synthetic minority oversampling using multivariate gaussian distribution

namagiri December 2023

1 Abstract

Highly imbalanced data is a common phenomenon from cases of fraudulent transactions to several medical datasets. Without rebalancing the dataset if it fits into any learning algorithm will result in a highly biased model towards the majority class to address this issue we have employed a technique of oversampling using a new smote(synthetic minority oversampling) strategy where k nearest neighbours are picked from the minority class and fitted to multivariate gaussian distribution to generate points which populate regions with high probability

2 Methodology

Partially guided synthetic minority oversampling using Gaussian kernel interpolation.

Let's consider the matrix X:

$$X = \begin{bmatrix} a_{11} & a_{12} & \cdots & a_{1n} \\ a_{21} & a_{22} & \cdots & a_{2n} \\ \vdots & \vdots & \ddots & \vdots \\ a_{a1} & a_{a2} & \cdots & a_{na} \end{bmatrix}$$

The dimensions of matrix X are $n \times a$, where n refers to rows/instances and a refers to attributes.

Let t be the target attribute and $X[t] = \{0, 1\}$, representing a binary classification task.

 $X = \text{Maj} \cup \text{Min}$, where Maj is the majority class and Min is the minority class.

Maj
$$[t] = \{0\}$$
 and Min $[t] = \{1\}$.
Let $\theta = \frac{n(\text{Min})}{n(\text{Maj})}$, where $\theta \in [0, 1]$.
 $n()$ denotes the cardinality of the class, and $\theta \ll 1$.

Algorithm 1 Selecting ensemble classifier

```
    Input: Minority data set Min ,Majority data set Maj, ratio balancer δ
    Output: Ensemble classifier E
    bounding the number of subsets b = ∫ n(Maj) / δ*n(Min) ∫
    Selecting b subsets from Maj | ∪ l=1 Maj(i) = Maj and ⋂ l=1 Maj(i) = ∅, ∀i, n(Maj(i)) > δ*n(Min)
    Number of synthetic samples to be generated
    sample number = min<sub>i∈[1,b]</sub> n(Maj(i)) - n(Min)
    Gen = Algorithm<sub>2</sub>(Min, Samplenumber)
    Min = Min ∪ GEN
    C<sub>i</sub>(Maj(i), Min)
    Getting b classifiers and trained to evaluate the best classifier BEST = decide({C<sub>1</sub>, C<sub>2</sub>, C<sub>3</sub>, ..., C<sub>b</sub>})
    Return: BEST
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Algorithm 2 Generating synthetic samples

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1: Input: Minority class Min, Sample number
 2: Output: Synthesized samples GEN
 3: Start s = sample \, size \, and GEN \leftarrow \phi
 4: for i \in \text{Min do}
          l = Sort(k \text{ nearest neighbours of } i)
         \begin{array}{l} l\_first \leftarrow \text{First } \frac{l}{4} \text{ elements of data} \\ l\_last \leftarrow \text{Last } \frac{l}{4} \text{ elements of data} \end{array}
 6:
 7:
          l_1 = l\_last \cup l\_first
 8:
 9:
          f \leftarrow \text{Multivariate gaussian distribution}(l_1)
                                                                                   ▶ Interpolated surface
10:
          Evaluation:
          l\_middle \leftarrow Middle \frac{l}{2} elements of data
11:
          for x in l-middle do
12:
               y \leftarrow \text{Evaluate}(f, x) \triangleright \text{Evaluate the function at the middle elements}
13:
14:
          end for
          s \leftarrow s - n(y)
15:
16:
          GEN \leftarrow GEN \cup y
          Min \leftarrow Min - l
17:
18: end for
19: Return: GEN
```

fitting the l1 vector into the distribution

Let $\mathbf{X} = (X_1, X_2, \dots, X_n)^T$ be a vector-valued random variable with a multivariate normal distribution. It has mean $\boldsymbol{\mu} \in R^n$ and covariance matrix $\boldsymbol{\Sigma} \in S_n^{++}$. The probability density function (pdf) of \mathbf{X} is given by:

$$p(\mathbf{x}; \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \frac{1}{(2\pi)^{n/2} |\boldsymbol{\Sigma}|^{1/2}} \exp\left(-\frac{1}{2} (\mathbf{x} - \boldsymbol{\mu})^T \boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu})\right)$$

To generate samples from this distribution, we can use the Cholesky decomposition. The Cholesky decomposition expresses the positive definite matrix Σ as the product of a lower triangular matrix L and its transpose:

$$\Sigma = \mathbf{L}\mathbf{L}^T$$

The algorithm for generating samples is as follows:

- 1. Ensure that Σ is positive definite.
- 2. Compute the Cholesky decomposition: $\mathbf{L} = \text{cholesky}(\Sigma)$.
- 3. Generate standard normal samples $\mathbf{z} \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$.
- 4. Transform to multivariate normal samples: $\mathbf{X} = \boldsymbol{\mu} + \mathbf{z} \mathbf{L}^T$.