CADE: Detecting and Explaining Concept Drift Samples for Security Applications

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<u>Limin Yang¹</u>, Wenbo Guo², Qingying Hao¹,

Arridhana Ciptadi³, Ali Ahmadzadeh³, Xinyu Xing², Gang Wang¹

¹University of Illinois at Urbana-Champaign ²Penn State ³Blue Hexagon Inc.







1. Train





2. Predict









Concept Drift! (Unseen family)

Another Type of Drift: In-class Evolution



the outdated model vulnerable to new attacks.

We envision that combating concept drift requires estab-

lishing a monitoring system to examine the relationship between the incoming data streams and the training data (and/or

the current classifier). The high-level idea is illustrated in

Figure 1. While the original classifier is working in the pro-

than traditional methods that focus on explaining a "decision

boundary" in this problem context. We evaluate CADE with

two case studies: Android malware classification and network intrusion detection. We further work with a security com-

pany to test CADE on its malware database. Our results show

that CADE can effectively detect drifting samples and provide

semantically meaningful explanations.

Why Concept Drift Matters?

• New attacks (zero-day) are NOT trivial



FireEye Annual Report 2020

"1.1 million malware samples per day" "41% malware families never seen before"

Why Concept Drift Matters?

• New attacks (zero-day) are NOT trivial

Both malware and goodware evolve over time





UserCode 3rdLib SDK

Functionality scope distribution of Android goodware and malware ^[1].

Why Concept Drift Matters?

• New attacks (zero-day) are NOT trivial

• Both malware and goodware evolve over time

• ML models' decision boundaries shift



GBDT malware classifier trained on Ember-2018; Tested a year later using malware samples from Blue Hexagon Inc. [DLS'21]

When NOT to Predict?



Existing Drifting Detection Solutions

- Solutions from ML community
 - JSTOR'54, SBIA'04, SDM'07, ICML'14, KDD'16, CIKM'19, etc.
 - Most require data labeling on the testing set \rightarrow *costly for security domain*
- Solutions from security community
 - Transcend [USENIX Sec'17]
 - Highly dependent on a good definition of "dissimilarity", scalability issues

Why It's Hard to Define a "Good" Distance Function?

• Distance loses effectiveness in high-dimensional data

– This sample feature space has 1,368 dimensions

 Drifting samples are not labeled, hard to differentiate from normal samples



T-SNE plot for the original space of an Android malware dataset (Unseen family: •)

Self-supervision: Contrastive Learning

- No knowledge about future drifting samples \rightarrow self-supervision
- Use contrastive learning to learn a compressed representation of the training data by contrasting with existing samples



• A sample is far away from ANY existing families' centroids, it's a potential drifting sample; rank for investigation

How to explain these drifting samples?



A Rich set of Explanation Methods

- Identify a small set of important features that make the drifting sample an outlier
- Naïve idea: boundary-based explanation
 - Approximate the decision boundary between drift and in-distribution
 - Explaining a supervised learning model
 - LIME [KDD'16], SHAP[NeurIPS'17], LEMNA [CCS'18], Perturbation [ICCV'17]
 - Using "crossing the boundary" as a signal to derive important features



"Dog"



"Cat"

Problems with Boundary-based Explanation

- Difficult to approximate the boundary
 - Drifting samples are limited
- Difficult to drag a drifting sample to cross the boundary
 - Drifting samples are far away from the boundary in the sparse area





Our Method: Distance-based Explanation

- Perturb the original features and observe the distance changes in latent space
- Perturbation strategy
 - Replace \boldsymbol{x}_t 's feature value with those of a reference sample \boldsymbol{x}_r
 - \mathbf{x}_r is closest to the centroid of nearest family
- Optimization goal
 - Minimize the distance between x_t and \textbf{C}_{A}
 - Use elastic-net regularization to minimize the number of selected important features



Drifting sample

Evaluation: Datasets

Drebin [NDSS'14]

- Top 8 malware families
- 3,317 malware samples
- Training set: 80% of 7 families
- Testing set: 20% of 7 families + unseen family



IDS2018 [<u>ICISSP'18</u>]

- Benign + 3 types of network intrusion
- 130,702 network flows
- Training set: 80% of 3 families
- Testing set: 20% of 3 families + unseen family



Drift Detection Results

Iteratively choose a family as the unseen family and report the average results here.

| Method | Drebin (Avg±Std) | | IDS2018 (Avg±Std) | |
|------------|------------------|-------------------|-------------------|-----------------|
| | F ₁ | Norm. Effort | F ₁ | Norm. Effort |
| Vanilla AE | 0.72 ± 0.15 | $1.48 {\pm} 0.31$ | 0.74±0.12 | 1.74 ± 0.40 |
| Transcend | 0.80 ± 0.12 | 1.29 ± 0.45 | 0.65 ± 0.46 | 1.45 ± 0.57 |
| CADE | 0.96±0.03 | 1.00±0.09 | 0.96±0.06 | 0.95±0.07 |

Real-world test: evaluate on Blue Hexagon PE malware dataset, still effective!

* Vanilla AE: Standard Autoencoder without contrastive learning.

Why CADE Works?



T-SNE visualization for Drebin dataset (Unseen family:
 Iconosys)

Drift Explanation: Case Study

Drifting sample family: FakeDoc; closest family: GingerMaster

• Key difference: FakeDoc usually subscribes to premium services via SMS.

[api_call::android/telephony/SmsManager;->sendTextMessage], [call::readSMS], [permission::android.permission.DISABLE_KEYGUARD], [permission::android.permission.RECEIVE_SMS], [permission::android.permission.SEND_SMS], [permission::android.permission.WRITE_SMS], [real_permission::android.permission.READ_CONTACTS], [permission::android.permission.READ_SMS], [feature::android.hardware.telephony], [api_call::android/location/LocationManager;->isProviderEnabled], [api_call::android/accounts/AccountManager;->getAccounts], [intent::android.intent.category.HOME], [feature::android.hardware.location.network], [real_permission::android.permission.RESTART_PACKAGES], [real_permission::android.permission.WRITE_SETTINGS], [api_call::android/net/ConnectivityManager;->getAllNetworkInfo], [api_call::android/net/wifi/WifiManager;->setWifiEnabled], [api_call::org/apache/http/impl/client/DefaultHttpClient], [url::https://ws.tapjoyads.com/], [url::https://ws.tapjoyads.com/set_publisher_user_id?], [permission::android.permission.CHANGE_WIFI_STATE], [real_permission::android.permission.ACCESS_WIFI_STATE], [real_permission::android.permission.BLUETOOTH], [real_permission::android.permission.BLUETOOTH_ADMIN], [call::setWifiEnabled].

Important features selected by CADE (avg # of selected features is 45 out of 1000+)

Takeaways

• Concept drift is a critical problem for ML/Security applications

• Contrastive Autoencoder is effective to detect concept drift

• Distance-based explanation is more suitable for explaining drifting samples



Thank you!

Homepage

https://liminyang.web.lllinois.edu



Code, features, and supplemental materials available

https://github.com/whyisyoung/CADE

A new PE malware dataset [DLS'21]

https://whyisyoung.github.io/BODMAS/



BODMAS: An Open Dataset for Learning based Temporal Analysis of PE Malware

Limin Yang*, Arridhana Ciptadi[†], Ihar Laziuk[†], Ali Ahmadzadeh[†], Gang Wang* *University of Illinois at Urbana-Champaign [†]Blue Hexagon liminy2@illinois.edu, {arri, ihar, ali}@bluehexagon.ai, gangw@illinois.edu

dataset called BODMAS to facilitate research efforts in machine learning based malware analysis. By closely examining existing open PE malware datasets, we identified two missing capabilities (i.e., recent/timestamped malware samples, and well-curated family information), which have limited researchers' ability to study pressing issues such as concept drift and malware family evolution. For these reasons, we release a new dataset to fill in the gaps. The BODMAS dataset contains 57,293 malware samples and 77,142 benign samples collected from August 2019 to September 2020, with carefully curated family information (581 families) We also perform a preliminary analysis to illustrate the impact of concept drift and discuss how this dataset can help to facilitate existing and future research efforts.

I. INTRODUCTION

Today, machine learning models (including deep neural networks) are broadly applied in malware analysis tasks, by researchers [30], [5], [11], [6] and antivirus vendors [1].

In this field of work, it is highly desirable to have public datasets and open benchmarks. On one hand, these datasets will be instrumental to facilitate new works to resolve open challenges (e.g., adversarial machine learning, interpretation techniques [28], [10]). On the other hand, public benchmarks and datasets can help researchers to easily compare their models and keep track of the progress as a community.

However, creating open malware datasets is highly challenging. For example, the authors of [5] have discussed many of such challenges including legal restrictions, costs and difficulty of labeling malware samples, and potential security liabilities. In addition to these factors, another key challenge is the dynamic evolving nature of malware (as well as benign software) [20]. As new malware families and variants appear over time, they constantly introduce changes to the underlying data distribution. As a result, there is a constant need for releasing new datasets and benchmarks over time.

Over the past decade, there were only a handful of open facilitate future research in our

Abstract-We describe and release an open PE malware malware detection and family attribution. First, most datasets mentioned above contain malware samples that appeared between 2017 to 2019. The data is slightly outdated to study recent malware behaviors. Second, most existing datasets do not contain well-curated family information. This limits researchers' ability to test learning-based family attribution methods and analyze family evolution patterns.

For these reasons, we compile a new dataset, called BODMAS, to complement existing datasets. Our dataset contains 57,293 malware samples and 77,142 benign samples (134,435 in total). The malware is randomly sampled each month from a security company's internal malware database, from August 29, 2019, to September 30, 2020 (one year). For each sample, we include both the original PE binary as well as a pre-extracted feature vector that shares the same format with existing datasets such as Ember [5] and SOREL-20M [11]. Researchers could easily combine our dataset with existing ones to use them together. More importantly, our dataset provides well-curated family labels (curated by security analysts) covering 581 malware families. The family label information is much richer than existing datasets (e.g., the Microsoft dataset [24] only has 9 families).

Preliminary Analysis. In this paper, we use our dataset (and existing datasets) to perform a preliminary analysis on the impact of concept drift (where the testing set distribution shifts away from the training set [8]) on binary malware classifiers and multi-class family attribution methods. We illustrate the impact of concept drift on different learning tasks. In particular, we highlight the challenges introduced by the arrival of previously unseen malware families, which have contributed to increasing false negatives of binary malware classifiers and crippled malware family classifiers in an "openworld" setting. In the end, we discuss the open questions related to our observations and how BODMAS could help to

Backup Slides

Drift Detection in the Latent Space

- Drift detection: if a sample is far away from ANY existing families' centroid, it's a potential drifting sample
- But different families' tightness vary, how to set distance thresholds?
- MAD (Median Absolute Deviation) ^[1]
 - Median of the median distance to the centroid
 - $-MAD = b * median(|X_i median(X)|)$
 - *X* is a set of distances to the centroid
 - Any new data outside $median(X) \pm A * MAD \rightarrow$ outlier
- Rank drifting samples for investigation





Drift Detection: Evaluation Metrics

| • | Procision _ detected unseen family samples | | | |
|---|--|--|--|--|
| | inspected samples | | | |
| • | Recall – detected unseen family samples | | | |
| | total # of unseen family samples | | | |
| | 2 * Precision * Recall | | | |
| • | $\Gamma_1 = -$ Precision + Recall | | | |
| • | Normalized Inspection effort = | | | |
| | inspected samples | | | |
| | total # of unseen family samples | | | |

Training set: A (200), B(200), C(200) Testing set: A(50), B(50), C(50), **D(10)**

| Family | Precision | Recall |
|--------|-----------------------|--|
| D | 1/1 = 1 | 1/10 = 0.1 |
| D | 2/2 = 1 | 2/10 = 0.2 |
| С | 2/3 = 0.67 | 2/10 = 0.2 |
| D | 3/4 = 0.75 | 3/10 = 0.3 |
| D | 4/5 = 0.8 | 4/10 = 0.4 |
| | D D C D D | PrecisionD $1/1 = 1$ D $2/2 = 1$ C $2/3 = 0.67$ D $3/4 = 0.75$ D $4/5 = 0.8$ |

A ranked list of detected samples

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Real-world Test on Blue Hexagon PE Malware

- 20,613 Windows PE malware, 395 families, Sept. 2019 Feb. 2020
- 2,381 features, training with top N families Sept. 2019 Jan. 2020
- Testing set: Feb. 2020
- CADE is still effective!

BLUEHEXAGON

| N (training families) | F ₁ | Norm. Effort | Detected Unseen Families |
|-----------------------------|----------------|-----------------|--------------------------------|
| 5 | 0.97 | 1.02 | 161/165 |
| 10 | 0.95 | 0.98 | 153/160 |
| 15 | 0.87 | 0.84 | 140/155 |

Drift Explanation: Evaluation Metrics and Results

- Metric: the latent distance between a perturbed sample and its closest centroid.
- Using CADE to select important features, $x_t \to x_p$, while baseline methods may $\to x_p'$, which is still far away from C_A .



| Method | Drebin-FakeDoc Avg±Std | IDS2018-Infiltration Avg±Std |
|--------------------------|---------------------------|---------------------------------|
| Original distance | 5.363 ± 0.568 | 11.715 ± 2.321 |
| Random | 5.422±1.773 | 11.546 ± 3.169 |
| Boundary-based | 3.960 ± 2.963 | 6.184 ± 3.359 |
| COIN [<u>IJCAI'18</u>] | 6.219 ± 3.962 | 8.921±2.234 |
| CADE | 0.065±0.035 | 2.349±3.238 |