

A Method of Choosing a Value of the Bending Constant in Huber's M-Estimation Function ¹

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Abstract

The shape of an M-estimation function is generally determined in the sense of either/both maximizing efficiency of an M-estimator at the model or/and bounding the influence function of an M-estimator. We propose an empirical method of choosing a value of the bending constant in Huber's ψ -function, which is the most widely used M-estimation function when estimating the location parameter.

Key Words and Phrases: Influential Observation; M-estimation; Robustness

1. Motivation and Algorithm

We assume that the observations X_1, X_2, \dots, X_n are i.i.d., each according to the distribution $F_\theta(x) = F(x - \theta)$, where F is known and θ is unknown. Any estimator $\hat{\theta}$, if σ is known, defined by a minimum problem of the form

$$\sum_{i=1}^n \rho \left(\frac{X_i - \hat{\theta}}{\sigma} \right) = \min!,$$

where ρ is an arbitrary differentiable function, is called an M-estimator. In practice, σ should be estimated by a proper estimator, say $\hat{\sigma}$.

We would like to measure the influence of a particular observation, say the i th observation, by comparing $\hat{\theta}$ by $\hat{\theta}_{-i}$, where $\hat{\theta}_{-i}$ is the M-estimate of θ computed without the i th observation. By a second order Taylor series, we have the followings;

$$\begin{aligned} \sum_{i=1}^n \rho \left(\frac{X_i - \hat{\theta}_{-i}}{\hat{\sigma}} \right) - \sum_{i=1}^n \rho \left(\frac{X_i - \hat{\theta}}{\hat{\sigma}} \right) \approx \\ - \left(\frac{\hat{\theta}_{-i} - \hat{\theta}}{\hat{\sigma}} \right) \sum_{i=1}^n \rho' \left(\frac{X_i - \theta}{\hat{\sigma}} \right) \Big|_{\theta=\hat{\theta}} + \frac{1}{2} \left(\frac{\hat{\theta}_{-i} - \hat{\theta}}{\hat{\sigma}} \right)^2 \sum_{i=1}^n \rho'' \left(\frac{X_i - \theta}{\hat{\sigma}} \right) \Big|_{\theta=\hat{\theta}}. \quad (1) \end{aligned}$$

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Let us define twice of the left-hand side of (1) as the measure of *Influence of i th observation for an M -estimation function in estimating a location parameter*, and denote it as IM_i . The first term on right-hand side of the equation (1) is in fact zero, so that we have

$$\begin{aligned} IM_i &= 2 \left\{ \sum_{i=1}^n \rho \left(\frac{X_i - \hat{\theta}_{-i}}{\hat{\sigma}} \right) - \sum_{i=1}^n \rho \left(\frac{X_i - \hat{\theta}}{\hat{\sigma}} \right) \right\} \\ &\approx \left(\frac{\hat{\theta}_{-i} - \hat{\theta}}{\hat{\sigma}} \right)^2 \sum_{i=1}^n \rho'' \left(\frac{X_i - \hat{\theta}}{\hat{\sigma}} \right). \end{aligned} \quad (2)$$

For Huber's proposal (Huber, 1981) such as $\rho'(t) = \psi_k(t)$, where

$$\psi_k(t) = \min\{k, \max\{t, -k\}\} = t \cdot \min \left\{ 1, \frac{k}{|t|} \right\}$$

for $0 < k < \infty$, the sign ' \approx ' in (2) can be replaced by ' $=$ ', so that we have

$$IM_i(k) = \left(\frac{\hat{\theta}_{-i} - \hat{\theta}}{\hat{\sigma}} \right)^2 \sum_{i=1}^n I_{[-k, k]} \left(\frac{X_i - \hat{\theta}}{\hat{\sigma}} \right).$$

We can notice that there is a relationship between IM_i and Cook's distance (Cook, 1977a; Cook and Weisberg, 1982), defined as

$$D_i \equiv D_i(\mathbf{X}^T \mathbf{X}, p\hat{\sigma}_n^2) = \frac{(\hat{\mathbf{b}}_{(i)} - \hat{\mathbf{b}})^T (\mathbf{X}^T \mathbf{X}) (\hat{\mathbf{b}}_{(i)} - \hat{\mathbf{b}})}{p\hat{\sigma}_n^2},$$

where

\mathbf{X} is an $n \times p$ full-rank matrix of known constants,
 \mathbf{Y} is an n -vector of observable responses,
 $\hat{\sigma}_n^2$ is a sample variance of residuals and $\hat{\mathbf{b}}$, and
 $\hat{\mathbf{b}}_{(i)}$ are the vectors of the least squares estimates of the vector of regression parameters, \mathbf{b} , with all observations and observation except the i th observation, respectively.

The D_i is a result of the second-order expansion of $\rho(\mathbf{Y} - \mathbf{X}\hat{\mathbf{b}})$ w.r.t $\hat{\mathbf{b}}_{(i)}$ if $\rho(\mathbf{Z}) = \mathbf{Z}^T \mathbf{Z}$ for a matrix \mathbf{Z} . Furthermore, if $p = 1$, D_i turns out

$$D_i = n \left(\frac{\bar{X}_{-i} - \bar{X}}{\hat{\sigma}_n} \right)^2 = 2 \left\{ \sum_{i=1}^n \left(\frac{X_i - \bar{X}_{-i}}{\hat{\sigma}_n / \sqrt{n}} \right)^2 - \sum_{i=1}^n \left(\frac{X_i - \bar{X}}{\hat{\sigma}_n / \sqrt{n}} \right)^2 \right\}, \quad (3)$$

where $\bar{X} = \sum X_i / n$, $\hat{\sigma}_n^2 = \sum (X_i - \bar{X})^2 / (n - 1)$, and \bar{X}_{-i} is a sample mean calculated without an i th observation. D_i in (3) is in fact a location parameter version of Cook's

distance. The above quantity is also a special case of IM_i when M-estimating function, $\rho'(t) = \psi(t) = t$ for all t . The (3) is in fact also a special case of the likelihood displacement in Cook, et al. (1988). The values of $D_i(\mathbf{X}^T\mathbf{X}, p\hat{\sigma}_n^2)$ can be converted to a familiar probability scale by comparing computed values to the $F(p, n-p)$ distribution. For example, if $D_i(\mathbf{X}^T\mathbf{X}, p\hat{\sigma}_n^2)$ equals the 0.5 value of the corresponding F distribution, then deletion of the i -th case would move the estimate of \mathbf{b} to the edge of a 50% confidence ellipsoid relative to $\hat{\mathbf{b}}$, a potentially important change (Cook, 1977b). If the largest D_i is substantially less than 1, deletion of a case will not change the estimate of \mathbf{b} by much.

Consider how to figure out a proper value of the bending constant for a particular data set by using IM_i . We want Huber's M-estimating function effectively to handle influential observations, so that no observation is considered as an influential observation; that is, we want IM_i for all i to be less than or equal to a proper value from F distribution. In order to attain our goal, we want the largest k , say k^* satisfying that

$$\max\{IM_i(k)\} \leq F(1-\alpha, 1, n-1), \quad (4)$$

where $F(1-\alpha, 1, n-1)$ is the $(1-\alpha) \times 100\%$ point of the F distribution with 1 and $n-1$ degrees of freedom, for a given α . The observations bounded by k^* and $-k^*$ are no longer influential observations in terms of Cook's distance, and in that sense Huber's estimate with that k^* is a robust alternative to a sample mean. Let see how Huber's estimates behaves according to the various values of α . We have generated 1000 samples of 20 random observations from the normal distributions in which samples are from $N(0, 1)$ with probability 1.0, 0.9, 0.8, 0.7, 0.6, respectively, and otherwise from $N(5, 1)$; that is, samples are simulated from $(1-\epsilon)N(0, 1) + \epsilon N(5, 1)$ for $\epsilon = 0.0, 0.1, 0.2, 0.3, \text{ and } 0.4$. For each sample, we picked up the largest k satisfying; $\max\{IM_i(k)\} \leq F(1-\alpha, 1, 19)$ for $\alpha = 0.5, 0.625, \text{ and } 0.75$, and calculated Huber's estimates for each level of α with bending constants obtained. The box-plots of the various estimates for a location parameter μ show how the estimates behave according to α 's under various contamination (Figure 1). There are five sections separated by the vertical lines according to the levels of contamination, and in each section. The box-plots of sample means, Huber's estimates according to the upper limit in (4) with $\alpha=0.5, 0.625, 0.75$, the estimates with $k = 1.28$ (suggested by Wilcox, 1997) and $k = 1.36$ (suggested by Hampel, et al., 1986), and medians are plotted from the left to the right. Under the normal density Huber's M-estimator attains its asymptotic efficiency 0.94 and 0.95 when $k=1.28$ and 1.36, respectively. When $\alpha = 0.5$, the Huber's M-estimates are closer to sample means, but when $\alpha = 0.75$, the estimates are closer to medians. When a level of contamination is 30% and more, Huber's M-estimates with $k = 1.28, 1.36$ seem to break down while Huber's M-estimates with $\alpha = 0.75$ are robust as the medians. However, Huber's M-estimates with $\alpha = 0.75$ and the medians have larger standardized errors than

Table 1: Means and Standardized errors in parentheses of the various M-estimates; $\alpha = \alpha_0$ stands for the M-estimates based on $(1 - \alpha_0)$ th quartile of F-distribution

	0%	10%	20%	30%	40%
sample mean	-0.0103 (0.2126)	0.5216 (0.2145)	0.9397 (0.2161)	1.4921 (0.2138)	1.9947 (0.2278)
Huber's with $\alpha=0.5$	0.0138 (0.2059)	0.3307 (0.2431)	0.7273 (0.7727)	1.2850 (0.4587)	1.7566 (0.4284)
Huber's with $\alpha=0.625$	0.0124 (0.2133)	0.2432 (0.2400)	0.5649 (0.3285)	1.0714 (0.4886)	1.5258 (0.5313)
Huber's with $\alpha=0.75$	0.0093 (0.2220)	0.1801 (0.2390)	0.4044 (0.2905)	0.7890 (0.3837)	1.1967 (0.4956)
Huber's with $k = 1.28$	0.0041 (0.2278)	0.1591 (0.2494)	0.4461 (0.2746)	0.9521 (0.3511)	1.7471 (0.3513)
Huber's with $k = 1.36$	0.0040 (0.2267)	0.1656 (0.2486)	0.4634 (0.2760)	0.9992 (0.3608)	1.7858 (0.3373)
median	-0.1394 (0.2587)	0.2571 (0.3126)	0.2571 (0.3126)	0.5118 (0.3742)	0.3821 (0.4497)

Huber's M-estimates with 1.28 and 1.36. That is, the plots are suggesting that there is a trade-off between robustness and efficiency as the level of contaminations increases. However, the ranges of Huber's M-estimates with $k=1.28$ and 1.36 are wider than Huber's M-estimates with $\alpha = 0.75$, that is, Huber's estimates with a fixed bending constant are less stable than Huber's M-estimates with $\alpha = 0.75$.

The fact is that when we consider robustness, 20% contamination at is a lot. We can notice that Huber's M-estimates with $\alpha = 0.75$ perform quite well over Huber's M-estimates with 1.28 and 1.36 at lower levels of contamination. Hence, we propose that Huber's M-estimator based on the algorithm in this article with $\alpha = 0.75$ should be a good alternative to those with $k = 1.28$ and 1.36.

2. Examples

In the following three examples, an M-estimate of location is a result of the first iteration of an iterative estimation procedure known as the Newton-Raphson method, that is,

$$T_n = T_n^0 + S_n^0 \sum \psi \left(\frac{X_i - T_n^0}{S_n^0} \right) / \sum \psi' \left(\frac{X_i - T_n^0}{S_n^0} \right),$$

where $T_n^0 = \text{median}$ and $S_n^0 = (\text{Median Absolute Deviation}) / 0.6745$.

Example 1: The Cushney and Peebles Data. In 1904, Cushney and Peebles published their experimental results on “The Action of Optimal Isomers” in the *Journal of Physiology*. The following data are copied from Table 4.1 of Staudte and Sheather (1990, p97), and written in order:

0.0, 0.8, 1.0, 1.2, 1.3, 1.3, 1.4, 1.8, 2.4, 4.6.

- The mean is 1.58 and the median is 1.3.
- The 20% two-sided trimmed mean is 1.40.
- The mean without {0.0, 2.4, 4.6}, which are considered outliers, is 1.26.
- F-distribution: when $k = 0.51$ ($\alpha=0.75$), 1.18 ($\alpha=0.625$), and 2.36 ($\alpha=0.5$), Huber’s estimates are 1.3, 1.37, and 1.40.
- Huber’s estimates with $k = 1.28$ and 1.36 are 1.338 and 1.343.

Example 2: Lifetimes of EMT6 Cells (Staudte, R. G. and Sheather, S. J., 1983). Mammalian cells in culture have lifetimes (cell cycle times) typically varying from 8 to 24 hours; the following is a typical data set of lifetimes in hours:

10.4, 10.9, 10.5, 8.8, 8.5, 8.7, 10.4, 7.8, 8.4, 9.1,
9.8, 10.3, 9.5, 10.4, 9.0, 8.9, 7.7, 8.2, 9.1, 22.2.

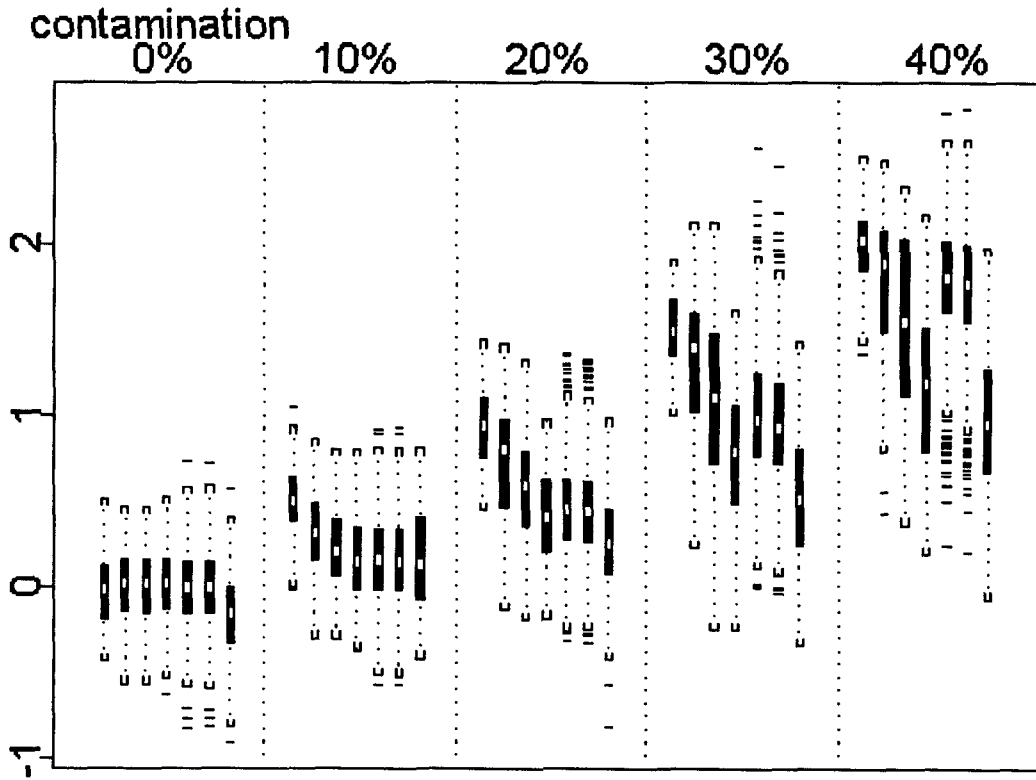
- The mean and median are 9.93 and 9.1, respectively.
- F-distribution: when $k = 0.53$ ($\alpha=0.75$), 2.43 ($\alpha=0.4$), and 3.07 ($\alpha=0.5$) Huber’s estimates are 9.25, 9.42, and 9.45.
- Huber’s estimates are 9.375 when both $k = 1.28$ and 1.36.

Example 3: Self-Awareness Data (Wilcox, 1997, p33). Dana (1990) conducted a study dealing with self-awareness and self-evaluation. One segment of his study measured the time subjects could keep a portion of an apparatus in contact with a specified target. The following shows some data for one of the groups:

77, 87, 88, 114, 151, 210, 219, 246, 253, 262,
296, 299, 306, 376, 428, 515, 666, 1310, 2611.

- The sample mean and the sample trimmed means with 10% and 20% trimmings are 448, 343, and 283.
- Huber's M-estimate with $k = 1.28, 1.36$ are 285.1 and 2.8853.
- F-distribution: when $k = 1.01$ ($\alpha=0.75$), 1.68 ($\alpha=0.4$), and 2.24 ($\alpha=0.5$) Huber's estimates are 227.56, 298.68, and 314.72.

Figure 1: Box-plots of the estimates under various contaminations; sample means, Huber's M-estimates based on $\alpha = 0.5, 0.625$ and 0.75 , and Huber's M-estimates with $k = 1.28$ and 1.36 , and medians in order from the left to the right in each section.



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