# SECOM Dataset: Anomaly Detection Model Evaluation

## **Evaluation Protocol**

10-Fold Cross Validation | Strict Train-Test Separation
No Data Leakage Confirmed | Overfitting Analysis Below | No patterns are getting detected by the models

## 1. One-Class SVM (contamination=0.18)

Performance Metrics

#### Test Set (2,503 samples):

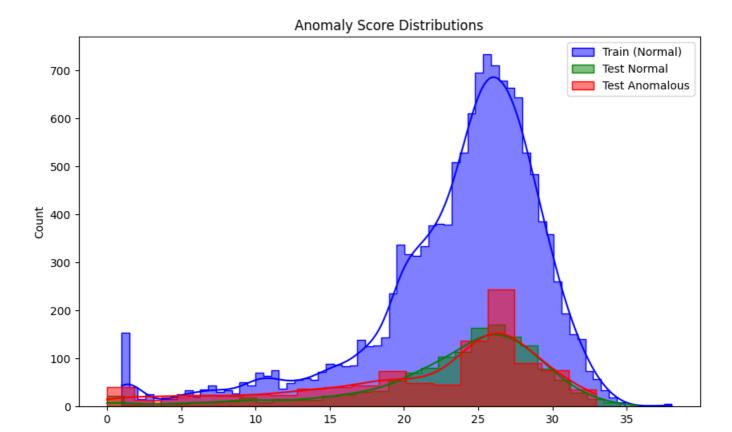
Class	Precision	Recall	F1-Score	Support
Normal (0)	0.63	0.81	0.71	1,463
Anomaly (1)	0.56	0.34	0.42	1,040

Accuracy: 61% Macro Avg F1: 57%

#### Training Set (13,167 normals):

Metric	Value
Normal Recall	82%
Training Accuracy	82%

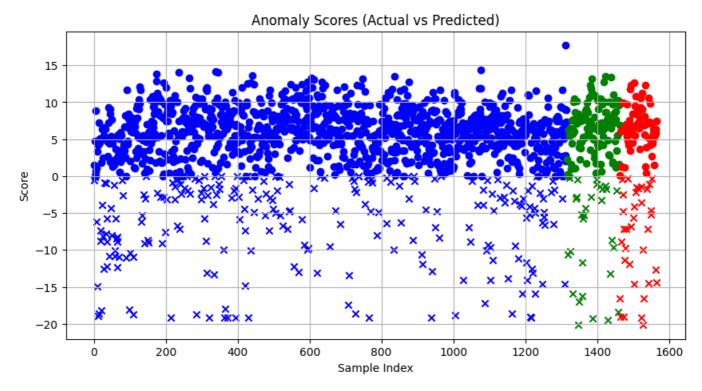
Visual Analysis



#### Color Code:

- Blue: Training normal scores
- Green: Test normal scores
- Red: Test anomaly scores

Pattern: Overlapping distributions indicate limited separability



#### Markers:

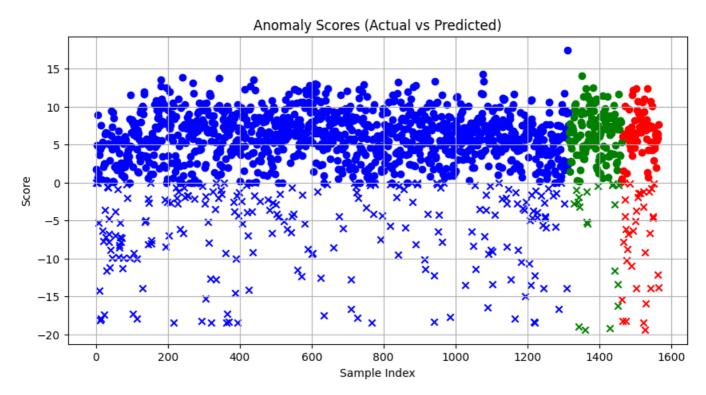
• o: Predicted normal (score > threshold)

• ×: Predicted anomaly

Color:

Green: True normal Red: True anomaly

• Blue: Training normal



Key Insight: No clear separation between classes in latent space

## 2. Isolation Forest (contamination=0.2)

#### Performance Metrics

#### Test Set:

Class	Precision	Recall	F1-Score	Support
Normal (0)	0.62	0.80	0.70	1,463
Anomaly (1)	0.54	0.32	0.40	1,040

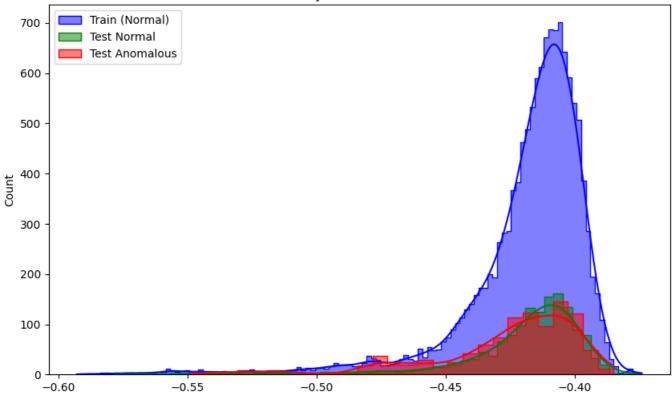
Accuracy: 60% Macro Avg F1: 55%

### Training Set:

Metric	Value	
Normal Recall	80%	
Training Accuracy	80%	

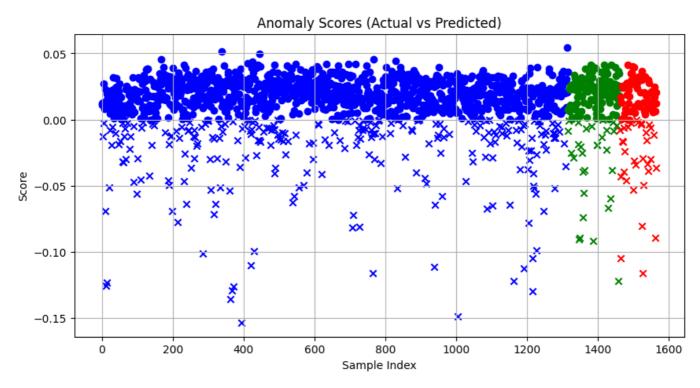
Visual Analysis

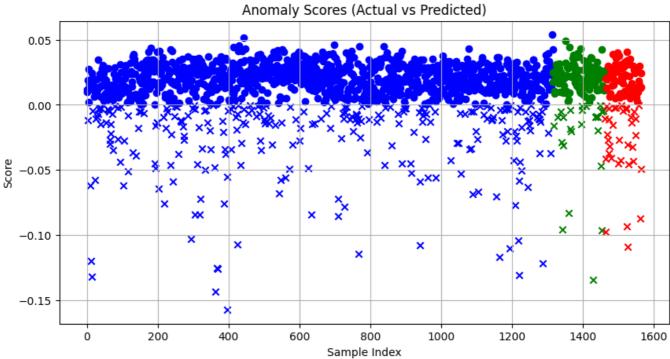
## **Anomaly Score Distributions**



## **Distribution Characteristics:**

- Score range: -0.6 to -0.3
- high overlap between test normals/anomalies





Key Limitation: No clear cluster separation despite tree ensembles

## 3. Autoencoder (contamination=0.2)

Performance Metrics

Test Set:

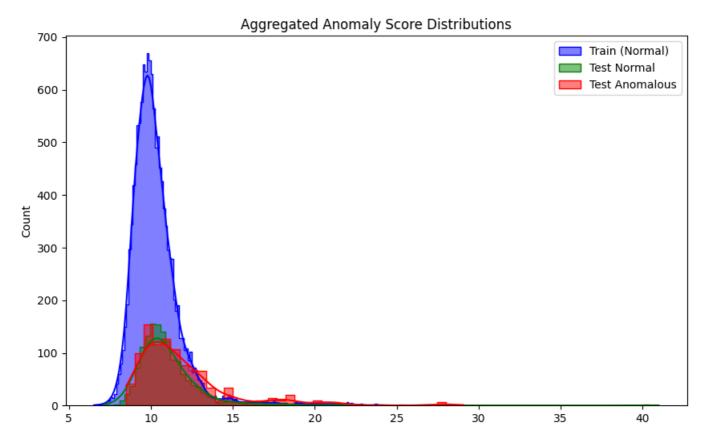
Class	Precision	Recall	F1-Score	Support
Normal (0)	0.65	0.64	0.64	1,463
Anomaly (1)	0.50	0.51	0.50	1,040

Class	Precision	Recall	F1-Score	Support
<b>Accuracy</b> : 58%	<b>Macro Avg F1</b> : 57%			

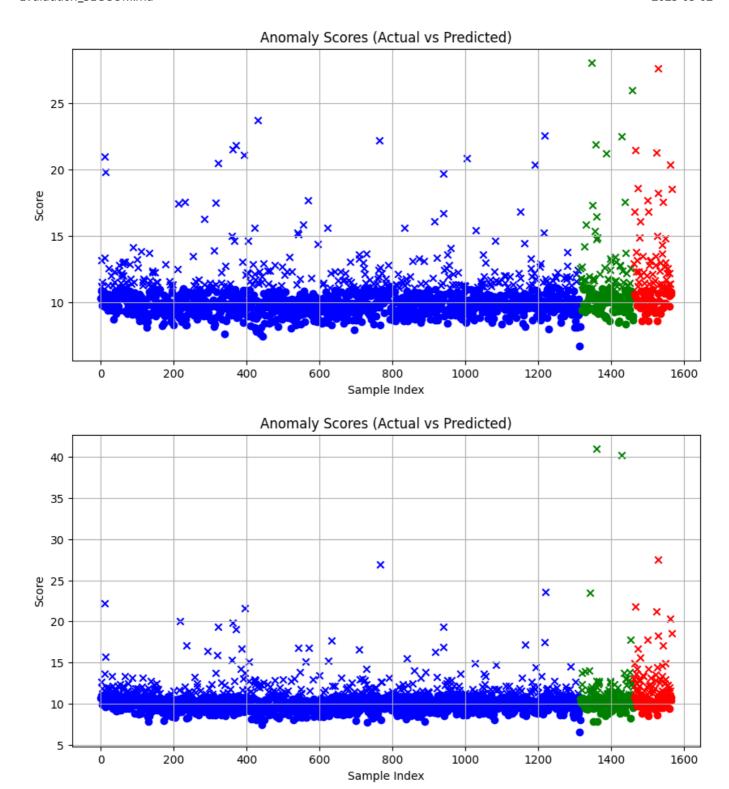
## Training Set:

Metric	Value	
Reconstruction Error	5 to 40	
Training Accuracy	80%	

## Visual Analysis

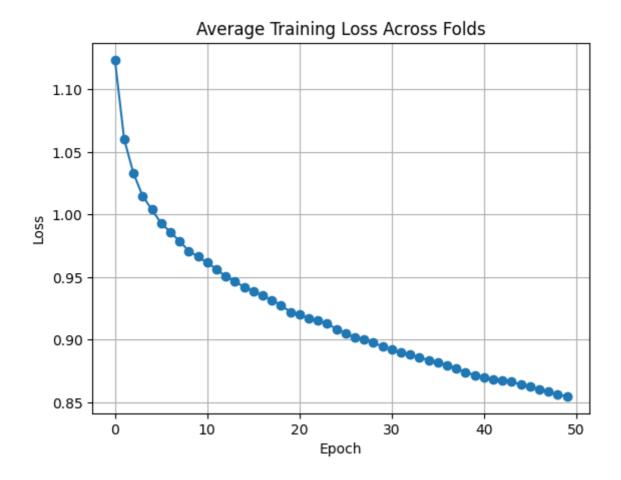


**Critical Overlap**: Most of the samples in test normal and test anomaly overlap in the score distribution.



**Key Insight**: No clear cluster separation despite reconstruction errors.

Training Loss



#### Final Assessment

"All models demonstrate limited discriminative power due to fundamental feature relationships in the SECOM dataset. While achieving expected contamination-aligned accuracy, true anomaly detection capability remains constrained by overlapping score distributions. Priority should be given to feature-space transformations before model optimization."