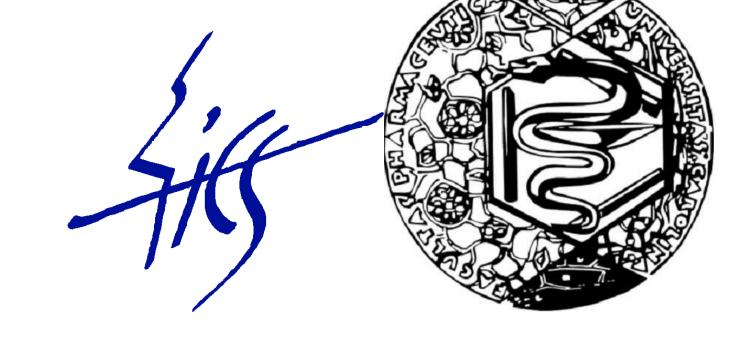
# Implementations for the minimum covariance determinant estimator J. Duintjer Tebbens<sup>1,2</sup> and J. Kalina<sup>1</sup>



Institute of Computer Science, Czech Academy of Sciences<sup>1</sup> and Faculty of Pharmacy in Hradec Králové, Charles University in Prague<sup>2</sup>

#### Robust estimation of location and scatter

In statistics, the term *robustness* is mostly used to indicate robustness with regards to outliers in the observed data. More precisely, a descriptive value is said to be robust of it is not significantly influenced by possible outliers in the data. The detection of outliers in p-dimensional data (i.e. observations with p recorded properties) is difficult if p > 3 because one can not rely on visual inspection. In the univariate case, a single outlier might still be relatively easily detected by measuring with a norm called Mahalanobis distance. This distance is in fact the energy norm for the inverse of the symmetric positive definite covariance matrix S and scales the p-dimensional space such that the variabilities of the individual properties are normalized. In a multivariate situation, with multiple outliers, the Mahalanobis distance itself is too strongly influenced by the outliers to give a reliable tool for their detection, a phenomenon called the masking effect.

If the aim is to estimate the location and scatter by robust estimators (i.e. to compute a robust mean vector and robust covariance matrix), one can compute the location and scatter for a subset of the observations which hopefully does not contain outliers. Assume we • have *n* observations  $x_i \in \mathbb{R}^p$  of *p* variables, given by the data matrix

$$X = [x_1, \ldots, x_n]^T \in \mathbb{R}^{n \times p},$$

• and look for a subset of size h of the indices  $\{1, 2, \ldots, n\}$ , where  $[(n+p+1)/2] \le h \le n$ , such that no index in the subset corresponds to an outlier.

A criterion to base the search of the subset on and that has been proved to lead to highly robust estimators of location and scatter is to *minimize the determinant* of the covariance matrix [2]. For a given subset H of size h of the indices  $\{1, 2, \ldots, n\}$ , let • the corresponding mean  $\bar{x}_H$  be

$$\bar{x}_H = \frac{\sum_{i \in H} x_i}{h} \in \mathbb{R}^p$$

 $\bullet$  and the corresponding covariance matrix  $S_H$  be

$$S_{H} = \frac{1}{h-1} \sum_{i \in H} (x_{i} - \bar{x}_{H})(x_{i} - \bar{x}_{H})^{T} = \frac{1}{h-1} (X_{H}^{T} - \bar{x}_{H} \mathbf{1}_{h}^{T})(X_{H} - \mathbf{1}_{h} \bar{x}_{H}^{T}) = \frac{1}{h-1} (X_{H}^{c})^{T} X_{H}^{c} \in \mathbb{R}^{p \times p},$$

where  $\mathbf{1}_h \in \mathbb{R}^h$  is the vector of ones,  $X_H$  the data matrix for the indices in H and  $X_H^c \in \mathbb{R}^{h \times p}$  is the corresponding centered data matrix. The Minimum Covariance Determinant Estimator [3] defines the optimal subset  $H_0$  of size h of  $\{1, 2, ..., n\}$  as

$$H_0 = \arg\min_{H} \det(S_H) = \arg\min_{H} \det(\sum_{i \in H} (x_i - \bar{x}_H)(x_i - \bar{x}_H)^T)$$

and defines the corresponding estimates of location and scatter as  $\bar{x}_{H_0}$  and  $S_{H_0}$ , respectively.

#### The fast MCD algorithm [4]

The computation of the Minimum Covariance Determinant Estimator requires minimization over all  $\binom{n}{h}$  h-subsets of  $\{1, 2, \ldots, n\}$ , thus has combinatorial complexity and becomes infeasible for very moderate numbers of observations *n*. In the widely used *fast MCD* [4] algorithm:

- one attempts to approximate the minimum determinant
- with several determinant minimizing steps for a large ( $\pm$  500) number of trial h-subsets • and selects the *h*-subset leading after the minimizing steps to the smallest determinant The determinant minimizing steps are called C-steps (concentration steps) and rely on the following theorem:

## Theorem (C-step [4])

Let  $H_1$  be an h-subset with corresponding location  $\bar{x}_{H_1}$  and scatter  $S_{H_1}$ . If  $\det(H_1) \neq 0$  compute the Mahalanobis distances

$$d(i) = \sqrt{(x_i - \bar{x}_{H_1})^T S_{H_1}^{-1}(x_i - \bar{x}_{H_1})}, \qquad i = 1, \dots, n$$

and find a re-ordering  $j_1, \ldots, j_n$  of  $\{1, 2, \ldots, n\}$  such that

$$d(j_1) \leq d(j_2) \leq \cdots \leq d(j_n).$$

Then if  $H_2$  is the h-subset consisting of the indices  $\{j_1, \ldots, j_h\}$  and  $S_{H_2}$  is the corresponding covariance matrix,

$$\det(S_{H_2}) \leq \det(S_{H_1})$$

with equality if and only if  $\bar{x}_{H_1} = \bar{x}_{H_2}$  and  $S_{H_1} = S_{H_2}$ .

The main computational costs of one C-step can be summarized as follows:

- construction of the current covariance matrix  $S_{H_1}$ :  $\mathcal{O}(np^2)$  flops.
- Cholesky- or eigendecomposition of  $S_{H_1}$  (this also yields  $\det(S_{H_1})$ ) :  $\mathcal{O}(p^3)$  flops.
- computation of the distances d(i):  $\mathcal{O}(np^2)$  flops.

We will consider C-steps based on eigendecomposition, that is, they compute

$$S_{H_1} = Z_1 D_1 Z_1^T$$

with  $D_1, Z_1$  the eigenvalue and eigenvector matrix, respectively, and find the Mahalanobis distances d(i) using

$$d(i) = \sqrt{(x_i - \bar{x}_{H_1})^T Z_1 D_1^{-1} Z_1^T (x_i - \bar{x}_{H_1})}, \qquad i = 1, \ldots, n.$$

Our contribution consists of two cheap,  $\mathcal{O}(np)$  permutations that can be added to the C-step to improve its power with regards to minimizing the determinant.

## An *a-posteriori* permutation

Suppose after a C-step, we have selected a new h-subset based on the ordered distances  $d(j_1) \le d(j_2) \le \cdots \le d(j_n)$  as described in the previous theorem. In other words, the new h-subset  $H_2$  consists of the indices  $\{j_1,\ldots,j_h\}$ . A natural question is whether among the discarded indices  $\{j_{h+1},\ldots,j_n\}$  there may be indices that, if included in  $H_2$ , would yield a covariance matrix with smaller determinant. This can be checked in a computationally inexpensive way as follows.

If instead of  $H_2$  we use the h-subset  $\{j_1,\ldots,j_{h-1},j_r\}\equiv H_r$  for some index  $j_r\in\{j_{h+1},\ldots,j_n\}$ , then the data matrix for  $H_r$  differs from the data matrix for  $H_2$  in one column only. Therefore, the corresponding covariance matrices are small rank updates from eachother.

#### Theorem (Low rank update of a covariance matrix ([1], Theorem 3.2.2))

Let  $d_r = x_{j_h} - x_{j_r} \in \mathbb{R}^p$ , let  $S_r$  denote the covariance matrix for  $H_r$  and let  $f = e_h - \mathbf{1}_h/h \in \mathbb{R}^h$ , where  $e_h$  denotes the hth unit vector. Then there holds

$$S_r = S_{H_2} - d_r f^T X_2^c - (X_2^c)^T f d_r^T + ||f||^2 d_r d_r^T.$$

All vectors involved in the low-rank update can be computed with  $\mathcal{O}(p)$  flops. Moreover, information on the determinant of  $S_r$  can be obtained from the determinant of  $S_{H_2}$  in  $\mathcal{O}(p)$ flops as well: Using the eigendecomposition  $S_{H_2} = Z_2 D_2 Z_2^T$ , the eigendecomposition of  $S_r$  for the modified h-subset  $H_r$  can be written as

$$S_r = S_{H_2} - d_r f^T X_2^c - (X_2^c)^T f d_r^T + ||f||^2 d_r d_r^T$$
  
=  $Z_2 \left( D_2 - Z_2^T d_r f^T X_2^c Z_2 - (X_2^c Z_2)^T f d_r^T Z_2 + ||f||^2 Z_2^T d_r d_r^T Z_2 \right) Z_2^T.$ 

Thus the eigenvalues of  $S_r$  are the eigenvalues of a symmetric rank-three update of the diagonal matrix  $D_2$  and each eigenvalue can be obtained, using (inverse) power iteration, in  $\mathcal{O}(p)$  flops. To keep the flop count at  $\mathcal{O}(p)$ , we propose to compute only the  $s, s \leq 5$ , largest eigenvalues of each covariance matrix  $S_r$ . After testing for all  $j_r \in \{j_{h+1}, \ldots, j_n\}$ , we select the index  $j_r$  for which the product of the s largest eigenvalues of  $S_r$  is minimal. The total flop count for this a posteriori permutation is of order (n - h)sp.

#### A look-ahead permutation

The weakness of the *a posteriori* permutation is that it tends to find, in numerical tests, an index  $j_r$  to exchange the index  $j_h$  of  $H_2$  with, which would have been found anyway in the next C-step, i.e. the index  $j_r$  often becomes a member of  $H_3$  anyway. The proposed a posteriori permutation is therefore mainly useful to add to the very last C-step to be performed.

To overcome this weakness, we propose a second permutation which looks ahead at the indices of  $H_3$  and attempts to add an index to  $H_2$  that will not be in  $H_3$ . Assume that with a candidate h-subset  $H_2$  we compute the Mahalanobis distances

$$d(i) = \sqrt{(x_i - \bar{x}_{H_2})^T S_{H_2}^{-1}(x_i - \bar{x}_{H_2})}, \qquad i = 1, \dots, n$$
(1)

and find a re-ordering  $k_1, \ldots, k_n$  of  $\{1, 2, \ldots, n\}$  such that

$$d(k_1) \leq d(k_2) \leq \cdots \leq d(k_n).$$

Then  $H_3$  would be defined as the indices  $\{k_1, \ldots, k_h\}$ . We can test whether indices in  $\{k_{h+1},\ldots,k_n\}\setminus H_2$  yield a lower determinant of  $S_{H_2}$  when interchanged with  $i_h$ . This can be done in  $\mathcal{O}((n-h)sp)$  flops as before. When the index for which the product of the s largest eigenvalues of  $S_r$  is minimal is found, we replace  $H_2$  with  $H_r$  and have to recompute the Mahalanobis distances

$$d(i) = \sqrt{(x_i - \bar{x}_{H_r})^T S_{H_r}^{-1}(x_i - \bar{x}_{H_r})}, \qquad i = 1, \ldots, n$$

to perform the next C-step. Fortunately, this does not require the full  $\mathcal{O}(np^2)$  flops for a regular C-step. Thanks to the fact that  $H_r$  is a small-rank update of  $H_2$ , it can be done in  $\mathcal{O}(np)$  flops using (1) and the Sherman-Morrison formula.

## Experiment

We generated 10 data sets  $X = [x_1, \dots, x_n]^T \in \mathbb{R}^{100 \times 10}$  each with 100 observations and 10 variables. 80 observations were normally distributed with mean vector 0 and covariance matix  $\Sigma = 0.6 \cdot I_{10} + 0.4 \cdot \mathbf{1}_p \cdot \mathbf{1}_p^T$  and 20 randomly placed outliers were normally distributed with mean vector  $3 \cdot \mathbf{1}_p$  and covariance matrix  $2 \cdot \Sigma$ . For 25 random initial choices of  $H_0$  and each of the ten datasets, we performed 4 regular C-steps and compared with 4 C-steps including the look-ahead permutations (dashed curve) and with 4 C-steps including both proposed permutations (solid curve). The quality of the results is measured by the squared norm of  $\bar{x}_{H_4}$ (left figure) and the Frobenius norm of  $\Sigma - S_{H_4}$  (right figure). The curves give the ratio of these measures for the improved vs. regular C-steps, averaged over all 25 random initial *h*-subsets.

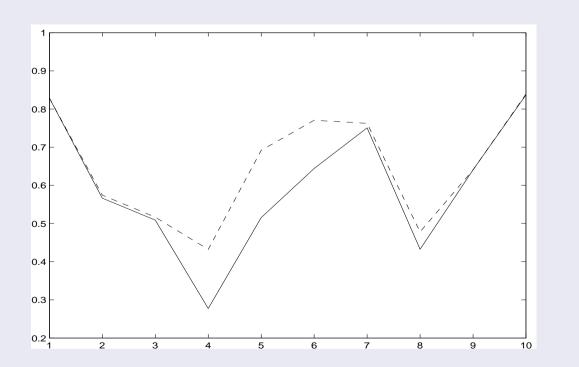


Figure 1: Ratios (averaged over 25 random initial choices of  $H_0$ ) of  $||\bar{x}_{H_4}||^2$  for look-ahead improved C-steps (dashed) or C-steps improved with both proposed permutations (solid) versus regular C-steps; the x-axis gives the data set number.

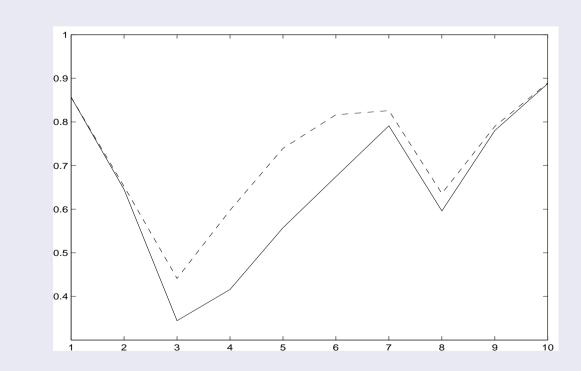


Figure 2: Ratios (averaged over 25 random initial choices of  $H_0$ ) of  $\|\Sigma - S_{H_4}\|_F$  for look-ahead improved C-steps (dashed) or C-steps improved with both proposed permutations (solid) versus regular C-steps; the x-axis gives the data set number.

http://www.cs.cas.cz/duintjertebbens

## Acknowledgements

The work of J. Kalina was financially supported by the Neuron Fund for Support of Science. The work of J. Duintjer Tebbens was supported by the grant GA13-06684S of the Czech Science Foundation.

- 1. Athanasiadis, S. *The small sample size problem in gene expression tasks*, Diploma thesis, Faculty of Pharmacy, Charles University, 2015.
- 2. Grübel, R. A minimal characterization of the covariance matrix Metrika, vol. 35, 49–52, 1988.
- 3. Hubert, M. and Debruyne, M. *Minimal covariance determinant* Metrika, Wiley Interdisciplinary Reviews: Computational Statistics, vol. 2, 36–43, 2010.
- 4. Rousseeuw, P. and Van Driessen, K. A fast algorithm for the minimum covariance determinant estimator Technometrics, vol. 34(3), 212-223, 1999.