Outlier Detection

Ninghao Liu

University of Georgia

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Overview

- Background and Problem Definition
- Outlier Detection (OD) Strategies
 - Supervised, Unsupervised, Semi-supervised
- Outlier Detection Methods
 - {Statistical, Proximity, Clustering, Classification} based
 - k-means for Outlier Detection
 - Isolation Forest
- Dealing with High-Dimensional Data in OD
 - Autoencoders for OD

1. A Real-World Application

Imagine you are a transaction auditor in a credit card company:

- Protect customers from credit card fraud.
- Pay special attention to card usages that are different from typical cases.
- Not-that-typical cases: the purchase occurs far from the owner's resident city, the purchase amount is much bigger than usual, very frequent purchases in a short time, ...
- Detect such transactions as soon as they occur and contact the card owner for verification.

1. Problem Definition

An **outlier** is a data object that deviates significantly from the rest of the objects, as if it were generated by a different mechanism.

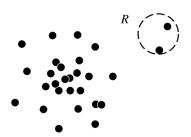


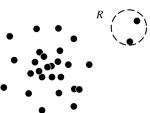
Figure 1: The objects in region R are outliers.

1. Problem Definition

Outlier detection (or **anomaly detection**) is the process of finding data objects with behaviors that are very different from expectation.

Please note that:

- Outliers are different from noisy data.
- It may be important to justify why the outliers are detected.
- Outlier detection is also related to *novelty detection* (but they are not the same).



Outlier detector training strategies:

- Supervised Learning
- Unsupervised Learning
- Semi-supervised Learning
- ...

2.1 Supervised Learning for OD

Given a dataset $\{(x_i, y_i)\}$, $1 \le i \le N$, train a classifier $f: \mathcal{X} \to \{0, 1\}$. For a new instance x, it is classified as an outlier if f(x) = 1, or a normal instance if f(x) = 0.

2.1 Supervised Learning for OD

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Challenges:

- Data imbalance: Methods are needed for handling data imbalance.
- Evaluation gap: Classification maximizes prediction accuracy, which may not be of interest in applications.
- Labeling difficulty: The sampled data for labeling may not sufficiently represent the outlier distribution.

2.2 Unsupervised Learning for OD

Given a dataset $\{x_i\}$, $1 \le i \le N$, train a detector $f: \mathcal{X} \to \{0, 1\}$. For a new instance x, it is regarded as an outlier if f(x) = 1, or a normal instance if f(x) = 0.

2.2 Unsupervised Learning for OD

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- Implicit assumption: The normal objects are "clustered".
- This assumption may not be true in some cases.
 - Outliers may also be "clustered".
 - Normal objects are diversely distributed.
- Computational challenge: We have to process a large population of non-target data entries (i.e., the normal objects) before one can touch the real meat (i.e., the outliers).

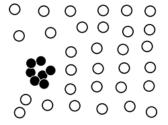


Figure 2: The black objects form outliers collectively.

2.3 Semi-supervised Learning for OD

Given a dataset $\{x_i\}$ and $\{y_j\}$, $1 \le i \le N$, $j \in \mathcal{J}$, $|\mathcal{J}| \ll N$, train a detector $f : \mathcal{X} \to \{0, 1\}$.

- Most of the instances are unlabeled, so effective unsupervised methods serve as the important basis.
- Both labeled normal/outlier instances are possible.

3. Outlier Detection Methods

- Statistical Methods
- Clustering-Based Methods
- Proximity-Based Methods
- ..

The general idea: Learn a probability density function p(x), denoting the probability that x is generated by the normal mechanism.

- Parametric methods: Assume p follows certain priori distribution (e.g., gaussian) with parameters θ to be estimated.
- Non-parametric methods: Do not assume a priori distribution.

Steps of parametric methods:

- Find a certain distribution to work with.
- ullet Estimate the parameters θ of the distribution.
- Calculate $p(x; \theta)$, and decide if the probability is low enough.

Example 1:

Univariate outlier detection using maximum likelihood. Suppose a city's average temperature values in July in the last 10 years are, in value-ascending order, 24.0°C, 28.9°C, 29.0°C, 29.1°C, 29.1°C, 29.2°C, 29.2°C, and 29.4°C.

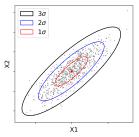
$$p(x) = \frac{1}{\sigma\sqrt{2\pi}} \cdot \exp\left(-\frac{1}{2}\frac{(x-\mu)^2}{\sigma^2}\right)$$

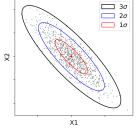
Steps of parametric methods:

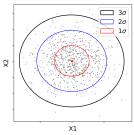
- Find a certain distribution to work with.
- ullet Estimate the parameters θ of the distribution.
- Calculate $p(x; \theta)$, and decide if the probability is low enough.

Example 2:

$$p(\mathbf{x}) = (2\pi)^{-\frac{k}{2}} \cdot \mathsf{det}(\Sigma)^{-\frac{1}{2}} \cdot \exp\left(-\frac{1}{2}(\mathbf{x} - \boldsymbol{\mu})^{\intercal} \Sigma^{-1}(\mathbf{x} - \boldsymbol{\mu})\right)$$





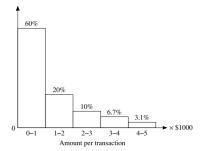


Commonly used non-parametric techniques:

- Histogram
- Kernel based density estimation

Example:

Outlier detection using a histogram. *AllElectronics* records the purchase amount for every customer transaction. Figure 12.5 uses a histogram (refer to Chapters 2 and 3) to graph these amounts as percentages, given all transactions.



Q: Should a transaction amounted \$6500 be regarded as an outlier?

Steps of histogram methods:

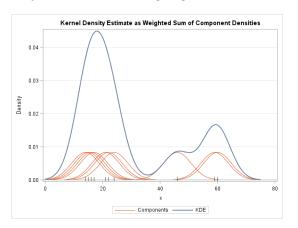
- Build the histogram from data.
- Given an instance x, check it against the histogram, and assign it with a outlier score of $\frac{1}{hist(x)}$.

Challenges

• How to choose the bin size?

Another Method: Treat an observed object as an indicator of high probability density in the surrounding region.

Another Method¹: Treat an observed object as an indicator of high probability density in the surrounding region.



¹https://blogs.sas.com/content/iml/2016/07/27/
visualize-kernel-density-estimate.html

Kernel density estimation

Given samples $\{x_1, x_2, ..., x_N\}$ and a test instance x,

$$p(x) = \frac{1}{Nh} \sum_{i=1}^{N} K(\frac{x - x_i}{h})$$
 (1)

where K() is the kernel function, h is the bandwidth parameter.

A kernel K() is a non-negative real-valued function that satisfies:

- $\int_{-\infty}^{+\infty} K(u) du = 1$,
- K(u) = K(-u).

A frequently used kernel is the RBF kernel: $K(x) = \exp(-\gamma \cdot ||x||_2^2)$.

Properties of statistical methods:

- Statistically justifiable
- Handling high-dimensional data is challenging
- Computational cost: For simple parametric models, it is linear time cost. For kernel density estimation, the model learning cost can be up to quadratic.

Possible scenarios:

- Does the instance belong to any cluster?
 - If no. then an outlier.
- Is there a large distance between the instance and the cluster to which it is closest?
 - If yes, then an outlier.
- Is the instance part of a small or sparse cluster?
 - If yes, then the whole cluster is outlying.

Usually **two steps**: (1) clustering, (2) outlier score measuring.

• Use existing outlier detection methods for the second step.

A review of k-mean clustering:

- Given a set of observations $(x_1, x_2, ..., x_N)$, k-means clustering aims to partition the N observations into $k \leq n$ sets $S = \{S_1, S_2, ..., S_k\}$.
- Formally, the objective is to find:

$$\underset{S}{\operatorname{argmin}} \sum_{i=1}^{k} \sum_{\mathbf{x} \in S_i} \|\mathbf{x} - \boldsymbol{\mu_i}\|^2, \tag{2}$$

where μ_i is the mean of S_i .

• This is equivalent to:

$$\underset{S}{\operatorname{argmin}} \sum_{i=1}^{k} \frac{1}{2|S_i|} \sum_{\mathbf{x}, \mathbf{y} \in S_i} \|\mathbf{x} - \mathbf{y}\|^2, \tag{3}$$

i.e., minimizing the pairwise squared deviations of points in the same cluster.

Equivalence between the two objectives:

$$\underset{S}{\operatorname{argmin}} \sum_{i=1}^{k} \sum_{\mathbf{x} \in S_i} \|\mathbf{x} - \boldsymbol{\mu_i}\|^2$$

$$\underset{S}{\operatorname{argmin}} \sum_{i=1}^{k} \frac{1}{2|S_i|} \sum_{\mathbf{x}, \mathbf{y} \in S_i} \|\mathbf{x} - \mathbf{y}\|^2$$

A review of k-means clustering:

- Initiate the k mean vectors $\{\mu_1, \mu_2, ..., \mu_k\}$.
- Alternate between the two steps as below.
 - **1 Assignment:** Assign each instance to the cluster with the nearest mean vector:

$$S_i^{(t)} = \{ \mathbf{x}_p : \|\mathbf{x}_p - \mu_i\| \le \|\mathbf{x}_p - \mu_j\|, \forall j, 1 \le j \le k \}$$
 (4)

Update: Recalculate mean vectors

$$u_{i} = \frac{1}{|S_{i}^{(t)}|} \sum_{\mathbf{x}_{p} \in S_{i}^{(t)}} \mathbf{x}_{p}$$
 (5)

 The algorithm converges when the assignments no longer change.

Some additional details in k-means:

- How to initialize?
 - The Forge method.
 - The Random Partition method.
- How to find the best *k*?
 - Through validation.
 - We could use the Gap Statistic ².
- Limitations of k-mean?

²" Estimating the number of clusters in a data set via the gap statistic." Journal of the Royal Statistical Society. 2001.

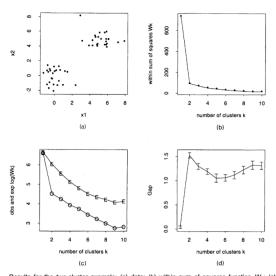


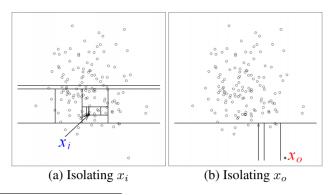
Fig. 1. Results for the two-cluster example: (a) data; (b) within sum of squares function W_k ; (c) functions $\log(W_k)$ (O) and $\hat{\mathcal{E}}_n^*\log(W_k)$) (E); (d) gap curve

Properties of clustering-based methods:

- Obtain a summary of data
- Good explainability
- The first step could be time consuming
- The detection performance largely depends on clustering quality
 - Gaussian Mixture Model
 - Hierarchical Clustering
 - Spectral Clustering
 - Deep Model Based Clustering
 - · ...

Isolation Forest (iForest) ³

- **Isolation** means "separating an instance from the rest of the instances".
- Easily being separated → More likely to be an outlier.



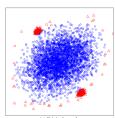
³"Isolation forest." ICDM. 2008.

Algorithm 1 : $iForest(X, t, \psi)$

Inputs: X - input data, t - number of trees, ψ - surpling size

Output: a set of t iTrees

- 1: Initialize Forest
- 2: set height limit $l = ceiling(\log_2 \psi)$
- 3: **for** i = 1 to t **do**
- 4: $X' \leftarrow sample(X, \psi)$
- 5: $Forest \leftarrow Forest \cup iTree(X', 0, l)$
- 6: end for
- 7: return Forest





```
Algorithm 2 : iTree(X, e, l)
Inputs: X - input data, e - current tree height, l - height
limit
Output: an iTree
 1: if e > l or |X| < 1 then
       return exNode\{Size \leftarrow |X|\}
 3: else
       let Q be a list of attributes in X
       randomly select an attribute q \in Q
 5:
       randomly select a split point p from max and min
 6.
       values of attribute q in X
      X_l \leftarrow filter(X, q < p)
       X_r \leftarrow filter(X, q > p)
       return inNode\{Left \leftarrow iTree(X_l, e+1, l),
 9:
                       Right \leftarrow iTree(X_r, e+1, l),
10:
11.
                       SplitAtt \leftarrow q,
                       SplitValue \leftarrow p
12:
13 end if
```

There are two types of nodes: inNode and exNode.

Not always separate each point.

Complexity $O(nt\log\psi)$

```
Algorithm 3 : PathLength(x, T, e)
Inputs: x - an instance, T - an iTree, e - current path length;
to be initialized to zero when first called
Output: path length of x
 1: if T is an external node then
      return e + c(T.size) \{c(.) \text{ is defined in Equation } 1\}
 3: end if
 4: a \leftarrow T.splitAtt
 5: if x_a < T.splitValue then
      return PathLength(x, T.left, e + 1)
 7: else \{x_a \geq T.splitValue\}
      return PathLength(x, T.right, e + 1)
 9: end if
```

The outlier score is an instance x is:

$$s(\mathbf{x}, \psi) = 2^{-\frac{E(h(\mathbf{x}))}{c(\psi)}} \tag{6}$$

- h(x): a single path length from root to x.
- $c(\psi)$: normalization
- iTrees have an equivalent structure to Binary Search Tree (BST).
- The estimation of E(h(x)) for external node terminations is the same as the unsuccessful search in BST.
- ullet $c(\psi)=$ the average path length of unsuccessful search in BST.

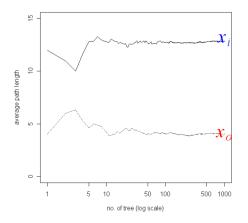
The outlier score is an instance x is:

$$s(\mathbf{x}, \psi) = 2^{-\frac{E(h(\mathbf{x}))}{c(\psi)}} \tag{7}$$

- If instances return $s \approx 1$, then they are outliers.
- If instances have $s \ll 0.5$, then they are quite safe to be seen as normal instances.
- If all the instances return $s \approx 0.5$, then the entire sample does not really have any distinct outliers.

Two variables to choose in iForest:

- number of trees to build t (default: 100)
- sub-sampling size ψ (default: 256)



4 Outlier Detection in High-Dimensional Data

Dimensionality Curse: Various phenomena that arise when analyzing data in high-dimensional spaces that do not occur in low-dimensional ones ⁴.

- The pairwise distances between different samples in the space converging to the same value as the dimensionality of the data increases.
- As the dimensionality increases, the distance between objects may be heavily dominated by noise.
- The *proximity* or *similarity* between samples may not be qualitatively relevant in higher dimensions.

⁴https://en.wikipedia.org/wiki/Curse_of_dimensionality

4 Outlier Detection in High-Dimensional Data

Dimensionality Reduction: compress data

- Classical approaches
 - E.g., Principle Component Analysis (PCA)
- Deep learning approaches.
 - E.g., Robust Deep Autoencoder (RDA) ⁵

⁵Zhou, Chong, and Randy C. Paffenroth. "Anomaly detection with robust deep autoencoders." KDD. 2017.

- A feature/dimensionality reduction technique.
- A relevant (but different) technique is feature selection.
- The goal of PCA is to transform the high-dimensional features to a lower-dimensional space.
- PCA is unsupervised.

What is a good data transformation?

What is a good data compression (transformation)?

- High compression rate (smaller data size).
- Key characteristics are maintained (keep more information).

In the deep learning era, the transformed/compressed data is also called as:

- the "representation" of data
- "latent representation"
- "embeddings"

- Let us start from the simple case where PCA projects the features to a 1-dimensional space.
- Suppose we have n p-dimensional (p > 1) instances, $x_1, x_2, ..., x_n \in \mathbb{R}^p$.
- We want to learn a mapper $a \in \mathbb{R}^p$ that $a^T x_i = z_i$. Here z_i is where x_i locates in the 1-dimensional space.
- PCA tries to solve:

$$\mathbf{a}^* = \operatorname{argmax}_{\|\mathbf{a}\|=1} \underbrace{\frac{1}{n} (z_i - \overline{z})^2}_{\text{variance}}, \tag{8}$$

where PCA tries to find the projection with the maximum variance in reduced data.

PCA tries to solve:

$$a^* = \operatorname{argmax}_{\|\mathbf{a}\|=1} \underbrace{\frac{1}{n} (z_i - \overline{z})^2}_{\text{variance}}.$$
 (9)

- Q1: Why ||a|| = 1?
- Q2: What does the objective imply?
 - Trying to make the data points distinguishable in the new space.
 - Why?

Since

$$\bar{z} = \frac{1}{n} \sum_{i=1}^{n} z_i = \frac{1}{n} \sum_{i=1}^{n} \boldsymbol{a}^T \boldsymbol{x}_i = \boldsymbol{a}^T (\frac{1}{n} \sum_{i=1}^{n} \boldsymbol{x}_i) = \boldsymbol{a}^T \bar{\boldsymbol{x}},$$

the problem can be written as

$$\mathbf{a}^* = \underset{||\mathbf{a}||=1}{\operatorname{arg max}} \frac{1}{n} \sum_{i=1}^n (z_i - \bar{z})^2$$

$$= \underset{||\mathbf{a}||=1}{\operatorname{arg max}} \frac{1}{n} \sum_{i=1}^n (\mathbf{a}^T \mathbf{x}_i - \bar{z})^2$$

$$= \underset{||\mathbf{a}||=1}{\operatorname{arg max}} \frac{1}{n} \sum_{i=1}^n (\mathbf{a}^T \mathbf{x}_i - \mathbf{a}^T \bar{\mathbf{x}})^2$$

$$= \underset{||\mathbf{a}||=1}{\operatorname{arg max}} \frac{1}{n} \sum_{i=1}^n \mathbf{a}^T (\mathbf{x}_i - \bar{\mathbf{x}}) (\mathbf{x}_i - \bar{\mathbf{x}})^T \mathbf{a}$$

$$= \underset{||\mathbf{a}||=1}{\operatorname{arg max}} \mathbf{a}^T \underbrace{\left(\frac{1}{n} \sum_{i=1}^n (\mathbf{x}_i - \bar{\mathbf{x}}) (\mathbf{x}_i - \bar{\mathbf{x}})^T\right)}_{p \times p \text{ covariance matrix}} \mathbf{a}$$

$$= \underset{||\mathbf{a}||=1}{\operatorname{arg max}} \mathbf{a}^T C \mathbf{a},$$

where $C = \frac{1}{n} \sum_{i=1}^{n} (x_i - \bar{x})(x_i - \bar{x})^T$, denotes the covariance matrix.

The problem is

$$\max_{\mathbf{a}} \mathbf{a}^{\mathsf{T}} \mathbf{C} \mathbf{a}$$

$$s.t., \|\mathbf{a}\|^2 = 1.$$
(10)

How to solve this?

Eigen-Decomposition

A square $n \times n$ matrix C with n linearly independent eigenvectors can be factorized as:

$$C = U\Lambda U^{-1} \tag{11}$$

- U is a $n \times n$ matrix whose columns are eigenvectors of C,
- $oldsymbol{\Lambda}$ is a diagonal matrix whose diagonal elements are the corresponding eigenvalues.

If C is symmetric, then

$$C = U\Lambda U^{\mathsf{T}} = \sum_{i=1}^{n} \lambda_{i} u_{i} u_{i}^{\mathsf{T}}. \tag{12}$$

Q: What about mapping data into a k-dimensional space? A: The PCA problem is then defined as:

$$\mathbf{A}^* = \underset{\mathbf{A} \in \mathbb{R}^{p \times k}: \mathbf{A}^{\mathsf{T}} \mathbf{A} = \mathbf{I}_k}{\operatorname{argmax}} \operatorname{trace}(\mathbf{A}^{\mathsf{T}} \mathbf{C} \mathbf{A})$$
 (13)

where $\mathbf{A} = [\mathbf{a}_1, \mathbf{a}_2, ..., \mathbf{a}_k] \in \mathbb{R}^{p \times k}$. PCA requires different projection vectors to be orthogonal.

$$\operatorname{trace}(\mathbf{A}^{\mathsf{T}}\mathbf{C}\mathbf{A}) = \sum_{i=1}^{k} (\mathbf{a}_{i}^{\mathsf{T}}\mathbf{C}\mathbf{a}_{i}), \tag{14}$$

which is a natural extension of 1-D PCA.

Q: How to solve this?

When k = 2, we want to solve

$$\max_{\mathbf{a}_{2}} \ \mathbf{a}_{2}^{\mathsf{T}} \mathbf{C} \mathbf{a}_{2}$$
 $s.t., \ \|\mathbf{a}_{i}\|^{2} = 1$
 $\mathbf{a}_{2}^{\mathsf{T}} \mathbf{a}_{1} = 0.$ (15)

How to solve this?

We then have the following theorem.

Theorem. (*Ky Fan*) Let $\mathbf{H} \in \mathbb{R}^{n \times n}$ be a symmetric matrix with eigenvalues

$$\lambda_1 \geq \lambda_2 \geq \cdots \geq \lambda_n$$

and the corresponding eigenvectors $\boldsymbol{U} = [\boldsymbol{u}_1, \dots, \boldsymbol{u}_n]$. Then

$$\lambda_1 + \cdots \lambda_k = \max_{m{A} \in \mathbb{R}^{n imes k} : m{A}^T m{A} = m{I}_k} \operatorname{trace}\left(m{A}^T m{H} m{A}
ight).$$

And the optimal A^* is given by $A^* = [u_1, ..., u_k]Q$ with Q an arbitrary orthogonal matrix.

• Note that in Eqn. (2), the covariance matrix C is a symmetric matrix. Given the above theorem, we directly obtain

$$\lambda_1 + \cdots \lambda_k = \underset{m{A} \in \mathbb{R}^{n \times k} : m{A}^T m{A} = m{I}_k}{\operatorname{arg \, max}} \operatorname{trace} \left(m{A}^T m{C} m{A} \right),$$
 $m{A}^* = [m{u}_1, \dots, m{u}_k] m{Q},$

where $\lambda_1, \ldots, \lambda_k$ are the k largest eigenvalues of the covariance matrix C, and the solution A^* is the matrix whose columns are corresponding eigenvectors.

② It also follows from the above theorem that solutions to PCA are not unique, and they differ by an orthogonal matrix. We used the special case where Q = I, i.e., $A^* = [u_1, \dots, u_k]$.

Using PCA, we obtain the mappers $\mathbf{A} = \{\mathbf{u}_i\}$, $1 \le i \le k$. An original data instance \mathbf{x} will be mapped to $\mathbf{z} = [z_1, z_2, ..., z_k]$. After mapping all the data, outlier detection could be done in the \mathbf{z} -space.

- Apply the detection methods we already discussed.
- Choose the mappers corresponding to higher or lower eigenvalues?
- How many k's to choose?

There is another way to understand PCA from the **data reconstruction** perspective.

- Assume X is centered.
- Data mapping: $Z = A^T X$.
- Reconstructed data: $\tilde{\mathbf{X}} = \mathbf{A}\mathbf{Z} = \mathbf{A}\mathbf{A}^{\mathsf{T}}\mathbf{X}$.
- Reconstruction error: $\|X \tilde{X}\|_F = \|X AA^{\mathsf{T}}X\|_F$.
- It has also been shown that:

$$\mathbf{A}\mathbf{A}^{\mathsf{T}}\mathbf{X} = \mathbf{B}^* = \underset{\mathbf{B}:\mathsf{rank}(\mathbf{B}) \leq k}{\mathsf{argmin}} \|\mathbf{X} - \mathbf{B}\|_{F}.$$
 (16)

PCA projects the high-dimensional features to a lower-dimensional space with minimum reconstruction error.

- Keep most of the information = Be able to recover.
- Instances with large reconstruction errors $||x AA^{T}x||_{2}$ are outliers.

In previous parts, we need to compute C first before doing PCA. This step could be costly if p or n is large. Can this step be avoided?

Singular Value Decomposition:

The singular value decomposition (SVD) of an $m \times n$ real matrix R (we assume $m \ge n$) can be written as:

$$R = U\Sigma V^{\mathsf{T}},\tag{17}$$

where $U \in \mathbb{R}^{m \times m}$, $\Sigma \in \mathbb{R}^{m \times n}$, and $V \in \mathbb{R}^{n \times n}$, $UU^{\mathsf{T}} = U^{\mathsf{T}}U = I_{m \times m}$, and $VV^{\mathsf{T}} = V^{\mathsf{T}}V = I_{n \times n}$.

1. Pre-process data.

$$egin{array}{lll} oldsymbol{X} &=& [oldsymbol{x}_1, oldsymbol{x}_2, \ldots, oldsymbol{x}_n] \in \mathbb{R}^{p imes n}, \ ar{oldsymbol{X}} &=& rac{1}{n} oldsymbol{X} oldsymbol{1}_n \in \mathbb{R}^{p imes 1}, \ ar{oldsymbol{X}} &=& (oldsymbol{X} - ar{oldsymbol{X}} oldsymbol{1}_n^T) \in \mathbb{R}^{p imes n}, \end{array}$$

- 2. We know $C = \frac{1}{n} \sum_{i=1}^{n} (x_i \overline{x})(x_i \overline{x})^{\mathsf{T}} = \frac{1}{n} \tilde{X} \tilde{X}^{\mathsf{T}}$.
- 3. If the SVD of \tilde{X} is $\tilde{X} = U\tilde{\Sigma}V$, the columns of U (left-singular vectors) are eigenvectors of $\tilde{X}\tilde{X}$.
- 4. We do not perform eigen-value decomposition of C, but do singular-value decomposition of \tilde{X} , to obtain the mappers U.

A more efficient way of computing PCA can be described as:

$$\underbrace{m{X}}_{p imes n} o \underbrace{m{ ilde{X}}}_{p imes n} o m{ ilde{X}} = m{U} m{ ilde{\Sigma}} m{V}^T o m{G} = m{U}_k \in \mathbb{R}^{p imes k}$$

$$oldsymbol{G}^T ilde{oldsymbol{X}} = oldsymbol{Z} \in \mathbb{R}^{k imes n}$$

4.1 Connection between PCA and Autoencoders

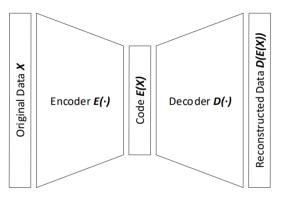


Figure 3: An autoendoer.

- What does each module in PCA correspond to?
- How to go deeper?

Deep Autoencoders

- A feed-forward multi-layer neural network.
- The desired output is the input itself.
 - Looks trivial?
 - Making hidden representation to be low-dimensional.
- The loss function of training an autoencoder:

$$\min_{D,E} \|\boldsymbol{X} - D(E(\boldsymbol{X}))\| \tag{18}$$

Robust PCA (RPCA)

- Try to reduce the sensitivity of PCA to outliers. Intuition?
- Split the data matrix into two parts

$$X = L + S, \tag{19}$$

where \boldsymbol{L} is a low-rank matrix, and \boldsymbol{S} is a sparse matrix.

Robust PCA (RPCA)

• The problem can be formulated as:

$$\min_{\boldsymbol{L},\boldsymbol{S}} \|\boldsymbol{L}\|_* + \lambda \|\boldsymbol{S}\|_1
s.t. \|\boldsymbol{X} - \boldsymbol{L} - \boldsymbol{S}\|_F^2 = 0,$$
(20)

where $\|\cdot\|_*$ is the nuclear norm, $\|\cdot\|_1$ is the L1 norm.

 If we let D and E to be linear mappers, then PCA is the optimal solution for recovering L.

Robust Deep Autoencoders (RDA)

• The problem can be formulated as:

$$\min_{\boldsymbol{L},\boldsymbol{S}} \|\boldsymbol{L} - D_{\theta}(\boldsymbol{E}_{\theta}(\boldsymbol{L}))\|_{2} + \lambda \|\boldsymbol{S}\|_{1}$$

$$s.t. \, \boldsymbol{X} - \boldsymbol{L} - \boldsymbol{S} = 0,$$
(21)

where both the encoder and the decoder are parameterized by $\theta.$

- The first term encourages **L** to contain the "main features" in data.
- ullet λ plays an essential role:
 - a smaller $\lambda \to \text{encourage more data into } \boldsymbol{S}$
 - a larger $\lambda \rightarrow$ include less data in ${\bf S}$

Robust Deep Autoencoders (RDA)

- How to choose D_{θ} and E_{θ} ?
 - Any commonly used deep autoencoder architectures could be used.
 - The paper choose a simple architecture:

$$E_{\theta}(\mathbf{x}) = g(\mathbf{W}\mathbf{x} + b_{E})$$

$$D_{\theta}(\mathbf{x}) = g(\mathbf{W}^{\mathsf{T}}E_{\theta}(\mathbf{x}) + b_{D})$$
(22)

• $g(x) = \frac{1}{1 + \exp(-x)}$, a nonlinear transformation.

Robust Deep Autoencoders (RDA)

- Is $||S||_1$ always a good choice?
- What if each anomaly point is not a matrix entry, but the whole row or column?

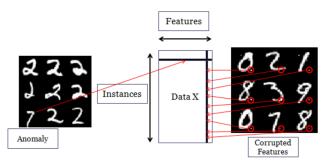


Figure 4: Two examples of group anomalies.

Regularization

- $\min \|x\|_1$ vs $\min \|x\|_2$
- Which one encourages a sparse x?
- An example of taxes.
- Can we combine them?

Robust Deep Autoencoders (RDA)

- $I_{2,1}$ normalization encourages structured/group sparsity.
- $\|X\|_{2,1} = \sum_{j=1}^{n} \|x_j\|_2 = \sum_{j=1}^{n} (\sum_{i=1}^{m} |x_{i,j}|^2)^{1/2}$.
- $\|X^{\mathsf{T}}\|_{2,1}$?

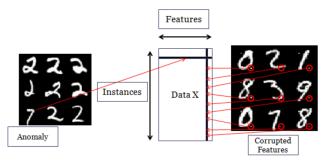


Figure 5: Two examples of group anomalies.

Robust Deep Autoencoders (RDA)

• The problem can be formulated as:

$$\min_{\boldsymbol{L},\boldsymbol{S}} \|\boldsymbol{L} - D_{\theta}(\boldsymbol{E}_{\theta}(\boldsymbol{L}))\|_{2} + \lambda(\|\boldsymbol{S}\|_{1} \text{ or } \|\boldsymbol{S}\|_{2,1} \text{ or } \|\boldsymbol{S}^{\mathsf{T}}\|_{2,1})$$
s.t. $\boldsymbol{X} - \boldsymbol{L} - \boldsymbol{S} = 0$,

- Just denoising: choose $\|S\|_1$.
- To find abnormal data instances: choose $\|S^{\mathsf{T}}\|_{2,1}$.

Robust Deep Autoencoders (RDA)

• How to solve the problem?

$$\min_{\boldsymbol{L},\boldsymbol{S}} \|\boldsymbol{L} - D_{\theta}(E_{\theta}(\boldsymbol{L}))\|_{2} + \lambda(\|\boldsymbol{S}\|_{1} \text{ or } \|\boldsymbol{S}\|_{2,1} \text{ or } \|\boldsymbol{S}^{\mathsf{T}}\|_{2,1})$$
s.t. $\boldsymbol{X} - \boldsymbol{L} - \boldsymbol{S} = 0$,

- Alternating Direction Method of Multipliers (ADMM):
 - Divide the objective into several pieces.
 - Optimizes one of the pieces while keeping the others fixed.
- How to update L with respect to $\min_{L} \|L D_{\theta}(E_{\theta}(L))\|_{2}$?
- How to update S with respect to $\min_{S} ||S||_1$?

Robust Deep Autoencoders (RDA)

The algorithm of solving

$$\min_{\boldsymbol{L},\boldsymbol{S}} \|\boldsymbol{L} - D_{\theta}(E_{\theta}(\boldsymbol{L}))\|_{2} + \lambda(\|\boldsymbol{S}\|_{1} \text{ or } \|\boldsymbol{S}\|_{2,1} \text{ or } \|\boldsymbol{S}^{\mathsf{T}}\|_{2,1})$$
s.t. $\boldsymbol{X} - \boldsymbol{L} - \boldsymbol{S} = 0$.

- Initialize θ , **S**
- while not converge:
 - 0 L = X S
 - **2** Minimize $\|\mathbf{L} D_{\theta}(E_{\theta}(\mathbf{L}))\|_2$ with gradient descent

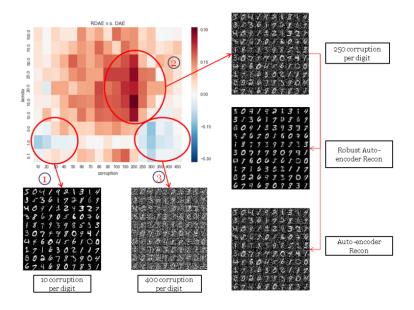
 - S = X L
 - Optimize S using a proximal operator
 - 6 Check convergence.

Experiment Design

- Datasets
- Research Problems
 - Baseline methods
 - Metrics
 - Visualization
- Effect of hyperparameters or modules

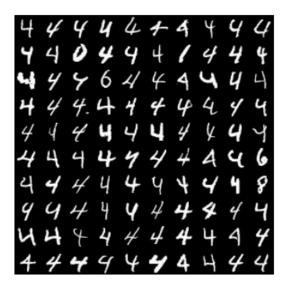
Experiment Design in RDA

- Datasets: MNIST
- Research Problem 1: Whether excluding from data an independent sparse matrix using $\|S\|_1$ could help the autoencoder in capturing the essential patterns in data.
 - Baseline methods: Autoencoder
 - Metrics: Classification error
- Experiment design:
 - Extract hidden features from $E_{\theta}(\mathbf{L})$.
 - Use the hidden features to build a classifier to classify digits.
 - Lower error \rightarrow Better model.



Experiment Design in RDA

- Datasets: MNIST
- Research Problem 2: Is $\|S\|_{2,1}$ regularized RDA effective in detecting outliers?
 - Baseline method: Isolation Forest (IS)
 - Evaluation metric: F1-score
- Experiment design:
 - Compare detected outliers with the ground truths.
 - 0.64 (RDA) vs 0.37 (IS)



Experiment Design in RDA

Datasets: MNIST

• Research Problem 3: Effect of λ ?

