

Anomaly Detection:

Clustering Based, Statistical Approaches and Reconstruction Based



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OUTLOOK

- Clustering Based
- Statistical Approaches
- Reconstruction Based

CLUSTERING BASED

■ ADVANTAGE

- unsupervised algorithm
- existing clustering algorithms can be plugged in

■ DRAWBACKS

- if the data object does not have a natural clustering or the clustering algorithm is not able to detect the natural clusters, the techniques may fail
- computationally expensive
 - using indexing structures (k-d tree, R* tree) may alleviate this problem
- in high dimensional spaces, data is sparse and distances between any two data objects may become quite similar
- can be difficult to decide on a clustering technique
- can be difficult to decide on number of clusters
- outliers can distort the clusters

CLUSTERING BASED

- **KEY ASSUMPTION:** normal data instances belong to large and dense clusters, while anomalies do not belong to any significant cluster.

- **GENERAL APPROACH:**
 - cluster data objects into a finite number of clusters
 - analyze each data object with respect to its closest cluster
 - anomalous data objects
 - do not fit into any cluster (residuals from clustering)
 - belong to small clusters
 - are located in low density clusters
 - are far from other data objects within the same cluster

CLUSTERING BASED: BASIC ALGORITHM

- Fixed-width clustering is first applied
 - the first data object is the center of first cluster
 - two data objects p_1 and p_2 are “near” if $d(p_1, p_2) < \varepsilon$ (ε is a user specified parameter)
 - if every subsequent data objects is “near”, add to the current cluster
 - otherwise create a new cluster

- Data objects in small clusters are anomalies

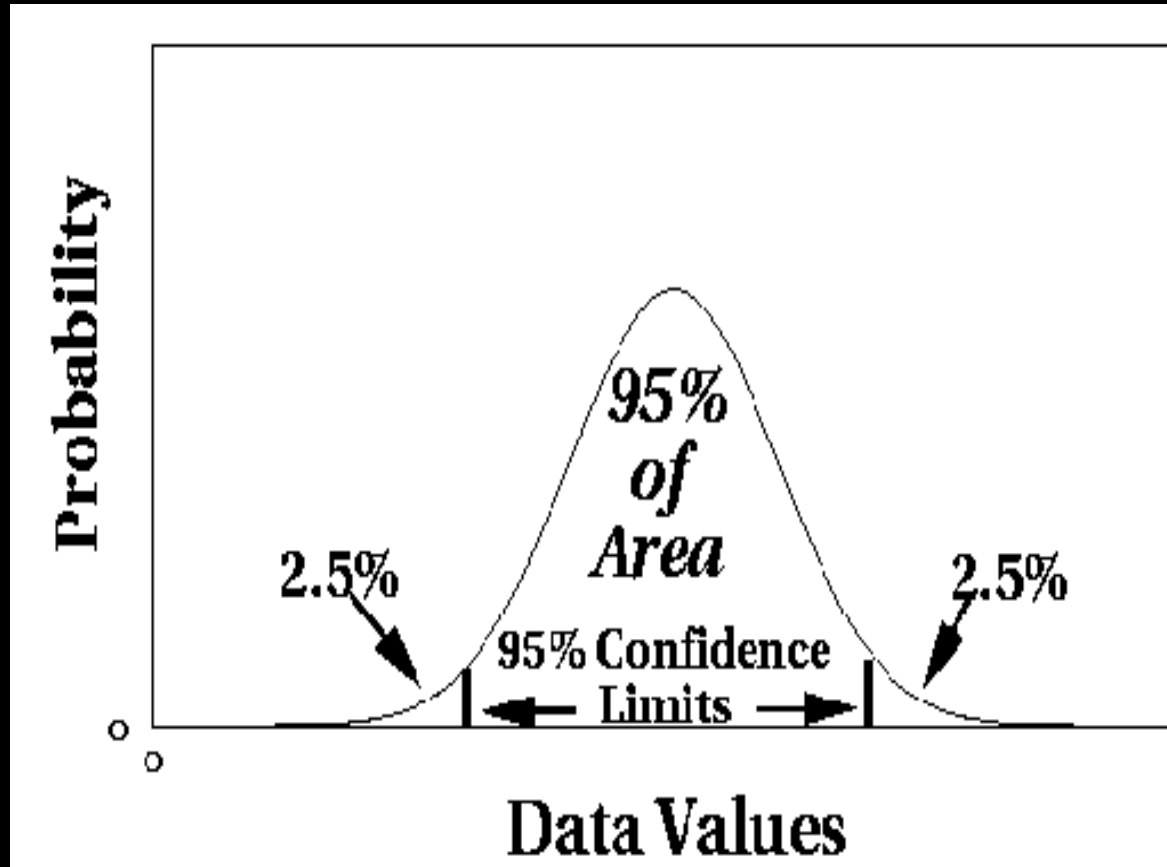
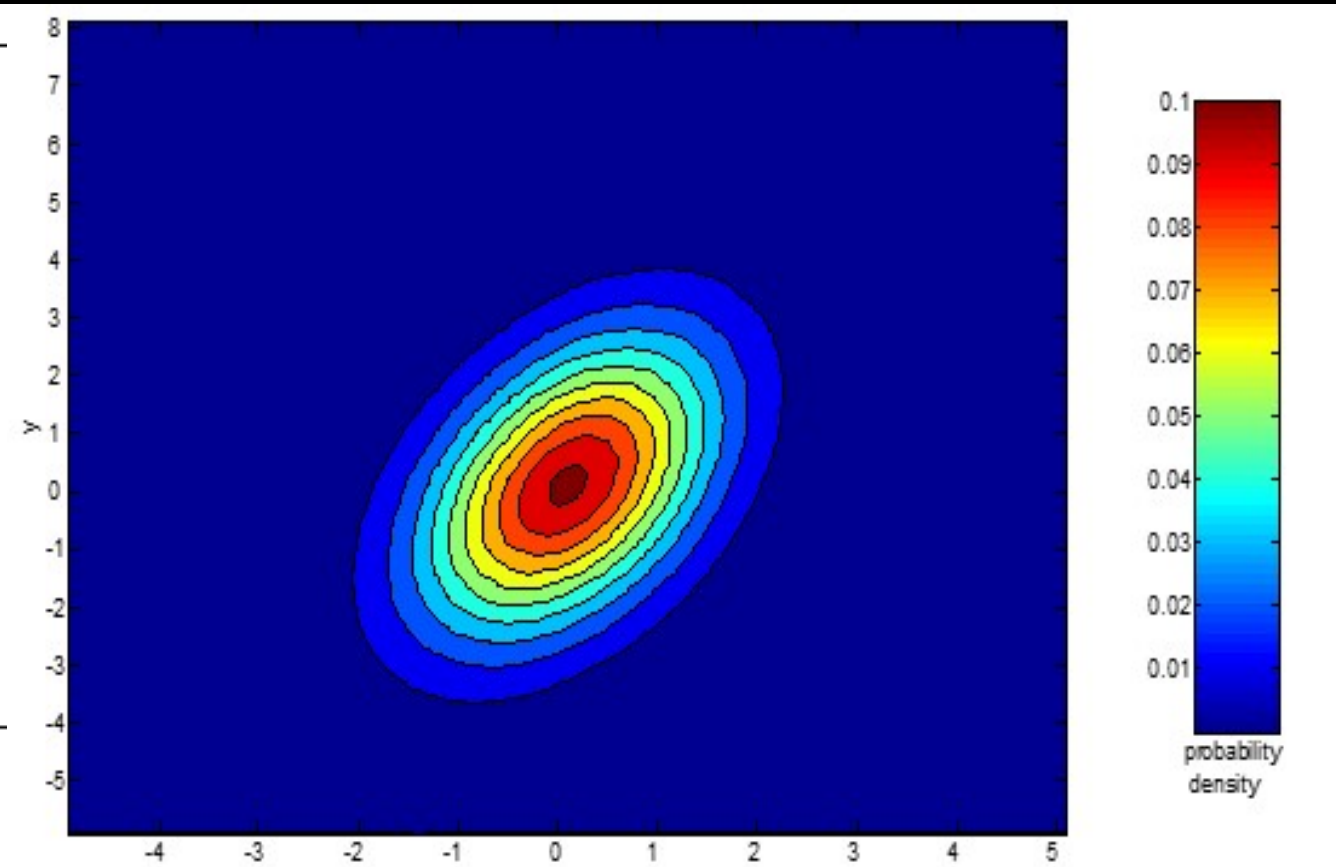
CLUSTERING BASED: CLUSTER BASED LOCAL OUTLIER FACTOR (CBLOF)

- An data object is a **cluster-based outlier** if it does not strongly belong to any cluster
 - for **prototype-based clusters**, an data object is an outlier if it is **not close enough to a cluster center**
 - outliers can impact the clustering produced
 - for **density-based clusters**, an data object is an outlier if **its density is too low**
 - can't distinguish between noise and outliers
 - for **graph-based clusters**, an data object is an outlier if **it is not well connected**

STATISTICAL APPROACHES

Probabilistic definition of an outlier: an outlier is an **data object that has a low probability with respect to a probability distribution model of the data.**

- Usually assume a **parametric model describing the distribution of the data** (e.g., normal distribution)
- Apply a statistical test that depends on
 - data distribution
 - parameters of distribution (e.g., mean, variance)
 - number of expected outliers (confidence limit)
- Issues
 - identifying the distribution of a data set
 - heavy tailed distribution
 - number of attributes
 - is the data a mixture of distributions?

STATISTICAL APPROACHES: NORMAL DISTRIBUTION**one-dimensional Gaussian****two-dimensional Gaussian**

STATISTICAL APPROACHES: GRUBBS'S TEST

- Detects outliers in univariate data
- Assumes data comes from normal distribution
- Detects one outlier at a time, remove the outlier, and repeat
 - H_0 : there is no outlier in data
 - H_1 : there is at least one outlier

- Grubbs's test statistic:
$$G = \frac{\max |X - \bar{X}|}{s}$$

- Reject H_0 if:
$$G > \frac{(N-1)}{\sqrt{N}} \sqrt{\frac{t^2_{(\alpha/(2N), N-2)}}{N-2 + t^2_{(\alpha/(2N), N-2)}}}$$

STATISTICAL APPROACHES: LIKELIHOOD APPROACH

- Assumes the data set D contains samples from a mixture of two probability distributions:
 - M (majority/non-anomalous distribution)
 - A (anomalous distribution)
- General Approach:
 - initially, assumes all the data objects belong to M
 - let $LL_t(D)$ be the log likelihood of D at time t
 - for each data object x_t that belongs to M , move it to A
- Let $LL_{t+1}(D)$ be the new log likelihood
- Computes the difference, $\Delta = LL_t(D) - LL_{t+1}(D)$
- If $\Delta > c$ (some threshold), then x_t is declared as an anomaly and moved permanently from M to A

STATISTICAL APPROACHES: LIKELIHOOD APPROACH

- Data distribution, $D = (1 - \lambda)M + \lambda A$
- M is a probability distribution estimated from data
 - can be based on any modeling method (naïve Bayes, maximum entropy, etc.)
- A is initially assumed to be uniform distribution

- Likelihood at time t :

$$L_t(D) = \prod_{i=1}^N P_D(x_i) = \left((1 - \lambda)^{|M_t|} \prod_{x_i \in M_t} P_{M_t}(x_i) \right) \left(\lambda^{|A_t|} \prod_{x_i \in A_t} P_{A_t}(x_i) \right)$$

$$LL_t(D) = |M_t| \log(1 - \lambda) + \sum_{x_i \in M_t} \log P_{M_t}(x_i) + |A_t| \log \lambda + \sum_{x_i \in A_t} \log P_{A_t}(x_i)$$

STATISTICAL APPROACHES: **STRENGTHS AND WEAKNESSES**

- Firm mathematical foundation
- Can be very efficient
- Good results if distribution is known
- In many cases, data distribution may not be known
- For high dimensional data, it may be difficult to estimate the true distribution
- Anomalies can distort the parameters of the distribution

RECONSTRUCTION BASED

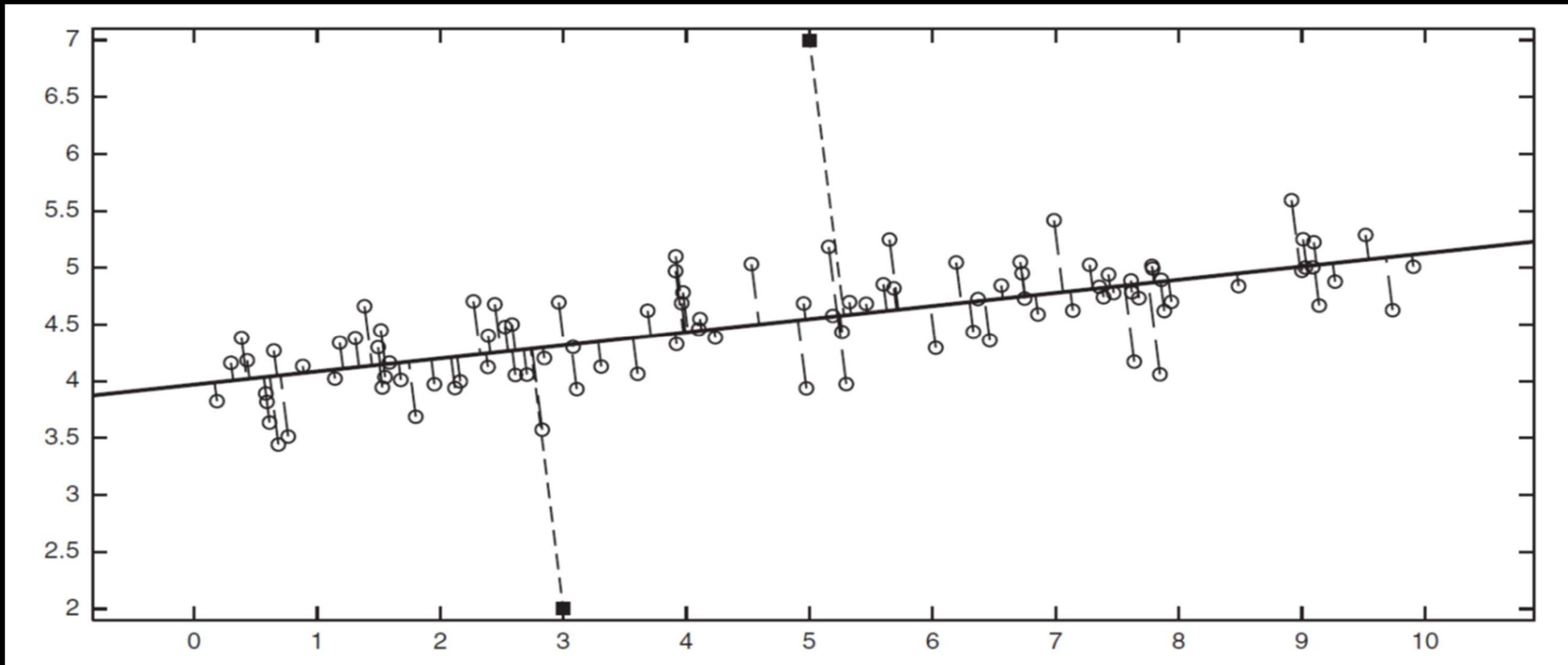
- Based on assumptions there are patterns in the distribution of the normal class that can be captured using lower-dimensional representations
- Reduce data to lower dimensional data
 - e.g. use Principal Components Analysis (PCA) or auto-encoders
- Measure the reconstruction error for each object
 - the difference between original and reduced dimensionality version

RECONSTRUCTION BASED: RECONSTRUCTION ERROR

- Let x be the original data object
- Find the representation of the data object in a lower dimensional space
- Project the object back to the original space
- Call this object \hat{x}

$$\text{Reconstruction Error} = \|x - \hat{x}\|$$

- Objects with large reconstruction error are anomalies

RECONSTRUCTION BASED: RECONSTRUCTION OF TWO DIMENSIONAL DATA

RECONSTRUCTION BASED: **PRINCIPAL COMPONENTS ANALYSIS**

- Compute the principal components of the dataset
- For each test data object, compute its projection on these components
- If y_i denotes the i^{th} component, then the following has a chi-squared distribution

$$\sum_{i=1}^q \frac{y_i^2}{\lambda_i} = \frac{y_1^2}{\lambda_1} + \frac{y_2^2}{\lambda_2} + \dots + \frac{y_q^2}{\lambda_q} \quad q < n$$

- an data object is anomalous, if for a given significance level α

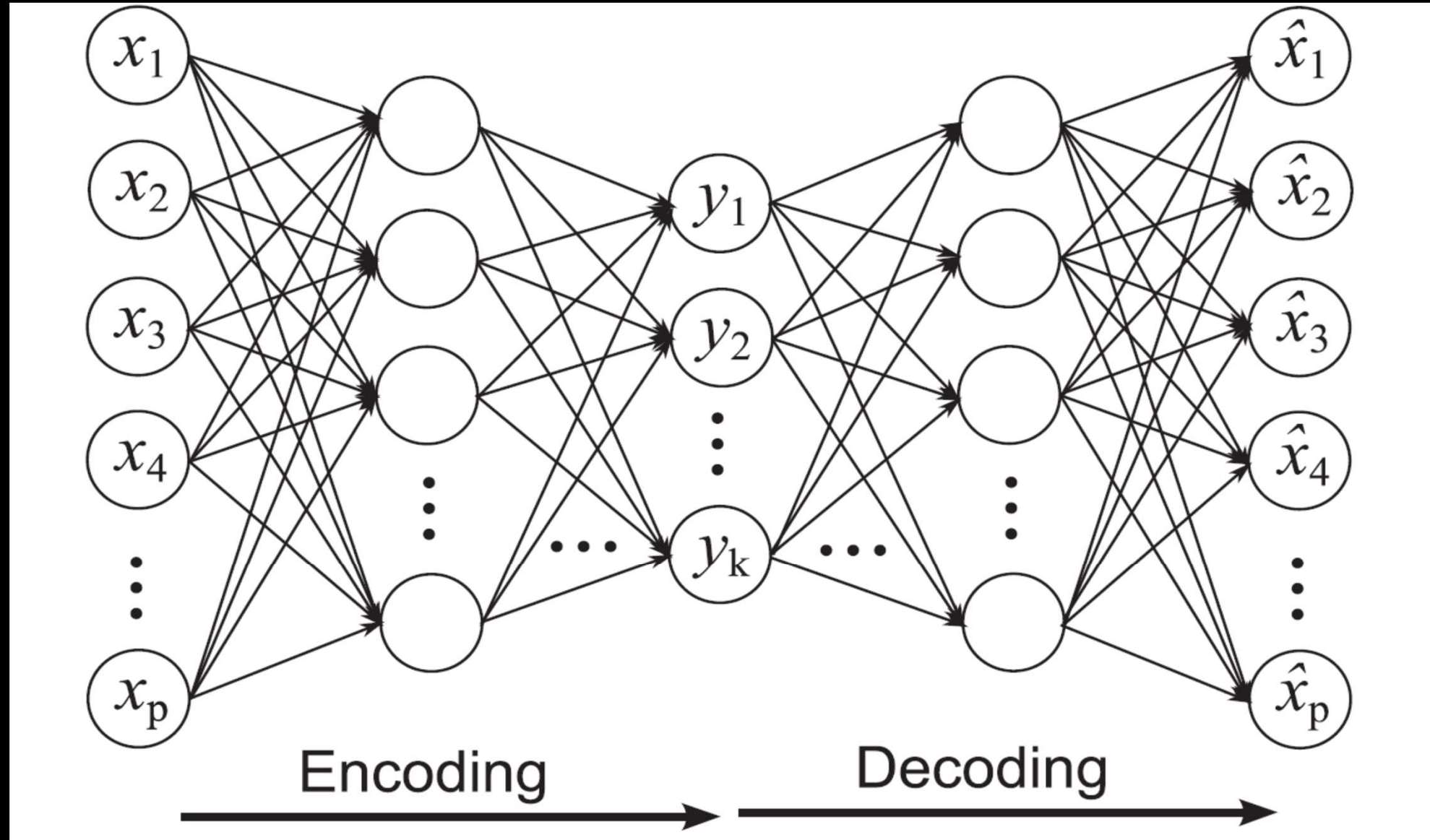
$$\sum_{i=1}^q \frac{y_i^2}{\lambda_i} > \chi_q^2(\alpha)$$

- Another measure is to observe last few principal components
 - anomalies have high value for the above quantity

$$\sum_{i=p-r+1}^p \frac{y_i^2}{\lambda_i}$$

RECONSTRUCTION BASED: AUTO-ENCODER

- An auto-encoder is a multi-layer neural network
- The number of input and output neurons is equal to the number of original attributes



RECONSTRUCTION BASED

■ STRENGTHS

- does not require assumptions about distribution of normal class
- can use many dimensionality reduction approaches

■ WEAKNESSES

- the reconstruction error is computed in the original space
 - this can be a problem if dimensionality is high

RECAP

- Clustering Based
- Statistical Approaches
- Reconstruction Based