Secure Data Aggregation for Sensor Networks in the Presence of Collusion Attack using Local Outlier Factor

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SECURE DATA AGGREGATION FOR SENSOR NETWORKS
IN THE PRESENCE OF COLLUSION ATTACK
USING LOCAL OUTLIER FACTOR

By
Anes Yessembayev

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Aggregation of data from multiple sensor nodes is usually done by simple methods such as averaging or, more sophisticated, iterative filtering methods. However, such aggregation methods are highly vulnerable to malicious attacks where the attacker has knowledge of all sensed values and has ability to alter some of the readings. In this work, we develop and evaluate algorithms that eliminate or minimize the influence of altered readings. The basic idea is to consider altered data as outliers and find algorithms that effectively identify altered data as outliers and remove them. Once the outliers have been removed, use some standard technique to estimate a true value.

We calculate local outlier factor (LOF) for each data point. We propose methods for computing threshold values from these LOF values. The data points that have LOF value above a calculated threshold value are removed before computing an estimated signal from the data points reported by the sensors. Thus, the proposed data aggregation algorithm operates in two phases: removal of outliers and computation of an estimated true value from the remaining sensor data. Extensive evaluation of the proposed algorithms show that they significantly outperform all existing methods. For simple cases, we have developed expressions for calculating LOF values.
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Chapter 1

Introduction

The present level of processes automation requires extensive use of various sensors. Due to unreliability of sensors they are deployed redundantly. Data from multiple sensor nodes is accumulated and combined by an aggregator node. An aggregator node not only collects readings from sensors, but also minimizes or eliminates the influence of readings from faulty or compromised sensors. Secure data aggregation algorithms for sensor networks aim to provide mechanisms for eliminating or resisting data distortion. These algorithms are usually run on an aggregator node or a base station.

1.1 Data Aggregation Methods

Probably the earliest and easiest method of data aggregation is simple averaging of readings from all sensors. However, simple averaging method has some major drawbacks, because it does not consider the existence of bias errors or faulty sensors and not to mention malicious attacks. Only one faulty sensor may reduce the accuracy of aggregated result significantly. In addition, the method does not doubt any sensor's reading. This makes the method highly vulnerable even to a simple attack, where the attacker skews reading of one or more sensors to a certain degree to alter estimated reading.

Iterative Filtering (IF) algorithms offer refined approaches (see [8] and [18] for examples). They initially assign one weight to the reading of each sensor and then weights are recalculated at each iteration based on the distance of the readings from the estimated value obtained in the previous iteration. They reduce the effect of a simple attack. But the weakest point of these iterative filtering algorithms is the use of a predetermined procedure for assigning an initial weight to each sensor’s reading.

An iterative algorithm is vulnerable to a malicious attacker who has knowledge of all readings and has power to alter two or more readings [18]. A malicious attacker can force the iterative filtering algorithm to converge to a desired value by altering readings of the compromised sensors (for details of an example situation see [18]).
To overcome weakness of the iterative filtering algorithm in [8], a *Robust Data Aggregation Method* (RDAM) was proposed in [18]. The main idea of the algorithm is to estimate a set of non-equal initial weights for the readings. The objective of the method is to calculate smaller initial weights for the readings of the compromised sensors. Results of extensive empirical evaluation of the RDAM method against other methods demonstrated highest accuracy for both simple and collusion attacks (for details see [18]).

However, any estimation of mean and standard deviation is extremely sensitive to presence of outliers [1],[14]. Thus, any aggregation algorithm that estimates true values from sensor-readings before removing outliers is susceptible to errors. In this work, we propose and evaluate a set of two phase data aggregation algorithms. In the first phase the outliers are identified and then, in the second phase an estimate is calculated.

### 1.2 Overview of the Thesis

In this work, we consider variations of the *collusion attack*. Our extensive evaluations discovered that in certain conditions the RDAM cannot overcome the influence of malicious attacker. In this work several two phase algorithms are proposed and evaluated.

The first phase of the proposed method employs a variant of *Local Outlier Factor* (LOF) calculation method [6] to estimate the degree that an object is an outlier because of collusion attack, sensor fault, noise, or a combination of them. This gives a flexible instrument to exclude suspected sensor-readings before estimation of a true value. Note that this method, unlike methods described previously, removes sensor-readings from compromised or bad sensors, which improve estimated values, while decrease the amount of calculations in the second phase of the proposed method. We present a set of different methods, including IF and RDAM, after LOF method. Also we test some variations of the algorithms in cooperation with *k-means clustering method* [9]. Finally, we introduce two LOF-based non-iterative algorithms that we found most accurate almost in all considered attacks. Moreover, the methods work without IF portion, that excludes disadvantages and vulnerabilities related to it.
The rest of thesis is organized as follows. Chapter 2 describes the problem statement. Here we briefly present most widely used topology of sensor networks and review their potential vulnerabilities. Chapter 3 briefly reviews work related to studies here. We describe existing data aggregation as well as outlier detection methods. Chapter 4 presents basic definitions and steps for computation of LOF. Chapter 5 presents our novel algorithms. We describe different combinations of using LOF method that we apply in our algorithms. Chapter 6 presents some theoretical results for computing LOF values when part of the data is from collusion attack. Chapter 7 shows our experimental results. Finally, the conclusion is provided in Chapter 8.
Chapter 2

Background

2.1 Model

The sensor network topology used for our work is an abstract model proposed in [20] (see Figure 2.1). A sensor network is built of a base station and a set of sensor clusters. Each cluster has a cluster head that gathers data from all sensor-nodes connected to it. A cluster head is also known as an aggregator node, because it gathers readings from multiple sensor nodes. The main functions of an aggregator node are collecting data from its sensor nodes, aggregating the raw data to produce an estimated reading, and communicating the processed data to the base station. Each sensor node has a micro-controller with one or multiple sensors. The micro-controller is equipped with relatively small memory and computing power, while an aggregator has bigger memory and higher computing power. A base station has larger memory and computing power, in addition to communication capabilities.

![Figure 2.1: Sensor network topology.](image-url)
2.2 Problem Statement

Sensor networks often operate in unattended environments and are deployed distributively, which makes them highly susceptible to failure and physical attacks. This creates threats to sensor networks security. Although in general, security can be defined as the combination of availability, confidentiality, and integrity, in our work we focus only on integrity. In addition, it is assumed that the aggregator nodes and the base station are not compromised. Assuming these limitations, we identify possible threats to the sensor network and propose solutions to overcome them (see Figure 2.2).

Classification of threats to sensor networks and their solutions

Nowadays, inexpensive sensors are embedded in numerous devices. These inexpensive sensors are not only unreliable, but they are also susceptible to failure. Moreover, they are usually insecure and their outputs are easily manipulated by malicious attackers.

Any sensor reading further from true value can be considered as an outlier, and in the literature they are categorized into several classes (see [13],[19], and [17]).

1. **Noise**: data with greater variance.
2. **Spike**: data with one or more out-of-bound readings.
3. **Stuck-at**: data with quasi-zero variance.

Consequently we cannot depend on readings from a single sensor. However, low cost of sensors allows one to compensate for their unreliability through use of multiple sensors. For example, after one receives readings from multiple sensors and apply a filtering technique, e.g. *Iterative Filtering*, the contributions of reading from unreliable sensors are reduced. Furthermore, this approach minimizes errors because of simple attacks mentioned in Chapter 1.

However, this technique cannot overcome a malicious attack, if and when the attacker has knowledge of all readings, details about filtering technique, and capability to alter some
of the readings. An example of a malicious attack is described in [18], where a *collusion attack* forces the *Iterative filtering algorithm* to converge to a skewed value. Rezvani et al. in [18] proposed an algorithm to improve *Iterative Filtering* and it is capable of reducing impacts of colluded readings. However, in our experiments we found conditions under which algorithms for reducing influence of colluded readings, such as RDAM method, are not effective. As a solution for such types of attacks we propose a set of variations of outlier detection algorithms that identify and, note, completely remove the impact of compromised readings. Considering that a micro-controller is very limited in its performance we set a criteria for the algorithms that they do not require high computational efforts.
Figure 2.2: Sensor network threats and solutions.
Chapter 3

Related Work

The extensive use of cheap and unreliable sensors mandates development of new algorithms for aggregating data securely, because these sensors are not only unreliable, but they are also easily compromised by malicious attackers. Also, it is necessary to take into account that new types of attacks continue to emerge with time. Thus, a robust algorithm that is resistant to known attacks, also should be resistant to emerging attacks. There are a number of research approaches and works related to our topic.

In our examples and experiments we consider a data model depicted in Table 3.1, where \( n \) - is total number of sensors, \( m \) - is number of readings from each sensor, \( x_{s}^{(t)} \) - data from sensor \( s \) at time \( t \).

### Table 3.1: Data Model

<table>
<thead>
<tr>
<th>Sensor readings</th>
<th>( s = 1 )</th>
<th>( s = 2 )</th>
<th>...</th>
<th>( s = n )</th>
</tr>
</thead>
<tbody>
<tr>
<td>( t = 1 )</td>
<td>( x_{1}^{(1)} )</td>
<td>( x_{2}^{(1)} )</td>
<td>...</td>
<td>( x_{n}^{(1)} )</td>
</tr>
<tr>
<td>( t = 2 )</td>
<td>( x_{1}^{(2)} )</td>
<td>( x_{2}^{(2)} )</td>
<td>...</td>
<td>( x_{n}^{(2)} )</td>
</tr>
<tr>
<td>...</td>
<td>...</td>
<td>...</td>
<td>...</td>
<td>...</td>
</tr>
<tr>
<td>( t = m )</td>
<td>( x_{1}^{(m)} )</td>
<td>( x_{2}^{(m)} )</td>
<td>...</td>
<td>( x_{n}^{(m)} )</td>
</tr>
</tbody>
</table>

3.1 Iterative Filtering Algorithms

Certainly the most studied method related to our research is Iterative Filtering Algorithms [16],[23],[26],[8],[15],[2]. In general, the majority of the work was dedicated for rating networks, where users give ratings to objects. Mizzaro proposed an algorithm for the assessment of scholarly papers [16]. Yu et al. [23] and, more recently, Zhou et al. [26] presented iterative algorithms for rating networks. However, these reputation-based algorithms may not always converge. Subsequently, a number of studies have proposed novel methods to overcome the convergence issue. C. de Kerchove and P. Van Dooren introduced a convergent Iterative Filtering algorithm using three discriminant functions: inverse, exponential, and Affine [8]. Li et al. in [15] presented six iterative algorithms, where users reputation is
calculated using the aggregated difference between the users rating and the corresponding objects ranking. Ayday et al. developed an iterative algorithm with probabilistic and belief propagation-based approach (see details in [2]). However, a fundamental problem of all IF algorithms is that they are primarily aimed against simple cheaters and do not consider severe malicious attacks such as a collusion attack that we mentioned in Chapter 1. Let us consider an example to illustrate this issue.

**Example 1.** Let us consider a dataset $X$ with $m = 3$ readings from each of the $n = 8$ sensors. Sensors $s_1 - s_5$ are providing their true readings to the aggregator, while $s_6 - s_8$ are under the influence of a malicious attacker and providing manipulated readings. The reading from $s_8$ at time $t$ is equal to the average of all readings of all sensors. Table 3.2 shows how the algorithm converges quickly to the average readings from $s_8$. This happens because the algorithm assigns equal initial weights for each sensor at the first round.

Table 3.2: Iterative Filtering Algorithm converges to colluded readings from $s_8$ sensor.

<table>
<thead>
<tr>
<th>instant</th>
<th>$s = 1$</th>
<th>$s = 2$</th>
<th>$s = 3$</th>
<th>$s = 4$</th>
<th>$s = 5$</th>
<th>$s = 6$</th>
<th>$s = 7$</th>
<th>$s = 8$</th>
</tr>
</thead>
<tbody>
<tr>
<td>$t = 1$</td>
<td>10.73</td>
<td>9.61</td>
<td>9.77</td>
<td>10.20</td>
<td>10.42</td>
<td>15.45</td>
<td>15.13</td>
<td>11.6157</td>
</tr>
<tr>
<td>$t = 2$</td>
<td>10.75</td>
<td>9.63</td>
<td>9.73</td>
<td>10.21</td>
<td>10.43</td>
<td>15.75</td>
<td>15.93</td>
<td>11.7757</td>
</tr>
<tr>
<td>$t = 3$</td>
<td>10.74</td>
<td>9.69</td>
<td>9.92</td>
<td>10.32</td>
<td>10.9</td>
<td>15.98</td>
<td>15.39</td>
<td>11.8486</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>round</th>
<th>Sensor weights</th>
<th>$t=1$</th>
<th>$t=2$</th>
<th>$t=3$</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td>11.6157</td>
</tr>
<tr>
<td>2</td>
<td>0.9786</td>
<td>0.2258</td>
<td>0.2652</td>
<td>0.4417</td>
</tr>
<tr>
<td>3</td>
<td>0.9786</td>
<td>0.2258</td>
<td>0.2652</td>
<td>0.4417</td>
</tr>
</tbody>
</table>

**3.2 Enhanced Iterative Filtering Algorithm**

To reduce effect of collusion attack, Rezvani et al. proposed a Robust Data Aggregation Method (RDAM), which is an improvement to IF method. In this method, unequal initial weights are calculated using the readings available to the aggregator [18]. This enhancement not only makes an IF algorithm collusion attack resistant, but it also reduces the number of required iterations to converge to an estimated value. The method is based on the assumption that the distribution of stochastic components of sensor errors is known or can be estimated.
Example (Cont.). With the same data the algorithm reduces the impact of readings from \( s_{6} - s_{8} \) (see Table 3.3). The algorithm estimates unequal initial weights for the readings at the first round. This allows to avoid converging to the values of \( s_{8} \). Although the algorithm in the example converges to the values of \( s_{4} \), in real data the algorithm usually converges to unique values, which actually occurred in our extensive simulations.

Table 3.3: RDAM avoids converging to colluded readings from \( s_{8} \) sensor.

<table>
<thead>
<tr>
<th>Sensor readings</th>
<th>aggregate values</th>
</tr>
</thead>
<tbody>
<tr>
<td>instant</td>
<td></td>
</tr>
<tr>
<td>( t = 1 )</td>
<td></td>
</tr>
<tr>
<td>( s = 1 )</td>
<td>10.73</td>
</tr>
<tr>
<td>( s = 2 )</td>
<td>9.61</td>
</tr>
<tr>
<td>( s = 3 )</td>
<td>9.77</td>
</tr>
<tr>
<td>( s = 4 )</td>
<td>10.20</td>
</tr>
<tr>
<td>( s = 5 )</td>
<td>10.42</td>
</tr>
<tr>
<td>( s = 6 )</td>
<td>15.45</td>
</tr>
<tr>
<td>( s = 7 )</td>
<td>15.13</td>
</tr>
<tr>
<td>( s = 8 )</td>
<td>11.6157</td>
</tr>
<tr>
<td>( t = 2 )</td>
<td></td>
</tr>
<tr>
<td>( s = 1 )</td>
<td>10.75</td>
</tr>
<tr>
<td>( s = 2 )</td>
<td>9.63</td>
</tr>
<tr>
<td>( s = 3 )</td>
<td>9.73</td>
</tr>
<tr>
<td>( s = 4 )</td>
<td>10.21</td>
</tr>
<tr>
<td>( s = 5 )</td>
<td>10.43</td>
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<tr>
<td>( s = 6 )</td>
<td>15.75</td>
</tr>
<tr>
<td>( s = 7 )</td>
<td>15.93</td>
</tr>
<tr>
<td>( s = 8 )</td>
<td>11.7757</td>
</tr>
<tr>
<td>( t = 3 )</td>
<td></td>
</tr>
<tr>
<td>( s = 1 )</td>
<td>10.74</td>
</tr>
<tr>
<td>( s = 2 )</td>
<td>9.69</td>
</tr>
<tr>
<td>( s = 3 )</td>
<td>9.92</td>
</tr>
<tr>
<td>( s = 4 )</td>
<td>10.32</td>
</tr>
<tr>
<td>( s = 5 )</td>
<td>10.9</td>
</tr>
<tr>
<td>( s = 6 )</td>
<td>15.98</td>
</tr>
<tr>
<td>( s = 7 )</td>
<td>15.39</td>
</tr>
<tr>
<td>( s = 8 )</td>
<td>11.8486</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>round</th>
<th>Sensor weights</th>
<th>( t=1 )</th>
<th>( t=2 )</th>
<th>( t=3 )</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>0.0722</td>
<td>0.3551</td>
<td>0.0707</td>
<td>0.4689</td>
</tr>
<tr>
<td>2</td>
<td>2.5168</td>
<td>4.5607</td>
<td>10.5808</td>
<td>57.2282</td>
</tr>
<tr>
<td>3</td>
<td>3.3182</td>
<td>3.2789</td>
<td>6.6131</td>
<td>408.4660</td>
</tr>
<tr>
<td>4</td>
<td>3.9912</td>
<td>2.7799</td>
<td>5.2341</td>
<td>2.07E+05</td>
</tr>
<tr>
<td>5</td>
<td>4.0060</td>
<td>2.7741</td>
<td>5.2145</td>
<td>1.38E+04</td>
</tr>
</tbody>
</table>

3.3 Statistical Estimation

If the readings from the sensors have no outliers, a statistical method for estimating true values is described here. The main assumption for the method is that every sensor has a certain error in its readings, because no sensor gives accurate readings continuously. This suggests to find a method for estimation of a true value in such a way that all the readings are considered, but weights vary based on temporal variation pattern of the readings. The sensor that has least variance is considered to have provided better readings. We propose to use the variance of readings of a sensor to determine a weight that is applied to the readings of the sensor when a true value is estimated from readings of all the sensors. A top-down description of the method starts from Equation (3.1).

\[
    w_{s} = \frac{1}{\nu_{s}} \cdot \frac{1}{\sum_{i=1}^{n} \frac{1}{\nu_{i}}}, \tag{3.1}
\]

where \( \nu_{s} \) is defined as

\[
    \nu_{s} = \frac{1}{m - 1} \cdot \sum_{t=1}^{m} (\mu_{s} - x_{s}^{(t)})^2, \tag{3.2}
\]
and $\mu_s$ is the mean of all readings from sensor $s$

$$
\mu_s = \frac{1}{m} \cdot \sum_{t=1}^{m} x_s^{(t)}
$$

(3.3)

After the weights are calculated, estimates of the true values are obtained using the equation below.

$$
\tilde{r}^{(t)} = \sum_{s=1}^{n} W_s \cdot x_s^{(t)}
$$

(3.4)

This method should be used only if the sensor readings are free of outliers or outliers have been identified and removed from the readings. Next we describe algorithms useful for outlier detection.

### 3.4 Outlier Detection Algorithms

Detection of outliers in a dataset is a well studied subject. But the information age, especially big-data and Internet of Everything (IoE), has seen renewed interest in the subject to fit the current needs. In our study, we use a recently proposed method and it is described in Chapter 4. In this section we provide a brief overview of the existing outlier detection method.

In general, outlier detection methods form five main classes: statistical, nearest neighbor, clustering, classification, and spectral decomposition [25]. In statistical methods outliers are defined as objects that do not fit in assumed distribution [3],[24],[22]. Nearest neighbor methods use a distance as a mean to distinguish outliers [5],[6].

Clustering algorithms use similarity metrics [12]. Most popular and probably old clustering algorithm is $k$-means [9], which we also apply in our experiments. Classification methods use training phase when they create mathematical models for regular data and outliers. After that they classify new data using these models [21],[11]. Spectral decomposition methods use principal component analysis to create a model structure. Readings that do not fit the structure are considered as outliers [7].
The methods in last three groups are characterized as requiring high computational efforts. However, sensor micro-controllers are limited in their performance. This motivates us to use in our algorithm a method that demands for relatively low computing power.

There is the *Local Outlier Factor* algorithm among *Nearest neighbor methods* group proposed by Breunig M. et al. for finding anomalous objects by measuring the local deviation of a given object with respect to its neighbors [6]. The method is able to detect outliers even if they form a dense cluster together regardless the data distribution. The method has a complexity $O(n^2)$, where $n$ is the number of data objects and is considered not a good choice for devices with limited computing power. However, our theoretical analysis in Chapter 6 shows that original equations described in Chapter 4 can be significantly simplified and adapted for the type of attacks considered here.
### Chapter 4

**An Algorithm for LOF and Its Application**

#### 4.1 Steps for Calculations of LOF

In this section we present basic definitions and steps for computation of *local outlier factor* (LOF) presented in [6]. The idea of LOF is based on the concept of local density. Computation of LOF involves six steps (see Figure 4.1). Since each step is essential for obtaining a final value of each data object and we will use these definitions for developing some novel computation algorithms for special cases, we formally define computation of each step. First we present an informal definition from [10].

![Figure 4.1: Outlier Detection algorithm.](image)

**Definition 1.** (Informal definition [10]): An outlier is an observation that deviates so much from other observations as to arouse suspicions that it was generated by a different mechanism.
For ease of communication, we use symbols \( o, p, \) and \( q \) to denote objects in a dataset \( X \). The notation \( d(p, q) \) is used to represent distance between two objects \( p \) and \( q \). A formal definition for \( d(p, q) \) is provided next.

**Definition 2.** (Distance \( d(p, q) \):) Distance between two points \( p \) and \( q \), denoted as \( d(p, q) \), is a nonnegative number, such that (i) \( d(p, p) = 0 \), (ii) \( d(p, q) > 0 \), if \( p \neq q \), and (iii) for three distinct points \( o, p, \) and \( q \) if \( d(o, p) = x \) and \( d(p, q) = y \), then \( d(o, q) \leq (x + y) \).

Above definition encompasses many methods for measuring the distance between two points, including Euclidean distance. For ease of conveying concepts, most of the definitions are accompanied by a running example.

**Example 2.** Let us consider four data points \( a(0) \), \( b(1) \), \( c(2) \), and \( d(5) \). The number in the parenthesis represents the position on the \( x \)-axis. These points are shown in Figure 4.2. We use Euclidean distance between points and the values of distance for all distinct pair of points are shown as following.

\[
\begin{align*}
&d(a, b) = 1; \quad d(a, c) = 2; \quad d(a, d) = 5; \quad d(b, c) = 1; \quad d(b, d) = 4; \quad d(c, d) = 3.
\end{align*}
\]

Figure 4.2: Four data points used to illustrate calculations in the algorithm that follows.

In the next step, the outlier-detection algorithm calculates \( d_k(p) \). It should be noted that \( d_k(p) \) is neither distance of the furthest point nor the number of points in the neighborhood of the point \( p \). It is true that the neighborhood of \( p \) will have at least \( k \) points, but the actual number may be much more than \( k \).
Definition 3. (k-distance $d_k(p)$ [6]) For any positive integer $k$, the $k$-distance of object $p$, denoted as $d_k(p)$, is defined as the distance $d(p,o)$ between $p$ and an object $o \in X$ such that:

(i) for at least $k$ objects $q \in X \setminus \{p\}$ it holds that $d(p,q) \leq d(p,o)$, and
(ii) for at most $(k-1)$ objects $q \in X \setminus \{p\}$ it holds that $d(p,q) < d(p,o)$.

\[ \square \]

In our example, $d_2(\cdot)$ of all the points are 2, but this is not true in general.

Example (Cont.). We consider four data points $a(0)$, $b(1)$, $c(2)$, and $d(5)$ from Example 4.2. We use $k = 2$ for this example.

$d_2(a) = d(a,c) = 2$; ($c$ is the second nearest neighbor);

$d_2(b) = d(b,a) = 1$; ($a/c$ is the second nearest neighbor);

$d_2(c) = d(c,a) = 2$; ($a$ is the second nearest neighbor);

$d_2(d) = d(d,b) = 4$; ($b$ is the second nearest neighbor);

\[ \square \]

After $d_k(\cdot)$, the $k$-distance, calculation of all the points, the next step is to compute $N_k(\cdot)$. Intuitively, $N_k(p)$ is the set of all points in the neighborhood of $p$ whose distance is less than or equal to $d_k(p)$. Formally, $N_k(p)$ is defined as

Definition 4. (k-distance neighbors $N_k(p)$ [6]) Given the $k$-distance $d_k(p)$ of $p$, $k$-distance neighbors of an object $p$ are the objects whose distance from $p$ is not greater than the $k$-distance of $p$, that is, $N_k(p) = \{q \in X \setminus \{p\} \mid d(p,q) \leq d_k(p)\}$.

\[ \square \]

Example (Cont.). We continue to use the same four points in Figure 4.2. Below are $2$-distance neighbors of all four points.

$N_2(a) = \{b,c\}$; $N_2(b) = \{a,c\}$; $N_2(c) = \{a,b\}$; $N_2(d) = \{b,c\}$
Reachability Distance of an object \( p \) with respect to another object \( o \), denoted as \( Rd_k(p \leftarrow o) \), is defined next. The definition is followed by computation of \( Rd_2(b \leftarrow a) \) for points \( b \) and \( a \) in our example. It is important to note that we have extended the scope of the definition in [6]. Since readings from all sensors could be identical in some cases, the \( k \)-distance could be zero in those cases. Our definition takes care of those special cases.

**Definition 5.** (Reachability Distance \( Rd_k(p \leftarrow o) \)) The Reachability Distance of object \( p \) with respect to another object \( o \) is defined as

\[
Rd_k(p \leftarrow o) = \max\{d_k(o), d(p, o), \epsilon\},
\]

where \( \epsilon \) is a small constant that is introduced to avoid division by zero operation in further calculations.

For the objects \( b \) and \( a \) in the Example 2, \( Rd_2(b \leftarrow a) = \max\{d_2(a), d(a, b), \epsilon\} = \max\{2, 1, \epsilon\} = 2 \). As will be clear later, computation of \( Rd_k(p \leftarrow o) \) for all pairs of points in \( X \) may not be necessary.

For an object \( p \in X \), the values of \( Rd_k(o) \in N_k(p) \) and cardinality of \( N_k(p) \) are used to compute Local Reachability Density, \( lrd_k(p) \).

**Definition 6.** (Local Reachability Density \( lrd_k(p) \) [6]) The local reachability density of an object \( p \) is defined as

\[
lrd_k(p) = \frac{|N_k(p)|}{\sum_{q \in N_k(p)} Rd_k(q \leftarrow p)},
\]

where \( |N_k(p)| \) is the number of objects in \( N_k(p) \).

**Example (Cont.).** Now we illustrate computation of \( lrd_2(a) \).

\[
\begin{align*}
lrd_2(a) &= \frac{|N_2(a)|}{\sum_{q \in N_2(a)} Rd_2(q \leftarrow a)} = \frac{|N_2(a)|}{Rd_2(b \leftarrow a) + Rd_2(c \leftarrow a)}.
\end{align*}
\]
Now,
\[ Rd_2(b \leftarrow a) = \max\{d_2(a), d(a, b), \epsilon\} \]
\[ = \max\{2, 1, \epsilon\} = 2, \]
and
\[ Rd_2(c \leftarrow a) = \max\{d_2(a), d(a, c), \epsilon\} \]
\[ = \max\{2, 2, \epsilon\} = 2 \]

Thus, \( lrd_2(a) = \frac{2}{2+2} = 0.5 \)

Following similar steps, we obtain values of \( lrd_2(b) \), \( lrd_2(c) \), and \( lrd_2(d) \):

\[ lrd_2(b) = \frac{|N_2(b)|}{Rd_2(a \leftarrow b) + Rd_2(c \leftarrow b)} = \frac{2}{1 + 1} = 1; \]
\[ lrd_2(c) = \frac{|N_2(c)|}{Rd_2(a \leftarrow c) + Rd_2(b \leftarrow c)} = \frac{2}{2 + 2} = 0.5; \]
\[ lrd_2(d) = \frac{|N_2(d)|}{Rd_2(b \leftarrow d) + Rd_2(c \leftarrow d)} = \frac{2}{4 + 4} = 0.25; \]

The final step in the algorithm is to compute Local Outlier Factor, \( LOF_k(p) \) of all the objects \( p \in X \). For defining \( LOF_k(p) \) of an object \( p \in X \), value of \( lrd_k(p) \), values of \( lrd_k(q) \in N_k(p) \), and cardinality of \( N_k(p) \) are used, but the expression can be simplified as shown below.

**Definition 7.** (Local Outlier Factor \( LOF_k(p) \) \[6\]) The local outlier factor of \( p \) is defined as

\[ LOF_k(p) = \frac{\sum_{q \in N_k(p)} \frac{lrd_k(q)}{lrd_k(p)}}{|N_k(p)|} = \left( \frac{\sum_{q \in N_k(p)} lrd_k(q)}{|N_k(p)|} \right), \quad (4.3) \]
Example (Cont.). We complete the example after calculating \( \text{LOF}_2(.) \) of all objects in the example.

\[
\begin{align*}
\text{LOF}_2(a) = \frac{\text{ld}_2(b) + \text{ld}_2(c)}{|N_k(a)| \ast \text{ld}_2(a)} &= \frac{1 + 0.5}{2 \ast 0.5} = 1.5; \\
\text{LOF}_2(b) = \frac{\text{ld}_2(a) + \text{ld}_2(c)}{|N_k(b)| \ast \text{ld}_2(b)} &= \frac{0.5 + 0.5}{2 \ast 1} = 0.5; \\
\text{LOF}_2(c) = \frac{\text{ld}_2(a) + \text{ld}_2(b)}{|N_k(c)| \ast \text{ld}_2(c)} &= \frac{0.5 + 1}{2 \ast 0.5} = 1.5; \\
\text{LOF}_2(d) = \frac{\text{ld}_2(b) + \text{ld}_2(c)}{|N_k(d)| \ast \text{ld}_2(d)} &= \frac{1 + 0.5}{2 \ast 0.25} = 3;
\end{align*}
\]

After sorting these local outlier factors we obtain,

\[
\text{LOF}_2(d) = 3; \quad \text{LOF}_2(a) = 1.5; \quad \text{LOF}_2(c) = 1.5; \quad \text{LOF}_2(b) = 0.5
\]

Obviously, top first outlier in this example is the object d.

4.2 Selecting \( k \) for Computing \( \text{LOF} \) in the Presence of Collusion Attack

Weaknesses of the Local Outlier Factor method are complexity \( O(n^2) \) and selecting the right \( k \) is not obvious. According to [6] the optimal value of \( k \) is defined as \( k = (n_g - 1) \), where \( n_g \) is the number of reliable objects. For our case the simplest way to select \( k \) is based on the assumption: there are considerably more ‘normal’ observations than ‘abnormal’ observations (outliers/anomalies) in the data. That hypothesis is enough to set

\[
k = \lceil n/2 \rceil \tag{4.4}
\]

to get decent results, where \( n \) is total number of observations.

Assume we have 10 data points, 3 of which are colluded (see Figure 4.3).

By calculating Local Outlier Factors (4.3) based on the average distance from every point \( p \) to the nearest \( k = \lceil n/2 \rceil = 5 \) points \( q \) we get that Local Outlier Factors of colluded data points are significantly higher, that gives a mathematical instrument to exclude them from further calculations. However, the simplicity has a drawback. Non-adaptive \( k \) leads to the same number of calculations for different number of colluders, which in some cases can make the value of \( \text{LOF} \) for outlier not so clearly prominent from reliable objects.
There is a better way to determine the optimal $k$ that uses $k$-means\textsuperscript{1} clustering method. The method splits all objects into a given number of clusters in which each object belongs to the cluster with the nearest mean, serving as a prototype of the cluster. Assuming that objects form two clusters: reliable $n_g$ and colluded $n_b$ objects, $k$-means clustering method allows to find the optimal $k$ defined in (4.5).

$$k = n_g - 1$$  \hfill (4.5)

Using this method allows us to consider the optimal $k$ in calculations (see Figure 4.4 and Section 5.1.4). In this case values of LOF for outliers reach the maximum, which makes them clearly distinguishable from reliable objects.

\textsuperscript{1}In this context symbol $k$ is a part of the name of the method and it is not related to the k-distance definition.
Chapter 5

Proposed Two Phase Data Aggregation Algorithms

In this chapter we present our algorithms. Logically and functionally our algorithms consist of two phases: 1) Detection and removal of outliers, 2) True value estimation with remaining data (see Figure 5.1). Table 5.1 contains notations used in descriptions for the algorithms.

Figure 5.1: Two phases of our algorithms.

<table>
<thead>
<tr>
<th>Notation</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>$r(t)$</td>
<td>true value of the signal at time $t$</td>
</tr>
<tr>
<td>$\hat{r}(t)$</td>
<td>estimated value of the signal at time $t$</td>
</tr>
<tr>
<td>$e_{s}(t)$</td>
<td>noise (error) of sensor $s$ at time $t$</td>
</tr>
<tr>
<td>$S$</td>
<td>a vector including numbers of all sensors</td>
</tr>
<tr>
<td>$S'$</td>
<td>a vector including numbers of reliable sensors</td>
</tr>
<tr>
<td>$X$</td>
<td>an array of source data</td>
</tr>
<tr>
<td>$X'$</td>
<td>an array of reliable data</td>
</tr>
<tr>
<td>$n$</td>
<td>number of sensors</td>
</tr>
<tr>
<td>$m$</td>
<td>number of readings from each sensor</td>
</tr>
<tr>
<td>$c$</td>
<td>number of compromised sensors</td>
</tr>
<tr>
<td>$x_{s}(t)$</td>
<td>data from sensor $s$ at time $t$</td>
</tr>
<tr>
<td>$w_{s}$</td>
<td>weight that reflects the trustworthiness of a sensor $s$</td>
</tr>
<tr>
<td>$\bar{X}$</td>
<td>a vector of average readings from each sensor</td>
</tr>
<tr>
<td>$\bar{x}_{s}$</td>
<td>an average reading from sensor $s$</td>
</tr>
<tr>
<td>$\bar{X}'$</td>
<td>a vector of average readings from reliable sensors</td>
</tr>
<tr>
<td>$\text{LOF}_{k}(X)$</td>
<td>an average of all $\text{LOF}<em>{k}(x</em>{s})$</td>
</tr>
<tr>
<td>$\text{LOF}_{k}(\bar{X})$</td>
<td>an average of all $\text{LOF}<em>{k}(\bar{x}</em>{s})$</td>
</tr>
</tbody>
</table>
5.1 Detection of Outliers

Following the logic of the construction of our algorithms, in this section we describe methods that we apply in the first phase of our algorithms. This phase is dedicated to the detection of outliers.

5.1.1 Detection of Outliers in a Single Step

As the name suggests the proposed algorithm detects outliers (that is, colluded data) and excludes them from further calculations in a single step. In Chapter 4 we described how to obtain a sequence of $LOF_k(x_1), LOF_k(x_2), ..., LOF_k(x_n)$ for a vector $X = x_1, x_2, ..., x_n$. As discussed earlier, higher the value of $LOF$ of an object, higher the probability of the object being a colluded object. The main challenge here is to set a criteria to split the sequence of $LOF$ values so that only colluded objects are excluded.

Example 3. For the data points depicted on the Figures 4.3 and 4.4, for $k = 5$ corresponding $LOF$ parameters are shown in the Table 5.2.

Table 5.2: Data points and corresponding LOF values

<table>
<thead>
<tr>
<th>value of $x_s$</th>
<th>$x_1$</th>
<th>$x_2$</th>
<th>$x_3$</th>
<th>$x_4$</th>
<th>$x_5$</th>
<th>$x_6$</th>
<th>$x_7$</th>
<th>$x_8$</th>
<th>$x_9$</th>
<th>$x_{10}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>$LOF_5(x_s)$</td>
<td>1.41</td>
<td>1.23</td>
<td>0.87</td>
<td>0.67</td>
<td>0.87</td>
<td>1.23</td>
<td>1.41</td>
<td>1.93</td>
<td>2.36</td>
<td>2.57</td>
</tr>
<tr>
<td>$LOF_5(X)$</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td>1.456</td>
<td></td>
</tr>
<tr>
<td>$LOF_5(x_s) &lt; LOF_5(X)$</td>
<td>true</td>
<td>true</td>
<td>true</td>
<td>true</td>
<td>true</td>
<td>true</td>
<td>true</td>
<td>false</td>
<td>false</td>
<td>false</td>
</tr>
</tbody>
</table>

Our empirical observations suggest that $LOF_k$ of an outlier is above $\overline{LOF}_k$, which is defined as the average of all $LOF_k$ values:

$$\overline{LOF}_k(X) = \frac{\sum_{s=1}^{n} LOF_k(x_s)}{n} \quad (5.1)$$

Example (Cont.). So, for the dataset in Table 5.2 $\overline{LOF}_5(X) = (1.41 + 1.23 + 0.87 + 0.67 + 0.87 + 1.23 + 1.41 + 1.93 + 2.36 + 2.57)/10 = 1.456$
Thus, data points $x_8(15), x_9(17),$ and $x_{10}(18),$ whose $LOF$ values are greater than $\overline{LOF}_S,$ are identified as outliers, and in reality are.

Using the notations introduced in Table 3.1, the Algorithm 1 provides a pseudo code description of the Outliers Detection in a Single Step.

**Algorithm 1:** Outliers Detection in a Single Step Algorithm

- **Input:** $X[m, n]$
- **Output:** $S'$

1. Compute $k = \lceil n/2 \rceil$
2. Compute $\overline{x}_s = \frac{\sum_{m=1}^{m}x_s^{(t)}}{m}$ for each sensor $1 \leq s \leq n$
3. Compute $LOF_k(\overline{x}_s)$ for each sensor $1 \leq s \leq n$
4. Compute $\overline{LOF}_k(\overline{X}) = \frac{\sum_{s=1}^{n}LOF_k(\overline{x}_s)}{n}$
5. for $s = 1 : n$ do
6.  if $LOF(\overline{x}_s) < \overline{LOF}_k(\overline{X})$ then
7.  S' = S' + {s};
8. end
9. end

On the input we have a dataset $X$ with $m$ observations for each of $n$ sensors. Output of the algorithm is a vector $S'$ that consists of indices of sensors that are found reliable. First, the algorithm calculates $k = \lceil n/2 \rceil$. Then the algorithm computes an average of all $m$ readings for each sensor $x_s, \overline{x}_s$. This is repeated for all $n$ sensors. Now we have a vector $\overline{X}[n]$ of $n$ data points. Third, compute $LOF_k(\overline{x}_s)$ for each element of $\overline{X}$. Next, calculate the average $\overline{LOF}_k(\overline{X})$ of all $LOF_k(\overline{x}_s)$. Then, compare each $LOF_k(\overline{x}_s)$ value with the $\overline{LOF}_k(\overline{X})$. Those sensors whose $LOF_k(\overline{x}_s)$ values are below the average are considered as reliable.

**5.1.2 Detection of Outliers Iteratively**

A variation of the previous method is the method in which outliers are found iteratively. Unlike the previous method, only one outlier with the maximum $LOF$ value is removed in every cycle.
Values of LOF parameter are recalculated considering the remaining data points after every iteration. A threshold and a ratio of the maximum LOF value to the minimum (see Equation (5.2)) are used for finding a loop exit-condition.

\[
\text{ratio} = \frac{\max(\text{LOF}_k(X))}{\min(\text{LOF}_k(X))} 
\]  

(5.2)

According to the [10] those elements that are deep inside a cluster have LOF values close to 1. On the other hand, all outliers have LOF values significantly greater than 1. This gives a basis to expect that if the ratio exceeds a certain threshold, then \(x_s\) is an outlier, where \(\text{LOF}_k(x_s) = \max(\text{LOF}_k(X))\). Initially we set a threshold equal 2. Our empirical observations and educated guess show that a value of 2 is good enough to detect obvious outliers. However, keeping the same threshold for all iterations often leads to false exclusion of reliable data. On the other hand, if we initially set the threshold too high it causes accepting outliers as a reliable data. Again, our extensive empirical evaluations has shown that if we increase the threshold every iteration using Equation (5.3) performance of the algorithm is excellent.

\[
\text{threshold}_{i+1} = \frac{\text{size}(X)}{\text{size}(X) - 1} \times \text{threshold}_i, 
\]  

(5.3)

where \(i\) is the current iteration number.

The Equation 5.3 changes the threshold value slowly, if the size of the set \(X\) large, but the threshold is increased faster if the size of \(X\) smaller. Thus, as we near the elimination of outliers, loop exit-condition is also close to be satisfied.
Example 4. For the example considered in Section 5.1.1 four iterations are needed.

Initialization:

threshold = 2;

\[ X = \{1.5, 2, 3, 4, 5, 6, 6.5, 15, 17, 18\} \]

\[ k = \left\lceil \frac{\text{size}(X)}{2} \right\rceil = 5 \]

Iteration 1:

\[ \text{LOF}_5(X) = \{1.41, 1.23, 0.87, 0.67, 0.87, 1.23, 1.41, 1.93, 2.36, 2.57\} \]

\[ \text{ratio} = \frac{2.57}{0.67} = 3.8 \]

\[ (\text{ratio} > \text{threshold}) \implies X = X - \{x_{10}\} = \{1.5, 2, 3, 4, 5, 6, 6.5, 15, 17\} \]

\[ \text{threshold} = \frac{9}{8} \times 2 = 2.25 \]

Iteration 2:

\[ \text{LOF}_5(X) = \{1.41, 1.23, 0.87, 0.67, 0.87, 1.23, 1.41, 2.82, 3.37\} \]

\[ \text{ratio} = \frac{3.37}{0.67} = 5 \]

\[ (\text{ratio} > \text{threshold}) \implies X = X - \{x_{9}\} = \{1.5, 2, 3, 4, 5, 6, 6.5, 15\} \]

\[ \text{threshold} = \frac{8}{7} \times 2.25 = 2.57 \]

Iteration 3:

\[ \text{LOF}_5(X) = \{1.41, 1.23, 0.87, 0.67, 0.87, 1.23, 1.41, 3.69\} \]

\[ \text{ratio} = \frac{3.69}{0.67} = 5.5 \]

\[ (\text{ratio} > \text{threshold}) \implies X = X - \{x_{8}\} = \{1.5, 2, 3, 4, 5, 6, 6.5\} \]

\[ \text{threshold} = \frac{7}{6} \times 2.57 = 3 \]

Iteration 4:

\[ \text{LOF}_5(X) = \{1.41, 1.23, 0.87, 0.67, 0.87, 1.23, 1.41\} \]

\[ \text{ratio} = \frac{1.41}{0.67} = 2.1 \]

\[ (\text{ratio} < \text{threshold}) \implies X' = X \]

Initially we set a threshold and calculate k, which we keep the same throughout whole the algorithm.
In first iteration we calculate values of LOF for each element of $X$. Then, find the ratio of the maximum value of LOF, which is $\text{LOF}_5(x_{10})$, to the minimum - $\text{LOF}_5(x_4)$. Next, we find that the value of the ratio is greater than threshold, which means that $x_{10}$ is an outlier. Exclude $x_{10}$ from the set $X$, recalculate a new value of threshold and continue iterations.

In every iteration we recalculate values of LOF for remaining elements of the $X$ set, find new values of the ratio and threshold until the exit-condition is satisfied.

After completing the last iteration on the input set $X$, the elements in the set $X'$ are considered to be good data points.

Using the notations introduced in Table 3.1, the Algorithm 2 provides a pseudo code description of the Detection of Outliers Iteratively.

### Algorithm 2: Outliers Detection Iteratively

**Input**: $X[m,n]$, threshold  
**Output**: $X'$

1. Compute $\bar{x}_s = \frac{1}{m} \sum_{t=1}^{m} x_{st}^{(t)}$ for each sensor $1 \leq s \leq n$
2. Compute $k = \lceil \frac{n}{2} \rceil$
3. do
4. Compute $\text{LOF}_k(\bar{x}_s)$ for each sensor $1 \leq s \leq n$
5. Compute $\text{ratio} = \frac{\max(\text{LOF}_k(\bar{x}))}{\min(\text{LOF}_k(\bar{x}))}$
6. if \text{ratio} $> \text{threshold}$ then
7. \quad $\bar{X} = \bar{X} - \bar{x}_s$ \hspace{5mm} $\text{LOF}_k(\bar{x}_s) = \max(\text{LOF}_k(\bar{X}))$
8. \quad $n = n - 1$
9. end
10. threshold $= \frac{\text{size}(\bar{X})}{\text{size}(\bar{X}) - 1} * \text{threshold}$
11. while \text{ratio} $> \text{threshold}$;
12. $X' = X$

Note that the example 4 excludes computation of average values for each sensor.

### 5.1.3 Detection of Outliers using Row-Wise Votes

Two outlier detection algorithms presented in previous sections, compute an average of all readings from a sensor. Then outlier detection algorithm identify outliers from these
averages. The Algorithm 3 presented in this section identifies outliers from the \( n \) readings that are reported at each time instance \( t \). In other words, the Algorithm 3 processes raw data in a row-wise manner. The outlier detection algorithm in Section 5.1.1 is applied to each row of \( X \) for identification outliers, that is, \( \text{LOF}_k(x^{(t)}_s) \) values are calculated for every observation \( x^{(t)}_s \) of sensor \( s \). The average \( \text{LOF}(x^{(t)}) \) value is calculated for every vector of observation \( x^{(t)} \). For those \( x^{(t)}_s \) whose LOF values indicate them as reliable corresponding elements of \( \text{votes}[m, n] \) array are assigned 1, otherwise it is assigned 0. This gives a table of votes (see example in the Table 5.3) with average values of all votes for each sensor, which is referred to reliability. Sensors that have reliability greater than a certain threshold are considered as reliable. After completing the algorithm the vector \( S' \) contains numbers of sensors that are found reliable.

### Table 5.3: Votes table

<table>
<thead>
<tr>
<th>Sensors</th>
<th>( s_1 )</th>
<th>( s_2 )</th>
<th>( s_3 )</th>
<th>( s_4 )</th>
<th>( s_5 )</th>
</tr>
</thead>
<tbody>
<tr>
<td>( t = 1 )</td>
<td>1</td>
<td>0</td>
<td>1</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>( t = 2 )</td>
<td>1</td>
<td>0</td>
<td>1</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>( t = 3 )</td>
<td>1</td>
<td>0</td>
<td>1</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>( t = 4 )</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>( t = 5 )</td>
<td>1</td>
<td>0</td>
<td>1</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>reliability</td>
<td>1</td>
<td>0.2</td>
<td>1</td>
<td>0.8</td>
<td>0.6</td>
</tr>
</tbody>
</table>

The advantage of this method is that it allows to set a threshold for excluding outliers; A higher threshold will eliminate readings from a larger number of sensors, but the readings from remaining sensors is expected to be highly reliable. The data shown in the Table 5.3 identify \( s_1, s_3, s_4, s_5 \) as reliable, if threshold is set to 0.5. But for same data sensors \( s_1, s_3, s_4 \) are found to be reliable data, if the threshold is set to 0.8.

#### 5.1.4 Detection of Outliers using K-means Clustering

The main challenge in using LOF based Outlier Detection algorithms is selection of an optimal value of \( k \). As will be shown in Chapter 7, using Equation (4.4) it is possible to obtain good results, but it is not very good if the number of colluded sensors is much less
Algorithm 3: Outliers Detection Row-Wise Algorithm

**Input**: \( X[m,n], \text{threshold} \)

**Output**: \( S' \)

1. **for** \( t = 1 : m \) **do**
   2. Compute \( \text{LOF}_k(x^{(t)}_s) \) for current observation, where \( 1 \leq s \leq n \)
   3. Compute \( \overline{\text{LOF}}_k(x^{(t)}) = \frac{\sum_{s=1}^{n} \text{LOF}_k(x^{(t)}_s)}{n} \)
   4. **for** \( s = 1 : n \) **do**
      5. if \( \text{LOF}_k(x^{(t)}_s) < \overline{\text{LOF}}_k(x^{(t)}) \) **then**
          6. \( \text{votes}^{(t)}_s = 1; \)
      7. **end**
      8. **else**
          9. \( \text{votes}^{(t)}_s = 0; \)
      10. **end**
   11. **end**
12. **end**

13. **for** \( s = 1 : n \) **do**
14. \( \frac{\sum_{t=1}^{m} \text{votes}^{(t)}_s}{m} > \text{threshold} \) **then**
15. \( S' = S' + \{ s \}; \)
16. **end**
17. **end**

than \( n/2 \) (see Figures 4.3 and 4.4). As we have already mentioned, the optimal value of \( k \) is \((n_g - 1)\), where \( n_g \) is the number of reliable objects.

For that task \( k \)-means clustering method has potential to produce better results [4]. The clustering algorithm splits data points into a given number of disjoint sets or clusters in which each data point belongs to one of the clusters. A data point closest to the mean value of all the data in a cluster serves as the **prototype** of the cluster.

**Example 5.** *Let us consider data points from previous examples\(^2\) \( a(1.5), b(2), c(3), d(4), e(5), f(6), g(6.5), h(15), i(17), \) and \( j(18) \) (shown in Table 5.2). The number in the parenthesis represents the value of an observation. First, we set the number of clusters \( k^2 \), which is in our case 2 (for reliable and colluded data objects).*

\(^2\)Since the symbol \( k \) is an inherent component of k-means method terminology, we do not use other symbols. Symbol \( k \) is used to denote the number of clusters only in this context. Further, we again use symbol \( k \) in the meaning related to \( k\)-distance.
Then select two data objects furthest apart a(1.5) and j(18). Next, define the initial cluster means (centroids): Mean\(_1\) and Mean\(_2\), which are initially equal to the values of the single data objects in the clusters.

The remaining data objects are now examined one at a time and assigned to the cluster whose centroid is closest to the data point. The cluster centroid of a cluster is recalculated every time a new data object is added to it. After nine steps the data objects are partitioned into two clusters (see Table 5.4).

Table 5.4: k-means clustering: splitting into clusters

<table>
<thead>
<tr>
<th>step</th>
<th>Cluster(_1) data objects</th>
<th>mean(_1)</th>
<th>Cluster(_2) data objects</th>
<th>mean(_2)</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>a(1.5)</td>
<td>1.5</td>
<td>j(18)</td>
<td>18</td>
</tr>
<tr>
<td>2</td>
<td>a(1.5), b(2)</td>
<td>1.75</td>
<td>j(18)</td>
<td>18</td>
</tr>
<tr>
<td>3</td>
<td>a(1.5), b(2), c(3)</td>
<td>2.17</td>
<td>j(18)</td>
<td>18</td>
</tr>
<tr>
<td>4</td>
<td>a(1.5), b(2), c(3), d(4)</td>
<td>2.63</td>
<td>j(18)</td>
<td>18</td>
</tr>
<tr>
<td>5</td>
<td>a(1.5), b(2), c(3), d(4), e(5)</td>
<td>3.1</td>
<td>j(18)</td>
<td>18</td>
</tr>
<tr>
<td>6</td>
<td>a(1.5), b(2), c(3), d(4), e(5), f(6)</td>
<td>3.6</td>
<td>j(18)</td>
<td>18</td>
</tr>
<tr>
<td>7</td>
<td>a(1.5), b(2), c(3), d(4), e(5), f(6), g(6.5)</td>
<td>4</td>
<td>j(18)</td>
<td>18</td>
</tr>
<tr>
<td>8</td>
<td>a(1.5), b(2), c(3), d(4), e(5), f(6), g(6.5)</td>
<td>4</td>
<td>h(15), j(18)</td>
<td>16.5</td>
</tr>
<tr>
<td>9</td>
<td>a(1.5), b(2), c(3), d(4), e(5), f(6), g(6.5)</td>
<td>4</td>
<td>h(15), i(17), j(18)</td>
<td>16.7</td>
</tr>
</tbody>
</table>

Next we verify that each data point has been allocated to the right cluster. To do that, we compare each data point’s distance to its current cluster centroid and to its distance the other cluster’s centroid. The procedure is illustrated in the Table 5.5.

The entries in the table confirms that the set of data points has been partitioned into two clusters correctly. Otherwise, data points that do not pass verification test are relocated to the other cluster and the centroids are recalculated. This iterative relocation continues until no more data movements are required.

To guarantee that the algorithm stops its execution, it is necessary to provide additional conditions.
After partitioning the data into two clusters, we know the number of elements in each, and assuming that reliable sensors are more than the colluded sensors, we can select the optimal $k$ using Equation (4.5), that is, $k = (n_g - 1)$, where $n_g$ is the number of elements in the largest cluster.

An obvious question is why one would need to use LOF algorithm, since the data has already been partitioned into two clusters. The answer is simple — $k$-means clustering algorithm partitions any set of data into a given number of clusters irrespective of their inherent property. But LOF values on a dataset give a measure for each data point that are useful for verification of the partitioning process.

**Example (Cont.).** After determined that the size of the largest cluster $\{1.5, 2, 3, 4, 5, 6, 6.5\}$ is 7 we set $k = 7 - 1 = 6$ and apply Detection of Outliers in a Single Step Algorithm (see Table 5.6)

<table>
<thead>
<tr>
<th>data point</th>
<th>Distance to the $mean_1 (4)$</th>
<th>Distance to the $mean_2 (16.7)$</th>
</tr>
</thead>
<tbody>
<tr>
<td>a(1.5)</td>
<td>2.5</td>
<td>15.2</td>
</tr>
<tr>
<td>b(2)</td>
<td>2</td>
<td>14.7</td>
</tr>
<tr>
<td>c(3)</td>
<td>1</td>
<td>13.7</td>
</tr>
<tr>
<td>d(4)</td>
<td>0</td>
<td>12.7</td>
</tr>
<tr>
<td>e(5)</td>
<td>1</td>
<td>11.7</td>
</tr>
<tr>
<td>f(6)</td>
<td>2</td>
<td>10.7</td>
</tr>
<tr>
<td>g(6.5)</td>
<td>2.5</td>
<td>10.2</td>
</tr>
<tr>
<td>h(15)</td>
<td>11</td>
<td>1.7</td>
</tr>
<tr>
<td>i(17)</td>
<td>13</td>
<td>0.3</td>
</tr>
<tr>
<td>j(18)</td>
<td>14</td>
<td>1.3</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>value of $x_s$</th>
<th>$x_1$</th>
<th>$x_2$</th>
<th>$x_3$</th>
<th>$x_4$</th>
<th>$x_5$</th>
<th>$x_6$</th>
<th>$x_7$</th>
<th>$x_8$</th>
<th>$x_9$</th>
<th>$x_{10}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>$LOF_6(x_s)$</td>
<td>1.35</td>
<td>1.2</td>
<td>0.89</td>
<td>0.59</td>
<td>0.89</td>
<td>1.2</td>
<td>1.35</td>
<td>2.3</td>
<td>2.75</td>
<td>2.98</td>
</tr>
<tr>
<td>$LOF_6(X)$</td>
<td>1.55</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>$LOF_6(x_s) &lt; LOF_6(X)$</td>
<td>true</td>
<td>true</td>
<td>true</td>
<td>true</td>
<td>true</td>
<td>true</td>
<td>true</td>
<td>false</td>
<td>false</td>
<td>false</td>
</tr>
</tbody>
</table>
After we applied Outlier Detection Algorithm in a Single Step we detected the same outliers as in Table 5.2, but now the values of outliers’ LOF are greater than using $k = 5$, which makes outliers to standout for detection.

5.2 Combinations of Algorithms

After the first phase of the algorithm is completed, readings suspected as outliers have been removed. In the second phase of the algorithm a true value is estimated from the remaining sensor-readings for each time instance. At this phase we can choose from IF, RDAM, and statistical estimation method. There are 12 possible combinations of four outlier detection algorithms described in Section 5.1, and three true value estimation algorithms in Chapter 3. In the following section, we describe some of them.

5.2.1 Detection of Outliers and Iterative Filtering

A combination of four methods for detection of outliers presented in Section 5.1 with IF gives four algorithms. Below is an example of combination of Detection of Outliers in a Single Step with IF. We call this algorithm LOF-single-IF.

Example 6. We use the same data from the Example 1 of Chapter 3 for this example. We have a dataset $X$ with $m = 3$ observations from each of $n = 8$ sensors. Sensors $s_1$ - $s_5$ are giving reliable data, while $s_6$ - $s_8$ are reporting false data, where readings from $s_8$ are equal to the average of all sensors. Table 5.7 illustrates results of LOF-single-IF. The algorithm excludes the impact of skewed readings from $s_6$ - $s_8$. First phase: calculate average readings $\bar{x}_s$ for each sensor. Next, calculate $\text{LOF}_k$ for each average reading using equation (4.4) $k = \lceil n/2 \rceil = 8/2 = 4$. Then, compute $\text{LOF}_4(\bar{X})$ - average of all $\text{LOF}_4(\bar{x}_s)$. After comparing each $\text{LOF}_4(\bar{x}_s)$ to $\text{LOF}_4(\bar{X})$ we detect that $s_6$, $s_7$, and $s_8$ are outliers and exclude them from further calculations. In the second phase we apply IF.
Table 5.7: Detection of Outliers in a Single Step and IF (LOF-single-IF)

<table>
<thead>
<tr>
<th>instant</th>
<th>$s = 1$</th>
<th>$s = 2$</th>
<th>$s = 3$</th>
<th>$s = 4$</th>
<th>$s = 5$</th>
<th>$s = 6$</th>
<th>$s = 7$</th>
<th>$s = 8$</th>
</tr>
</thead>
<tbody>
<tr>
<td>$t = 1$</td>
<td>10.73</td>
<td>9.61</td>
<td>9.77</td>
<td>10.20</td>
<td>10.42</td>
<td>15.45</td>
<td>15.13</td>
<td>11.6157</td>
</tr>
<tr>
<td>$t = 2$</td>
<td>10.75</td>
<td>9.53</td>
<td>9.75</td>
<td>10.21</td>
<td>10.45</td>
<td>15.75</td>
<td>15.35</td>
<td>11.7757</td>
</tr>
<tr>
<td>$t = 3$</td>
<td>10.74</td>
<td>9.69</td>
<td>9.92</td>
<td>10.32</td>
<td>10.98</td>
<td>15.98</td>
<td>15.39</td>
<td>11.8486</td>
</tr>
<tr>
<td>$\bar{X}$</td>
<td>10.7400</td>
<td>9.6433</td>
<td>9.8067</td>
<td>10.2433</td>
<td>10.5833</td>
<td>15.7267</td>
<td>15.4833</td>
<td>11.7467</td>
</tr>
<tr>
<td>$LOF_{4}(\bar{X})$</td>
<td>1.0865</td>
<td>1.3148</td>
<td>1.0816</td>
<td>0.6061</td>
<td>1.0912</td>
<td>3.5704</td>
<td>3.3897</td>
<td>2.3257</td>
</tr>
<tr>
<td>$LOF_{4}(X)$</td>
<td>1.8083</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>$LOF_{4}(\bar{X}) &lt; LOF_{4}(X)$</td>
<td>true</td>
<td>true</td>
<td>true</td>
<td>true</td>
<td>false</td>
<td>false</td>
<td>false</td>
<td></td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>round</th>
<th>Sensor weights</th>
<th>$t=1$</th>
<th>$t=2$</th>
<th>$t=3$</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1</td>
<td>1</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>2</td>
<td>3.3993</td>
<td>3.1677</td>
<td>6.3423</td>
<td>4.7585</td>
</tr>
<tr>
<td>3</td>
<td>3.9715</td>
<td>2.7916</td>
<td>5.2642</td>
<td>1.3E+05</td>
</tr>
<tr>
<td>4</td>
<td>4.0059</td>
<td>2.7742</td>
<td>5.2146</td>
<td>1.4E+10</td>
</tr>
</tbody>
</table>

5.2.2 Detection of Outliers and RDAM

We created four combinations of methods of detection of outliers described in Section 5.1 with RDAM. Below is an example of combination showing Detection of Outliers in a Single Step with RDAM. We call this algorithm LOF-single-RDAM.

Example 7. Combination of Detection of Outliers in a Single Step and RDAM similarly detects and excludes outlier readings from sensors $s_6$ - $s_8$ using LOF technique at the first phase. After removing skewed data, RDAM method is applied in the second phase. Table 5.8 shows how the algorithm excludes the impact of readings from $s_6$ - $s_8$.

Table 5.8: Detection of Outliers in a Single Step and RDAM (LOF-single-RDAM)

<table>
<thead>
<tr>
<th>instant</th>
<th>$s = 1$</th>
<th>$s = 2$</th>
<th>$s = 3$</th>
<th>$s = 4$</th>
<th>$s = 5$</th>
<th>$s = 6$</th>
<th>$s = 7$</th>
<th>$s = 8$</th>
</tr>
</thead>
<tbody>
<tr>
<td>$t = 1$</td>
<td>10.73</td>
<td>9.61</td>
<td>9.77</td>
<td>10.20</td>
<td>10.42</td>
<td>15.45</td>
<td>15.13</td>
<td>11.6157</td>
</tr>
<tr>
<td>$t = 2$</td>
<td>10.75</td>
<td>9.63</td>
<td>9.75</td>
<td>10.21</td>
<td>10.43</td>
<td>15.75</td>
<td>15.35</td>
<td>11.7757</td>
</tr>
<tr>
<td>$t = 3$</td>
<td>10.74</td>
<td>9.69</td>
<td>9.92</td>
<td>10.32</td>
<td>10.98</td>
<td>15.98</td>
<td>15.39</td>
<td>11.8486</td>
</tr>
<tr>
<td>$\bar{X}$</td>
<td>10.7400</td>
<td>9.6433</td>
<td>9.8067</td>
<td>10.2433</td>
<td>10.5833</td>
<td>15.7267</td>
<td>15.4833</td>
<td>11.7467</td>
</tr>
<tr>
<td>$LOF_{4}(\bar{X})$</td>
<td>1.0865</td>
<td>1.3148</td>
<td>1.0816</td>
<td>0.6061</td>
<td>1.0912</td>
<td>3.5704</td>
<td>3.3897</td>
<td>2.3257</td>
</tr>
<tr>
<td>$LOF_{4}(X)$</td>
<td>1.8083</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>$LOF_{4}(\bar{X}) &lt; LOF_{4}(X)$</td>
<td>true</td>
<td>true</td>
<td>true</td>
<td>true</td>
<td>false</td>
<td>false</td>
<td>false</td>
<td></td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>round</th>
<th>Sensor weights</th>
<th>$t=1$</th>
<th>$t=2$</th>
<th>$t=3$</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>0.0789</td>
<td>0.4362</td>
<td>0.081</td>
<td>0.3997</td>
</tr>
<tr>
<td>2</td>
<td>1.7568</td>
<td>8.4680</td>
<td>29.2168</td>
<td>15.1868</td>
</tr>
<tr>
<td>3</td>
<td>1.6380</td>
<td>9.6996</td>
<td>41.1109</td>
<td>1.2574E+01</td>
</tr>
<tr>
<td>4</td>
<td>1.4588</td>
<td>13.2041</td>
<td>83.6730</td>
<td>9.2561E+00</td>
</tr>
<tr>
<td>5</td>
<td>1.2570</td>
<td>22.0700</td>
<td>485.0123</td>
<td>6.4817E+00</td>
</tr>
<tr>
<td>6</td>
<td>1.1477</td>
<td>32.9083</td>
<td>5.30E+04</td>
<td>5.3023E+00</td>
</tr>
<tr>
<td>7</td>
<td>1.1388</td>
<td>33.9074</td>
<td>5.78E+08</td>
<td>5.2144E+00</td>
</tr>
</tbody>
</table>
5.2.3 Algorithm Based on Combination of Weights and Votes

This algorithm is obtained by combining row-wise votes method described in 5.1.3 and IF. In this combination, actually, there is no clear separation between detection of outliers phase and estimation of true values phase. Having a table of votes (see Table. 5.3) the algorithm uses its values as an indicator to use the corresponding reading in calculations or not to use (see Equation (5.4)).

$$\bar{r}(t) = \sum_{s=1}^{n} w_s \cdot x_s^{(t)} \cdot votes_s^{(t)}$$ \hspace{1cm} (5.4)

If an indicator $votes_s^{(t)}$ is 1 then the algorithm uses the corresponding reading in the same way as in IF. Otherwise the reading is simply ignored. Pseudo-code of the algorithm is shown in Algorithm 4.

---

**Algorithm 4: Weights and Votes Algorithm**

<p>| Input : $X[m, n]$ |</p>
<table>
<thead>
<tr>
<th>Output: $\bar{r}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>for $t = 1 : m$ do</td>
</tr>
<tr>
<td>Compute $LOF_k(x_s^{(t)})$ for current observation, where $1 \leq s \leq n$</td>
</tr>
<tr>
<td>Compute $\overline{LOF}<em>k(x^{(t)}) = \frac{\sum</em>{s=1}^{n} LOF_k(x_s^{(t)})}{n}$</td>
</tr>
<tr>
<td>for $s = 1 : n$ do</td>
</tr>
<tr>
<td>if $LOF(x_s^{(t)}) &lt; \overline{LOF}_k(x^{(t)})$ then</td>
</tr>
<tr>
<td>votes_s^{(t)} = 1;</td>
</tr>
<tr>
<td>else</td>
</tr>
<tr>
<td>votes_s^{(t)} = 0;</td>
</tr>
<tr>
<td>end</td>
</tr>
<tr>
<td>end</td>
</tr>
</tbody>
</table>

$W \leftarrow 1$;

while $\bar{r}$ notConverged do

| for $t = 1 : m$ do |
| $\bar{r}(t) = \sum_{s=1}^{n} w_s \cdot x_s^{(t)} \cdot votes_s^{(t)}$ |
| end |

Compute $W$;

end
5.2.4 Detection of Outliers and Statistical Estimation

For our studies, we combined Detection of Outliers in a Single Step and Detection of Outliers using K-means clustering method, described in Sections 5.1.1 and 5.1.4, with Statistical Estimation, described in Section 3.3. As presented in Chapter 7, these two combinations produce best results in our experiments. There we call these algorithms LOF-single-weightedSum and k-means-LOF-single-weightedSum, respectively.

Example 8. Let us apply LOF-single-weightedSum for the same data set used in previous examples. At the first phase the algorithm similarly detects and excludes outlier readings from sensors s_6 - s_8. At the second phase Statistical estimation method is applied (see Table 5.9).

Table 5.9: Detection of Outliers in a Single Step and Statistical estimation
Chapter 6

Estimation of LOF for Simple Datasets

Let \( G = \{g_1, g_2, \ldots, g_{n_g}\} \) be a set of readings from \( n_g \) ‘good’ sensors. In our study, a reading from a ‘good’ sensor means that the sensor, the embedded device, and communication from the embedded device to the destination are normal; there is no hardware, software, and network issues have changed or altered the reading. Similarly, let \( B' = \{b_1, b_2, \ldots, b_{n_b+1}\} \) be a set of ‘bad’ readings from \( b_b + 1 \) sensors. A ‘bad’ reading may be due to several reasons, including but not limited to, a faulty sensor or a reading altered by an adversary or a colluder. If a collusion attack of the type described in [18] has occurred, then one of readings \( b_i \in B' \) is the average of the remaining \( n_b \) from \( B' \) and all \( n_g \) readings from \( G \).

Without loss of generality, let us assume that \( b_{n_b+1} \) is the average reading introduced by the colluder. Let us define \( B = B' \setminus \{b_{n_b+1}\} \), and \( av = b_{n_b+1} \), where

\[
    av = \frac{\sum_{i=1}^{n_g} g_i + \sum_{j=1}^{n_b} b_j}{n_g + n_b}.
\]

(6.1)

Figure 6.1: Collusion Attack Model.
For simplifying presentation, initially we assume that all readings in $G$ are same and equal to $\mu_g$ and similarly, we also assume that readings in $B$ are equal to $\mu_b$. Then the value of colluded reading $b_{n_b+1} = av$ is given by,

$$\mu = \frac{n_g \cdot \mu_g + n_b \cdot \mu_b}{n_g + n_b}$$  \hspace{1cm} (6.2)

Later, we will discuss how this restriction can be eliminated. By these assumptions, $d(g_i, g_j) = 0$ for $1 \leq i, j \leq n_g$ and also, $d(b_i, b_j) = 0$ for $1 \leq i, j \leq n_b$. For real-world situations, there should be considerably more ‘good’ observations than ‘bad’, that is, $n_g > (n_b + 1)$.

In this situation, we want to theoretically determine the value of $LOF_k$ for different range of values for $k$. For good sensors and $av$, we need to consider only one case, but for bad sensors, we have to consider three cases: $k < \min\{n_g, n_b\}$, $k = \min\{n_g, n_b\}$, and $\min\{n_g, n_b\} < k < \max\{n_g, n_b\}$.

**Lemma 1.** For $g_i, g_j \in G$, and $1 \leq k < \max\{n_g, n_b\}$,

1. $d_k(g_i) = 0$,
2. $N_k(g_i) = G \setminus \{g_i\}$,
3. $|N_k(g_i)| = n_g - 1$,
4. $Rd_k(g_j \leftarrow g_i) = \epsilon$,
5. $lr d_k(g_i) = \frac{1}{\epsilon}$.

**Proof.** The proofs of all items of the Lemma start from definitions. Since $k$ is less than $n_g$, the number of objects in $G$, a k-nearest neighbor $d_k(g_i)$ of any object $g_i \in G$ is another object in $G$. Because of the assumption that all readings in $G$ are equal and by Definition 3, we have $d_k(g_i) = 0$.

By Definition 4 all objects in $G$ except the object $g_i$ are in $N_k(g_i)$, the set of the k-nearest neighbors of $g_i$. 
The observation just made leads us to the fact that there are \((n_g - 1)\) objects in \(N_k(g_i)\).

By Definition 5, and the fact that \(d_k(g_i) = d(g_i, g_j)\), for all \(g_i, g_j \in G\),

\[
Rd_k(g_j \leftarrow g_i) = \max(d_k(g_i), d(g_i, g_j), \epsilon) = \epsilon.
\]

By Definition 6,

\[
lrd_k(g_i) = \frac{|N_k(g_i)|}{\sum_{g_j \in N_k(g_i)} Rd_k(g_j \leftarrow g_i)}.
\]

Now using the results we have already proved,

\[
lrd_k(g_i) = \frac{n_g - 1}{(n_g - 1)\epsilon} = \frac{1}{\epsilon}.
\]

This completes the proofs. \(\square\)

**Theorem 1.** For all \(g_i \in G\), and \(1 \leq k < \max\{n_g, n_b\}\), \(LOF_k(g_i) = 1\).

\(\square\)

**Proof.** By Definition 7, \(LOF_k(g_i)\) for \(g_i \in G\)

\[
LOF_k(g_i) = \frac{\sum_{q \in N_k(g_i)} lrd_k(q)}{|N_k(g_i)|} \cdot \frac{lrd_k(g_i)}{\sum_{q \in G \setminus g_i} lrd_k(q)} = \frac{(n_g - 1)}{(n_g - 1)} \cdot \frac{1}{\epsilon} = 1
\]

(by Lemma 1) \(\square\)
Next we present a lemma for the $av$.

**Lemma 2.** For $g_i \in G$, $av$, and $1 \leq k < \max\{n_g, n_b\}$,

1. $d_k(av) = \frac{n_b \cdot |\mu_b - \mu_g|}{n_g + n_b}$,
2. $N_k(av) = G$,
3. $|N_k(av)| = n_g$,
4. $Rd_k(g_i \leftarrow av) = \frac{n_g \cdot |\mu_g - \mu|}{n_g + n_b}$,
5. $lr_d_k(av) = \frac{n_g + n_b}{n_b |\mu_b - \mu|}$.

□

**Proof.** A $k$-nearest object of $av$ can be either in $G$ or in $B$, which are at distances $|\mu - \mu_g|$ and $|\mu_b - \mu|$, respectively. After substituting expression for $\mu$ from Equation (6.2) in $|\mu - \mu_g|$ and $|\mu_b - \mu|$, and then some simplifications we get,

$$|\mu - \mu_g| = \frac{n_b \cdot |\mu_b - \mu_g|}{n_g + n_b},$$

and

$$|\mu_b - \mu| = \frac{n_g \cdot |\mu_b - \mu_g|}{n_g + n_b}.$$

Since it was assumed that $n_G > (n_b + 1)$, we have $|\mu - \mu_g| < |\mu_b - \mu|$. Thus, all object in $G$ are at $k$-distance from $av$, and

$$d_k(av) = \frac{n_b \cdot |\mu_b - \mu_g|}{n_g + n_b}.$$

Also, a $k$-nearest neighbor of $av$ is in $G$ and distance between any two objects in $G$ is 0, by Definition 4 all $n_G$ objects in $G$ are in the set $N_k(av)$, that is, $N_k(av) = G$.

From above discussion, it is clear the value of $|N_k(av)|$ is $n_g$.

By Definition 5,

$$Rd_k(g_i \leftarrow av) = \max(d_k(av), d(av, g_i), \epsilon).$$
We have already proved that \( d(av, g_i) = |\mu - \mu_g| \) and \( d_k(av) = |\mu - \mu_g| \). Hence,

\[
Rd_k(g_i \leftarrow av) = |\mu - \mu_g| = \frac{n_b \cdot |\mu_b - \mu_g|}{n_g + n_b}.
\]

Finally, by Definition 6,

\[
lrd_k(av) = \frac{|N_k(av)|}{\sum_{g_i \in N_k(av)} Rd_k(g_i \leftarrow av)}.
\]

After substitution of values of \( Rd_k(g_i \leftarrow av) \) in the equation above, we get

\[
lrd_k(av) = \frac{n_g}{n_g \cdot |\mu - \mu_g|} = \frac{1}{|\mu - \mu_g|} = \frac{n_g + n_b}{n_b \cdot |\mu_b - \mu_g|}.
\]

\[\square\]

**Theorem 2.** For \( g_i \in G \) and \( 1 \leq k < \max\{n_g, n_b\} \), \( LOF_k(av) = \frac{n_b \cdot |\mu_b - \mu_g|}{(n_g + n_b) \cdot \epsilon} \)

\[\square\]

**Proof.** By Definition 7,

\[
LOF_k(av) = \frac{\sum_{q \in N_k(av)} lrd_k(q)}{|N_k(av)| \cdot lrd_k(av)} = \frac{\sum_{q \in G} lrd_k(q)}{n_g \cdot lrd_k(av)} \quad \text{(by Lemma 2)}
\]

\[
= \frac{n_b \cdot |\mu_b - \mu_g|}{(n_g + n_b) \cdot \epsilon} \quad \text{(by Lemmas 1 and 2)}
\]

\[\square\]
Lemma 3. For \( b_i, b_j \in B \), and \( 1 \leq k < \min\{n_g, n_b\} \),

1. \( d_k(b_i) = 0 \),
2. \( N_k(b_i) = B \setminus \{b_i\} \),
3. \( |N_k(b_i)| = n_b - 1 \),
4. \( R_d(k) = b_j \leftarrow b_i \),
5. \( l_r d_k(b_i) = \frac{1}{\epsilon} \).

\[ \square \]

Proof is similar to the proof of Lemma 1, and omitted for avoiding duplicity and conserving space.

Theorem 3. For \( b_i \in B \) and \( 1 \leq k < \min\{n_g, n_b\} \), \( LOF_k(b_i) = 1 \),

Proof. By Definition 7, \( LOF_k(b_i) \) for \( b_i \in B \)

\[
LOF_k(b_i) = \frac{\sum_{q \in N_k(b_i)} l_r d_k(q)}{|N_k(b_i)| \cdot l_r d_k(b_i)}
\]

(6.9)

\[ = \frac{\sum_{q \in B \setminus \{b_i\}} l_r d_k(q)}{(n_b - 1) \cdot l_r d_k(b_i)} \quad \text{(by Lemma 3)} \]

(6.10)

\[ = \frac{(n_b - 1) \cdot \frac{1}{\epsilon}}{(n_b - 1) \cdot \frac{1}{\epsilon}} = 1 \quad \text{(by Lemma 3)} \]

(6.11)

Lemma 4. For \( b_i, b_j \in B, av, \) and \( k = \min(n_g, n_b) \),

1. \( d_k(b_i) = \left| \frac{n_g \cdot (\mu_b - \mu_g)}{n_g + n_b} \right| \),
2. \( N_k(b_i) = \{av\} \cup B \setminus \{b_i\} \),
3. \( |N_k(b_i)| = n_b \),
4. \( R_d(k) = b_j \leftarrow b_i \),
5. \( R_d(k) = \frac{n_g \cdot (\mu_b - \mu_g)}{n_g + n_b} \),
6. \( R_d(k) = \frac{n_g \cdot (\mu_b - \mu_g)}{n_g + n_b} \),
7. \( l_r d_k(b_i) = \left| \frac{n_g + n_b}{n_g \cdot (\mu_b - \mu_g)} \right| \).
Proof. Since $k$ is equal to the number of objects in $B$, the k-nearest neighbor of any object in $B$ must be an object that is not in $B$. Since the object $av$ is closest to the objects in $B$, it is the k-nearest neighbor of all objects in $B$. The distance between an object $b_i \in B$ and $av$ is $\frac{n_g \cdot |\mu_b - \mu_g|}{n_g + n_b}$, and equals to $d_k(b_i)$.

Since distance between any objects $b_i \in B$ is 0 and the k-nearest neighbor of $b_i$ is $av$, then by Definition 4 object $av$ and all objects in $B$ except the object $b_i$ are the k-nearest neighbors. Thus, $N_k(b_i) = \{av\} \cup B \setminus \{b_i\}$.

From the proof above, it is clear that $b_i \in B$, $|N_k(b_i)| = n_b$.

By Definition 5, $Rd_k(b_j \leftarrow b_i) = \max(d_k(b_i), d(b_i, b_j), \epsilon)$ and $Rd_k(av \leftarrow b_i) = \max(d_k(b_i), d(b_i, av), \epsilon)$. Already we have proved that $d_k(b_i) = \frac{n_g \cdot |\mu_b - \mu_g|}{n_g + n_b}$. Hence, $Rd_k(b_j \leftarrow b_i) = \frac{n_g \cdot |\mu_b - \mu_g|}{n_g + n_b}$ and $Rd_k(av \leftarrow b_i) = \frac{n_g \cdot |\mu_b - \mu_g|}{n_g + n_b}$.

By Definition 6,

$$lr \text{d}_{k}(b_i) = \frac{|N_k(b_i)|}{\sum_{p \in N_k(b_i)} Rd_k(p \leftarrow b_i)} = \frac{|N_k(b_i)|}{\sum_{p \in N_k(b_i) \setminus \{av\}} Rd_k(p \leftarrow b_i) + Rd_k(av \leftarrow b_i)}.$$ (6.12)

Substitution of all values of $Rd_k(b_i)$ and simplification leads to the desired result,

$$lr \text{d}_{k}(b_i) = \frac{n_g + n_b}{n_g \cdot |\mu_b - \mu_g|}.$$ (6.13)

Theorem 4. For $g_i \in G$, $b_i \in B$, $av$ and $k = \min\{n_g, n_b\}$

$$LOF_k(b_i) = 1 + \frac{n_g - n_b}{(n_b)^2}.$$
Proof of Theorem 4. By definition of \( \text{LOF}_k(b_i) \) we have,

\[
\text{LOF}_k(b_i) = \frac{\sum_{q \in N_k(b_i)} lrd_k(q)}{|N_k(b_i)| \cdot lrd_k(b_i)} \tag{6.15}
\]

\[
= \frac{lrd_k(\text{av}) + \sum_{q \in B \setminus \{b_i\}} lrd_k(q)}{n_b \cdot lrd_k(b_i)} \quad \text{(by Lemma 4)} \tag{6.16}
\]

\[
= \frac{lrd_k(\text{av}) + \sum_{q \in B \setminus \{b_i\}} lrd_k(q)}{n_b \cdot lrd_k(b_i)} + \frac{(n_b - 1) \cdot lrd_k(b_i)}{n_b \cdot lrd_k(b_i)} \tag{6.17}
\]

\[
= \frac{lrd_k(\text{av})}{n_b \cdot lrd_k(b_i)} + \frac{(n_b - 1) \cdot lrd_k(b_i)}{n_b \cdot lrd_k(b_i)} \tag{6.18}
\]

Now using results from Lemmas 2 and 4, and doing some simplifications we get,

\[
\text{LOF}_k(b_i) = 1 + \frac{n_g - n_b}{n_b}
\]

\( \square \)

Lemma 5. For \( b_i, b_j \in B, \text{av}, \text{and, } \min\{n_g, n_b\} < k < \max\{n_g, n_b\} \)

(1) \( d_k(b_i) = |\mu_b - \mu_g| \),

(2) \( N_k(b_i) = G \cup \{\text{av}\} \cup B \setminus \{b_i\} \),

(3) \( |N_k(b_i)| = n_g + n_b \),

(4) \( R_k(b_j \leftarrow b_i) = |\mu_b - \mu_g| \),

\[ R_k(\text{av} \leftarrow b_i) = |\mu_b - \mu_g|, \]

\[ R_k(g_i \leftarrow b_i) = |\mu_b - \mu_g|, \]

(5) \( lrd_k(b_i) = \frac{1}{|\mu_b - \mu_g|} \)

\( \square \)

Proof. Since \( k \) is greater than the number of objects in \( B \), a k-nearest neighbor of any object in \( B \) is an object in \( G \), which gives k-distance, \( d_k(b_i) = |\mu_b - \mu_g| \).
Since distance between any two objects in $B$ is 0, a $k$-nearest neighbor is in $G$, and distance between any objects in $G$ is also 0, then by Definition 4 all objects in $G$, object $a_v$, and all objects in $B$ except objects $b_i$ are in the $k$-nearest objects set denoted as $N_k(b_i)$.

Since all objects in $G$, object $a_v$, and all objects in $B$ except the subject $b_i$ are in $N_k(b_i)$, the number of members of $N_k(b_i)$ is calculated as $n_g + 1 + (n_b - 1) = n_g + n_b$.

By Definition 5,

\[ Rd_k(b_j \leftarrow b_i) = \max(d_k(b_i), d(b_i, b_j), \epsilon), \quad (6.19) \]
\[ Rd_k(a_v \leftarrow b_i) = \max(d_k(b_i), d(b_i, a_v), \epsilon), \quad (6.20) \]
\[ Rd_k(g_i \leftarrow b_i) = \max(d_k(b_i), d(b_i, g_i), \epsilon). \quad (6.21) \]

We have already proved that $d_k(b_i) = |\mu_b - \mu_g|$. Hence, $Rd_k(b_j \leftarrow b_i) = |\mu_b - \mu_g|$, $Rd_k(a_v \leftarrow b_i) = |\mu_b - \mu_g|$, and $Rd_k(g_i \leftarrow b_i) = |\mu_b - \mu_g|$.

By Definition 6,

\[ lrd_k(b_i) = \frac{|N_k(b_i)|}{\sum_{p \in N_k(b_i)} Rd_k(p \leftarrow b_i)} \quad (6.22) \]
\[ = \frac{|N_k(b_i)|}{\sum_{p \in B \setminus b_i} Rd_k(p \leftarrow b_i) + Rd_k(a_v \leftarrow b_i) + \sum_{q \in G} Rd_k(q \leftarrow b_i)}. \quad (6.23) \]

Now substitution of expressions we have already found for $Rd_k(b_j \leftarrow b_i)$, $Rd_k(a_v \leftarrow b_i)$, and $Rd_k(g_i \leftarrow b_i)$ in the expression above, we get the desired result:

\[ lrd_k(b_i) = \frac{1}{|\mu_b - \mu_g|}. \quad (6.24) \]

\[ \Box \]

**Theorem 5.** For $k > \min\{n_g, n_b\}$, $g_i \in G$, $b_j \in B$, and $a_v$

\[ LOF_k(b_i) = \frac{n_g |\mu_b - \mu_g|}{(n_g + n_b) \epsilon} + \frac{1}{n_b} + \frac{n_b - 1}{n_g n_b}. \]
**Proof of Theorem 5.** By definition of $LOF_k(b_i)$ we have,

\[
LOF_k(b_i) = \frac{\sum_{q \in N_k(b_i)} lrd_k(q)}{|N_k(b_i)| \cdot lrd_k(b_i)}
\]

\[
= \frac{\sum_{q \in G} lrd_k(q) + lrd_k(\text{av}) + \sum_{q \in B \setminus \{b_i\}} lrd_k(q)}{(n_g + n_b) \cdot lrd_k(b_i)} \quad \text{(by Lemma 5)}
\]

\[
= \frac{\sum_{q \in G} lrd_k(q)}{(n_g + n_b) \cdot lrd_k(b_i)} + \frac{lrd_k(\text{av})}{(n_g + n_b) \cdot lrd_k(b_i)}
\]

\[
+ \frac{\sum_{q \in B \setminus \{b_i\}} lrd_k(q)}{(n_g + n_b) \cdot lrd_k(b_i)} \quad \text{(by Lemmas 1, 2, and 5)}
\]

\[
= \frac{n_g \cdot |\mu_b - \mu_g|}{(n_g + n_b) \cdot \epsilon} + \frac{1}{n_b} + \frac{n_b - 1}{n_g + n_b}
\]

(6.25) (6.26) (6.27) (6.28) (6.29)

\(\square\)
Chapter 7
Experimental Results

The experimental evaluation of the proposed algorithms aims to verify their reliability and efficiency in the presence of a collusion attack. Since the RDAM performs best in the presence of collusion attack [18], we use this algorithm’s performance as a benchmark for measuring that of the proposed algorithms. In order to evaluate the accuracy of the investigated algorithms in the presence of collusion attack, we assume that an attacker compromises $c$ sensor nodes out of $n$ nodes in a cluster and $c < n/2$.

7.1 Experimental Settings

Unless otherwise stated, we reconstruct the same conditions that are used in [18] for generating datasets for evaluation of the proposed algorithms. We used Matlab 9.0 (R2015b) as our programming platform. We model a cluster of $n = 20$ sensor nodes and assume that these 20 nodes reports their readings to a cluster head or data aggregator. Each sensor gathers $m = 400$ readings before sending them to the cluster head. For consistency and fair comparisons, the original signal is generated using the Equation (7.1), which has been used in [18].

$$r^{(t)} = 5 \cdot n \cdot \sin \left( \frac{\sqrt{2} \cdot t}{2 \cdot \pi} \right), \quad (7.1)$$

where $t \in \{1..m\}$.

To generate synthetic data sets $X$, we add noise $e_s^{(t)}$ to the original signal $r^{(t)}$ as shown by Equation (7.2)

$$x_s^{(t)} = r^{(t)} + e_s^{(t)}. \quad (7.2)$$

We use Gaussian distribution for both stochastic noise for readings from reliable sensors and correlated noise for readings from colluded sensors. In our experiments for the sensors that are considered reliable we add the same biased error as in [18] and is defined by Equations (7.3) and (7.4).
Moreover, we conduct experiments with another type of additive noise signals for the reliable sensors. Equation (7.5) describes this type of noise signal. Note that the baseline variance of the noise is increased by a multiplicative factor of $\sqrt{s}$ for the sensor with id $s$.

$$
e^{(t)}_s \sim \mathcal{N}(b_s, \sigma^2)
$$

(7.5)

For errors of colluded readings we use the same models, which are described by Equations (7.6) and (7.7) as in [18]. For noise we also use a modified model described by Equation (7.8). In this modified model the multiplicative factor $\sqrt{s}$ is removed (or set to 1).

$$
b'_s \sim \mathcal{N}(0, 3 \cdot \sigma^2_b),
$$

(7.6)

$$
e^{(t)}_s \sim \mathcal{N}(b'_s, 5 \cdot \sqrt{s} \cdot \sigma^2)
$$

(7.7)

$$
e^{(t)}_s \sim \mathcal{N}(b'_s, 5 \cdot \sigma^2)
$$

(7.8)

Parameters used in our experiments:

- each experiment is repeated 200 times and results are averaged;
- for all experiments $\sigma^2_b = 4$;
- value of standard deviation $\sigma$ in each experiment is varied from 1 to 5;
- number of compromised sensors $c$ in each experiment is varied from 3 to 8.

---

3We noted in Chapter 2 that sensor networks are often used in hostile environment. So it could happen that every next sensor is exposed to greater external influence and therefore noise increases accordingly.
We use only one discriminant function in our experiments \( g(d) = d^{-1} \) (see details in [18]) for iterative filtering phase of RDAM algorithm. The selection of this discriminant functions is compelled by the fact that out of four discriminant functions used for evaluations in [18], this function had shown the best performance.

For fair comparisons performance of the RDAM and our proposed algorithms, we used same sets of generated data for both methods.

### 7.2 Performance Evaluation Metric

To measure performance of the proposed algorithms we use two metrics: Root Mean Squared error (RMSE) and the Maximum Error (ME) defined by Equations (7.9) and (7.10), respectively. The RDAM algorithm used RMSE for evaluating performance of the algorithm against existing algorithms. We added ME to the metric for capturing worst-case performance.

\[
RMSE = \sqrt{\frac{\sum_{t=1}^{m} (r^{(t)} - \bar{r}^{(t)})^2}{m}},
\]

where \( r^{(t)} \) and \( \bar{r}^{(t)} \) are true values and estimate of true values of the signal at time \( t \), respectively.

\[
ME = \max_{\forall t}(|r^{(t)} - \bar{r}^{(t)}|),
\]

where max is the function that returns maximum value for all \( t \) and that \( |.| \) is absolute value function. The efficiency is evaluated by the number of iterations needed for the IF algorithm to converge.

### 7.3 Results

We present some of our typical observations from our extensive simulation results. For ease of comparison we illustrate our results as line plots. Moreover, we included corresponding data in the table format for those who want to examine the results more closely and critically.
We evaluated all the proposed algorithms, but we report results only for those algorithms that have produced best results. It should be noted that with rare exceptions, all our algorithms performed better than the RDAM algorithm.

The first set of results is for datasets that are (statistically) identical to those used for experiments in [18]. For visual display of the results, Figure 7.1 show plots RSME vs standard deviation for our proposed algorithms and RDAM algorithms. For detailed comparisons data is shown in Table 7.1. It is clear from the table and the plots that performance of our algorithm is as good or better than that of RDAM. For the case with 8 colluders the RMSE for our methods are 3-13% lower than that of the RDAM algorithm.

Figure 7.1: RMSE with original conditions of collusion attack
Table 7.1: Evaluation parameters for algorithms with original conditions of collusion attack

<table>
<thead>
<tr>
<th>Parameter</th>
<th>$\sigma = 1$</th>
<th>$\sigma = 2$</th>
<th>$\sigma = 3$</th>
<th>$\sigma = 4$</th>
<th>$\sigma = 5$</th>
</tr>
</thead>
<tbody>
<tr>
<td>RMSE</td>
<td>0.6574 1.4048 30.1</td>
<td>0.7505 2.054 18.2</td>
<td>0.8034 2.617 13.2</td>
<td>1.1022 3.963 11.6</td>
<td>1.3304 4.121 10.5</td>
</tr>
<tr>
<td>Maximum Error</td>
<td>0.8577 2.4223 20</td>
<td>1.0750 3.2259 15.4</td>
<td>1.3499 4.063 13.6</td>
<td>1.6044 4.996 12.2</td>
<td>1.9162 5.572 11.3</td>
</tr>
<tr>
<td>Iterations</td>
<td>0.7086 1.9174 2.2</td>
<td>0.9609 2.6835 2.2</td>
<td>1.1309 3.4292 2.2</td>
<td>1.3899 4.2388 2.2</td>
<td>1.6513 4.8549 2.2</td>
</tr>
</tbody>
</table>

Assessing performances of the algorithms in terms of the maximum error (defined by (7.10)), it can be concluded that all our proposed methods give approximately the
Figure 7.2: RMSE with changed conditions of collusion attack with 3 colluders

Figure 7.3: RMSE with changed conditions of collusion attack with 5 colluders
Figure 7.4: RMSE with changed conditions of collusion attack with 7 colluders

Figure 7.5: RMSE with changed conditions of collusion attack with 8 colluders
same level of maximum error, except \textit{LOF-single-weightedSum} and \textit{k-means-LOF-single-weightedSum}, which show significantly lower ME as can be seen in Figure 7.7 and Table 7.2.

Figures 7.2, 7.3, 7.4, 7.5, and 7.7 also show that two of our methods — \textit{LOF-single-weightedSum} and \textit{k-means-LOF-single-weightedSum} — outperform all others. For a given level of added noise level and number of colluders, their RMSE and ME are almost identical. The reason for their success is their ability to identify colluded readings with fairly high degree of accuracy (see Tables 7.1 and 7.2). For original conditions of collusion attack for \( \sigma = \{3, 4, 5\} \) and 8 colluders the estimated number of colluders is exactly 8.

Recall that algorithms \textit{LOF-single-weightedSum} and \textit{k-means-LOF-single-weightedSum} do not use iterative filtering. Thus, they reduce computation time and do not suffer from weaknesses of the iterative filtering algorithms.
(a) The maximum error for collusion attack with 3 colluders

(b) The maximum error for collusion attack with 5 colluders

(c) The maximum error for collusion attack with 7 colluders

(d) The maximum error for collusion attack with 8 colluders

Figure 7.7: The maximum error with changed conditions of collusion attack
Chapter 8

Conclusion

In this work, we concentrated on the development of outlier detection methods. We described the conception of local outlier factor, which plays a key role in our methods. Then we proposed several two phase outlier detection algorithms.

The main feature of our algorithms is detection and removal of outliers before estimation true value. First, this increases the accuracy by removing the influence of outliers in aggregated result. Second, having only reliable data true values can be estimated non-iteratively, which decreases computational cost.

We used the RDAM [18] as the benchmark for comparison since it has shown best results against original collusion attack. We tested RDAM and our algorithms against collusion attacks considered in [18]. Moreover, we created examples of collusion attack scenario that were not considered in [18]. Under these novel attack scenario the proposed algorithms performed much better than the RDAM.

We have presented experimental results that show that (1) the estimates have higher accuracy than RDAM and (2) the algorithms have better efficiency than that of the RDAM. We observed that detecting local outliers for sensor readings using LOF is efficient.

Since computation of local outlier factor is not a low computational-cost method, we have found expressions that are useful for calculating outlier factors for simple data. While conducting this research we changed some original definitions and discovered promising opportunities that can lead to new clustering methods, which will be a scope for future work.
References


