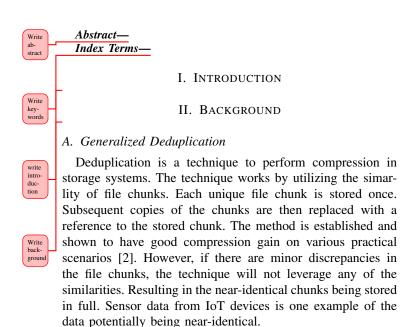
Outlier Detection of Generalized Deduplication Compressed IoT Data

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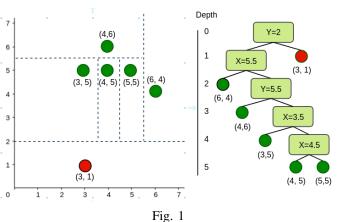
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To utilize the similarities in the almost identical data, a generalization of deduplication has been studied. This method consider the chunks at the bit level and splits them into two parts, the *base* and *deviation*. The *base* is the identical part that is to be stored once and herafter referenced with pointers. The *deviation* is the disparity between the chunks. Looking at a simple example with four 6-bit numbers, 100000, 100001, 100010 and 100011. It can be identified that the four most significant bits of the numbers are identical. Hence, leading to all having a shared *base* of 1000. The two least significant bits are then the *deviation*[4].

B. Isolation Forest

Anomaly detection is a combination of outlier- and novelty detection. Including both identifying outliers in the training data and determining if unseen observations are outliers. Isolation Forest (iForest) is an anomaly detection method. It differs from other popular techniques in the way that it identifies anomalies explicitly instead of profiling ordinary



data points[1]. IForest utilizes decision trees similar to other tree ensemble methods. The main principle is to recursively split each data point, and then evaluate the amount of splits necessary to split each data point. The logic is that anomalies will requires less splits to be isolated than an ordinary point. Trees are built by selecting a random feature and then selecting a random value between the minimum and maximum value of that feature. The process is then repeated untill all data points are isolated or a maximum height of the tree is reached. An illustration can be seen on Figure 1. The graphic shows an example of a decision tree and how an anomaly is at a lower depth of the tree. When determining if an observation is an outlier iForest calculates a score, it is defined as:

$$s(x,n) = 2^{-\frac{E(h(x))}{c(n)}}$$
 (1)

E(h(x)) is the average length from the root node to the specific data point. This is the average over a group of trees. c(n) is the average length from the root node to an external node. The anomaly score s is between 0 and 1. Scores close to 1 is seen as anomalies while values close to 0 is seen as normal data points.

Write related work

III. RELATED WORK

Performing analytics on compressed data is not an untouched subject.

A collection of models and algorithms are developed to perform classification and anomaly detection within network communication on compressed data[3].

Another paper looks into anomaly detection based on compressed data. They do it on the edge of the cloud on compressed data. Lots of formulas regarding rate vs. distortion. Dont know compression or anomaly detection method[7].

Direct analytics on data compressed using generalized deduplication has been carried out. It was studied how clustering(K-Means, IMM, DTC) could be performed on synthetic, synthetic with noise and a power consumption data set [6].

Since isolation forest only performs horizontal and vertical splits certain anomalies will not be detected. Imagining a two dimensional data set. Then the ones having the same x and y values might not be isolated correctly. This is extended by allowing diagonal splits in the extended isolation forest[5].

write method

IV. METHODS

A. Isolation Forest on GD compressed Data

Data compressed with generalized deduplication results in having a set of bases, deviation and references linking a data point to its base and deviation. In the following example the deviation will be omitted. Say we have the data set S where $S \in \mathbb{R}^2$. Performing GD on S will result in each feature of the points being mapped to their bases. Having an point $x = [x_1, x_2]$ where $x \in S$ and some computed bases ids $b_0, b_1, ..., b_n$, then the transformed version $x^* = [x_1^*, x_2^*]$ will hold the computed bases. This is depicted on Figure 2. The bases are computed on the raw data and referenced in the features x_1^* and x_2^* .

Isolation forest is then to be performed on the transformed version of the data set. The isolation forest splits before compression could be seen on Figure 1. Figure 3 is similar but is instead performing the splits on the bases. It is seen that certain data points will map to identical bases on both features.

B. DupRes Isolation Forest

The bases of GD compressed data will inherintly be grouped. Stripping the deviation of each data point will result in data points being placed in bins. This binning is illustrated on figure 4. The graphic shows the bins created with different amount of deviation bits. The circles are the bases. The dotted lines are enclosing areas where data points in an area will be mapped to the closest base in the negative direction. Having a larger amount of deviation bits is leading to larger bins.



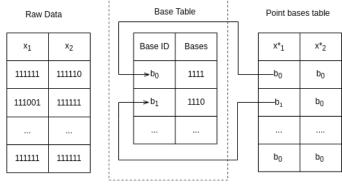


Fig. 2

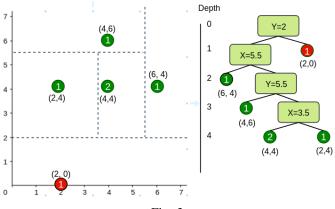
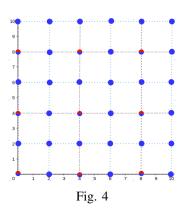


Fig. 3

Having larger bins might lead to a better compression rate however it could lead to undesired behaviour when trying to detect anomalies with iForest. An outlier could be mapped to the same base as an inlier on all or some of it features. Having the same base on some features will make it harder to isolate the outlier meanwhile having identical bases on all features makes it impossible. The binning aswell leads to inliers being grouped on fewer points. This causes them to be isolated more easily, and thus labeled as outliers.

Isolation Forest is not fit for the large amount duplicates that is potentially created by compressing with generalized



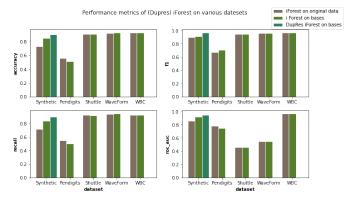


Fig. 5

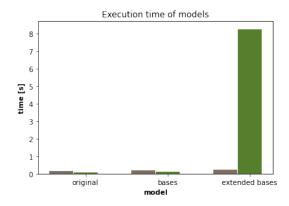


Fig. 6

deduplication. Therefore, a more duplicate resistant version is proposed. The core idea of the new version is to utilize the amount of duplicates when building the tree. The amount is then used to adjust the score of an observation. ...



VI. RESULTS

VII. CONCLUSION

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