Advanced Techniques for Mitigating Abnormal Instances and Class Imbalance in High-Dimensional Data Classification

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Machine Learning for Complex Data

- The rapid advancement of science and technology has resulted in increasingly complex datasets
- Predictive Modeling
- Make data-driven decisions
- Challenges in Predictive Modeling
 - Class Imbalance Issue
 - Abnormal Instances
 - Curse of Dimensionality
 - Categorical Features

RQ 1

Class Imbalance Issue

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Class Imbalance Issue

- Occurs when the number of instances in different classes is significantly disproportionate.
- Issue: Leads to biased models and decreases predictive accuracy.
- Abnormal Instances



Synthetic Minority Oversampling Technique (SMOTE)

- Resampling
- Balancing the Dataset:
 - Create new samples for the minority class.
- Technique:
 - Interpolate between randomly chosen minority class samples and their nearest neighbors.

•
$$p_{new} = p_0 + \alpha (p_3 - p_0)$$



Figure: SMOTE data generation

Limitation with SMOTE

• Challenged by outliers within the minority class.



Figure: Original Data



Figure: Re-sampled data with SMOTE

Proposed Solution [7]

- Technique:
 - Use a weighted average of neighbouring instances.
 - $p_{new} = \frac{\sum_{j=1}^{k} (w_j \times p_j)}{\sum_{j=1}^{k} w_j}, j = 1, \dots, k$
 - Improve robustness against outliers and noisy data.
 - Learn from a more extensive set of nearest neighbours.
- Challenge:
 - Selecting suitable weights to enhance resilience to outliers and noisy data.



Figure: Proposed method data generation

How to Define Weights?

- Distance-based approach: Higher weights for closer instances in feature space.
- Use inverse distance to the median centroid of the minority class.
- Developing new SMOTE extensions:
 - Distance extSMOTE
 - ② Dirichlet extSMOTE [1]
 - **③** FCRP SMOTE SMOTE with Chinese Restaurant Process Idea
 - BGMM SMOTE SMOTE with Bayesian Gaussian Mixture Model

References

Handling imbalanced data with abnormal minority instances

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Enhancing SMOTE for imbalanced data with abnormal minority instances



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1. Distance extSMOTE

• $d_j \in \mathbb{R}$ is the Euclidean distance between the median centroid of the minority class and the nearest neighbours

•
$$w_j = d_{j,norm}^{-1} = Normalized inverse distance$$

Algorithm Distance ExtSMOTE

Require: $X \in \mathbb{R}^{n \times p}$ the features, $Y \in \{0, 1\}^n$ the binary class label outputs.

Require: $k \in \mathbb{N}$ the number of neighbors to select for the k-Nearest Neighbors.

Ensure: Generated data $X_{new} \in \mathbb{R}^{q \times p}$ and $Y_{new} \in \{0, 1\}^q$ with q points created.

- 1: Denote by S_1 the number of points labelled as the minority class and S_0 the number of points labelled as the majority class.
- 2: Initialize Xnew and Ynew as empty vectors.
- 3: Obtain the median centroid (μ) of the minority class.
- 4: while $S_1 < S_0$ do
- 5: Filter $\mathcal{D} = \{X_i | Y_i = 1\}$, the set of points labeled as minority class 1.
- 6: Randomly choose $r \in D$ and find the indices of its k nearest neighbors, r_1, \ldots, r_k .
- 7: Consider the inverse distances, from μ , to each nearest neighbour as weights, $w_i = d_i^{-1}$

8:
$$x^{new} \leftarrow \frac{\sum (w_j \times x_{r_j})}{\sum w_j}$$
 for all j from 1 to k .
9: $y^{new} \leftarrow 1$
10: $S_1 = S_1 + 1$
11: Append x^{new} to X_{new} , append y^{new} to Y_{new}
12: end while
13: return X_{new} , Y_{new}
14: X_{new} , Y_{new}
15: X_{new} , Y_{new}
16: X_{new} , Y_{new}
17: X_{new} , Y_{new}
18: X_{new} , Y_{new}
19: X_{new} , Y_{new}
10: X_{new} , Y_{new}
11: X_{new} , Y_{new}
12: Y_{new} , Y_{new}
13: Y_{new} , Y_{new} , Y_{new}
14: Y_{new} , Y_{new} , Y_{new} , Y_{new}
15: Y_{new} , Y

1. Distance extSMOTE



(a) This scenario occurs when an outlier is chosen as a neighbouring point.

(b) The values within parentheses indicate (d_j, w_j) .

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Figure: An example of creating a sample - Distance extSMOTE

2. Dirichlet extSMOTE

- The Dirichlet distribution is defined by a set of parameters
 *α*₁, *α*₂,..., *α*_K, where K is the dimensionality of the probability
 simplex.
- The distribution is parameterized by α = [α₁, α₂,..., α_K], which can be considered pseudo-counts or prior observations.
- Let $p = [p_1, p_2, \dots, p_k]$ be a *K*-dimensional vector such that for all $j = 1, \dots, k$ we have $p_j \ge 0, j = 1, 2, \dots, k$ and $\sum_{j=1}^{K} p_j = 1$.
- The pdf of the Dirichlet distribution for a point *p* on the simplex [2]:

$$w_j = P(p|\alpha) \sim Dir(\alpha_1, \alpha_2, \dots, \alpha_K) \stackrel{\text{def}}{=} \frac{\Gamma(\sum_j \alpha_j)}{\prod_j \Gamma(\alpha_j)} \prod_{j=1}^K p_j^{\alpha_j - 1}$$
(1)

Dirichlet extSMOTE

Algorithm Dirichlet ExtSMOTE

Require: $X \in \mathbb{R}^{n \times p}$ the features, $Y \in \{0, 1\}^n$ the binary class label outputs, $k \in \mathbb{N}$ the number of neighbors to select for the k-Nearest Neighbors, $m > 0 \in \mathbb{R}$, the multiplier of the parameter of the distribution.

Ensure: Generated data $X_{new} \in \mathbb{R}^{q \times p}$ and $Y_{new} \in \{0, 1\}^q$ with q points created.

- 1: Denote by S_1 the number of points labelled as the minority class and S_0 the number of points labelled as the majority class.
- 2: Initialize X_{new} and Y_{new} as empty vectors.
- 3: Obtain the median centroid (μ) of the minority cluster.
- 4: while $S_1 < S_0$ do
- 5: Filter $\mathcal{D} = \{X_i | Y_i = 1\}$, the set of points labeled as minority class 1.
- 6: Randomly choose $r \in D$ and find the indices of its k nearest neighbors, r_1, \ldots, r_k .
- 7: if Type is 'Inverse distance (D)' then

8: Calculate the distances, $D = [d_1, \ldots, d_k]$ from μ to each nearest neighbour and obtain the reciprocal of each distance $D^{-1} = [\frac{1}{d_1}, \ldots, \frac{1}{d_k}]$. Then $\alpha = D^{-1} \times m$

9: else if Type is 'Uniform Vector (UV)' then 10: Generate a vector $\alpha = 1$, $\chi = m$ when

0: Generate a vector
$$\alpha = \mathbf{1_k} \times m$$
, where $\mathbf{1_k} = [1, \dots, 1]$

11: else if Type is 'Uniform Distribution (UD)' then

12: Generate vector U of size k from uniform(0, 1) distribution, then $\alpha = U \times m$.

- 13: end if
- 14: Use α as parameters to the Dirichlet Distribution and generate random weights $w_i \sim Dir(\alpha)$

15:
$$x^{new} \leftarrow \sum w_j x_{r_j}$$
 for all j from 1 to k , as $\sum w_j = 1$

16:
$$y^{new} \leftarrow 1, S_1 = S_1 + 1$$

17: Append
$$x^{new}$$
 to X_{new} , append y^{new} to Y_{new}

18: end while

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2. Dirichlet extSMOTE (Inverse Distance)



(a) This scenario occurs when an outlier is chosen as a neighbouring point.

Synthetic point generation Dirichlet SMOTE - Inverse Distance Nearest Neighbours 1,0.46) (1.83,0.42) 2.65. 0.421 New Data Point 0.5 Random Data Point 0.0 -0.5 -1.0 -1.5-----(4:90% 0.13) -2.0 -2.5(2.3)2.25 2.50 3.75 4.00 4.25 3.00 3.50 X-axis

(b) The values within parentheses indicate (d_j, w_j) .

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Figure: An example of creating a sample - Dirichlet extSMOTE

Synthetic Data Generation

•
$$X_{minority-outliers} \sim \mathcal{N}_{2\times 2}(\mu_{2\times 1}^{(1)}, \Sigma_{2\times 2}^{(1)})$$

• $X_{majority} \sim \mathcal{N}_{2\times 2}(\mu_{2\times 1}^{(2)}, \Sigma_{2\times 2}^{(2)})$
• $X_{outliers} \sim \text{Uniform}([-10, 10]^2)$

$$\mu_{2\times1}^{(1)} = \begin{bmatrix} 0\\0 \end{bmatrix}_{2\times1}, \Sigma_{2\times2}^{(1)} = \begin{bmatrix} 2 & 0\\0 & 2 \end{bmatrix}_{2\times2}$$
$$\mu_{2\times1}^{(2)} = \begin{bmatrix} 3\\4 \end{bmatrix}_{2\times1}, \Sigma_{2\times2}^{(2)} = \begin{bmatrix} 2 & 0\\0 & 2 \end{bmatrix}_{2\times2}$$

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Synthetic Data Generation



Figure: Comparison of resampled data

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Synthetic Data Generation



Figure: F1 Scores for 100 simulated datasets with 5-fold cross validation

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Application Data

Table: Characteristics of the binary class datasets used in the computational study.

No	Dataset	Instances	Features	Minority class	Majority class	%Minority	%Majority	IR	Presence of LOF Outliers
1	yeast6	1484	8	EXC	Remaining classes	2.36	97.64	41.40	Yes
2	yeast5	1484	8	EXC, ERL	Remaining classes	2.70	97.30	36.10	Yes
3	yeast-1289vs7	947	8	VAC	NUC, CYT, ERL, POX	3.17	96.83	30.57	Yes
4	yeast4	1484	8	ME2	Remaining classes	3.44	96.56	28.10	Yes
5	yeast-2vs8	483	8	POX	CYT	4.14	95.86	23.15	Yes
6	glass12357vs6	214	9	6	Remaining classes	4.21	95.79	22.78	Yes
7	yeast-1458vs7	693	8	VAC	NUC, ME3, ME2, POX	4.33	95.67	22.10	Yes
8	oil	937	49	minority	majority	4.38	95.62	21.85	No
9	abalone9_18	731	7	9, 18	Remaining classes	5.75	94.25	16.40	Yes
10	glass12367vs5	214	9	5	Remaining classes	6.07	93.93	15.46	Yes
11	thyroid_sick	3772	52	sick	healthy	6.12	93.88	15.33	Yes
12	yeast-1vs7	459	8	VAC	NUC	6.54	93.46	14.30	Yes
13	us_crime	1994	100	>0.65	<=0.65	7.52	92.48	12.29	Yes
14	glass12vs5	159	9	5	1, 2	8.18	91.82	11.23	Yes
15	spectrometer	531	93	>=44	<44	8.47	91.53	10.80	Yes
16	landsat_satellite	6435	36	2	Remaining classes	9.73	90.27	9.28	Yes
17	mfeatmor0	2000	6	0, 1	Remaining classes	10.00	90.00	9.00	Yes
18	yeast3	1484	8	ME3	Remaining classes	10.98	89.02	8.10	Yes
19	mfeatmor01	2000	6	0	Remaining classes	20.00	80.00	4.00	Yes
20	glass123vs567	214	9	5, 6, 7	Remaining classes	23.83	76.17	3.20	Yes
21	parkinsons	195	22	1	0	24.62	75.38	3.06	Yes
22	habermans_survival	306	3	2	1	26.47	73.53	2.78	Yes
23	glass23567vs1	214	9	1	Remaining classes	32.71	67.29	2.06	Yes
24	breast_cancer	569	30	M	В	37.26	62.74	1.68	Yes
25	banknote	1372	4	1	Remaining classes	44.46	55.54	1.25	Yes

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Application Results



Figure: F1 Score Ranks for the datasets with 100 \times 5-fold cross validation

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RQ 2

High-Dimensional Data and Presence of Categorical Data

Matharaarachchi, S., Domaratzki, M. and Muthukumarana, S. (2025). Deep-ExtSMOTE: Integrating Autoencoders for Advanced Mitigation of Class Imbalance in High-Dimensional Data Classification. The Manuscript is submitted for the Journal of Big Data Research.

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High-Dimensional Data

- Curse of Dimensionality
 - A large number of features relative to the available data, "large p, small n" problem [4].
 - Challenges:
 - Data Sparsity
 - Increased Model Complexity and Overfitting
 - Computational Challenges
- Feature Reduction
 - A critical strategy to address the challenges of high dimensionality in class imbalance [3, 5, 6].

Autoencoders

- Neural network models are designed to learn a compressed, lower-dimensional input data representation.
- Composed of two parts:
 - **Encoder:** Takes the input data and maps it to a lower-dimensional representation.
 - **Decoder:** Takes the low-dimensional representation produced by the encoder and reconstructs the original input data.



- The encoder consists of several layers that sequentially reduce the dimensionality of the input **x**⁽ⁱ⁾, where *i* denotes the *i*th observation.
- The transformation at each layer can be mathematically expressed as:

$$\mathbf{h}_{1}^{(i)} = f_{1}(\mathbf{W}_{1}^{T} \cdot \mathbf{x}^{(i)} + \mathbf{b}_{1})$$

$$\mathbf{h}_{2}^{(i)} = f_{2}(\mathbf{W}_{2}^{T} \cdot \mathbf{h}_{1}^{(i)} + \mathbf{b}_{2})$$

$$\dots$$

$$\mathbf{z}^{(i)} = f_{n}(\mathbf{W}_{n}^{T} \cdot \mathbf{h}_{n-1}^{(i)} + \mathbf{b}_{n}),$$

where $\mathbf{h}_1^{(i)}, \mathbf{h}_2^{(i)}, \ldots$ are the intermediate representations at each layer, $\mathbf{W}_1, \mathbf{W}_2, \ldots, \mathbf{W}_n$ are the weight matrices for each layer, $\mathbf{b}_1, \mathbf{b}_2, \ldots, \mathbf{b}_n$ are the biased vectors for each layer, $\mathbf{z}^{(i)}$ is the latent vector, representing the lower-dimensional feature space and f_1, f_2, \ldots, f_n represent the non-linear activation functions used in each layer.

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Decoder

- The decoder takes the latent vector $\mathbf{z}^{(i)}$ and maps it back to a reconstruction of the original input.
- The process at each layer of the decoder can be expressed as:

$$\begin{split} \mathbf{h}_{n-1}^{(i)} &= f_{n-1}^{\prime} (\mathbf{W}_{n-1}^{\prime T} \cdot \mathbf{z}^{(i)} + \mathbf{b}_{n-1}^{\prime}) \\ \mathbf{h}_{n-2}^{(i)} &= f_{n-2}^{\prime} (\mathbf{W}_{n-2}^{\prime T} \cdot \mathbf{h}_{n-1}^{\prime(i)} + \mathbf{b}_{n-2}^{\prime}) \\ & \cdots \\ \mathbf{\hat{x}}^{(i)} &= f_{1}^{\prime} (\mathbf{W}_{1}^{\prime T} \cdot \mathbf{h}_{2}^{\prime(i)} + \mathbf{b}_{1}^{\prime}), \end{split}$$

where $\hat{\mathbf{x}}^{(i)}$ is the constructed output for the i^{th} observation, which approximates $\mathbf{x}^{(i)}$. $\mathbf{h}'_{n-1}^{(i)}$, $\mathbf{h}'_{n-2}^{(i)}$, \ldots are the intermediate representations in the decoder. $\mathbf{W}'_1, \mathbf{W}'_2, \ldots, \mathbf{W}'_{n-1}$ are the weight matrices for each decoder layer, where $\mathbf{b}'_1, \mathbf{b}'_2, \ldots, \mathbf{b}'_{n-1}$ are the biased vectors for each decoder layer. $\mathbf{z}'^{(i)}$ is the latent vector, representing the lower-dimensional feature space and $f'_1, f'_2, \ldots, f'_{n-1}$ represent the non-linear activation functions used in each decoder layer.

Train Multi-layer Autoencoder

 To train the multi-layer autoencoder, we use a loss function that measures the difference between the input x⁽ⁱ⁾ and the reconstructed output x̂⁽ⁱ⁾. The most common loss function for this purpose is Mean Squared Error (MSE):

$$\mathcal{L}(\mathbf{x}, \hat{\mathbf{x}}) = \frac{1}{n} \sum_{i=1}^{n} ||\mathbf{x}^{(i)} - \hat{\mathbf{x}}^{(i)}||^2$$
(2)

5. Deep-ExtSMOTE

- Autoencoder + Dirichlet ExtSMOTE
- Step 1: Train the Autoencoder



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5. Deep-ExtSMOTE





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5. Deep-ExtSMOTE

• Step 3: Resampling and Classification



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Simulation Results



Figure: F1-Score distribution for 100 trials using simulated datasets with 1000 samples and 5000 features (2000 informative), with an imbalance ratio (IR) of 3.

Application Results

- Application 1: Isolet (Isolated Letter Speech Recognition) -Continuous Binary Classification
 - **Objective:** Accurately recognize spoken letters based on high-dimensional acoustic features extracted from voice recordings of multiple speakers.
 - The dataset includes 617 continuous features, representing processed characteristics of the audio signals.
 - Scenario 1: Original Isolet Dataset
 - Dataset includes 7797 samples, resulting in a feature-to-sample ratio of approximately 0.0791.
 - Scenario 2: Reduced Isolet Dataset
 - Selected a subset of 1000 samples from the original 7797 samples. This adjustment resulted in a feature-to-sample ratio of 0.617.

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Application Results



Figure: F1 Scores for the Isolet dataset across 50 training and test splits.



Figure: F1 Scores for the reduced Isolet dataset across 50 training and test

splits.

Application Results

- Application 2: Chile (Categorical Binary Classification)
 - **Objective:** Predict the yield of 204 chile pepper genotypes from multi-environment trials in New Mexico, USA.
 - Conduct experiment by starting with 2,500 features and increasing the number of features to 7,500.
 - Feature-to-sample ratio ranging from approximately 12.25 to 37.7.

Application Results



Figure: F1 score comparison with varying feature numbers.

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Application Results



Figure: F1 score comparison with varying imbalance ratios.

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Thank You!

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