_1}\} = 1 - O(n^{-B_6}),
\]

(5.5)

\[
\inf_{j \in [n^{c_1}, n^{c_2}]} P\{B_4 \leq j \zeta_j Z_{j+1}^{-1} (\log n)^{1-(1/\alpha)} \leq B_5\} = 1 - O(n^{-B_6}).
\]

(5.6)

Note further that the constraint on \(c_2\) permits \(n^{c_2}\) to be of size \(\nu \exp n^{\varepsilon_1}\) (where \(\varepsilon_1 > 0\)).

Proof. If \(U(1) \leq \cdots \leq U(p)\) denote the order statistics of a sample of size \(p\) drawn from the uniform distribution on \([0, 1]\) then, for each \(p\), we can construct a collection of independent random variables \(Z_1, \ldots, Z_p\) with the standard negative exponential distribution on \([0, \infty]\), such that, for \(1 \leq j \leq p\),

\[
U(j) = 1 - \exp(-V_j)
\]

where

\[
V_j = \sum_{k=1}^{j} \frac{Z_k}{p - k + 1} = w_j + W_j.
\]

For details, see Rényi (1953). Further, uniformly in \(1 \leq j \leq \frac{1}{2} p\) and \(2 \leq p < \infty\),

\[
w_j = \sum_{k=p-j+1}^{p} \frac{1}{k} = \frac{j}{p} + O(j^2/p^2) = O(j/p),
\]

(5.7)

\[
W_j = \sum_{k=p-j+1}^{p} k^{-1}(Z_{p-k+1} - 1), \quad \sup_{1 \leq j \leq p/2} j^{-1/2}|W_j| \leq p^{-1}W(p),
\]

(5.8)

\[
\sup_{1 \leq j \leq p/2} j^{-3/2}\left|W_j - \frac{1}{p} \sum_{k=p-j+1}^{p} (Z_{p-k+1} - 1)\right| \leq p^{-2}W(p),
\]

(5.9)

where the nonnegative random variable \(W(p)\), which without loss of generality, we take to be common to (5.8) and (5.9), satisfies the expression \(P\{W(p) > p^\varepsilon\} = O(p^{-C})\) for each \(C, \varepsilon > 0\).

Using the second identity in (5.7), and (5.8), we deduce that

\[
U_{j+1} - U_j = (V_{j+1} - V_j)\left\{1 - \frac{1}{2}(V_{j+1} + V_j)
\right\}
\]

\[
+ \frac{1}{6}(V_{j+1}^2 + V_j V_{j+1} + V_j^2) - \cdots
\]

(5.10)

\[
= \frac{Z_{j+1}}{p - j}\left\{1 + \Psi_{j_1}\left(\frac{j}{p} + \frac{S_{j_1}}{p^{1/2}}\right)\right\},
\]
uniformly in $1 \leq j \leq \frac{1}{2} p$, where the random variable $\Psi_{j1}$ satisfies, for $k = 1$,

$$P\left( \max_{1 \leq j \leq p/2} |\Psi_{jk}| \leq A \right) = 1,$$

(5.11)

$A > 0$ is an absolute constant, and for each $C, \varepsilon > 0$ the nonnegative random variable $S_{j1}$ satisfies, with $k = 1$,

$$P\left( \sup_{1 \leq j \leq p/2} S_{jk} > p^\varepsilon \right) = O\left( p^{-C} \right).$$

(5.12)

Using the third identity in (5.7) and (5.9), we deduce that

$$0 \leq U(j) = w_j + W_j - \frac{1}{2} (w_j + W_j)^2 + \cdots$$

(5.13)

$$= \frac{j}{p} + \Psi_{j2} \left( \frac{j^2}{p^2} + \frac{j^{1/2} S_{j2}}{p} \right),$$

where $\Psi_{j2}$ and $S_{j2} \geq 0$ satisfy (5.11) and (5.12), respectively.

Define $D_j = U(j+1) - U(j)$ and without loss of generality, $C_0 = 1$ in (3.1). If the common distribution function of the $\Theta_1j$’s is $F$, then by Taylor’s expansion,

$$\zeta_j = F^{-1}(U(j) + D_j) - F^{-1}(U(j))$$

(5.14)

$$= D_j (F^{-1})'(U(j) + \omega_j D_j)$$

$$= \Psi_j \frac{D_j}{U(j) + \omega_j D_j} \left\{ - \log(U(j) + \omega_j D_j) \right\}^{(1/\alpha) - 1},$$

where $0 \leq \omega_j \leq 1$ and the last line makes use of (3.1). The random variable $\Psi_j$ satisfies, for constants $B_1, B_2$ and $B_3$ satisfying $0 < B_1 < B_2 < \infty$ and $0 < B_3 < 1$,

$$P\left( B_1 \leq \Psi_j \leq B_2 \text{ for all } j \text{ such that } U(j+1) < B_3 \right) = 1.$$

The required result then follows from (5.10), (5.13) and (5.14). □

**Proof of Theorem 1.** Take $j_0 < p$ a positive integer. Note that, taking $\mathcal{E}(j_0), \mathcal{E}_j, \tilde{\mathcal{E}}(j_0), \tilde{\mathcal{E}}_j, \mathcal{O}$ and $\mathcal{J}$ as for Lemma 1,

$$\{ \tilde{R}_j = R_j \text{ for } 1 \leq j \leq j_0 \}$$

$$\supset \{ |\tilde{Q}_{R_j}| \leq \frac{1}{2} \min(\zeta_{j-1}, \zeta_j) \text{ for } 1 \leq j \leq j_0 \} \cap \mathcal{E}(j_0),$$

where we define $\Theta_{(j-1)} = -\infty$ if $j = 1$ as before. Therefore, defining $\pi(j_0) = P(\tilde{R}_j = R_j \text{ for } 1 \leq j \leq j_0)$, we deduce that

$$\pi(j_0) \geq 1 - \sum_{j=1}^{j_0} P \left\{ |\tilde{Q}_{R_j}| > \frac{1}{2} \min(\zeta_{j-1}, \zeta_j) \right\} - P\{\tilde{\mathcal{E}}(j_0)\}.$$
Also,
\[
\{ \hat{R}_j = R_j \text{ for } 1 \leq j \leq j_0 \}
\]
\[
= \{ \bar{X}_{R_1} \leq \cdots \leq \bar{X}_{R_{j_0}} \text{ and } \bar{X}_j > \bar{X}_{R_{j_0}} \text{ for } j \notin \{R_1, \ldots, R_{j_0}\} \}
\]
\[
= \{ \zeta_j \geq - (\bar{Q}_{R_{j+1}} - \bar{Q}_{R_j}) \text{ for } 1 \leq j \leq j_0 \}
\]
and so
\[
(5.16) \quad \pi(j_0) \leq P(\zeta_j \geq - (\bar{Q}_{R_{j+1}} - \bar{Q}_{R_j}) \text{ for } 1 \leq j \leq j_0).
\]

Letting \( \pi_1(j_0) \) denote the probability that \( \mathcal{E}_j \) holds for all \( j \in \mathcal{J} \), by (5.16),
\[
(5.17) \quad \pi(j_0) \leq \pi_1(j_0).
\]

Note that if the components of each \( Q_i \) are independent, then the events \( \mathcal{E}_j \), for \( j \in \mathcal{J} \), are independent conditional on \( \mathcal{O} \). Therefore,
\[
\pi_1(j_0) = E \left\{ P \left( \bigcap_{j \in \mathcal{J}} \mathcal{E}_j \mid \mathcal{O} \right) \right\} = E \left[ \prod_{j \in \mathcal{J}} \left\{ 1 - P(\mathcal{E}_j \mid \mathcal{O}) \right\} \right]
\]
\[
(5.18) \quad \leq E \left[ \exp \left\{ - \sum_{j \in \mathcal{J}} P(\bar{\mathcal{E}}_j \mid \mathcal{O}) \right\} \right].
\]

Using Lemma 1, we have the following inequalities regarding \( \pi(j_0) \):
\[
(5.19) \quad \pi(j_0) \geq 1 - 2\{1 + o(1)\} \sum_{j=1}^{j_0} P(|N| > T_{1j}) - P(\tilde{\mathcal{E}}(j_0)) + o(1),
\]
\[
(5.20) \quad \pi(j_0) \leq \{1 + o(1)\} E \left[ \exp \left\{ -(1 + \Delta) \sum_{j \in \mathcal{J}} P(N > T_{2j} \mid \mathcal{O}) \right\} \right].
\]

To show that (3.6) implies (2.5), by (5.19) it is sufficient to show that \( P(\tilde{\mathcal{E}}(j_0)) \) and \( \sum_{j=1}^{j_0} P(|N| > T_{1j}) \) are both \( o(1) \), which we shall do in turn.

Define \( \ell = (\log n)^{(1/\alpha)-1} \), let \( N \) be a standard normal random variable independent of \( \mathcal{O} \), and let \( Z \) be independent of \( N \) and have the standard negative exponential distribution. Let \( K_1 \) be a positive constant. If \( a_n \) is a sequence of positive numbers and \( f_n \) is a sequence of nonnegative functions, write \( a_n \approx f_n(K) \) to mean that, for constants \( L_1, L_2 > 1 \), either (a) \( a_n \leq L_1 f_n(K) \) whenever \( K \geq L_2 \) and \( n \) is sufficiently large, and \( a_n \geq L_1^{-1} f_n(K) \) whenever \( K \leq L_2^{-1} \) and \( n \) is sufficiently large, or (b) \( a_n \geq L_1^{-1} f_n(K) \) whenever \( K \geq L_2 \) and \( n \) is sufficiently large, and \( a_n \leq L_1 f_n(K) \) whenever \( K \leq L_2^{-1} \) and \( n \) is sufficiently large. Let \( 0 < c_1 < c_2 < \frac{1}{2} \) and \( c_1 < \frac{1}{4} \), and let \( j_0 \) and \( j_1 \) denote integers satisfying \( |j_1 - n^{c_1}| \leq 1 \), \( j_1 \leq j_0 \leq n^{c_2} \) and \( j_1/j_0 \to 0 \).
When (3.1) holds with $C_0 = 1$, Lemma 2 implies that, for each $B_6 > 0$ and letting $\gamma_j = n^{-1/2} j^{\ell-1}$,

$$s(n) \equiv \sum_{j_1}^{j_0} P(\{|N| > K_1 n^{1/2}(\xi_j)\})$$

\[\equiv O\left\{ j_1 P(\{|N| > K_2 Z \gamma_{j_1}^{-1}\}) + n^{-B_6} \right\} + \sum_{j_1 < j \leq j_0} P(\{|N| > K Z \gamma_j^{-1}\})\]

\[\equiv O\left\{ j_1 \left( P(Z \leq \gamma_{j_1}) + E\left[ Z^{-1} \gamma_{j_1} \exp\left\{-\frac{1}{2}(K Z \gamma_{j_1}^{-1})^2 I(Z > \gamma_{j_1})\right\}\right]\right\} + \sum_{j_1 < j \leq j_0} \left( P(Z \leq \gamma_j) + E\left[ Z^{-1} \gamma_j \exp\left\{-\frac{1}{2}(K Z \gamma_j^{-1})^2 I(Z > \gamma_j)\right\}\right]\right)\]

\[\equiv O\left\{ j_1 \left( \gamma_{j_1} + E\left[ Z^{-1} \gamma_{j_1} \exp\left\{-\frac{1}{2}(K Z \gamma_{j_1}^{-1})^2 I(Z > \gamma_{j_1})\right\}\right]\right\} + \sum_{j_1 < j \leq j_0} \left( \gamma_j + E\left[ Z^{-1} \gamma_j \exp\left\{-\frac{1}{2}(K Z \gamma_j^{-1})^2 I(Z > \gamma_j)\right\}\right]\right)\] (5.21)

Now,

$$E\left[ Z^{-1} \gamma_j \exp\left\{-\frac{1}{2}(K Z \gamma_j^{-1})^2 I(Z > \gamma_j)\right\} \right]$$

\[= \int_0^\infty z^{-1} \gamma_j \exp\left\{-\frac{1}{2}(K Z \gamma_j^{-1})^2 - z\right\} dz\]

$$= \gamma_j \int_1^\infty u^{-1} \exp\left\{-\frac{1}{2}(K u)^2 - \gamma_j u\right\} du \approx \gamma_j \]

$$= n^{-1/2} j^{\ell}.\]

(Here, we have used the fact that $j \leq j_0 \leq n^{c_2}$ where $c_2 < \frac{1}{2}$.) Therefore,

$$s(n) \asymp j_1 \cdot n^{-1/2} j_1^{\ell-1} + \sum_{j_1 < j \leq j_0} n^{-1/2} j^{\ell-1}$$

\[\asymp n^{-1/2} j_1^2 j_1^{\ell-1} + n^{-1/2} j_0^{2/\ell-1}\]

\[\asymp n^{-1/2} j_0^2 j_0^{\ell-1}.\]

(Here, we have used the fact that $j_1/j_0 \rightarrow 0$.)

The right-hand side of (5.22) converges to zero if and only if (3.6) holds. Moreover, in view of the fact that

$$P(|N| > T_{1,j}) \leq P\left( |N| > \frac{\xi_{j-1}}{2(\text{var } Q_{R_j})^{1/2}} \right) + P\left( |N| > \frac{\xi_j}{2(\text{var } Q_{R_j})^{1/2}} \right),$$
and depending on the choice of \( K_1 \) in the definition of \( s(n) \) at (5.21), \( s(n) \) can be an upper bound to the series \( \sum_{j=1}^{j_0} P(|N| > T_{1j}) \) on the right-hand side of (5.1). Hence,

(5.23) \[ \sum_{j=1}^{j_0} P(|N| > T_{1j}) = o(1). \]

This deals with the second term on the right-hand side of (5.19). Similarly, if \( r \in [2, \infty) \) is a fixed integer, and if \( j_0 = o(n^{1/4} \ell^{1/2}) \), then

(5.24) \[ s_1(n) \equiv \sum_{j=j_0+1}^{j_0+r-1} P(|N| > K_1 n^{1/2}(\xi_j)) = o(1). \]

Moreover, if \( j_1 \) denotes the integer part of \( n^{r/2} - j_0 \) then, for constants \( K_2 \) and \( K_3 \) satisfying \( K_1 > K_2 > K_3 > 0 \), and for any \( B > 0 \),

\[
\begin{align*}
\sum_{j=j_0+1}^{j_0+r-1} P\{ |N| > K_1 n^{1/2}(\Theta_{(j+1)} - \Theta_{(j_0)}) \}
\leq \sum_{j=j_0+1}^{j_0+r-1} P\left\{ |N| > K_2 n^{1/2} \ell \sum_{k=1}^{j_0+k} Z_k \right\} + O(n^{-B}) \\
\leq j_1 P\{ |N| > K_2 n^{1/4} \ell^{1/2} (Z_1 + \cdots + Z_r) \} + O(n^{-B}) \\
= O\{ j_1 (n^{1/2} \ell^{2})^{-r} \},
\end{align*}
\]

where we have assumed that \( j_0 = o(n^{1/4} \ell^{1/2}) \) and also used the fact that \( Z_1 + \cdots + Z_r \) has a gamma\((r, 1)\) distribution. If we choose \( r \) so large that \( pn^{-r/2} = O(n^{-\varepsilon}) \) for some \( \varepsilon > 0 \), then we can deduce from (5.24) and (5.25) that \( s_1(n) + s_2(n) \to 0 \), and hence, by (5.6), that

(5.26) \[ \sum_{j=j_0+1}^{n^{r/2}} P(\tilde{Q}_{R_j} + \Theta_{R_j} > \tilde{Q}_{R_{j_0}} + \Theta_{R_{j_0}}) \to 0. \]

A more crude argument can be used to prove that if \( r \) is so large that \( p^2 n^{-r/2} = O(n^{-\varepsilon}) \) for some \( \varepsilon > 0 \), and if \( j_0 = o(n^{1/4} \ell^{1/2}) \), then

(5.27) \[ \sum_{n^{r/2} < j \leq p} P(\tilde{Q}_{R_j} + \Theta_{R_j} > \tilde{Q}_{R_{j_0}} + \Theta_{R_{j_0}}) \to 0. \]

Together, (5.26) and (5.27) imply that if \( j_0 = o(n^{1/4} \ell^{1/2}) \) then

(5.28) \[ P(\tilde{E}(j_0)) \to 0. \]
Thus, in light of (5.19), we see (5.23) and (5.28) imply that (3.6) is sufficient for (2.5).

We next show that (2.5) implies (3.6) in the independent case. If (2.5) holds, then by (5.20),
\[ \sum_{j \in J} P(N > T_{2j} \mid \mathcal{O}) \to 0 \]
in probability. Therefore, by Lemma 2, with \( j_0 \) and \( j_1 \) as above, there exists \( K_1 > 0 \) such that
\[ \sum_{j_1 < j \leq j_0} P(|N| > n^{1/2} K_1(\xi_j) \mid \mathcal{O}) \to 0 \]
in probability. (We can take the sum over all \( j \in [j_1 + 1, j_0] \), rather than just over even \( j \), since (5.2) holds for sums over odd \( j \) as well as over even \( j \).) Hence, arguing as in the lines below (5.21), we deduce that for sufficiently large \( K_2 > 0 \),
\[ \sum_{j_1 < j \leq j_0} f(Z_j / \delta_j) \to 0 \]
in probability, where the random variables \( Z_j \) are independent and have a common exponential distribution, \( \delta_j = n^{-1/2} j \ell^{-1} \) and
\[ f(z) = z^{-1} \exp(-K_2 z^2)I(z > 1). \]

We claim that this implies that the expected value of the left-hand side of (5.29) also converges to 0:
\[ \sum_{j_1 < j \leq j_0} E\{f(Z_j / \delta_j)\} \to 0 \]
or equivalently that \( \sum_{j_1 < j \leq j_0} \delta_j \to 0 \), and hence [using the argument leading to (5.22)] that \( s(n) \asymp n^{-1/2} J_0^2 \ell^{-1} \to 0 \), which is equivalent to (3.6). Therefore, if we establish (5.30) then we shall have proved that (2.5) implies (3.6).

It remains to show that (5.29) implies (5.30). This we do by contradiction. If (5.30) fails then, along a subsequence of values of \( n \), the left-hand side of (5.30) converges to a nonzero number. For notational simplicity, we shall make the inessential assumptions that the number is finite and that the subsequence involves all \( n \), and we shall take \( K_2 = 1 \) in the definition of \( f \). In particular,
\[ t(n) \equiv \sum_{j_1 < j \leq j_0} E\{f(Z_j / \delta_j)\} \to t(\infty), \]
where \( t(\infty) \) is bounded away from 0. Now, \( t(n) = (1 + o(1)) \mu(1) \delta(n) \), where \( \delta(n) = \sum_{j_1 < j \leq j_0} \delta_j \) and, for general \( \lambda \geq 1 \), \( \mu(\lambda) = \int_{z > \lambda} z^{-1} \exp(-z^2) \, dz \). There-
Therefore,\( \delta(n) \to \delta(\infty) \equiv t(\infty)/\mu(1) \).

For each \( \lambda > 1 \) the left-hand side of (5.29) equals \( \Delta_1 + \Delta_2 \), where, in view of (5.31),

\[
E(\Delta_2) = \sum_{j_1 < j \leq j_0} E\{f(Z_j/\delta_j)I(Z_j > \lambda \delta_j)\}
\]

(5.33)

\[
= \{1 + o(1)\} \mu(\lambda) \delta(n)
\]

and

\[
\Delta_1 = \sum_{j_1 < j \leq j_0} f(Z_j/\delta_j)I(Z_j \leq \lambda \delta_j) = \sum_{j_1 < j \leq j_0} f(W_j)I_j
\]

with \( W_j = Z_j/\delta_j \) and \( I_j = I(\delta_j \leq Z_j \leq \lambda \delta_j) \). However,

\[
\sum_{j_1 < j \leq j_0} P(I_j = 1) = \mu_1(\lambda) \delta(n) + o(1) = \delta(\infty) \mu(\lambda) + o(1),
\]

where \( \mu_1(\lambda) = \int_{-\infty}^{\lambda} \exp(-z^2) \, dz \). Therefore, in the limit as \( n \to \infty \), \( \Delta_1 \) equals a sum, \( S_\lambda \), say, of \( N \) independent random variables each having the distribution of \( f(W) \), where \( W \) is uniformly distributed on \([1, \lambda]\), \( N \) has a Poisson distribution with mean \( \delta(\infty) \mu_1(\lambda) \), and \( N \) and the summands are independent. The distribution of \( S_\lambda \) is stochastically monotone increasing, in the sense that \( P(S_\lambda > s) \) increases with \( \lambda \). On the other hand, since \( \mu(\lambda) \to 0 \) as \( \lambda \to \infty \) then, by (5.32) and (5.33),

\[
\lim_{\lambda \to \infty} \limsup_{n \to \infty} E(\Delta_2) = 0.
\]

Combining these results, we deduce that \( \Delta_1 + \Delta_2 \), that is, the left-hand side of (5.29), does not converge to zero in probability. This contradicts (5.29) and so establishes that \( t(\infty) \) must equal zero; that is, (5.30) holds.

Comments on proving the polynomial case: the proof for the case of polynomial tails proceeds similarly. The main difference is that in the proof of Lemma 2 we use (3.2) instead of (3.1), which forces a factor of \( p^{-1/\alpha} \) into the results of the lemma, rather than \( (\log n)^{1-(1/\alpha)} \). This in turn implies that \( s(n) \asymp n^{-1/2} j_0^{2+1/\alpha} p^{-1/\alpha} \), entailing that convergence occurs if (and, in the case of independence, only if) \( j_0 = o(\nu_{\text{pol}}) \), as required. \( \square \)

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