

MULTI-INSTANCE LEARNING BASED ANOMALY DETECTION METHOD FOR SEQUENCE DATA WITH APPLICATION TO THE CREDIT CARD DELINQUENCY RISK CONTROL

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ABSTRACT

1 Anomaly detection in sequence data is widely applicable to many fields and has
2 significant commercial value to the financial industry. The focus of this paper is its
3 utility as means to control credit card delinquency risk. Transactions that deviate
4 from the typical data sequence are a common precursor of payment difficulty. Cur-
5 rent detection methods do not effectively use transaction data to detect abnormal
6 transactions. This makes it difficult to control the overdue payment risk. We pro-
7 pose a Multi-Instance Learning based anomaly detection (MILAD) method with
8 well designed learning networks to address this problem. MILAD analyze users
9 monthly transactions and payment history, and detect exceptions through well de-
10 signed deep learning networks. By comparing the performance of MILAD and
11 DAGMM, which is currently the most commonly used unsupervised deep learn-
12 ing algorithm for credit card risk control, MILAD best controls overdue risk by
13 utilizing both transaction and payment information.

14 1 INTRODUCTION

15 In recent years, research on the anomaly detection of the sequential data has gradually become
16 a hot topic. It has a very wide range of applications in many industry fields. Especially in the
17 financial field, sequence data anomaly detection has a great commercial value. Traditional sequence
18 data anomaly detection searches for the changes in several parameters of temporal data sequences,
19 such as the time series data. For example, [Gao et al. \(2019; 2020\)](#) proposed methods to detect
20 the anomaly position of the variance structure for the data sequence with smoothly changing mean
21 function. Different from the traditional ones, this research focuses on analyzing the anomaly status
22 of a multivariate time series data sequence by studying the influence of anomaly samples on the
23 abnormal state of the whole data sequence in high dimensional space. The motivation of this paper
24 is to the common credit default problem in the financial field. Effectively controlling the overdue
25 risk of credit cards is a key issue. However, there is no effective algorithm which can effectively
26 analyze the overdue risk by utilizing transaction samples in credit card transaction sequence so far.

27 For these overdue credit card users, most of their transactions are normal, and only a few transactions
28 are abnormal, such as impulse purchase, fraudulent purchase, etc. These abnormal transactions
29 are the main reasons to cause the overdue problem. However, current credit card overdue risk
30 control approaches ([Lucas & Jurgovsky, 2020](#); [Chen & Guestrin, 2016](#); [Liu et al., 2019](#); [Bolton
31 et al., 2001](#)) having little power to utilize transaction information, and relying too much on business
32 experience when conducting risk control, and being relatively cumbersome to use the model in
33 practice, etc. A big challenge of utilizing these abnormal transactions is that there are no obvious
34 post event features for assigning the abnormality labels to these anomaly transactions. The only label
35 information we could use is the users monthly overdue information. [Zong et al. \(2018\)](#) proposed the
36 DAGMM algorithm combining traditional unsupervised methods and deep auto encoders together,
37 and achieved some good results. However, in practice, sample features are constructed artificially,
38 which makes the representation of samples is not comprehensive enough. Therefore, the difference
39 between abnormal samples and normal samples is limited, and the model can not distinguish them
40 very well. Further more, this unsupervised algorithm cannot use the overdue information effectively.

41 At present, this method only has few applications in the cold-start businesses because of the absence
42 of abnormal labels.

43 The characteristic of the credit card bill overdue risk detection is that the monthly bill has a label, but
44 transactions on the bill do not have labels, which also happens in other application scenarios. The
45 Multiple Instance Learning is a good solution to solve this kind of problem. There has been a lot
46 of work done in this field, such as [Carbonneau et al. \(2018\)](#). Under the Multiple Instance Learning
47 framework, samples are grouped into sets, which are defined as Bags. An abnormal status label is
48 assigned to the entire bag. But no label is assigned to the samples in the bag. Then the relationship
49 between the bag label and sample labels is determined based on the assumption of the Multiple
50 Instance Learning. [Ilse et al. \(2018\)](#) proposed an Attention based the Multiple Instance Learning
51 algorithm (ABMIL). ABMIL uses the Attention Neural Network to learn the attention weights of
52 samples in a bag. Then the attention weights are used to aggregate samples in the bag, followed by
53 the subsequent classification analysis. This aggregation method can assign weights to the samples
54 in a bag, and then detect important samples based on sample weights. Inspired by their method,
55 if we can utilize both individual transaction information and the overall bill overdue information
56 simultaneously, we can improve the existing methods.

57 In this article, we propose a new anomaly detection algorithm based on the Multiple Instance Learning
58 technique, named as Multiple Instance Learning for anomaly detection (MILAD). MILAD is a
59 sequence sample information based anomaly detection method, which can make full use of sample
60 information and sequence information. In the experiments studied in this paper, MILAD can control
61 the overdue risk from the transaction perspective, which can provide more accurate and effective
62 results. MILAD outperforms the most commonly used algorithms in terms of several major model
63 evaluation criteria and provides a better performance in model interpretation.

64 The rest of the paper is organized as follows. The model and its proposed algorithm MILAD are in-
65 troduced in Section 2, with the computation details of each module and their parameter optimization
66 techniques. Section 3 is the experiment data analysis. In this section we conduct several experiments
67 on the application data set and compare the results against those based on the DAGMM algorithm,
68 which is the most commonly used method in the financial field. Section 4 is the summary of the
69 paper.

70 2 METHODOLOGY

71 2.1 MODEL AND NOTATION

72 Suppose $X = \{\mathbf{x}_1, \dots, \mathbf{x}_j, \dots, \mathbf{x}_J\}$ is a time dependent multivariate sequence, where $\mathbf{x}_j \in \mathbb{R}^d$ is a
73 sample of the sequence at time $j = 1, \dots, J$. $y_j \in \{0, 1\}$ is the hidden label indicating the status of
74 the sample \mathbf{x}_j , and 1 means abnormal. y_j is the hidden state of the sample has to be predicted from
75 the following model,

$$y_j = \begin{cases} 1 & \text{if } f(x_j) \geq \delta \\ 0 & \text{else} \end{cases}, \quad j = 1, \dots, J, \quad (1)$$

76 where $f : \mathbb{R}^d \rightarrow \mathbb{N}$ is a classifier based on feature mapping. We need to estimate the hidden state
77 y_j of each sample. δ is the threshold parameter discriminating the abnormal status of samples. An
78 appropriate δ should be chosen according to the practical situation. Then the abnormal state label Y
79 of the sequence X is modeled as

$$Y = \begin{cases} 1 & \text{if } \mathcal{F}(y_1, \dots, y_j, \dots, y_J) \geq \Delta \\ 0 & \text{else} \end{cases}, \quad (2)$$

80 where $\mathcal{F} : \mathbb{R}^J \rightarrow \mathbb{R}$ is a function used to estimate the overall anomaly state of a data sequence. Δ is
81 the threshold to discriminate the overall anomaly state of the data sequence.

82 Figure 1 is the flowchart of our entire modeling framework. The framework is composed of three
83 parts. The first part is the Multiple Instance Learning based on the sample information and the

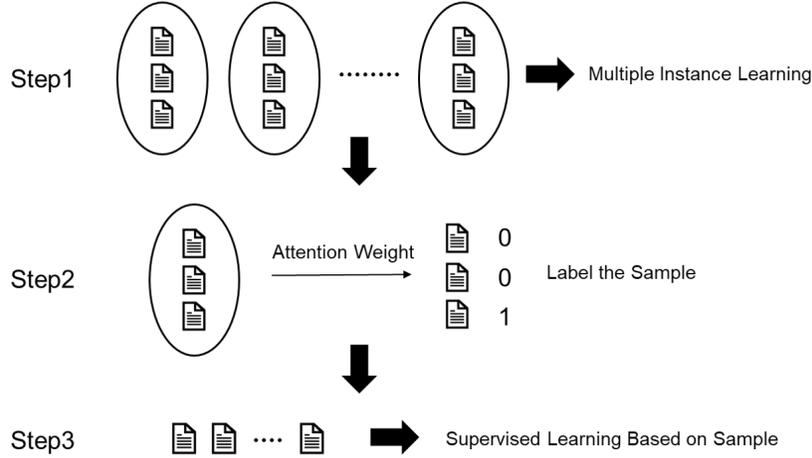


Figure 1: The framework of the MILAD algorithm

84 sequence information using the Attention mechanism. The second part is the anomaly label estimation
 85 of all samples in the data sequence according to the result from the previous Multiple Instance
 86 Learning procedure. The third part is the sequence anomaly detection procedure based on the esti-
 87 mated abnormal labels of samples using binary supervised learning method. Models are trained by
 88 the common optimization algorithm Adam (Kingma & Ba, 2014).

89 Algorithm 1 is the computational flow of our proposed sequence anomaly detection method MILAD.
 90 It is a Multiple Instance Learning based method, which can effectively associates the unknown
 91 sample label y_j with the known sequence label Y through the Attention network mechanism and
 92 achieve efficient modeling processes eventually. The MILAD algorithm constructs a risk analysis
 93 model \mathcal{F} based on sample anomaly detection in a data sequence. In practice, taking the credit card
 94 overdue risk prediction businesses as an example, we can use the model \mathcal{F} to evaluate card holders'
 95 overdue risk based on their transaction vector $\mathbf{x}' \in \mathbb{R}^d$. We can predict the overdue risk probability
 96 p' through the model \mathcal{F} , and finally determine whether to intercept the transaction \mathbf{x}' based on the
 97 actual needs of the business. In this way we can directly control the overdue risk in the transaction
 98 dimension. Comparing with traditional approach, controlling overdue risk based on the MILAD
 99 algorithm is much more convenient in practice.

Algorithm 1: MILAD

Input: The Multi time series sample bag $X = \{\mathbf{x}_1, \dots, \mathbf{x}_j, \dots, \mathbf{x}_J\}$ which is a collection of time series of length J , $\mathbf{x}_j \in \mathbb{R}^d$

Step 1 Multiple Instance Learning: Use the Algorithm 2 to estimate the classification probability $P = \text{Sigmoid}(W_C^\top Z + b)$ of the bag, the attention weight $w_j^* = \mathbf{w}_j / \sum_{m=1}^J \mathbf{w}_m$ of samples in the bag, and the abnormal probability $p_j = Pw_j^*$ of samples in the bag.

Step 2 Sample Anomaly Detection: Based on the Multiple Instance Learning results from Step 1, detect the abnormal state of each sample \mathbf{x}_j in the bag, and get the sample anomaly state set $S = \{\hat{y}_1, \hat{y}_2, \dots, \hat{y}_J\}$.

Step 3 Sequence Anomaly Detection: Based on the abnormal sample detection results from the Step 2, use the classification (e.g., Xgboost) method to estimate The abnormal state Y of the sample bag X .

Output: Y the abnormal state of the sample bag, and S the hidden abnormal state set.

100 2.2 MULTIPLE INSTANCE LEARNING

101 In model equation 1, the classification model f is built upon the feature information of sequence
 102 samples in the bag. However, the anomaly state label y_j of the sequence samples is generally
 103 unknown. Therefore, we cannot perform any supervised learnings directly. To effectively solve
 104 this problem, we use the Multiple Instance Learning approach. The Multiple Instance Learning
 105 model is composed of the following four parts: the transformer network T , attention network W ,
 106 aggregation network A , and classification network C . The transformer network T is designed to
 107 conduct a feature extraction and transformation on the original features. As mentioned in [Foulds
 108 & Frank \(2010\)](#), there are two different assumptions: the Standard Assumption and the Collective
 109 Assumption. The Standard Assumption is that each sample in the bag has its own label, the label
 110 of the bag is negative if all samples in the bag are negative, and the label of the bag is positive
 111 if there is at least one positive sample in the bag. The Collective Assumption states that the label
 112 of a bag cannot be determined by any single sample, but by the interactions between samples and
 113 the cumulative effect of some samples in the bag. Therefore, we propose two types of designs
 114 for the network T : the **Basic method** and the **Self-Attention based method**. The Basic method
 115 is adaptive to the standard assumption, while the Self-Attention based method is designed for the
 116 collective assumption, which has more practical usages. The attention network W is used to learn
 117 attention weights of samples in the bag, and the attention weights are estimated through a module
 118 conducted by a two-layer gated neural network. The aggregation network A is used to aggregate
 119 all samples in the bag. After calculating the attention weight of each sample through the attention
 120 network W , we can estimate the feature vector Z_i of the bag by calculating the weighted sum of the
 121 sample vectors in the bag. The classification network C is a network that classifies the bag vector.
 122 After the aggregation of samples in the bag, the classification problem is turned into a traditional
 123 binary supervised learning problem. To deal with features extracted from the neural network, a fully
 124 connection (FC) layer network together with the sigmoid activation function is used to calculate the
 125 anomaly classification probability P_i of the i^{th} bag. The details of the proposed multiple instance
 126 learning networks are listed in [Appendix A](#).

Algorithm 2: The Multiple Instance Learning Algorithm

Input: The multi time series sample bag $X = \{\mathbf{x}_1, \mathbf{x}_2 \dots, \mathbf{x}_J\}$

Step 1 Randomly initialize the weights of the parameters in the T, W, A, C network;

Step 2 Transform the original sample through the network T and obtain the transformed
 vector $\mathbf{h}_j = T(\mathbf{x}_j)$ through the Basic method or the Self-Attention method ([Algorithm 3](#)
 in [Appendix A](#)).

Step 3 Calculate the attention weight w_j^* for samples in the bag through the network W .

Step 4 Aggregate the samples through the network A : $Z = \sum_{j=1}^J w_j^* \mathbf{h}_j$.

Step 5 Obtain the abnormal probability of the bag through the network C :
 $P = \text{Sigmoid}(W_C^\top Z + b)$.

Output: The classification probability P of the sample bag

127 [Algorithm 2](#) is our designed Multiple Instance Learning Algorithm. We use the Attention mecha-
 128 nism to adaptively aggregate the samples in the bag Y_i . Since the feature extractor and classifier
 129 are conducted with neural networks, it allows to establish an end-to-end model to make the whole
 130 model to be more auto adaptive. Meanwhile, each step of the model is built upon neural networks,
 131 which makes the back propagation algorithm available for parameters optimization. All parameters
 132 are optimized by minimizing the Logarithmic loss

$$L(\Theta) = \frac{1}{N} \sum_{i=1}^N (Y_i \ln P_i + (1 - Y_i) \ln(1 - P_i)),$$

133 where N is the sample size of the training data, P_i is the anomaly probability of the i^{th} bag.

134 2.3 SAMPLE ANOMALY DETECTION

135 After the Attention based Multiple Instance Learning, we obtain the probability P_i of the label of
 136 the bag to be 1, and the attention weight w_{ij}^* of each sample in the bag. Unlike the traditional
 137 Multiple Instance Learning, the estimated attention weights of the samples are more important here,
 138 which can be used to detect the key samples in the bag. That is, which sample in the bag has a
 139 significant impact on the abnormal status of the bag. The samples with larger attention weights have
 140 a greater impact on bags, and these samples are likely to be the key samples which lead to the bag
 141 abnormality. Therefore, we can combine the prediction results of the bag and the estimated attention
 142 weights together to predict the anomaly status of each samples in the bag. Let

$$p_j = P_i w_{ij}^*$$

143 be the probability of sample \mathbf{x}_j being abnormal. The probability is used to rank samples in the
 144 bag, rather than discriminating samples with respect to the sample abnormality. By choosing a
 145 appropriate threshold δ , the abnormal state of the sample is $y_{ij} = 1$ if $p_{ij} \geq \delta$.

146 2.4 SEQUENCE ANOMALY DETECTION

147 After using the Multiple Instance Learning for abnormal sample detection, we get the anomaly set
 148 $S = \{\hat{y}_1, \dots, \hat{y}_J\}$, which contains the pseudo labels of all samples in the bag, and then we can do the
 149 binary supervised learning to estimate the sequence abnormal state Y in model equation 2 using the
 150 classification approaches. In this paper, we adopt the Xgboost algorithm. However in practice, there
 151 are only a few sequences or sample bags which are abnormal. Therefore, the binary classification
 152 problem we are dealing with is a highly imbalanced data analysis problem. We should adopt the
 153 imbalance data analysis techniques. To evaluate the model performance for the imbalanced data,
 154 AUC will be a good choice.

155 3 EMPIRICAL ANALYSIS

156 3.1 DATA PREPROCESSING

157 Since payment data often contains sensitive private information about individuals or institutions,
 158 and only banks and other related institutions have access to it. Therefore the acquisition of such
 159 public data set is quite limited. The lack of available effective public datasets is also an challenge for
 160 researches in this field. In this work, we evaluate the performance of the proposed MILAD algorithm
 161 on a commonly used real data set, which is the Credit Card Fraud Detection (CCFD) (Dal Pozzolo
 162 et al., 2015) data. The CCFD data composed of transactions of credit card users in Europe on
 163 September 2013. This dataset includes 284,807 transactions, where 492 are abnormal transactions.
 164 It is a highly imbalanced dataset, which only has 0.17% of abnormal transactions. To deal with this
 165 highly imbalance problem of the data, we use the common undersampling method to sample 10%
 166 of normal transactions. Then the abnormal rate increases to 1.70%. Due to the privacy issues in this
 167 field, this dataset cannot provide the original transaction features and the user information. The data
 168 contains 28 principal component features, $\{V_1 \dots V_{28}\}$, which are transformed from the original
 169 features, the transaction amount, and the anomaly label of each transaction.

170 In order to make the dataset suitable for solving our problem, we have to generate new dataset
 171 through the following data generating mechanism based on the original CCFD data. We randomly
 172 select a certain number of transactions from the original data set to form a sample bag, then take
 173 each sample bag as the user’s transaction set X_i , and then label the bag according to the sample label
 174 in the bag. The labeling process of the sample bag is based on these two assumptions of the Multiple
 175 Instance Learning, which are the standard assumption and collective assumption. In the subsequent
 176 data analysis we assume that there are no available labels for the samples in the bag. Generating
 177 the dataset in this way can effectively mimic our desired scenario in which we have the label for
 178 user’s transaction set, but lack of the labels for each transactions in the bag. The labeling rules
 179 for the sample bags are as follows. Under the Standard Assumption, as long as there is a sample
 180 $\mathbf{x}_{ij} \in \mathbb{R}^d$ whose label y_{ij} is abnormal in the sequence set $X_i = \{\mathbf{x}_{i1}, \dots, \mathbf{x}_{ij}, \dots, \mathbf{x}_{iJ}\}$, the label
 181 of the overall sequence $Y_i = \min\{1, \sum_{j=1}^J y_{ij}\}$. Under the Collective Assumption, only when the

182 sum of the amount of abnormal samples reaches a certain threshold, the label of the overall sequence
 183 $Y_i = 1$, if $\frac{\sum_{j=1}^J y_{ij} \nu_{ij}}{\sum_{j=1}^J \nu_{ij}} \geq \Delta$, where ν_{ij} is the transaction amount of the sample $\mathbf{x}_{ij} \in \mathbb{R}^d$, that is
 184 $\nu_j \in \{x_{j1}, \dots, x_{jd}\}$. The parameter $\Delta \in (0, 1)$ is determined according to the specific application
 185 scenario.

186 For the convenience of the experiment, we assume the sample size in each bag is the same when
 187 generating the data set. Under the standard assumption, since we use the probability, which reflects
 188 the anomaly status of the sample, to rank the samples in the bag, rather than discriminating the
 189 samples. In the subsequent discrimination analysis, the threshold δ in model equation 1 is chosen to
 190 be the one which makes the highest $F1$ score in the training set. Under the collective assumption,
 191 we need to consider the proportion of the abnormal transaction amount among all transactions in the
 192 bag. When the proportion of abnormal transaction amount reaches the threshold $\Delta = 0.1$, we will
 193 consider the user’s transaction bag to be overdue. Under each assumption, we have $N_1 = 200$ bags
 194 for training, $N_2 = 50$ bags for test. The size of the bag is $J = 10$. The anomaly rate of bags is
 195 16.6% for the standard assumption and 5.8% for the collective assumption.

196 We show the performance of the proposed method on both the standard assumption and collective
 197 assumption. We first evaluate the sample anomaly detection performance, and then analyze the
 198 performance of the sequence anomaly detection. In the sample anomaly detection part, we compare
 199 MILAD with the most commonly used unsupervised anomaly detection algorithm DAGMM in the
 200 financial field in terms of common model evaluation criteria (Precision, Recall, F1 score, AUC),
 201 as well as the interpretability of these two methods. In the Sequence Anomaly Detection part, we
 202 first built an idealized model, which is a model constructed based on the ideal assumption that the
 203 hidden labels are all available, hereafter denoted by the **Ideal** model. We use the Ideal model as the
 204 benchmark since it always has the best performance among all possible methods. We use AUC to
 205 evaluate model performances.

206 The computational resources of our experiments are *Windows 10, Intel(R) Core(TM) i5-9300H,*
 207 *GeForce GTX 1650 GPU, 16GB Ram.* We use *Python 3.8* under *Tensorflow 2.5.0* environment.

208 3.1.1 SAMPLE ANOMALY DETECTION UNDER THE STANDARD AND COLLECTIVE 209 ASSUMPTIONS

210 Table 1 is the network structures of the experiments. For DAGMM, we use the same network
 211 structure under both assumptions, and we also use the same hyperparameter settings in [Zong et al.](#)
 212 (2018) ($\lambda_1 = 0.1, \lambda_2 = 0.005$). For MILAD, the Basic method is adopted under the standard
 213 assumption, and the Self-Attention method is adopted under the collective assumption. $FC(a, b, c)$
 214 is a full connection network, where a and b are the number of input and output neurons, c is the
 215 activation function.

Table 1: Network structures under the standard and collective assumptions

Method	Layer	Structure
DAGMM	Encoder	FC(28, 16, tanh) \rightarrow FC(16, 4, tanh) \rightarrow FC(4, 1, none)
	Decoder	FC(1, 4, tanh) \rightarrow FC(4, 16, tanh) \rightarrow FC(16, 28, none)
	Estimate Network	FC(3, 10, tanh) - Dropout(0.2) \rightarrow FC(10, 2, Softmax)
MILAD (Standard)	Network T	FC(28, 16, ReLU) \rightarrow FC(16, 8, ReLU)
	Network W	FC(8, 8, tanh) \odot FC(8, 8, Sigmoid) \rightarrow FC(8, 1)
	Network C	FC(8, 1, Sigmoid)
MILAD (Collective)	Network T	Refer to the structure in Algorithm 3, where $d = 8$
	Network W	FC(8, 8, tanh) \odot FC(8, 8, Sigmoid) \rightarrow FC(8, 1)
	Network C	FC(8, 1, Sigmoid)

216 Figure 2 shows the loss function curves of these two algorithms with respect to 1,000 epochs in the
 217 training processes. We can see that the DAGMM algorithm converges after 1,000 epochs. Therefore
 218 we choose the model after the 1,000 epochs of training as the final DAGMM model. For the MILAD
 219 algorithm, since we treat each bag as a sample group, the sample size is relatively small. It can be
 220 seen that the model is overfitted after 30 epochs of training under the standard assumption, and 10
 221 epochs of training under the collective assumption. Therefore, under the standard assumption, we

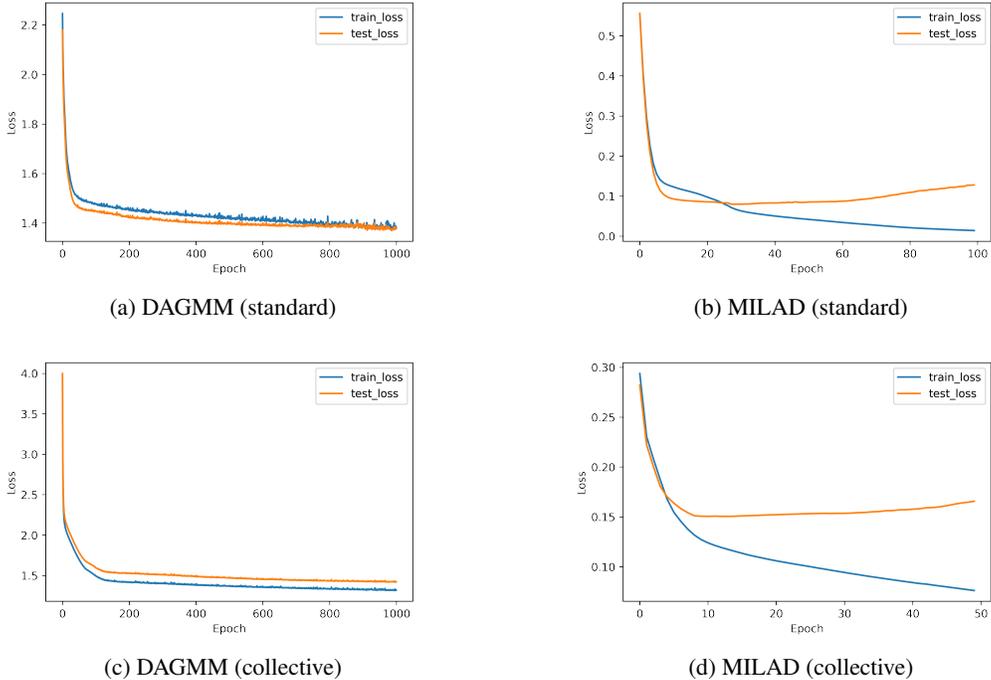


Figure 2: Loss curves under two assumptions

222 choose the model after 30 epochs of training as the final MILAD model, while under the collective
 223 assumption, we choose the model after 10 epochs of training as the final MILAD model.

224 The final output of the DAGMM model is the probability density of the sample. In order to compare
 225 it with the anomaly probability computed from the MILAD method, we use the function $f(x) =$
 226 $1 - \frac{2}{\pi} \arctan(x)$ to convert it into the probability ranged in (0,1). Figure 3 is the ROC curve of
 227 the model trained by the DAGMM algorithm and the MILAD algorithm. It can be seen that the
 228 performance of the MILAD algorithm is significantly better than that of the DAGMM algorithm
 229 under both assumptions.

230 Table 2 is the comparison matrix in several common model evaluation criteria. It can be seen that
 231 MILAD is significantly better than DAGMM in terms of these common model evaluation criteria,
 232 such as Precision, Recall, F1 score and AUC. This is because DAGMM is an unsupervised learning
 233 method, while MILAD is a supervised learning algorithm, which can effectively utilize the label
 234 information of the bag for complex data through the attention based Multiple Instance Learning ap-
 235 proach, and outperforms the unsupervised learning method. Therefore it is reasonable that MILAD
 236 achieves better performance, and is more useful in practice.

Table 2: Model comparison under two assumptions

Assumption	Type	Method	Precision	Recall	F1-score	AUC
Standard	Training	DAGMM	0.0899	0.3722	0.1449	0.8397
		MILAD	1.0000	0.8639	0.9270	0.9717
	Test	DAGMM	0.0971	0.4545	0.1600	0.8769
		MILAD	0.9302	0.9091	0.9195	0.9627
Collective	Training	DAGMM	0.0580	0.2232	0.0921	0.7725
		MILAD	0.6273	0.4000	0.4885	0.8854
	Test	DAGMM	0.0561	0.2526	0.0918	0.7856
		MILAD	0.5781	0.3895	0.4654	0.8878

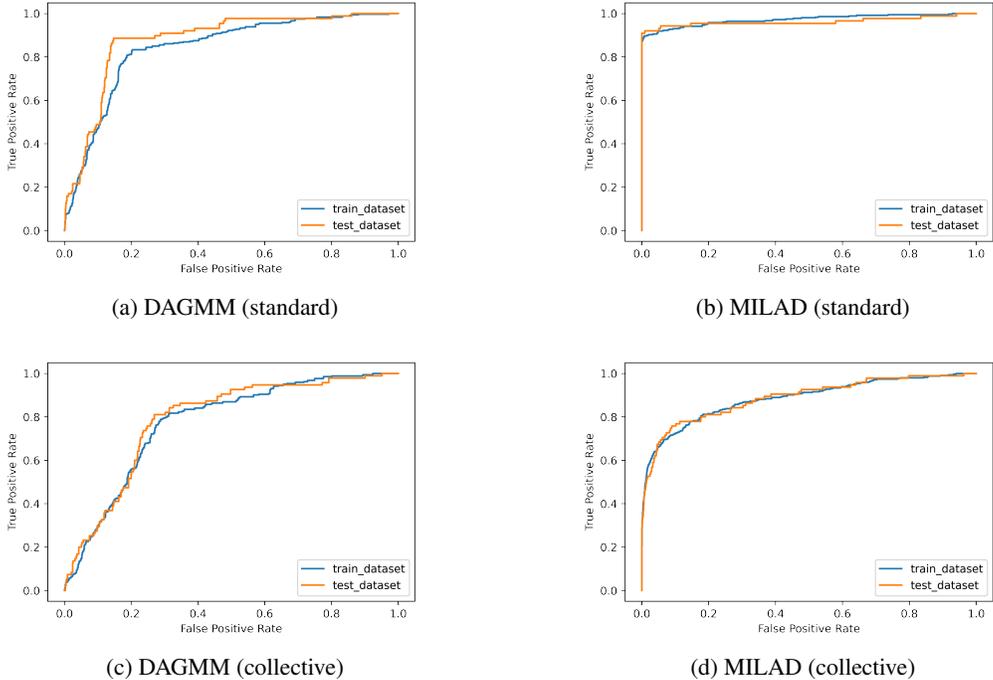


Figure 3: ROC curves under two assumptions

237 To show the interpretability of the MILAD algorithm, we also check these abnormal sample bags to
 238 see whether the method can identify the abnormal samples in the bag that cause the bag abnormality.
 239 Table 3 is the samples’ anomaly state (y_{ij}) and their attention weights (w_{ij}^*) of four ($i = 1, \dots, 4$)
 240 randomly selected anomaly sample bags under the standard and collective assumptions. The attention
 241 weights $\{0.75; 0.78; (0.30, 0.27); (0.20, 0.21)\}$ of the abnormal samples that cause the abnor-
 242 mality of the entire bag are significantly larger than other samples in the same bag. This result is
 243 consistent with our experiment setups, which fully demonstrates the outstanding interpretability of
 244 our MILAD method.

Table 3: Attention weights of two randomly selected cases under two assumptions

Assumption		x_1	x_2	x_3	x_4	x_5	x_6	x_7	x_8	x_9	x_{10}
Standard	y_{1j}	0	0	0	0	0	1	0	0	0	0
	w_{1j}^*	0.03	0.04	0.02	0.00	0.02	0.75	0.05	0.02	0.04	0.03
	y_{2j}	0	0	0	1	0	0	0	0	0	0
	w_{2j}^*	0.03	0.01	0.03	0.78	0.00	0.00	0.00	0.01	0.10	0.03
Collective	y_{3j}	1	1	0	0	0	0	0	0	0	0
	w_{3j}^*	0.30	0.27	0.03	0.05	0.03	0.07	0.06	0.05	0.04	0.10
	y_{4j}	0	0	0	0	1	0	0	0	0	1
	w_{4j}^*	0.07	0.04	0.12	0.06	0.20	0.06	0.11	0.11	0.02	0.21

245 3.1.2 SEQUENCE ANOMALY DETECTION

246 For sequence anomaly detection we adopt the Xgboost algorithm, which is a commonly used binary
 247 supervised learning method in this field. All models are trained to achieve their best performances.
 248 The AUC results are shown in Table 4. It can be seen that in the absence of transaction labels, our
 249 MILAD algorithm still can achieve the similar performance as the Ideal model with respect to the
 250 AUC criterion, which is significantly better than DAGMM. We can conclude that MILAD is more
 251 feasible than DAGMM under both standard and collective assumption.

Table 4: AUC under the standard and collective assumptions

Assumption	Type	Ideal	DAGMM	MILAD
Standard	Training	1	1	1
	Test	0.98	0.87	0.98
Collective	Training	1	1	1
	Test	0.98	0.80	0.94

252

4 CONCLUSION

253 In this paper, we focus on the anomaly state evaluation of the data sequence caused by the abnormal
254 samples contained in it. We propose a anomaly detection algorithm MILAD based on the Multiple
255 Instance Learning techniques. We apply the proposed method to the delinquency risk detection in the
256 credit card industry. The empirical results demonstrate that MILAD overcomes many short-comings
257 that existing methods have through its use of the sample information and the sequence anomaly
258 information simultaneously to effectively identify abnormal samples. The proposed method can
259 help financial institutions to control the overdue risk based on transactions directly and effectively.

260 AUTHOR CONTRIBUTIONS

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300 APPENDIX

301 A MULTIPLE INSTANCE LEARNING NETWORK

302 In model equation 1, the classification model $f(\cdot)$ is built upon the feature information of sequence
 303 samples in the bag. However, the anomaly state label y_j of the sequence samples is generally
 304 unknown. Therefore, we can not perform any supervised learnings directly. To effectively solve this
 305 problem, we use the Multiple Instance Learning approach.

306 The Multiple Instance Learning model is composed of the following four parts: the Transformer
 307 Network T, Attention Network W, Aggregation Network A, and Classification Network C. Accord-
 308 ing to the structure of the Transformer Network, we have two types of designs: the Basic method
 309 and the Self-Attention based method. The Basic method is adaptive to the Standard Assumption,
 310 while the Self-Attention method is designed for the Collective Assumption, which has more practical
 311 usages. Figure 4 and Figure 5 are the network structures of the Multiple Instance Learning
 312 model corresponding to the Basic method and the Self-Attention method.

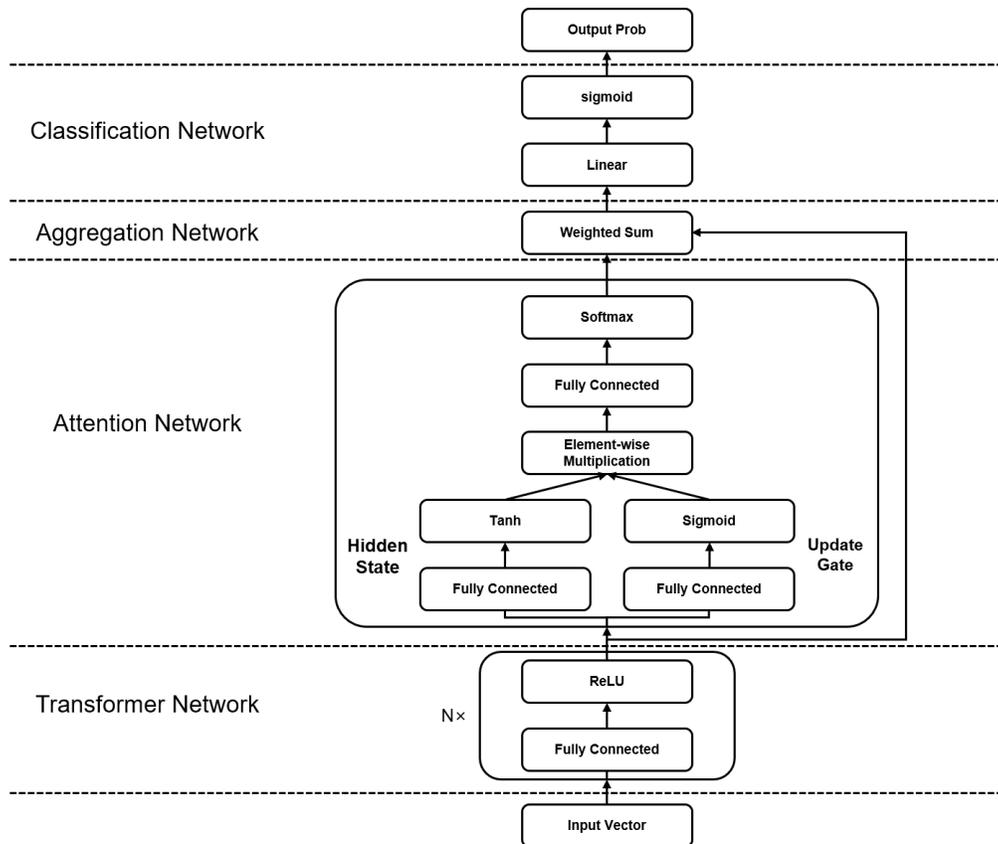


Figure 4: Multiple Instance Learning Network Structure (Basic Version)

313 The Algorithm 3 is the detailed transformation algorithm based on the Self-Attention method.

314 A.0.1 TRANSFORMER NETWORK T

315 The function of the Transformer Network T is to conduct a feature extraction and transformation on
 316 the original features. There are two approaches based on different assumptions: the Basic method
 317 and the Self-Attention method. Figure 4 and Figure 5 are the network structures of the Multiple
 318 Instance Learning model corresponding to the Basic method and the Self-Attention method.

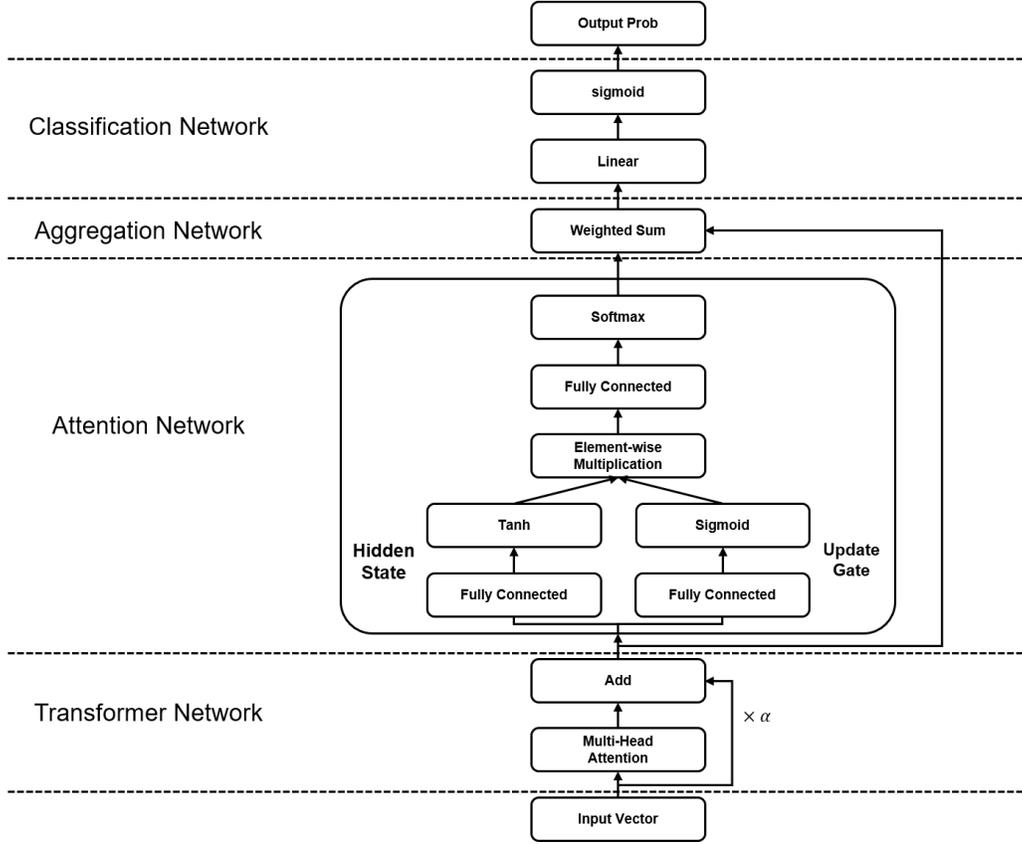


Figure 5: Multiple Instance Learning Network Structure (Self-Attention Version)

Algorithm 3: The Feature Transformation Algorithm Based on the Self-Attention Method

Input: The multi time series sample bag $X = \{\mathbf{x}_1, \dots, \mathbf{x}_J\}$

Step1 Rewrite the input sample vectors in matrix form $X = [\mathbf{x}_1, \dots, \mathbf{x}_J]^T$;

Step2 Calculate Q, K, V matrix:

$$Q = XW_Q$$

$$K = XW_K$$

$$V = XW_V$$

Step3 Calculate attention weight:

$$W_A = \text{Softmax}\left(\frac{QK^T}{\sqrt{d_1}}\right)$$

Step4 Calculate transformed matrix:

$$T = [\mathbf{t}_1, \dots, \mathbf{t}_J]^T = W_A V$$

Step5 Output the transformed feature vectors through Soft-Transformer \mathbf{h}_j :

$$\mathbf{h}_j = \mathbf{x}_j + \alpha \mathbf{t}_j, \forall j = 1 \dots J$$

Output: The transformed vectors $H = \{\mathbf{h}_1, \dots, \mathbf{h}_J\}$

319 **The Basic Method** As mentioned in Ilse et al. (2018), we use a double-layer neural network to
 320 calculate the attention weight. Let \mathbf{x}_j be the sample feature vector, then $X = \{\mathbf{x}_1, \dots, \mathbf{x}_J\}$ will be
 321 the corresponding sequence bag. The attention weight is

$$w_j^* = \frac{\exp\{W_2^\top \tanh(W_1^\top \mathbf{x}_j)\}}{\sum_{i=1}^J \exp\{W_2^\top \tanh(W_1^\top \mathbf{x}_i)\}}.$$

322 Then, the feature vector $Z \in \mathbb{R}^d$ for the bag will be estimated as the weighted average of the sample
323 vectors. That is

$$Z = \sum_{j=1}^J w_j^* \mathbf{x}_j.$$

324 The advantages of this approach are all weight parameters are able to be optimized in the training
325 process and the each sample in the bag has its own weight. Therefore, the feature vector conducted
326 in this way has more interpretability. Meanwhile, we could also present a exception sample detection
327 based on the their weights. The sample with a larger weight will have a larger chance to be a critical
328 sample in the bag.

329 For the Basic approach, since it has been assumed that there is no interaction effect and no structural
330 information between samples in the bag, sample \mathbf{x}_j can be transformed into a feature vector \mathbf{h}_j
331 directly using a two-layer fully connected network,

$$\mathbf{h}_j = W_2^\top \sigma(W_1^\top \mathbf{x}_j + \mathbf{b}_1) + \mathbf{b}_2,$$

332 where $\sigma(\cdot)$ is the activation function.

333 **The Self-attention Method** The Basic method assumes that the samples are independent when
334 calculating the Attention mechanism. However, in practice, there exists various interaction effects
335 between samples. In order to learn the interaction effectively, Vaswani et al. (2017) introduced a
336 Transformer framework based on the Attention mechanism to obtain the interaction information be-
337 tween sequences composed of words. Based on this method, Rymarczyk et al. (2021) proposed a
338 Soft-Transformer framework. The Soft-Transformer transforms samples into feature vectors first
339 before the Attention based Multiple Instance Learning, which can explore the interactive informa-
340 tion between samples more effectively. Let \mathbf{x}_j be the sample vector, and $X = [\mathbf{x}_1, \dots, \mathbf{x}_J]^\top$ be the
341 sample matrix composed of vectors. Firstly, we use weight matrix $W_Q^{d \times d_1}, W_K^{d \times d_1}, W_V^{d \times d_2}$ to calcu-
342 late the corresponding matrices $Q^{J \times d_1}$ (*Query*), $K^{J \times d_1}$ (*Key*), $V^{J \times d_2}$ (*Value*), where $Q = XW_Q$,
343 $K = XW_K$, $V = XW_V$. We usually make $d_1 = d_2$. Then we have

$$W_A = \text{Softmax}\left(\frac{QK^\top}{\sqrt{d_1}}\right).$$

344 Finally we get the transformed sample matrix:

$$T = W_A V = \text{Softmax}\left(\frac{QK^\top}{\sqrt{d_1}}\right) X W_V,$$

345 where $T = [\mathbf{t}_1, \dots, \mathbf{t}_J]^\top$ is a transformed sample matrix, and \mathbf{t}_j is the transformed sample vector.
346 Then we can perform the Multiple Instance Learning in the same way. Using this method we can
347 get the transformed sample vector and the interaction information between samples. However, after
348 transformation, the actual meaning of the vector is different from those for the original samples. The
349 subsequent sample weights do not represent the importance of the samples anymore, and cannot be
350 used to discriminate critical samples. Therefore, we use the Soft-Transformer to transform the
351 output vector \mathbf{t}_j into $\mathbf{x}_j + \alpha \mathbf{t}_j$. The Soft-Transformer we use makes the weight of the transformed
352 vector still has the ability to reflect the importance of the samples after considering the interaction
353 information in the analysis. In the subsequent analysis, we perform a critical sample discrimination
354 based on attention weights. Figure 6 is the schematic diagram of the transformation process.

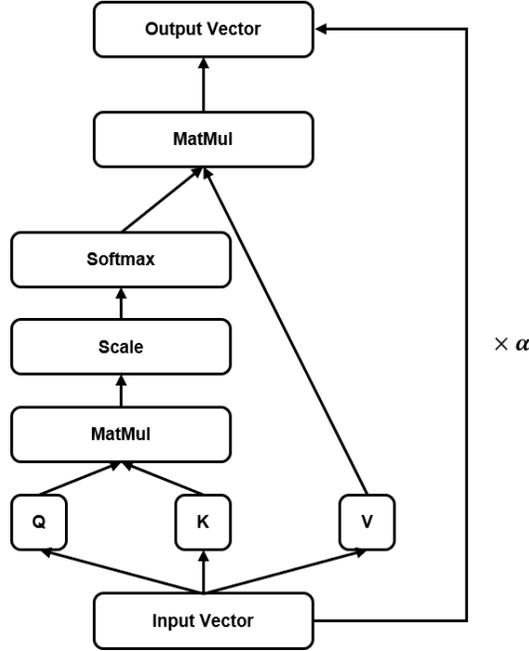


Figure 6: The structure of the Soft-Transformer

355 A.0.2 ATTENTION NETWORK W

356 The Attention Network is used to learn attention weights of samples in the bag, and the attention
 357 weights are estimated through a module conducted by a two-layer gated neural network:

$$\mathbf{v}_j = \tanh(V^\top \mathbf{h}_j),$$

$$\mathbf{u}_j = \text{Sigmoid}(U^\top \mathbf{h}_j),$$

$$w_j = W_a^\top (\mathbf{v}_j \odot \mathbf{u}_j),$$

$$w_j^* = \frac{\exp\{w_j\}}{\sum_{i=1}^J \exp\{w_i\}},$$

358 where \mathbf{v}_j is the hidden state, \mathbf{u}_j is the updated gate state, \odot represents the element-wise multi-
 359 plication of the vector, w_j is the attention weight, and w_j^* is the normalized attention weight by
 360 Softmax.

361 A.0.3 AGGREGATION NETWORK A

362 The Aggregation Network is used to aggregate all samples in the bag. After calculating the attention
 363 weight of each sample through the Attention Network W , we can estimate the feature vector Z_i of
 364 the bag by calculating the weighted sum of the sample vectors in the bag,

$$Z_i = \sum_{j=1}^J w_j^* \mathbf{h}_j.$$

365 where w_j^* is the attention weight of each sample in the bag, \mathbf{h}_j is the feature vector obtained from
 366 the Transformer Network T .

367 A.0.4 CLASSIFICATION NETWORK C

368 The Classification Network is a network that classifies the bag vector. After the aggregation of
369 samples in the bag, the classification problem is turned into a traditional binary supervised learning
370 problem. To deal with features extracted from the Neural Network, a fully connected (FC) layer
371 network together with the Sigmoid activation function is used to calculate the anomaly classification
372 probability P_i of the i^{th} bag, where

$$P_i = \text{Sigmoid}(W_C^T Z_i + b).$$