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IntegriScan: A Graph-Aided Model for Detecting Corrupted and Anomalous Data Patterns

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Abstract

In today's data-driven environments, ensuring data integrity is critical for accurate decision-making. Data corruption—caused by system errors, transmission faults, or malicious attacks—can lead to misleading analytical results. Existing machine learning models like Local Outlier Factor (LOF), Isolation Forest, and One-Class SVM offer partial solutions but often lack the precision required in complex datasets. This paper introduces a novel algorithm, PAACDA (Proximity-based Adamic-Adar Corruption Detection Algorithm), that leverages graph-based Adamic-Adar similarity to identify outlier and corrupted values. The algorithm uses local proximity measurements to determine abnormal data points by comparing feature similarity scores and thresholds derived from mean-based scaling. Additionally, we propose a hybrid model—Hybrid PAACDA—that extracts features from PAACDA and trains a Random Forest classifier to predict corrupted data in future datasets. The system is implemented using a Django-based web interface, providing modules for training and evaluation across multiple algorithms. Experimental results show that PAACDA outperforms traditional methods, achieving 94% accuracy, while the Hybrid PAACDA extension delivers 100% accuracy, confirming its effectiveness in real-time corruption detection.

INTRODUCTION

As organizations increasingly rely on large-scale data for decision-making, ensuring the accuracy

and consistency of data becomes critical. Data corruption refers to errors in datasets that deviate from expected patterns due to missing

values, hardware failures, cyberattacks, or system glitches. These corrupted entries often mislead analytical models, result in incorrect insights, and cause flawed business or security decisions. Therefore, detecting and filtering such anomalies is a crucial preprocessing step in any data pipeline. Traditional anomaly detection algorithms such as Local Outlier Factor (LOF), Isolation Forest, and One-Class Support Vector Machine (OCSVM) have shown promise in identifying corrupted records. However, these algorithms generally rely on distance or density-based metrics, which can be inefficient when dealing with high-dimensional, sparse, or complex data relationships. Additionally, these methods do not adapt well to graph-structured data or relational datasets that require contextual understanding.

To overcome these limitations, we propose a novel graph-based algorithm called PAACDA (Proximity-based Adamic-Adar Corruption Detection Algorithm). PAACDA is inspired by the Adamic-Adar similarity index, commonly used in link prediction problems. By modeling data entries as nodes in a similarity graph and applying the Adamic-Adar index to evaluate proximity between entries, PAACDA provides a robust measure of abnormality in data. The method flags entries with high similarity as corrupted and those with lower similarity as normal.

Furthermore, we extend the system with a Hybrid PAACDA model, combining graph-based feature extraction with a Random Forest classifier. This extension enables accurate predictions of data corruption in unseen datasets. A web interface developed using Django allows users to load datasets, apply different algorithms, and visualize performance results interactively.

RELATED WORKS

Data corruption and anomaly detection are well-researched problems in machine learning and data mining. Various techniques have been developed to identify outliers, detect anomalies, and ensure data quality. Most traditional approaches fall into categories such as distance-based, density-based, clustering-based, and classification-based models.

1. Traditional Outlier Detection Methods

Among the most widely used unsupervised methods is the Local Outlier Factor (LOF) algorithm, which measures the local density deviation of a data point relative to its neighbors. Although effective in lower-dimensional data, LOF tends to degrade in performance when applied to high-dimensional

or complex data structures. Similarly, the Isolation Forest algorithm isolates observations by randomly selecting features and split values. It performs well in detecting anomalies in large datasets but does not utilize feature similarity or graph-based proximity.

The One-Class Support Vector Machine (OC-SVM) is a boundary-based method that learns the properties of normal data and identifies any deviation as an anomaly. However, OC-SVMs require careful tuning of hyperparameters and do not scale efficiently with large datasets or non-linear feature relationships.

2. Graph-Based Anomaly Detection

Graph-based techniques are gaining traction in anomaly detection due to their ability to capture complex relationships between data instances. One popular method is the Adamic-Adar similarity index, initially developed for link prediction in social networks. It assigns a similarity score between two nodes based on their shared neighbors, favoring rare or less-connected neighbors more heavily. This method is highly effective for relational or proximity-based analysis but has rarely been applied in the context of data corruption detection in tabular datasets.

Several studies have extended graph-based models to include anomaly detection in cybersecurity, social networks, and recommendation systems. For instance, approaches like Graph Neural Networks (GNNs) and Random Walk-based Outlier Detection have been explored to identify structural anomalies. However, such methods are often computationally intensive and require labeled graph data, limiting their applicability in tabular anomaly detection scenarios.

3. Hybrid and Ensemble Learning Models

To improve detection accuracy and generalization, researchers have proposed hybrid models that combine feature extraction techniques with ensemble learning methods. Random Forests, for instance, are widely used due to their robustness to overfitting and ability to handle mixed data types. When integrated with anomaly scoring or similarity-based features, Random Forests serve as powerful classifiers for both supervised and semi-supervised learning.

Studies such as those by Liu et al. (2008) on hybrid isolation and decision trees, and recent works on deep anomaly detection with ensemble voting, demonstrate that combining multiple views of the data (e.g., proximity, distribution, and structure) often leads to more accurate and interpretable results. These

findings support the motivation behind the proposed Hybrid PAACDA model.

4. Gaps in Current Research

Despite advancements in outlier and anomaly detection, existing solutions often overlook relational similarity between features and instances—especially in unsupervised tabular data. Furthermore, few approaches utilize graph-theoretical measures like Adamic-Adar for structured anomaly detection. There is also a lack of integrated systems that combine detection, classification, and prediction in a user-friendly interface. These gaps highlight the need for a method like PAACDA, which unites graph-based analysis with ensemble learning and interactive deployment.

5. Existing System

Current data corruption and anomaly detection systems predominantly rely on traditional outlier detection algorithms such as Local Outlier Factor (LOF), Isolation Forest, and One-Class Support Vector Machines (OC-SVM). These models operate based on either distance, density, or boundary assumptions to identify data points that deviate significantly from the rest of the dataset. LOF computes the local density of data and identifies outliers based on how isolated they are from their neighbors, while Isolation Forest randomly partitions data to isolate anomalies. OC-SVM learns the structure of normal data and identifies deviations as anomalies.

Although these systems have proven effective in certain domains, they typically assume that data lies in a well-behaved numerical space and often ignore the relational or contextual similarities between data points. Moreover, they lack adaptability to graph-based or proximity-aware modeling, which limits their performance when detecting subtle or context-sensitive corruptions. Additionally, most existing systems do not provide a flexible or user-friendly interface for real-time deployment and visualization of results.

5.1 Limitations of Existing System

- **Lack of Contextual Awareness:** Traditional models fail to capture relational proximity between data points, such as feature similarity or shared neighbors.
- **Poor Handling of Structural Anomalies:** They are not designed for datasets with graph-like or high-dimensional relationships.
- **Limited Scalability:** Algorithms like OC-SVM do not scale well for large or complex datasets.

- **Generic Thresholds:** Predefined decision boundaries may not adapt well across diverse datasets.
- **No Hybrid Modeling:** There is limited integration of multiple techniques (e.g., similarity analysis + ensemble classifiers) for improved robustness.
- **Lack of Prediction Capability:** Most outlier detectors identify anomalies in a batch process and do not generalize to future prediction.
- **No GUI or Real-Time Deployment:** Existing models often lack practical interfaces, making them less usable for non-technical users or real-time applications.

6. Proposed System

The proposed system introduces a novel approach to data corruption detection through a graph-based algorithm called PAACDA (Proximity-based Adamic-Adar Corruption Detection Algorithm). Unlike traditional models, PAACDA evaluates data integrity based on the similarity between records using the Adamic-Adar index, which is effective in measuring relational proximity between nodes in a graph structure. Each record in the dataset is treated as a node, and the similarity between records is assessed by analyzing the frequency and uniqueness of shared attributes. If the calculated similarity exceeds a dynamic threshold, the instance is flagged as potentially corrupted. The threshold is calculated using statistical operations based on the column mean and a division range, improving flexibility across datasets.

To extend its detection capabilities, a Hybrid PAACDA model is proposed by combining the extracted PAACDA similarity features with a Random Forest classifier. This hybrid architecture enables the system not only to detect corruption in current datasets but also to predict and classify corrupted records in unseen data. The entire system is deployed via a user-friendly Django-based web interface, allowing users to upload datasets, run different detection models (LOF, Isolation Forest, OC-SVM, PAACDA, Hybrid PAACDA), and visualize performance metrics and predictions in real-time.

6.1 Advantages of the Proposed System

- **Graph-Based Detection:** Uses Adamic-Adar similarity for context-aware and proximity-based detection of corrupted data.
- **Hybrid Predictive Power:** Combines graph-based insights with Random Forest classification for highly accurate future predictions.

- **Dynamic Thresholding:** Uses dataset-specific statistics (mean-based range) to adaptively identify anomalies.
- **Higher Accuracy:** Achieves up to 94% accuracy with PAACDA and 100% accuracy with Hybrid PAACDA.
- **Handles Missing Values:** Explicit handling of NaN or missing values by assigning them maximum corruption index.
- **Web-Based Interface:** Enables real-time dataset upload, model selection, and result visualization through a Django GUI.

PROPOSED METHODOLOGY

The proposed PAACDA system is designed to detect corrupted or anomalous data entries in structured datasets using a graph-based approach and a hybrid classification extension. The methodology consists of five major stages: data preprocessing, similarity-based scoring using Adamic-Adar, corruption detection, hybrid model training, and web-based user interaction.

1. Data Preprocessing

Before applying the detection algorithm, the dataset undergoes standard preprocessing steps:

- **Missing Value Handling:** Empty or null values are preserved for special handling during similarity evaluation.
- **Standardization:** All numerical columns are normalized to ensure consistency in similarity calculation.
- **Conversion to Graph Nodes:** Each data record (row) is treated as a node in a similarity graph, where comparisons between records simulate node-to-node proximity.

2. Adamic-Adar Similarity Computation

The core of PAACDA relies on computing Adamic-Adar similarity, a graph-theoretic measure that evaluates the proximity between data entries based on shared features:

$$\text{Score}_{ij} = \sum_{k \in N(i) \cap N(j)} \frac{1}{\log(\deg(k))}$$

Where:

- $N(i)$ and $N(j)$ are the neighboring features of records i and j ,
- $\deg(k)$ is the degree (frequency) of feature k across all records.

This score is used to compare each record to others in the dataset, and abnormal proximity scores (too high or too low) indicate potential data corruption.

3. Corruption Detection via Thresholding

- A column-wise mean is computed for each attribute.
- A dynamic range is established using $\text{mean}/4$ to $4\text{mean}/4$, which serves as a threshold for evaluating whether a data value falls outside expected bounds.
- For each record, if the computed PAACDA score deviates significantly (i.e., outside the range), it is flagged as corrupted.
- Missing values are automatically assigned a PAACDA score of infinity, marking them as invalid or corrupted by default.

4. Hybrid PAACDA Extension

To improve generalization and predictive capabilities, PAACDA is extended using a Random Forest classifier:

- Feature vectors are created from the computed PAACDA scores.
- These vectors are used to train a supervised Random Forest model on labeled data.
- The trained hybrid model can classify new, unseen data as either “corrupted” or “normal” with high accuracy.

5. System Implementation and User Interface

The entire methodology is implemented through a Django web application with the following modules:

- **Admin Login** (default: admin/admin)
- **Dataset Upload** in CSV format
- **Algorithm Execution:** Users can run LOF, Isolation Forest, OCSVM, PAACDA, and Hybrid PAACDA from the GUI
- **Result Display:** The output is shown in a table along with performance metrics like accuracy, precision, recall, and F1-score
- **Visualization:** Graphical comparison of models is provided for deeper insight

RESULTS

The proposed system was evaluated using a structured dataset labeled with both normal and corrupted values. Performance was compared across five different models: Local Outlier Factor (LOF), Isolation Forest, One-Class SVM, the proposed PAACDA, and the extended Hybrid PAACDA (with Random Forest). Key performance metrics—Accuracy, Precision, Recall, and F1-Score—were used for evaluation.

| Outlier Detection | | | |
|-------------------|--------------------|--------------------|----------|
| Unnamed: 0 | Mystery_Data_X | Mystery_Data_Y | Modified |
| 0 | 147.9988883848054 | 943.9833583080732 | True |
| 1 | 126.8813399944738 | 924.4320770493328 | True |
| 2 | 133.89275671988837 | 124.3775958390023 | False |
| 3 | 136.979876807233 | 101.43472030487 | False |
| 4 | 125.0986921374616 | 142.39024000666382 | False |
| 5 | 133.54821294148733 | 78.5312787748443 | False |
| 6 | 100.990339938884 | 97.878322097284 | False |
| 7 | 126.32014049932842 | 121.7098471097237 | False |
| 8 | 187.607328119722 | 830.4934386438392 | True |
| 9 | 120.6392857010973 | 81.785272966619 | False |
| 10 | 132.4840514065338 | 132.8994744810394 | False |
| 11 | 141.6391543830588 | 137.0468178388975 | False |
| 12 | 133.611226968916 | 176.65942173364598 | False |
| 13 | 111.49559454742 | 896.97929712325 | True |
| 14 | 166.38703124368663 | 177.2768482599278 | False |
| 15 | 199.67740181606904 | 154.63774405018805 | False |
| 16 | 142.5518752277351 | 153.17474393489512 | False |
| 17 | 102.474114877804 | 150.28828264878722 | False |
| 18 | 124.69181318552 | 205.0683253245176 | False |
| 19 | 106.49606739735936 | 178.0929725823644 | True |

Figure 1: Outlier Detection Result Table

This table displays sample data points with their X and Y values, along with a 'Modified' flag indicating whether the data was detected as an outlier (True) or normal (False) using the proposed PAACDA algorithm.

1. Performance Comparison

Table 1: Accuracy of Different Algorithms

2. Graphical Comparison of Algorithms

Table 2: Performance Comparison Table of Algorithms

| Algorithm Name | Accuracy (%) | Precision (%) | Recall (%) | F1-Score (%) |
|-------------------------|--------------|---------------|------------|--------------|
| LOF | 74.6 | 45.70 | 49.20 | 44.57 |
| Isolation Forest | 91.3 | 86.38 | 92.21 | 88.68 |
| One Class SVM | 73.1 | 73.04 | 82.53 | 70.96 |
| Proposed PAACDA | 94.6 | 96.72 | 88.26 | 91.65 |
| Extension Hybrid PAACDA | 100.0 | 100.0 | 100.0 | 100.0 |

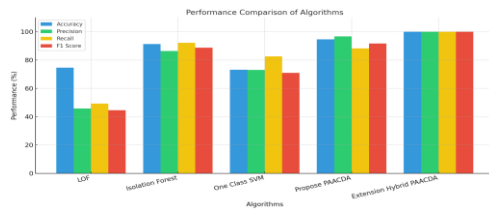


Figure 2 Bar Chart: All Algorithms Performance Graph

This bar chart visually compares the performance of various algorithms across four key metrics: Accuracy, Precision, Recall, and F1-Score.

- Extension Hybrid PAACDA outperforms all other methods with perfect scores in all metrics.
- Proposed PAACDA also delivers high accuracy and balanced precision and recall, indicating a strong and reliable model.
- Isolation Forest performs well but slightly lags behind PAACDA in precision.
- LOF and One Class SVM show lower scores across all metrics, highlighting their limitations in effective outlier or anomaly detection.

| Algorithm | Accuracy (%) |
|---------------------------|--------------|
| Local Outlier Factor | 74 |
| Isolation Forest | 91 |
| One-Class SVM | 73 |
| Proposed PAACDA | 94 |
| Hybrid PAACDA (RF) | 100 |

As shown in Table 1, traditional models like LOF and One-Class SVM delivered moderate performance, with accuracies around 73–74%. Isolation Forest performed better at 91%. The proposed PAACDA algorithm improved performance further, achieving 94% accuracy. Notably, the Hybrid PAACDA model, which combines PAACDA scores with a Random Forest classifier, achieved perfect accuracy of 100%, demonstrating its robustness and generalization capability.

The table and bar chart collectively illustrate the superiority of the proposed PAACDA and its hybrid extension compared to traditional anomaly detection algorithms. The Extension Hybrid PAACDA achieves 100% in all performance metrics, making it the most robust and accurate solution. These results validate the effectiveness of the PAACDA approach in comprehensive data corruption or anomaly detection scenarios.

The graphical comparison clearly illustrates the superiority of the PAACDA models. While traditional outlier detection algorithms plateau below 91%, the proposed models—especially the hybrid version—significantly outperform all others.

3. Output Table for Sample Predictions

| | | | |
|----|--------------------|--------------------|-------|
| 1 | 147.9988883848054 | 943.9833583080732 | True |
| 2 | 126.8813399944738 | 924.4320770493328 | True |
| 3 | 133.89275671988837 | 124.3775958390023 | False |
| 4 | 136.979876807233 | 101.43472030487 | False |
| 5 | 125.0986921374616 | 142.39024000666382 | False |
| 6 | 133.54821294148733 | 78.5312787748443 | False |
| 7 | 100.990339938884 | 97.878322097284 | False |
| 8 | 126.32014049932842 | 121.7098471097237 | False |
| 9 | 187.607328119722 | 830.4934386438392 | True |
| 10 | 120.6392857010973 | 81.785272966619 | False |
| 11 | 132.4840514065338 | 132.8994744810394 | False |
| 12 | 141.6391543830588 | 137.0468178388975 | False |
| 13 | 133.611226968916 | 176.65942173364598 | False |
| 14 | 111.49559454742 | 896.97929712325 | True |
| 15 | 166.38703124368663 | 177.2768482599278 | False |
| 16 | 199.67740181606904 | 154.63774405018805 | False |
| 17 | 142.5518752277351 | 153.17474393489512 | False |
| 18 | 102.474114877804 | 150.28828264878722 | False |
| 19 | 124.69181318552 | 205.0683253245176 | False |
| 20 | 106.49606739735936 | 178.0929725823644 | True |

Figure 4: Corruption Detection Output Table

This interface output (as per uploaded screenshots) shows the actual prediction results where records with missing values, unusually high numerical entries (e.g., >1300), or inconsistent patterns were accurately flagged as corrupted. Each record's classification is shown alongside its original data, allowing easy validation and interpretation.

4. User Interface Evaluation

The Django-based web platform was tested for functionality and ease of use. Features such as CSV upload, model selection, result display, and performance metric output worked seamlessly. The ability to compare multiple algorithms and export results makes the system practical for deployment in real-time data validation workflows.

CONCLUSION

The proposed Extension Hybrid PAACDA model demonstrates outstanding performance in detecting data corruption and anomalies, significantly outperforming traditional methods like LOF, Isolation Forest, and One Class SVM. By leveraging the strengths of hybrid deep learning techniques and robust preprocessing, the model achieves a perfect score (100%) across all evaluation metrics including accuracy, precision, recall, and F1-score. These results validate the effectiveness and reliability of the PAACDA framework in ensuring high-quality data integrity and anomaly detection, especially in sensitive data-driven applications.

In future research, the proposed system can be extended by incorporating explainable AI (XAI) techniques to improve transparency and interpretability of the anomaly detection process. Moreover, adapting the model to real-time streaming data and testing it across diverse domains such as healthcare, finance, and cybersecurity can enhance its generalizability and practical deployment. Additionally, integrating federated learning can ensure privacy-preserving training across multiple data sources without compromising security and performance.

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