

# Robust estimations from distribution structures: III. Invariant Moments

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1 **Descriptive statistics for parametric models are currently highly sensitive to departures, gross errors, and/or random errors. Here, leveraging the structures of parametric distributions and their central moment kernel distributions, a class of estimators, consistent simultaneously for both a semiparametric distribution and a distinct parametric distribution, is proposed. These efficient estimators are robust to both gross errors and departures from parametric assumptions, making them ideal for estimating the mean and central moments of common unimodal distributions. This article also illuminates the understanding of the common nature of probability distributions and the measures of them.**

moments | invariant | unimodal | adaptive estimation |  $U$ -statistics

1 **T**he potential biases of robust location estimators in estimating the population mean have been noticed for more than two centuries (1), with numerous significant attempts made to address them. In calculating a robust estimator, the procedure of identifying and downweighting extreme values inherently necessitates the formulation of distributional assumptions. Previously, it was demonstrated that, due to the presence of infinite-dimensional nuisance shape parameters, the semiparametric approach struggles to consistently address distributions with shapes more intricate than symmetry. Newcomb (1886) provided the first modern approach to robust parametric estimation by developing a class of estimators that gives "less weight to the more discordant observations" (2). In 1964, Huber (3) used the minimax procedure to obtain  $M$ -estimator for the contaminated normal distribution, which has played a pre-eminent role in the later development of robust statistics. However, as previously demonstrated, under growing asymmetric departures from normality, the bias of the Huber  $M$ -estimator (HM) increases rapidly. This is a common issue in parametric robust statistics. For example, He and Fung (1999) constructed (4) a robust  $M$ -estimator (HFM) for the two-parameter Weibull distribution, from which the mean and central moments can be calculated. Nonetheless, it is inadequate for other parametric distributions, e.g., the gamma, Perato, lognormal, and the generalized Gaussian distributions (SI Dataset S1). Another interesting approach is based on  $L$ -estimators, such as percentile estimators. For examples of percentile estimators for the Weibull distribution, the reader is referred to the works of Menon (1963) (5), Dubey (1967) (6), Marks (2005) (7), and Boudt, Caliskan, and Croux (2011) (8). At the outset of the study of percentile estimators, it was known that they arithmetically utilize the invariant structures of parametric distributions (5, 6). An estimator is classified as an  $I$ -statistic if it asymptotically satisfies  $I(LE_1, \dots, LE_l) = (\theta_1, \dots, \theta_q)$  for the distribution it is consistent, where LEs are calculated with the use of  $LU$ -statistics (defined in Subsection ??),  $I$  is defined using arithmetic operations and constants but may also incorporate transcendental functions and quantile functions, and  $\theta$ s are the population parameters it estimates. In this article, two subclasses of  $I$ -

statistics are introduced, recombined  $I$ -statistics and quantile  $I$ -statistics. Based on  $LU$ -statistics,  $I$ -statistics are naturally robust. Compared to probability density functions (pdfs) and cumulative distribution functions (cdfs), the quantile functions of many parametric distributions are more elegant. Since the expectation of an  $L$ -estimator can be expressed as an integral of the quantile function,  $I$ -statistics are often analytically obtainable. However, it is observed that even when the sample follows a gamma distribution, which belongs to the same larger family as the Weibull model, the generalized gamma distribution, a misassumption can still lead to substantial biases in Marks percentile estimator (MP) for the Weibull distribution (7) (SI Dataset S1).

On the other hand, while robust estimation of scale has also been intensively studied with established methods (9, 10), the development of robust measures of asymmetry and kurtosis lags behind, despite the availability of several approaches (11–15). The purpose of this paper is to demonstrate that, in light of previous works, the estimation of central moments can be transformed into a location estimation problem by using  $U$ -statistics, the central moment kernel distributions possess desirable properties, and by utilizing the invariant structures of unimodal distributions, a suite of robust estimators can be constructed whose biases are typically smaller than the variances (as seen in Table 1 for  $n = 5184$ ).

**A. Invariant Moments.** Most popular robust location estimators, such as the symmetric trimmed mean, symmetric Winsorized mean, Hodges-Lehmann estimator, Huber  $M$ -estimator, and median of means, are symmetric. As shown in REDS I, a symmetric weighted Hodges-Lehmann mean ( $SWHLM_{k,\epsilon}$ ) can achieve consistency for the population mean in any symmetric distribution with a finite mean. However, it falls considerably short of consistently handling other parametric distributions that are not symmetric. Shifting from semiparametrics to parametrics, consider a robust estimator with a non-sample-dependent breakdown point (defined in

## Significance Statement

Bias, variance, and contamination are the three main errors in statistics. Consistent robust estimation is unattainable without distributional assumptions. In this article, invariant moments are proposed as a means of achieving near-consistent and robust estimations of moments, even in scenarios where moderate violations of distributional assumptions occur, while the variances are sometimes smaller than those of the sample moments.

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Subsection ??) which is consistent simultaneously for both a semiparametric distribution and a parametric distribution that does not belong to that semiparametric distribution, it is named with the prefix ‘invariant’ followed by the name of the population parameter it is consistent with. Here, the recombined  $I$ -statistic is defined as

$$RI_{d,h_{\mathbf{k}},\mathbf{k}_1,\mathbf{k}_2,k_1,k_2,\epsilon=\min(\epsilon_1,\epsilon_2),\gamma_1,\gamma_2,n,LU_1,LU_2} := \lim_{c \rightarrow \infty} \left( \frac{(LU_{1h_{\mathbf{k}},\mathbf{k}_1,k_1,\epsilon_1,\gamma_1,n} + c)^{d+1}}{(LU_{2h_{\mathbf{k}},\mathbf{k}_2,k_2,\epsilon_2,\gamma_2,n} + c)^d} - c \right),$$

where  $d$  is the key factor for bias correction,  $LU_{h_{\mathbf{k}},\mathbf{k},k,\epsilon,\gamma,n}$  is the  $LU$ -statistic,  $\mathbf{k}$  is the degree of the  $U$ -statistic,  $k$  is the degree of the  $LL$ -statistic,  $\epsilon$  is the upper asymptotic breakdown point of the  $LU$ -statistic. It is assumed in this series that in the subscript of an estimator, if  $\mathbf{k}$ ,  $k$  and  $\gamma$  are omitted,  $\mathbf{k} = 1$ ,  $k = 1$ ,  $\gamma = 1$  are assumed, if just one  $\mathbf{k}$  is indicated,  $\mathbf{k}_1 = \mathbf{k}_2$ , if just one  $\gamma$  is indicated,  $\gamma_1 = \gamma_2$ , if  $n$  is omitted, only the asymptotic behavior is considered, in the absence of subscripts, no assumptions are made. The subsequent theorem shows the significance of a recombined  $I$ -statistic.

**Theorem A.1.** Define the recombined mean as  $rm_{d,k_1,k_2,\epsilon=\min(\epsilon_1,\epsilon_2),\gamma_1,\gamma_2,n,WL_1,WL_2} := RI_{d,h_{\mathbf{k}}=x,\mathbf{k}_1=1,\mathbf{k}_2=1,k_1,k_2,\epsilon=\min(\epsilon_1,\epsilon_2),\gamma_1,\gamma_2,n,LU_1=WL_1,LU_2=WL_2}$ . Assuming finite means,  $rm_{d,k_1,k_2,\epsilon=\min(\epsilon_1,\epsilon_2),\gamma_1,\gamma_2,n,WL_1,WL_2} = \frac{\mu - WL_{1k_1,\epsilon_1,\gamma_1}}{WL_{1k_1,\epsilon_1,\gamma_1} - WL_{2k_2,\epsilon_2,\gamma_2}}$ , is a consistent mean estimator for a location-scale distribution, where  $\mu$ ,  $WL_{1k_1,\epsilon_1,\gamma_1}$ , and  $WL_{2k_2,\epsilon_2,\gamma_2}$  are different location parameters from that location-scale distribution. If  $\gamma_1 = \gamma_2 = 1$ ,  $WL = SWHLM$ ,  $rm$  is also consistent for any symmetric distributions.

*Proof.* Finding  $d$  that make  $rm_{d,k_1,k_2,\epsilon=\min(\epsilon_1,\epsilon_2),\gamma_1,\gamma_2,n,WL_1,WL_2}$  a consistent mean estimator is equivalent to finding the solution of  $rm_{d,k_1,k_2,\epsilon=\min(\epsilon_1,\epsilon_2),\gamma_1,\gamma_2,n,WL_1,WL_2} = \mu$ . First consider the location-scale distribution. Since  $rm_{d,k_1,k_2,\epsilon=\min(\epsilon_1,\epsilon_2),\gamma_1,\gamma_2,n,WL_1,WL_2} = \lim_{c \rightarrow \infty} \left( \frac{(WL_{1k_1,\epsilon_1,\gamma_1} + c)^{d+1}}{(WL_{2k_2,\epsilon_2,\gamma_2} + c)^d} - c \right) = (d+1)WL_{1k_1,\epsilon_1,\gamma_1} - dWL_{2k_2,\epsilon_2,\gamma_2} = \mu$ . So,  $d = \frac{\mu - WL_{1k_1,\epsilon_1,\gamma_1}}{WL_{1k_1,\epsilon_1,\gamma_1} - WL_{2k_2,\epsilon_2,\gamma_2}}$ . In REDS I, it was established that any  $WL(k, \epsilon, \gamma)$  can be expressed as  $\lambda WL_0(k, \epsilon, \gamma) + \mu$  for a location-scale distribution parameterized by a location parameter  $\mu$  and a scale parameter  $\lambda$ , where  $WL_0(k, \epsilon, \gamma)$  is a function of  $Q_0(p)$ , the quantile function of a standard distribution without any shifts or scaling, according to the definition of the weighted  $L$ -statistic. The simultaneous cancellation of  $\mu$  and  $\lambda$  in  $\frac{(\lambda\mu_0 + \mu) - (\lambda WL_{10}(k_1, \epsilon_1, \gamma_1) + \mu)}{(\lambda WL_{10}(k_1, \epsilon_1, \gamma_1) + \mu) - (\lambda WL_{20}(k_2, \epsilon_2, \gamma_2) + \mu)}$  assures that the  $d$  in  $rm$  is always a constant for a location-scale distribution. The proof of the second assertion follows directly from the coincidence property. According to Theorem 19 in REDS I, for any symmetric distribution with a finite mean,  $SWHLM_{1k_1} = SWHLM_{2k_2} = \mu$ . Then  $rm_{d,k_1,k_2,\epsilon_1,\epsilon_2,SWHLM_1,SWHLM_2} = \lim_{c \rightarrow \infty} \left( \frac{(\mu+c)^{d+1}}{(\mu+c)^d} - c \right) = \mu$ . This completes the demonstration.  $\square$

For example, the Pareto distribution has a quantile function  $Q_{Par}(p) = x_m(1-p)^{-\frac{1}{\alpha}}$ , where  $x_m$  is the minimum possible value that a random variable following the Pareto distribution

can take, serving a scale parameter,  $\alpha$  is a shape parameter. The mean of the Pareto distribution is given by  $\frac{\alpha x_m}{\alpha-1}$ . As  $WL(k, \epsilon, \gamma)$  can be expressed as a function of  $Q(p)$ , one can set the two  $WL_{k,\epsilon,\gamma}$ s in the  $d$  value of  $rm$  as two arbitrary quantiles  $Q_{Par}(p_1)$  and  $Q_{Par}(p_2)$ . For the Pareto distribution,

$$d_{Per,rm} = \frac{\mu_{Per} - Q_{Par}(p_1)}{Q_{Par}(p_1) - Q_{Par}(p_2)} = \frac{\frac{\alpha x_m}{\alpha-1} - x_m(1-p_1)^{-\frac{1}{\alpha}}}{x_m(1-p_1)^{-\frac{1}{\alpha}} - x_m(1-p_2)^{-\frac{1}{\alpha}}}$$

$x_m$  can be canceled out. Intriguingly, the quantile function of exponential distribution is  $Q_{exp}(p) = \ln\left(\frac{1}{1-p}\right)\lambda$ ,  $\lambda \geq 0$ .  $\mu_{exp} = \lambda$ . Then,  $d_{exp,rm} = \frac{\mu_{exp} - Q_{exp}(p_1)}{Q_{exp}(p_1) - Q_{exp}(p_2)} = \frac{\lambda - \ln\left(\frac{1}{1-p_1}\right)\lambda}{\ln\left(\frac{1}{1-p_1}\right)\lambda - \ln\left(\frac{1}{1-p_2}\right)\lambda} = -\frac{\ln(1-p_1)+1}{\ln(1-p_1)-\ln(1-p_2)}$ . Since

$$\lim_{\alpha \rightarrow \infty} \frac{\frac{\alpha}{\alpha-1} - (1-p_1)^{-1/\alpha}}{(1-p_1)^{-1/\alpha} - (1-p_2)^{-1/\alpha}} = -\frac{\ln(1-p_1)+1}{\ln(1-p_1)-\ln(1-p_2)},$$

$d_{Per,rm}$  approaches  $d_{exp,rm}$ , as  $\alpha \rightarrow \infty$ , regardless of the type of weighted  $L$ -statistic used. That means, for the Weibull, gamma, Pareto, log-normal and generalized Gaussian distribution,

$$rm_{d,k_1,k_2,\epsilon=\min(\epsilon_1,\epsilon_2),SWHLM_1,SWHLM_2} = \frac{\mu - SWHLM_{1k_1,\epsilon_1}}{SWHLM_{1k_1,\epsilon_1} - SWHLM_{2k_2,\epsilon_2}}$$

is consistent for at least one particular case, where  $\mu$ ,  $SWHLM_{1k_1,\epsilon_1}$ , and  $SWHLM_{2k_2,\epsilon_2}$  are different location parameters from an exponential distribution. Let  $SWHLM_{1k_1,\epsilon_1,\gamma} = BM_{\nu=3,\epsilon=\frac{1}{24}}$ ,

$$SWHLM_{2k_2,\epsilon_2,\gamma} = m, \text{ then } \mu = \lambda, m = Q\left(\frac{1}{2}\right) = \ln 2\lambda, BM_{\nu=3,\epsilon=\frac{1}{24}} = \lambda \left( 1 + \ln \left( \frac{26068394603446272 \sqrt[6]{\frac{7}{247}} \sqrt[3]{11}}{3915^{5/6} 101898752449325\sqrt{5}} \right) \right),$$

$$\frac{\mu - BM_{\nu=3,\epsilon=\frac{1}{24}}}{BM_{\nu=3,\epsilon=\frac{1}{24}} - m} = \frac{\lambda - \lambda \left( 1 + \ln \left( \frac{26068394603446272 \sqrt[6]{\frac{7}{247}} \sqrt[3]{11}}{3915^{5/6} 101898752449325\sqrt{5}} \right) \right)}{\lambda \left( 1 + \ln \left( \frac{26068394603446272 \sqrt[6]{\frac{7}{247}} \sqrt[3]{11}}{3915^{5/6} 101898752449325\sqrt{5}} \right) \right) - \ln 2\lambda} = \frac{\ln \left( \frac{26068394603446272 \sqrt[6]{\frac{7}{247}} \sqrt[3]{11}}{3915^{5/6} 101898752449325\sqrt{5}} \right)}{1 - \ln(2) + \ln \left( \frac{26068394603446272 \sqrt[6]{\frac{7}{247}} \sqrt[3]{11}}{3915^{5/6} 101898752449325\sqrt{5}} \right)} \approx 0.103.$$

The biases of  $rm_{d \approx 0.103, \nu=3, \epsilon=\frac{1}{24}, BM, m}$  for distributions with skewness between those of the exponential and symmetric distributions are tiny (SI Dataset S1).  $rm_{d \approx 0.103, \nu=3, \epsilon=\frac{1}{24}, BM, m}$  exhibits excellent performance for all these common unimodal distributions (SI Dataset S1).

The recombined mean is a recombined  $I$ -statistic. Consider an  $I$ -statistic whose LEs are percentiles of a distribution obtained by plugging  $LU$ -statistics into a cumulative distribution function,  $I$  is defined with arithmetic operations, constants, and quantile functions, such an estimator is classified as a quantile  $I$ -statistic. One version of the quantile  $I$ -statistic can be defined as  $QI_{d,h_{\mathbf{k}},\mathbf{k},k,\epsilon,\gamma,n,LU} :=$

$$\begin{cases} \hat{Q}_{n,h_{\mathbf{k}}} \left( \left( \hat{F}_{n,h_{\mathbf{k}}}(LU) - \frac{\gamma}{1+\gamma} \right) d + \hat{F}_{n,h_{\mathbf{k}}}(LU) \right) & \hat{F}_{n,h_{\mathbf{k}}}(LU) \geq \frac{\gamma}{1+\gamma} \\ \hat{Q}_{n,h_{\mathbf{k}}} \left( \hat{F}_{n,h_{\mathbf{k}}}(LU) - \left( \frac{\gamma}{1+\gamma} - \hat{F}_{n,h_{\mathbf{k}}}(LU) \right) d \right) & \hat{F}_{n,h_{\mathbf{k}}}(LU) < \frac{\gamma}{1+\gamma} \end{cases}$$

where  $LU$  is  $LU_{\mathbf{k},k,\epsilon,\gamma,n}$ ,  $\hat{F}_{n,h_{\mathbf{k}}}(x)$  is the empirical cumulative distribution function of the  $h_{\mathbf{k}}$  kernel distribution,  $\hat{Q}_{n,h_{\mathbf{k}}}$  is the quantile function of the  $h_{\mathbf{k}}$  kernel distribution.

Similarly, the quantile mean can be defined as  $qm_{d,k,\epsilon,\gamma,n,WL} := QI_{d,h_{\mathbf{k}}=x,\mathbf{k}=1,k,\epsilon,\gamma,n,LU=WL}$ . Moreover, in extreme right-skewed heavy-tailed distributions, if the calculated percentile exceeds  $1 - \epsilon$ , it will be adjusted to  $1 - \epsilon$ . In a left-skewed distribution, if the obtained percentile is smaller than  $\gamma\epsilon$ , it will also be adjusted to  $\gamma\epsilon$ . Without loss of generality, in the following discussion, only the case where  $\hat{F}_n(WL_{k,\epsilon,\gamma,n}) \geq \frac{\gamma}{1+\gamma}$  is considered. The most popu-

lar method for computing the sample quantile function was proposed by Hyndman and Fan in 1996 (16). Another widely used method for calculating the sample quantile function involves employing linear interpolation of modes corresponding to the order statistics of the uniform distribution on the interval  $[0, 1]$ , i.e.,  $\hat{Q}_n(p) = X_{[h]} + (h - [h])(X_{[h]} - X_{[h-1]})$ ,  $h = (n-1)p + 1$ . To minimize the finite sample bias, here, the inverse function of  $\hat{Q}_n$  is deduced as  $\hat{F}_n(x) := \frac{1}{n} \left( \frac{x - X_{cf}}{X_{cf+1} - X_{cf}} + cf \right)$ , based on Hyndman and Fan's definition, or  $\hat{F}_n(x) := \frac{1}{n-1} \left( cf - 1 + \frac{x - X_{cf}}{X_{cf+1} - X_{cf}} \right)$ , based on the latter definition, where  $cf = \sum_{i=1}^n \mathbf{1}_{X_i \leq x}$ ,  $\mathbf{1}_A$  is the indicator of event  $A$ .

The quantile mean uses the location-scale invariant in a different way, as shown in the subsequent proof.

**Theorem A.2.**  $qm_{d = \frac{F(\mu) - F(WL_{k,\epsilon,\gamma})}{F(WL_{k,\epsilon,\gamma}) - \frac{\gamma}{1+\gamma}}, k, \epsilon, \gamma, WL}$  is a consistent mean estimator for a location-scale distribution provided that the means are finite and  $F(\mu)$ ,  $F(WL_{k,\epsilon,\gamma})$  and  $\frac{\gamma}{1+\gamma}$  are all within the range of  $[\gamma\epsilon, 1 - \epsilon]$ , where  $\mu$  and  $WL_{k,\epsilon,\gamma}$  are location parameters from that location-scale distribution. If  $WL = SWHLM$ ,  $qm$  is also consistent for any symmetric distributions.

*Proof.* When  $F(WL_{k,\epsilon,\gamma}) \geq \frac{\gamma}{1+\gamma}$ , the solution of  $(F(WL_{k,\epsilon,\gamma}) - \frac{\gamma}{1+\gamma})d + F(WL_{k,\epsilon,\gamma}) = F(\mu)$  is  $d = \frac{F(\mu) - F(WL_{k,\epsilon,\gamma})}{F(WL_{k,\epsilon,\gamma}) - \frac{\gamma}{1+\gamma}}$ . The  $d$  value for the case where  $F(WL_{k,\epsilon,\gamma}) < \frac{\gamma}{1+\gamma}$  is the same. The definitions of the location and scale parameters are such that they must satisfy  $F(x; \lambda, \mu) = F(\frac{x-\mu}{\lambda}; 1, 0)$ , then  $F(WL(k, \epsilon, \gamma); \lambda, \mu) = F(\frac{\lambda WL_0(k, \epsilon, \gamma) + \mu - \mu}{\lambda}; 1, 0) = F(WL_0(k, \epsilon, \gamma); 1, 0)$ . It follows that the percentile of any weighted  $L$ -statistic is free of  $\lambda$  and  $\mu$  for a location-scale distribution. Therefore  $d$  in  $qm$  is also invariably a constant. For the symmetric case,  $F(SWHLM_{k,\epsilon}) = F(\mu) = F(Q(\frac{1}{2})) = \frac{1}{2}$  is valid for any symmetric distribution with a finite second moment, as the same values correspond to same percentiles. Then,  $qm_{d,k,\epsilon,SWHLM} = F^{-1} \left( (F(SWHLM_{k,\epsilon}) - \frac{1}{2})d + F(\mu) \right) = F^{-1} \left( 0 + F(\mu) \right) = \mu$ . To avoid inconsistency due to post-adjustment,  $F(\mu)$ ,  $F(WL_{k,\epsilon,\gamma})$  and  $\frac{\gamma}{1+\gamma}$  must reside within the range of  $[\gamma\epsilon, 1 - \epsilon]$ . All results are now proven.  $\square$

The cdf of the Pareto distribution is  $F_{Par}(x) = 1 - \left(\frac{x_m}{x}\right)^\alpha$ . So, set the  $d$  value in  $qm$  with two arbitrary percentiles  $p_1$  and  $p_2$ ,  $d_{Par,qm} =$

$$1 - \left(\frac{x_m}{\frac{x_m}{\alpha-1}}\right)^\alpha - \left(1 - \left(\frac{x_m}{x_m(1-p_1)^{-\frac{1}{\alpha}}}\right)^\alpha\right)^\alpha = \frac{\left(1 - \left(\frac{x_m}{x_m(1-p_1)^{-\frac{1}{\alpha}}}\right)^\alpha\right)^\alpha - \left(1 - \left(\frac{x_m}{x_m(1-p_2)^{-\frac{1}{\alpha}}}\right)^\alpha\right)^\alpha}{1 - \left(\frac{\alpha-1}{\alpha}\right)^\alpha - p_1}.$$

The  $d$  value in  $qm$  for the exponential distribution is always identical to  $d_{Par,qm}$  as  $\alpha \rightarrow \infty$ , since  $\lim_{\alpha \rightarrow \infty} \left(\frac{\alpha-1}{\alpha}\right)^\alpha = \frac{1}{e}$  and the cdf of the exponential distribution is  $F_{exp}(x) = 1 - e^{-\lambda^{-1}x}$ , then

$$d_{exp,qm} = \frac{(1-e^{-1}) - \left(1 - e^{-\ln\left(\frac{1}{1-p_1}\right)}\right)}{\left(1 - e^{-\ln\left(\frac{1}{1-p_1}\right)}\right) - \left(1 - e^{-\ln\left(\frac{1}{1-p_2}\right)}\right)} = \frac{1 - \frac{1}{e} - p_1}{p_1 - p_2}.$$

So, for the Weibull, gamma, Pareto, lognormal and generalized

Gaussian distribution,  $qm_{d = \frac{F_{exp}(\mu) - F_{exp}(SWHLM_{k,\epsilon})}{F_{exp}(SWHLM_{k,\epsilon}) - \frac{1}{2}}, k, \epsilon, SWHLM}$

is also consistent for at least one particular case, provided that  $\mu$  and  $SWHLM_{k,\epsilon}$  are different location parameters from an exponential distribution and  $F(\mu)$ ,  $F(SWHLM_{k,\epsilon})$  and  $\frac{1}{2}$  are all within the range of  $[\epsilon, 1 - \epsilon]$ . Also let  $SWHLM_{k,\epsilon,\gamma} = BM_{\nu=3,\epsilon=\frac{1}{24}}$

and  $\mu = \lambda$ , then  $d = \frac{F_{exp}(\mu) - F_{exp}(BM_{\nu=3,\epsilon=\frac{1}{24}})}{F_{exp}(BM_{\nu=3,\epsilon=\frac{1}{24}}) - \frac{1}{2}} =$

$$\frac{-e^{-1} + e^{-\left(1 + \ln\left(\frac{26068394603446272 \sqrt[3]{\frac{7}{247}} \sqrt[3]{11}}{391^5/6 \cdot 101898752449325\sqrt{5}}\right)\right)}}{-e^{-\frac{1}{2}} - e^{-\left(1 + \ln\left(\frac{26068394603446272 \sqrt[3]{\frac{7}{247}} \sqrt[3]{11}}{391^5/6 \cdot 101898752449325\sqrt{5}}\right)\right)}} = \frac{101898752449325\sqrt{5} \sqrt[3]{\frac{247}{7}} 391^5/6 - \frac{1}{e}}{26068394603446272 \sqrt[3]{11} e - \frac{1}{e}} \approx 0.088. F_{exp}(\mu),$$

$F_{exp}(BM_{\nu=3,\epsilon=\frac{1}{24}})$  and  $\frac{1}{2}$  are all within the range of  $[\frac{1}{24}, \frac{23}{24}]$ .  $qm_{d \approx 0.088, \nu=3, \epsilon=\frac{1}{24}, BM}$  works better in the fat-tail scenarios (SI Dataset S1). Theorem A.1 and A.2 show that  $rm_{d \approx 0.103, \nu=3, \epsilon=\frac{1}{24}, BM}$  and  $qm_{d \approx 0.088, \nu=3, \epsilon=\frac{1}{24}, BM}$  are both consistent mean estimators for any symmetric distribution and the exponential distribution with finite second moments. It's obvious that the asymptotic breakdown points of  $rm_{d \approx 0.103, \nu=3, \epsilon=\frac{1}{24}, BM}$  and  $qm_{d \approx 0.088, \nu=3, \epsilon=\frac{1}{24}, BM}$  are both  $\frac{1}{24}$ . Therefore they are all invariant means.

To study the impact of the choice of WLS in  $rm$  and  $qm$ , it is constructive to recall that a weighted  $L$ -statistic is a combination of order statistics. While using a less-biased weighted  $L$ -statistic can generally enhance performance (SI Dataset S1), there is a greater risk of violation in the semiparametric framework. However, the mean- $WA_{\epsilon,\gamma}$ -median inequality is robust to slight fluctuations of the QA function of the underlying distribution. Suppose for a right-skewed distribution, the QA function is generally decreasing with respect to  $\epsilon$  in  $[0, u]$ , but increasing in  $[u, \frac{1}{1+\gamma}]$ , since all quantile averages with breakdown points from  $\epsilon$  to  $\frac{1}{1+\gamma}$  will be included in the computation of  $WA_{\epsilon,\gamma}$ , as long as  $\frac{1}{1+\gamma} - u \ll \frac{1}{1+\gamma} - \gamma\epsilon$ , and other portions of the QA function satisfy the inequality constraints that define the  $\nu$ th  $\gamma$ -orderliness on which the  $WA_{\epsilon,\gamma}$  is based, if  $0 \leq \gamma \leq 1$ , the mean- $WA_{\epsilon,\gamma}$ -median inequality still holds. This is due to the violation of  $\nu$ th  $\gamma$ -orderliness being bounded, when  $0 \leq \gamma \leq 1$ , as shown in REDS I and therefore cannot be extreme for unimodal distributions with finite second moments. For instance, the SQA function of the Weibull distribution is non-monotonic with respect to  $\epsilon$  when the shape parameter  $\alpha > \frac{1}{1-\ln(2)} \approx 3.259$  as shown in the SI Text of REDS I, the violation of the second and third orderliness starts near this parameter as well, yet the mean- $BM_{\nu=3,\epsilon=\frac{1}{24}}$ -median inequality retains valid when  $\alpha \leq 3.387$ . Another key factor in determining the risk of violation of orderliness is the skewness of the distribution. In REDS I, it was demonstrated that in a family of distributions differing by a skewness-increasing transformation in van Zwet's sense, the violation of orderliness, if it happens, only occurs as the distribution nears symmetry (12). When  $\gamma = 1$ , the over-corrections in  $rm$  and  $qm$  are dependent on the  $SWA_\epsilon$ -median difference, which can be a reasonable measure of skewness after standardization (11, 13), implying that the over-correction is often tiny with moderate  $d$ . This qualitative analysis suggests the general reliability of  $rm$  and  $qm$  based on the mean- $WA_{\epsilon,\gamma}$ -median inequality, es-

pecially for unimodal distributions with finite second moments when  $0 \leq \gamma \leq 1$ . Extending this rationale to other weighted  $L$ -statistics is possible, since the  $\gamma$ - $U$ -orderliness can also be bounded with certain assumptions, as discussed previously.

Consider two continuous distributions belonging to the same location-scale family, according to Theorem ??, their corresponding  $k$ th central moment kernel distributions only differ in scaling. Define the recombined  $k$ th central moment as  $rkmd_{d,k_1,k_2,\epsilon=\min(\epsilon_1,\epsilon_2),\gamma_1,\gamma_2,n,WHLkm_1,WHLkm_2} := RL_{d,hk=\psi_k,k_1=k,k_2=k,k_1,k_2,\epsilon_1,\epsilon_2,\gamma_1,\gamma_2,n,LU_1=WHLkm_1,LU_2=WHLkm_2}$ . Then, assuming finite  $k$ th central moment and applying the same logic as in Theorem A.1,  $rkmd_{d,k_1,k_2,\epsilon=\min(\epsilon_1,\epsilon_2),\gamma_1,\gamma_2,WHLkm_1,WHLkm_2}$  is a consistent  $k$ th central moment estimator for a location-scale distribution, where  $\mu_k$ ,  $WHLkm_{1k_1,\epsilon_1,\gamma_1}$ , and  $WHLkm_{2k_2,\epsilon_2,\gamma_2}$  are different  $k$ th central moment parameters from that location-scale distribution. Similarly, the quantile will not change after scaling. The quantile  $k$ th central moment is thus defined as

$$qkm_{d,k,\epsilon,\gamma,n,WHLkm} := QI_{d,hk=\psi_k,k=k,\epsilon,\gamma,n,LU=WHLkm}$$

$qkm_{d,k,\epsilon,\gamma,n,WHLkm}$  is also a consistent  $k$ th central moment estimator for a location-scale distribution provided that the  $k$ th central moment is finite and  $F_{\psi_k}(\mu_k)$ ,  $F_{\psi_k}(WHLkm_{k,\epsilon,\gamma})$  and  $\frac{\gamma}{1+\gamma}$  are all within the range of  $[\gamma\epsilon, 1 - \epsilon]$ , where  $\mu_k$  and  $WHLkm_{k,\epsilon,\gamma}$  are different  $k$ th central moment parameters from that location-scale distribution. According to Theorem ??, if the original distribution is unimodal, the central moment kernel distribution is always a heavy-tailed distribution, as the degree term amplifies its skewness and tailedness. From the better performance of the quantile mean in heavy-tailed distributions, the quantile  $k$ th central moments are generally better than the recombined  $k$ th central moments regarding asymptotic bias.

Finally, the recombined standardized  $k$ th moment is defined to be

$$rskm_{\epsilon=\min(\epsilon_1,\epsilon_2),k_1,k_2,k_3,k_4,\gamma_1,\gamma_2,\gamma_3,\gamma_4,n,WHLkm_1,WHLkm_2,WHLvar_1,WHLvar_2} = \frac{rkmd_{d,k_1,k_2,\epsilon_1,\gamma_1,\gamma_2,n,WHLkm_1,WHLkm_2}}{(rvar_{d,k_3,k_4,\epsilon_2,\gamma_3,\gamma_4,n,WHLvar_1,WHLvar_2})^{k/2}}$$

The quantile standardized  $k$ th moment is defined similarly,

$$qskm_{\epsilon=\min(\epsilon_1,\epsilon_2),k_1,k_2,\gamma_1,\gamma_2,n,WHLkm,WHLvar} := \frac{qkm_{d,k_1,\epsilon_1,\gamma_1,n,WHLkm_i=i+1}}{(qvar_{d,k_2,\epsilon_2,\gamma_2,n,WHLvar})^{k/2}}$$

## B. A shape-scale distribution as the consistent distribution.

In the last section, the parametric robust estimation is limited to a location-scale distribution, with the location parameter often being omitted for simplicity. For improved fit to observed skewness or kurtosis, shape-scale distributions with shape parameter ( $\alpha$ ) and scale parameter ( $\lambda$ ) are commonly utilized. Weibull, gamma, Pareto, lognormal, and generalized Gaussian distributions (when  $\mu$  is a constant) are all shape-scale unimodal distributions. Furthermore, if either the shape parameter  $\alpha$  or the skewness or kurtosis is constant, the shape-scale distribution is reduced to a location-scale distribution. Let  $D(|skewness|, kurtosis, \mathbf{k}, etype, dtype, n) = d_{i\mathbf{k}m}$  denote the function to specify  $d$  values, where the first input is the absolute value of the skewness, the second input is the kurtosis,

the third is the order of the central moment (if  $\mathbf{k} = 1$ , the mean), the fourth is the type of estimator, the fifth is the type of consistent distribution, and the sixth input is the sample size. For simplicity, the last three inputs will be omitted in the following discussion. Hold in awareness that since skewness and kurtosis are interrelated, specifying  $d$  values for a shape-scale distribution only requires either skewness or kurtosis, while the other may be also omitted. Since many common shape-scale distributions are always right-skewed (if not, only the right-skewed or left-skewed part is used for calibration, while the other part is omitted), the absolute value of the skewness should be the same as the skewness of these distributions. This setting also handles the left-skew scenario well.

For recombined moments up to the fourth ordinal, the object of using a shape-scale distribution as the consistent distribution is to find solutions for the system of equations

$$\begin{cases} rm(WHLM, \gamma_m, D(|rskew|, rkurt, 1)) = \mu \\ rvar(WHLvar, \gamma mvar, D(|rskew|, rkurt, 2)) = \mu_2 \\ rtm(WHLtm, \gamma mtm, D(|rskew|, rkurt, 3)) = \mu_3 \\ rfm(WHLfm, \gamma mfm, D(|rskew|, rkurt, 4)) = \mu_4 \\ rskew = \frac{\mu_3}{\mu_2} \\ rkurt = \frac{\mu_4}{\mu_2^2} \end{cases}, \quad 349$$

where  $\mu_2$ ,  $\mu_3$  and  $\mu_4$  are the population second, third and fourth central moments.  $|rskew|$  and  $rkurt$  should be the invariant points of the functions  $\zeta(|rskew|) = \left| \frac{rtm(WHLtm, \gamma mtm, D(|rskew|, 3))}{rvar(WHLvar, \gamma mvar, D(|rskew|, 2))^{3/2}} \right|$  and  $\varkappa(rkurt) = \frac{rfm(WHLfm, \gamma mfm, D(rkurt, 4))}{rvar(WHLvar, \gamma mvar, D(rkurt, 2))^2}$ . Clearly, this is an overdetermined nonlinear system of equations, given that the skewness and kurtosis are interrelated for a shape-scale distribution. Since an overdetermined system constructed with random coefficients is almost always inconsistent, it is natural to optimize them separately using the fixed-point iteration (see Algorithm 1, only  $rkurt$  is provided, others are the same).

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### Algorithm 1 $rkurt$ for a shape-scale distribution

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**Input:**  $D$ ;  $WHLvar$ ;  $WHLfm$ ;  $\gamma mvar$ ;  $\gamma mfm$ ;  $maxit$ ;  $\delta$   
**Output:**  $rkurt_{i-1}$   
1:  $rkurt_i \leftarrow \varkappa(kurtosis_{max})$   $\triangleright$  Using the maximum kurtosis available in  $D$  as an initial guess.  
**repeat**  
2:  $rkurt_i \leftarrow \varkappa(kurtosis_{max})$   $\triangleright$  Using the maximum kurtosis available in  $D$  as an initial guess.  
**repeat**  
3:  $rkurt_i \leftarrow \varkappa(kurtosis_{max})$   $\triangleright$  Using the maximum kurtosis available in  $D$  as an initial guess.  
**until**  $i > maxit$  or  $|rkurt_i - rkurt_{i-1}| < \delta$   $\triangleright$   $maxit$  is the maximum number of iterations,  $\delta$  is a small positive number.

---

The following theorem shows the validity of Algorithm 1.

**Theorem B.1.** Assuming  $\gamma = 1$  and  $m\mathbf{k}m$ s, where  $2 \leq \mathbf{k} \leq 4$ , are all equal to zero,  $|rskew|$  and  $rkurt$ , defined as the largest attracting fixed points of the functions  $\zeta(|rskew|)$  and  $\varkappa(rkurt)$ , are consistent estimators of  $\tilde{\mu}_3$  and  $\tilde{\mu}_4$  for a shape-scale distribution whose  $k$ th central moment kernel distributions are  $U$ -congruent, as long as they are within the domain of  $D$ , where  $\tilde{\mu}_3$  and  $\tilde{\mu}_4$  are the population skewness and kurtosis, respectively.

370 *Proof.* Without loss of generality, only  $rkurt$  is considered, 424  
371 while the logic for  $|rskew|$  is the same. Additionally, the 425  
372 second central moments of the underlying sample distribu- 426  
373 tion and consistent distribution are assumed to be 1, with 427  
374 other cases simply multiplying a constant factor according 428  
375 to Theorem ???. From the definition of  $D$ ,  $\frac{\alpha(rkurt_D)}{rkurt_D} =$  429  
376  $\frac{f_{m_D} - SWHLf_{m_D}}{SWHLf_{m_D} - m_{f_{m_D}}} \frac{(SWHLf_{m_D} - m_{f_{m_D}}) + SWHLf_{m_D}}{rkurt_D \left( \frac{var_D - SWHLvar_D}{SWHLvar_D - m_{var_D}} (SWHLvar - m_{var}) + SWHLvar \right)^2}$ , where 430  
377 the subscript  $D$  indicates that the estimates are from the 431  
378 central moment kernel distributions generated from the consis- 432  
379 tent distribution, while other estimates are from the underlying 433  
380 distribution of the sample. 434

381 Then, assuming the  $mkms$  are all equal to zero and 435  
382  $var_D = 1$ ,  $\frac{\alpha(rkurt_D)}{rkurt_D} = \frac{f_{m_D} - SWHLf_{m_D}}{SWHLf_{m_D}} \frac{(SWHLf_{m_D}) + SWHLf_{m_D}}{rkurt_D \left( \frac{SWHLvar}{SWHLvar_D} \right)^2} =$  436  
383  $\frac{(f_{m_D} - SWHLf_{m_D} + 1)(SWHLf_{m_D})}{f_{m_D} \left( \frac{SWHLvar}{SWHLvar_D} \right)^2} = \frac{SWHLf_{m_D} SWHLvar_D^2}{SWHLf_{m_D} SWHLvar^2} =$  437  
384  $\frac{SWHLf_{m_D}}{SWHLvar^2} = \frac{SWHLkurt}{SWHLkurt_D}$ . Since  $SWHLf_{m_D}$  are from the 438  
385 same fourth central moment kernel distribution as  $f_{m_D} =$  439  
386  $rkurt_D var_D^2$ , according to the definition of  $U$ -congruence, 440  
387 an increase in  $f_{m_D}$  will also result in an increase in 441  
388  $SWHLf_{m_D}$ . Combining with Theorem ??,  $SWHLkurt$  is 442  
389 a measure of kurtosis that is invariant to location and scale, 443  
390 so  $\lim_{rkurt_D \rightarrow \infty} \frac{\alpha(rkurt_D)}{rkurt_D} < 1$ . As a result, if there is at 444  
391 least one fixed point, let the largest one be  $fix_{max}$ , then 445  
392 it is attracting since  $\left| \frac{\partial(\alpha(rkurt_D))}{\partial(rkurt_D)} \right| < 1$  for all  $rkurt_D \in$  446  
393  $[fix_{max}, kurtosis_{max}]$ , where  $kurtosis_{max}$  is the maximum 447  
394 kurtosis available in  $D$ . 448  
395  $\square$  449

396 As a result of Theorem B.1, assuming continuity,  $mkms$  455  
397 are all equal to zero, and  $U$ -congruence of the central moment 456  
398 kernel distributions, Algorithm 1 converges surely provided 457  
399 that a fixed point exists within the domain of  $D$ . At this 458  
400 stage,  $D$  can only be approximated through a Monte Carlo 459  
401 study. The continuity of  $D$  can be ensured by using linear 460  
402 interpolation. One common encountered problem is that the 461  
403 domain of  $D$  depends on both the consistent distribution and 462  
404 the Monte Carlo study, so the iteration may halt at the bound- 463  
405 ary if the fixed point is not within the domain. However, 464  
406 by setting a proper maximum number of iterations, the algo- 465  
407 rithm can return the optimal boundary value. For quantile 466  
408 moments, the logic is similar, if the percentiles do not exceed 467  
409 the breakdown point. If this is the case, consistent estimation 468  
410 is impossible, and the algorithm will stop due to the maximum 469  
411 number of iterations. The fixed point iteration is, in principle, 470  
412 similar to the iterative reweighing in Huber  $M$ -estimator, but 471  
413 an advantage of this algorithm is that it is solely related to 472  
414 the inputs in Algorithm 1 and is independent of the sample 473  
415 size. Since they are consistent for a shape-scale distribution, 474  
416  $|rskew|$  can specify  $d_{rm}$  and  $d_{tm}$ ,  $rkurt$  can specify  $d_{var}$  and 475  
417  $d_{rfm}$ . Algorithm 1 enables the robust estimations of all four 476  
418 moments to reach a near-consistent level for common unimodal 477  
419 distributions (Table 1, SI Dataset S1), just using the Weibull 478  
420 distribution as the consistent distribution. 479

421 **C. Critical points, lines, and Multiple consistent distributions.** 480  
422 In 1895, Pearson considered how to construct probability distri- 481  
423 butions in which the skewness and kurtosis could be adjusted 482

424 equally freely. The Pearson distribution is a family of unimodal 425  
426 continuous probability distribution functions that satisfy the 427  
428 following differential equation  $\frac{dP(x)}{dx} = -\frac{(a_0 + a_1x + a_2x^2)P(x)}{b_0 + b_1x + b_2x^2 + b_3x^3}$ , 429  
430 where  $P(x)$  is the pdf of the Pearson distribution, and  $a_0$ , 431  
432  $a_1$ ,  $a_2$ ,  $b_0$ ,  $b_1$ ,  $b_2$ , and  $b_3$  are constants that determine the 433  
434 specific type of Pearson distribution. This differential equa- 435  
436 tion ensures that the distribution is unimodal and that the 436  
437 density function is continuous. The Pearson family subsumes 437  
438 many common unimodal distributions, e.g., beta distribution, 438  
439 Cauchy distribution, Chi-squared distribution, continuous uni- 439  
440 form distribution, gamma distribution, inverse-chi-squared 440  
441 distribution, inverse-gamma distribution, normal distribution, 441  
442 and tudent's t-distribution. Plotting the kurtosis-skewness 442  
443 lines of the five shape-scale unimodal distributions used in 443  
444 this series into the Pearson diagram. One can immediately 444  
445 identify two critical points, one is 9-2, which is the exponential 445  
446 distribution or the limiting form of the Pareto distribution, 446  
447 another is the normal distribution. This further explains why 447  
448 using the exponential distribution as the consistent distribu- 448  
449 tion has excellent performance even better than using the 449  
450 Weibull distribution as the consistent distribution, since it 450  
451 lays in this critical point. Then, we further consider the criti- 451  
452 cal lines in the Pearson diagram (Figure 1). Consider if the 452  
453 underlying distribution is not the Weibull distribution, but 453  
454 the gamma distribution, then, using the Weibull distribution 454  
455 as an initial guess, calculating the Euclidean distance of the 455  
456 obtained kurtosis-skewness to the kurtosis-skewness lines of 456  
457 other consistent distributions, choosing the one that has the 457  
458 smallest distance, then recalculate the kurtosis-skewness ac- 458  
459 cording to the new consistent distribution, iterately, we can 459  
460 use multiple distributions as consistent distributions. 460  
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484 **D. Root mean square error** .Lai, Robbins, and Yu (1983) 455  
485 proposed an estimator that adaptively chooses the mean or 456  
486 median in a symmetric distribution and showed that the choice 457  
487 is typically as good as the better of the sample mean and 458  
488 median regarding variance (17). Another approach which can 459  
489 be dated back to Laplace (1812) (18) is using  $w\bar{x} + (1-w)m_n$  as 460  
490 a location estimator and  $w$  is deduced to achieve optimal vari- 461  
491 ance. Inspired by Lai et al's approach (17), in this study, for 462  
492  $rkurt$ , there are 364 combinations based on 14  $SWHLfms$  and 463  
493 26  $SWHLvars$  (SI Text). Each combination has a root mean 464  
494 square error (RMSE) for a single-parameter distribution, which 465  
495 can be inferred using a Monte Carlo study. For  $qkurt$ , there 466  
496 are another 364 combinations, but if the percentiles of quantile 467  
497 moments exceed the breakdown point, that combination is 468  
498 excluded. Then, the combination with the smallest RMSE, cal- 469  
499 ibrated by a two-parameter distribution, is chosen. Similar to 470  
500 Subsection B, let  $I(kurtosis, dtype, n) = ikurt_{WHLfm, WHLvar}$  471  
501 represent these relationships. In this article, the breakdown 472  
502 points of the  $SWHLms$  in  $SWHLkm$  were adjusted to ensure 473  
503 the overall breakdown points were  $\frac{1}{24}$ , as detailed in Theorem 474  
504 ???. There are two approaches to determine  $ikurt$ . The first 475  
505 one is computing all 364+364  $rkurt$  and  $qkurt$ , and then, since 476  
506  $\lim_{ikurt \rightarrow \infty} \frac{I(ikurt)}{ikurt} < 1$ , the same fix point iteration algorithm 477  
507 as Algorithm 1 can be used to choose the RMSE-optimum 478  
508 combination. The only difference is that unlike  $D$ ,  $I$  is defined 479  
509 to be discontinuous but linear interpolation can also ensure 480  
510 continuity. The second approach is shown in SI Algorithm 481  
511 2. The RMSEs of these  $ikurt$  from the two approaches can 482  
512 be further determined by a Monte Carlo study. Algorithm 1 483

**Table 1. Evaluation of invariant moments for five common unimodal distributions in comparison with current popular methods**

Errors	$\bar{x}$	TM	H-L	SM	HM	WM	SQM	BM	MoM	MoRM	$m$ HLM	$rm_{exp,BM}$	$qm_{exp,BM}$
WASAB	0.000	0.107	0.088	0.078	0.078	0.066	0.048	0.048	0.034	0.035	0.034	0.002	0.003
WRMSE	0.014	0.111	0.092	0.083	0.083	0.070	0.053	0.053	0.041	0.041	0.038	0.017	0.018
WASB <sub><math>n=5184</math></sub>	0.000	0.108	0.089	0.078	0.079	0.066	0.048	0.048	0.034	0.036	0.033	0.002	0.003
WSE $\vee$ WSSE	0.014	0.014	0.014	0.015	0.014	0.014	0.014	0.015	0.017	0.014	0.014	0.017	0.017

Errors	HFM <sub><math>\mu</math></sub>	MP <sub><math>\mu</math></sub>	$rm$	$qm$	$im$	$var$	$var_{bs}$	Tsd <sup>2</sup>	HFM <sub><math>\mu_2</math></sub>	MP <sub><math>\mu_2</math></sub>	$rvar$	$quar$	$ivar$
WASAB	0.037	0.043	0.001	0.002	0.001	0.000	0.000	0.200	0.027	0.042	0.005	0.018	0.003
WRMSE	0.049	0.055	0.015	0.015	0.014	0.017	0.017	0.198	0.042	0.062	0.019	0.026	0.019
WASB <sub><math>n=5184</math></sub>	0.038	0.043	0.001	0.002	0.001	0.000	0.001	0.198	0.027	0.043	0.005	0.018	0.003
WSE $\vee$ WSSE	0.018	0.021	0.015	0.015	0.014	0.017	0.017	0.015	0.024	0.032	0.018	0.017	0.018

Errors	$tm$	$tm_{bs}$	HFM <sub><math>\mu_3</math></sub>	MP <sub><math>\mu_3</math></sub>	$rtm$	$qtm$	$itm$	$fm$	$fm_{bs}$	HFM <sub><math>\mu_4</math></sub>	MP <sub><math>\mu_4</math></sub>	$rfm$	$qfm$	$ifm$
WASAB	0.000	0.000	0.052	0.059	0.006	0.083	0.034	0.000	0.000	0.037	0.046	0.024	0.038	0.011
WRMSE	0.019	0.018	0.063	0.074	0.018	0.083	0.044	0.026	0.023	0.049	0.062	0.037	0.043	0.029
WASB <sub><math>n=5184</math></sub>	0.001	0.003	0.052	0.059	0.007	0.082	0.038	0.001	0.009	0.037	0.047	0.024	0.036	0.013
WSE $\vee$ WSSE	0.019	0.018	0.021	0.091	0.015	0.012	0.017	0.024	0.021	0.020	0.027	0.021	0.020	0.022

The first table presents the use of the exponential distribution as the consistent distribution for five common unimodal distributions: Weibull, gamma, Pareto, lognormal, and generalized Gaussian distributions. Popular robust mean estimators discussed in REDS 1 were used as comparisons. The breakdown points of mean estimators in the first table, besides H-L estimator and Huber  $M$ -estimator, are all  $\frac{1}{8}$ . The second and third tables present the use of the Weibull distribution as the consistent distribution not plus/plus using the lognormal distribution for the odd ordinal moments optimization and the generalized Gaussian distribution for the even ordinal moments optimization. SQM is the robust mean estimator used in recombined/quantile moments. Unbiased sample central moments ( $var$ ,  $tm$ ,  $fm$ ),  $U$ -central moments with quasi-bootstrap ( $var_{bs}$ ,  $tm_{bs}$ ,  $fm_{bs}$ ), and other estimators were used as comparisons. The generalized Gaussian distribution was excluded for He and Fung  $M$ -Estimator and Marks percentile estimator, since the logarithmic function does not produce results for negative inputs. The breakdown points of estimators in the second and third table, besides  $M$ -estimators and percentile estimator, are all  $\frac{1}{24}$ . The tables include the average standardized asymptotic bias (ASAB, as  $n \rightarrow \infty$ ), root mean square error (RMSE, at  $n = 5184$ ), average standardized bias (ASB, at  $n = 5184$ ) and variance (SE  $\vee$  SSE, at  $n = 5184$ ) of these estimators, all reported in the units of the standard deviations of the distribution or corresponding kernel distributions.  $W$  means that the results were weighted by the number of Google Scholar search results on May 30, 2022 (including synonyms). The calibrations of  $d$  values and the computations of ASAB, ASB, and SSE were described in Subsection D. ?? and SI Methods. Detailed results and related codes are available in SI Dataset S1 and [GitHub](#).

can also be used to determine the optimum choice among the two approaches. The  $364+364 rkurt$  and  $qkurt$  can form a vector,  $Vkurt$ , where the  $Q_{Vkurt}(\frac{1}{5})$  to  $Q_{Vkurt}(\frac{4}{5})$  can be used to determine the  $d$  values of  $rkms$  and  $qkms$ . The RMSEs of those  $rkms$  and  $qkms$  can also be estimated by a Monte Carlo study and the estimator with the smallest RMSE of each ordinal is named as  $ikm$ . When  $k$  is even, the  $ikurt$  determined by  $ism$  (detailed in the SI Text) is used to determine  $ikm$ . This approach yields results that are often nearly optimal (SI Datasets S1). The estimations of skewness and  $ikm$ , when  $k$  is odd, follow the same logic.

In general, the variances of invariant central moments are much smaller than those of corresponding unbiased sample central moments (deduced by Cramér (19, 20)), except that of the corresponding second central moment (Table 1).

## Discussion

Statistics, encompassing the collection, analysis, interpretation, and presentation of data, has evolved over time, with various approaches emerging to meet challenges in practice. Among these approaches, the use of probability models and measures of random variables for data analysis is often considered the core of statistics. While the early development of statistics was focused on parametric methods, there were two main approaches to point estimation. The Gauss–Markov theorem (1, 21) states the principle of minimum variance unbiased estimation which was further enriched by Neyman (1934) (22), Rao (1945) (23), Blackwell (1947) (24), and Lehmann and Scheffé (1950, 1955) (25, 26). Maximum likelihood was first introduced by Fisher in 1922 (27) in a multinomial model and

later generalized by Cramér (1946), Hájek (1970), and Le Cam (1972) (19, 28, 29). In 1939, Wald (30) combined these two principles and suggested the use of minimax estimates, which involve choosing an estimator that minimizes the maximum possible loss. Following Huber’s seminal work (3),  $M$ -statistics have dominated the field of parametric robust statistics for over half a century. Nonparametric methods, e.g., the Kolmogorov–Smirnov test, Mann–Whitney–Wilcoxon Test, and Hoeffding’s independence test, emerged as popular alternatives to parametric methods in 1950s, as they do not make specific assumptions about the underlying distribution of the data. In 1963, Hodges and Lehmann proposed a class of robust location estimators based on the confidence bounds of rank tests (31). In REDS I, when compared to other semiparametric mean estimators with the same breakdown point, the H-L estimator was shown to be the bias-optimal choice, which aligns Devroye, and Lerasle, Lugosi, and Oliveira’s conclusion that the median of means is near-optimal in terms of concentration bounds (32) as discussed. The formal study of semiparametric models was initiated by Stein (33) in 1956. Bickel, in 1982, simplified the general heuristic necessary condition proposed by Stein (33) and derived sufficient conditions for this type of problem, adaptive estimation (34). These conditions were subsequently applied to the construction of adaptive estimates (34). It has become increasingly apparent that, in robust statistics, many estimators previously called "nonparametric" are essentially semiparametric as they are partly, though not fully, characterized by some interpretable Euclidean parameters. This approach is particularly useful in situations where the data

do not conform to a simple parametric distribution but still have some structure that can be exploited. In 1984, Bickel addressed the challenge of robustly estimating the parameters of a linear model while acknowledging the possibility that the model may be invalid but still within the confines of a larger model (35). He showed by carefully designing the estimators, the biases can be very small. The paradigm shift here opens up the possibility that by defining a large semiparametric model and constructing estimators simultaneously for two or more very different semiparametric/parametric models within the large semiparametric model, then even for a parametric model belongs to the large semiparametric model but not to the semiparametric/parametric models used for calibration, the performance of these estimators might still be near-optimal due to the common nature shared by the models used by the estimators. Maybe it can be named as comparometrics. Closely related topics are "mixture model" and "constraint defined model," which were generalized in Bickel, Klaassen, Ritov, and Wellner's classic semiparametric textbook (1993) (36) and the method of sieves, introduced by Grenander in 1981 (37). As the building blocks of statistics, invariant moments can reduce the overall errors of statistical results across studies and thus can enhance the replicability of the whole community (38, 39).

## Methods

Methods of generating the Table 1 are summarized below, with details in the SI Text. The  $d$  values for the invariant moments of the Weibull distribution were approximated using a Monte Carlo study, with the formulae presented in Theorem A.1 and A.2. The computation of  $I$  functions is summarized in Subsection D and further explained in the SI Text. The computation of ASABs and ASBs is described in Subsection ???. The SEs and SSEs were computed by approximating the sampling distribution using 1000 pseudorandom samples for  $n = 5184$  and 50 pseudorandom samples for  $n = 2654208$ . The impact of the bootstrap size, ranging from  $n = 2.7 \times 10^2$  to  $n = 2.765 \times 10^4$ , on the variance of invariant moments and  $U$ -central moments was studied using the SEs and SSEs methods described above. A brute force approach was used to estimate the maximum biases of the robust estimators discussed for the five unimodal distributions. The validity of this approach is discussed in the SI Text.

**Data and Software Availability.** Data for Table 1 are given in SI Dataset S1-S4. All codes have been deposited in [GitHub](#).

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