

Learning and Classification of Malware Behavior

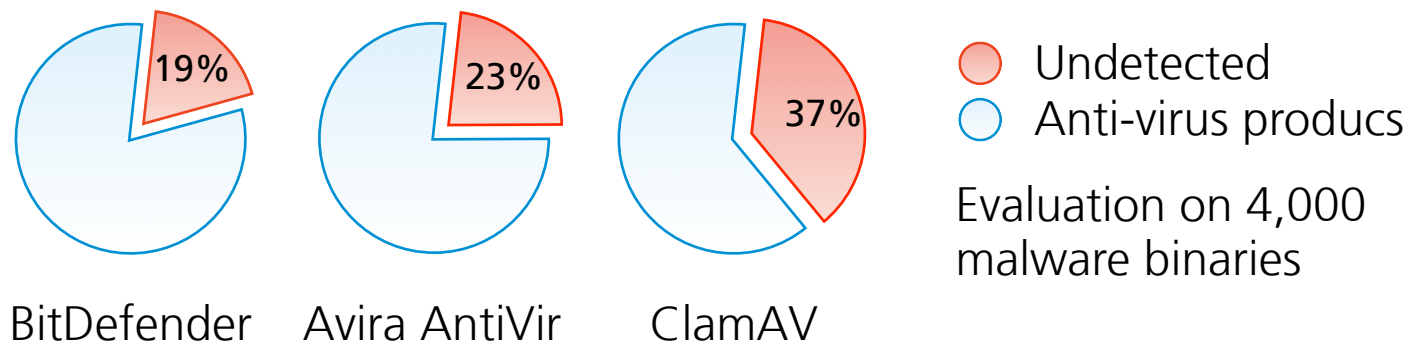
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DIMVA 2008, Paris, France

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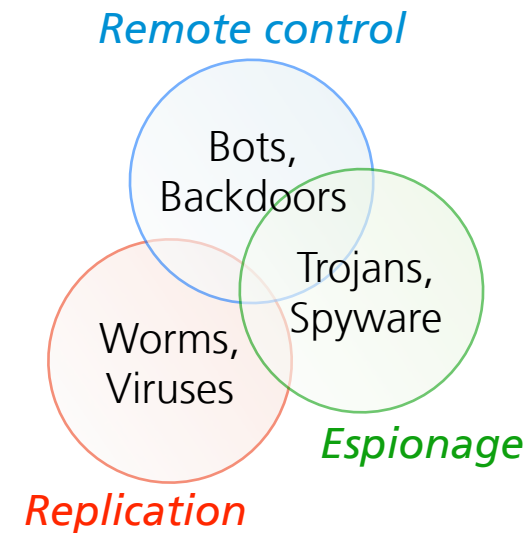
² University of Mannheim, Germany

- ▶ **Malicious software: A vivid threat**
 - ▶ Plethora of worms, trojans, bots, backdoors
 - ▶ Exponential growth of malware in the wild
 - ▶ Emergence of criminal “industries”
- ▶ **Conventional static defenses insufficient**
 - ▶ High degree of polymorphy and obfuscation



- ▶ **Malware behavior**

- ▶ Malware differs in purpose and functionality
- ▶ Typical and discriminative behavioral patterns

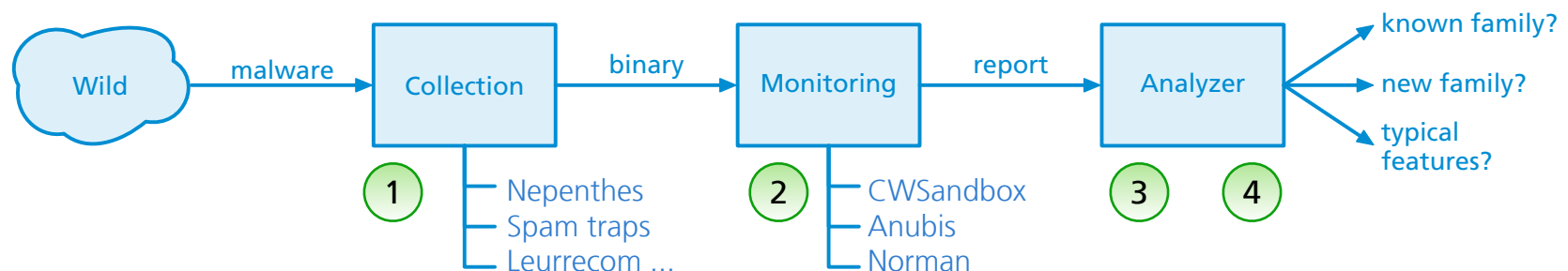


- ▶ **Behavior-based analysis**

- ▶ Monitoring and detection of malicious behavior
- ▶ AV products: manually generated behavior rules
- ▶ Alternative, fully automated approaches?

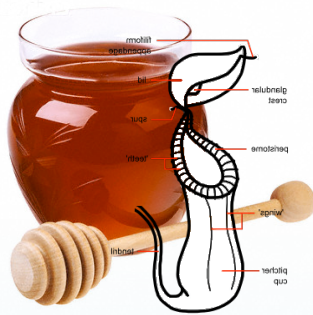
Learning-based approaches

- ▶ **Clustering of malware behavior** (e.g. Bailey et al., RAID 2007)
 - ▶ Difficult to control cluster models (many vs. few)
 - ▶ Clustering often non-predictive, e.g. linkage clustering
- ▶ **Idea: Generalize from behavior and prior knowledge**
 - ▶ Incorporate (noisy) labels, e.g. by anti-virus tool
 - ▶ Learn classification of malware families using labels



- ▶ Automatic collection of current malware families
 - ▶ Broad range of malware using diverse methods, e.g. honeypots, spam traps, honeyclients

*Vulnerability
emulation*



Nepenthes

Self-replicating
malware



Spam traps

Trojans and
backdoors

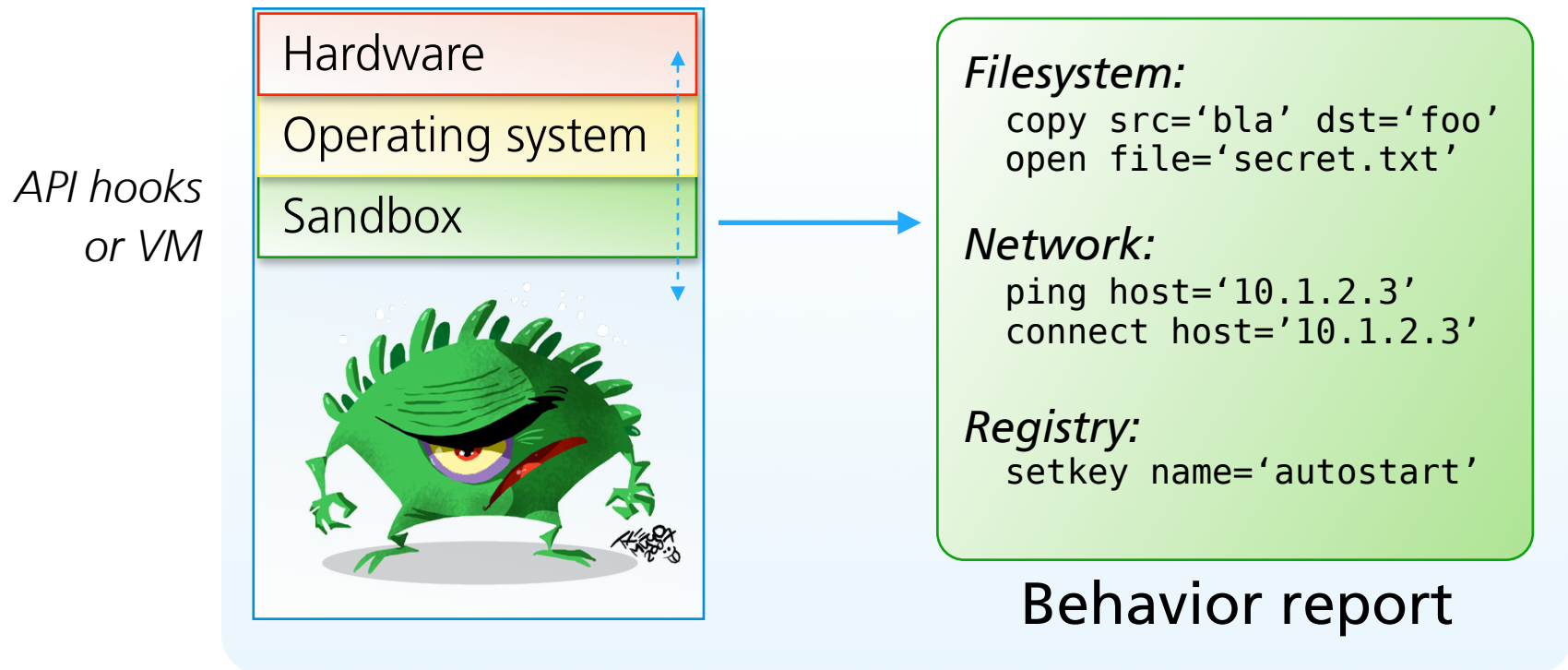


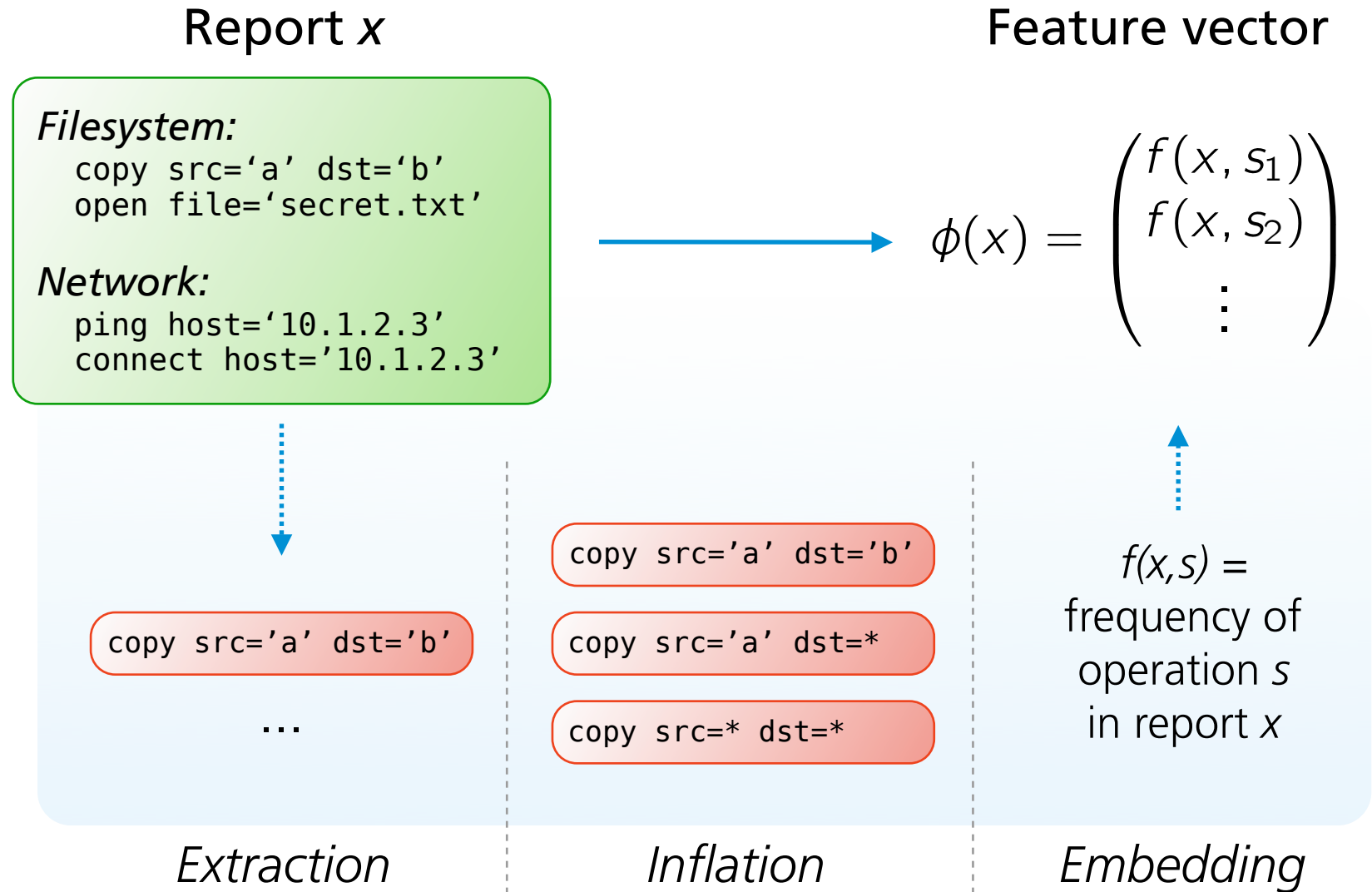
Honeyclients

Drive-by
malware

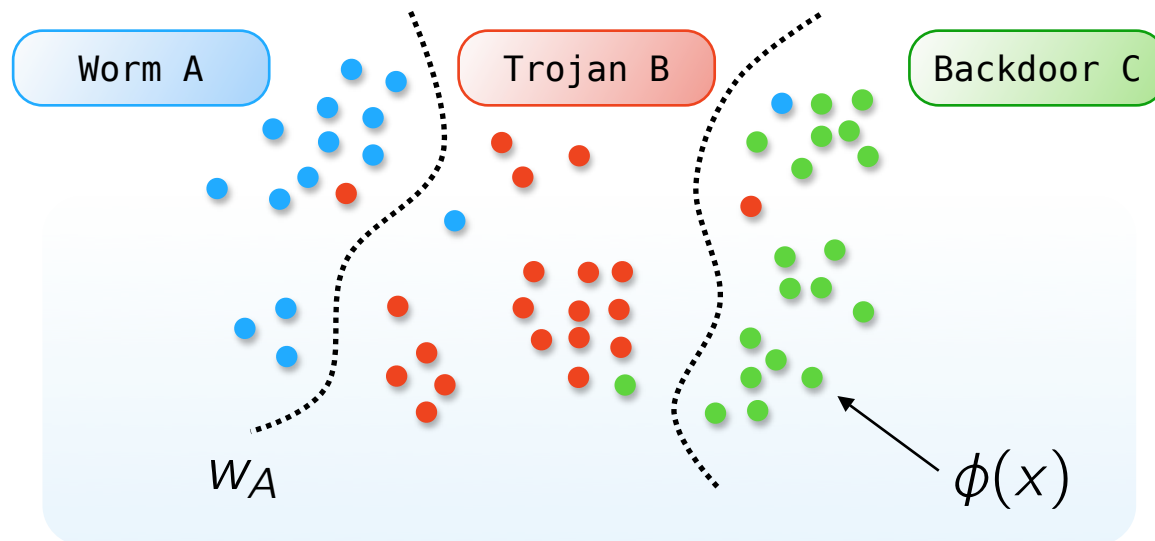
*Client-side
emulation*

- ▶ Sandbox for malware
 - ▶ Protected execution environment (e.g. CWSandbox)
 - ▶ Monitors and reports observed behavior





- ▶ Discrimination of malware families in feature space
 - ▶ Assign family label to embedded reports, e.g. AV label



- ▶ Learn maximum-margin hyperplane w for each family
- ▶ Incorporation of non-linearity using kernel functions

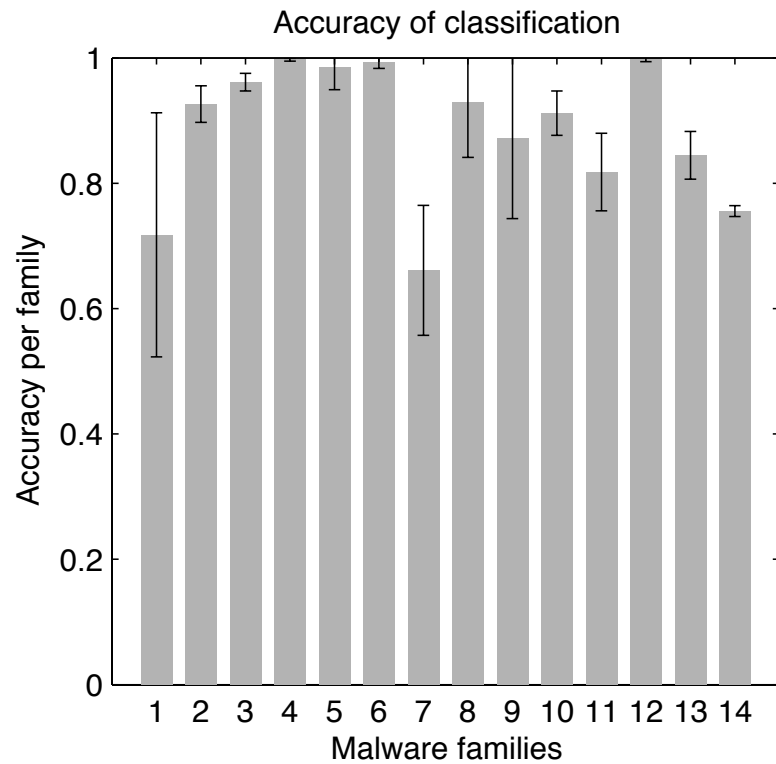
Experimental evaluation

- ▶ Malware collection labeled using AV tool (AntiVir)
 - ▶ #k: 10,072 malware binaries from 14 families
 - ▶ #u: 3,139 unknown variants (detected 4 weeks later)

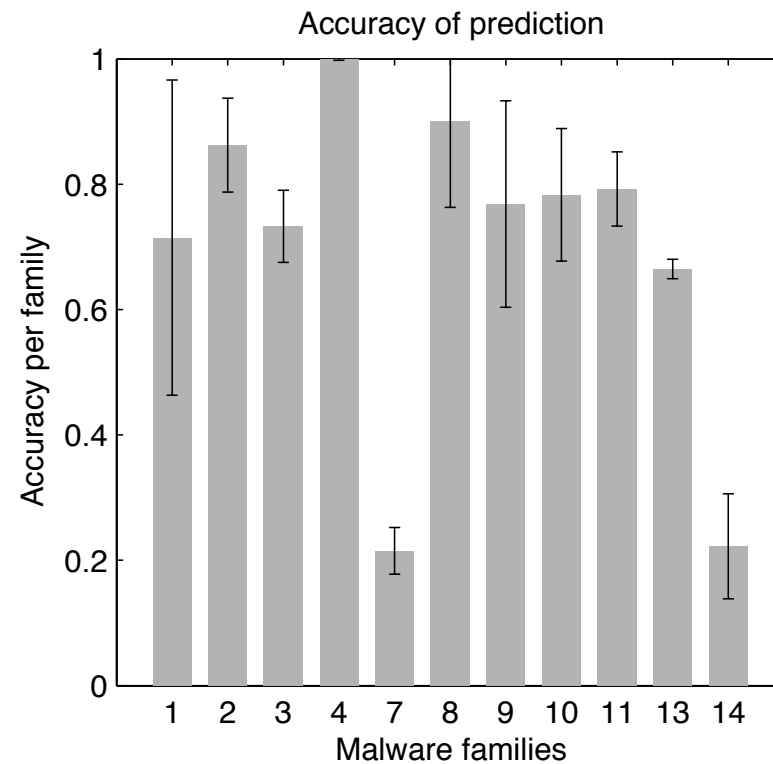
Malware family	#k	#u	Malware family	#k	#u
1: Backdoor.VanBot	91	169	8: Worm.Korgo	244	4
2: Trojan.Bancos	279	208	9: Worm.Parite	1215	19
3: Trojan.Banker	834	185	10: Worm.PoeBot	140	188
4: Worm.Allaple	1500	614	11: Worm.Rbot	1399	904
5: Worm.Doomber	426	0	12: Worm.Sality	661	0
6: Worm.Gobot	777	0	13: Worm.SdBot	777	597
7: Worm.IRCBot	229	107	14: Worm.Virut	1500	144

Results: Classification

- ▶ Learning on known, prediction on unknown variants



Known variants, avg. 88%



Unknown variants, avg. 69%

- ▶ High detection accuracy (Note: random guessing = 7%)

Results: Explanation

- ▶ Explanation of learned malware behavior classifier
 - ▶ Most discriminative dimensions in hyperplane vectors

Worm.Sality

```
0.0142: create_file ... srcpath="C:\windows\system32\" src=*
0.0073: create_file ... srcpath="C:\windows\system32\" src="vcmgcd32.dl_"
0.0068: delete_file ... srcpath="C:\windows\system32\" src=*
0.0051: create_mutex name="kuku_joker_v3.09"
0.0035: enum_processes apifunction="Process32First"
```

Worm.Doomber

```
0.0084: create_mutex name="GhostBOT0.58c"
0.0073: create_mutex name="GhostBOT0.58b"
0.0052: create_mutex name="GhostBOT0.58a"
0.0014: enum_processes apifunction="Process32First"
0.0011: query_value key="HKEY_LOCAL_MACHINE\...\run" value="GUARD"
```

- ▶ **Behavior-based malware analysis**
 - ▶ Extension of current AV tools *(see Oberheide et al., USENIX 2008)*
 - ▶ Hinders simple obfuscation and polymorphy
- ▶ **Supervised learning on malware behavior**
 - ▶ Detection accuracy: *69% unknown malware variants*
 - ▶ No black box: *Explanation via hyperplane vectors*
 - ▶ Further extension: *Rejection of unknown behavior*
- ▶ **Perspectives**
 - ▶ Semi-supervised learning: Best of both worlds.

Thanks. *Questions?*

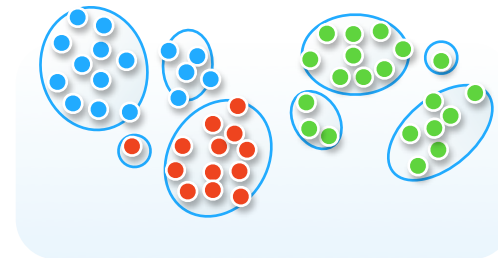


- ▶ **Evasion attacks**
 - ▶ Detection of honeypot or sandbox environment
 - ▶ Obfuscated and polymorphic behavior
 - ▶ Mimic behavior of benign programs or other malware
- ▶ **Consequences & defenses**
 - ▶ Run multiple honeypots and sandboxes in parallel
 - ▶ Obfuscation and polymorphy: Discriminative features?
 - ▶ Fruitless to mimic benign program = No real activity

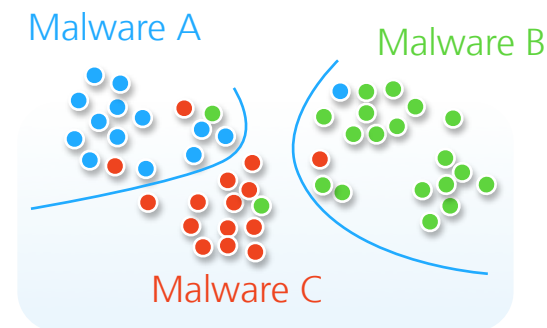
- ▶ Bächer, Kötter, Holz, Dornseif, Freiling. *The Nepenthes platform: An efficient approach to collect malware*. RAID 2006.
- ▶ Bailey, Oberheide, Andersen, Mao, Jahanian, Nazario. *Automated classification and analysis of Internet malware*. RAID 2007.
- ▶ Burges. *A Tutorial on Support Vector Machines for Pattern Recognition*. Knowledge Discovery and Data Mining 2(2), 1998.
- ▶ Oberheide, Cooke, Jahanian. *N-Version Antivirus in the Network Cloud*. USENIX 2008.
- ▶ Rieck, Laskov. *Linear-Time Computation of Similarity Measure for Sequential Data*. Journal of Machine Learning Research 9(1), 2008.
- ▶ Willems, Holz, Freiling. *Towards automated dynamic binary analysis*, IEEE Magazine Security & Privacy 5(2), 2007.

Unsupervised vs. Supervised

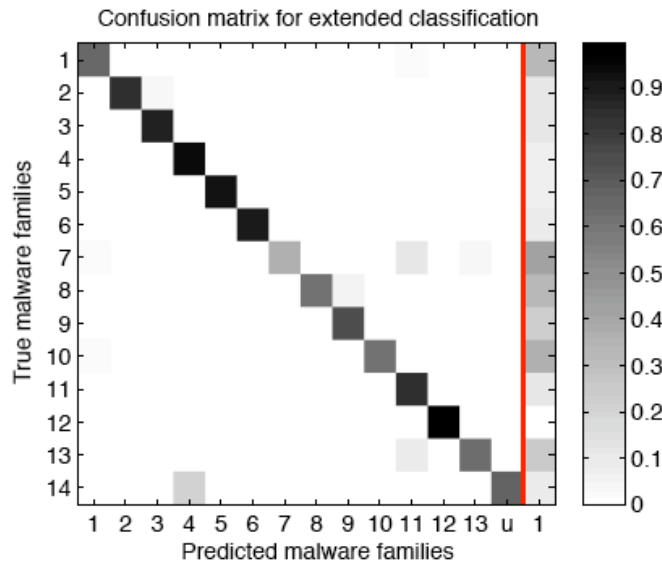
- ▶ **Clustering (unsupervised)**
 - ▶ Determine malware families from structure only
 - ▶ Difficult to control model complexity without labels



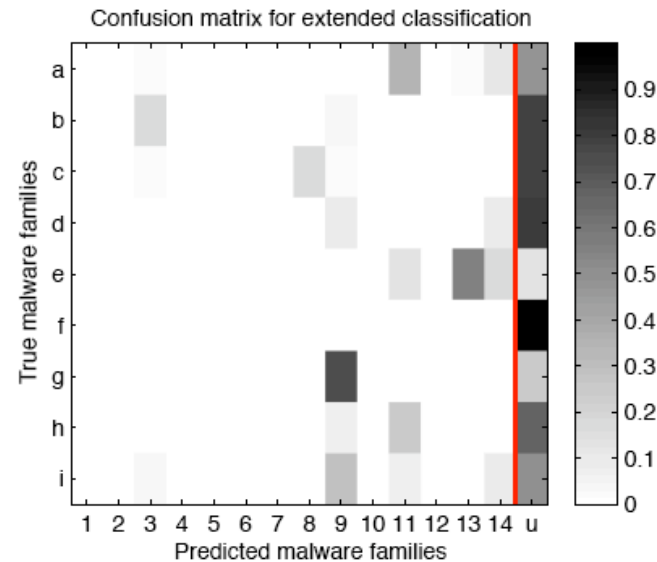
- ▶ **Classification (supervised)**
 - ▶ Determine malware families using structure *and* labels
 - ▶ Generalization beyond noisy labels



- ▶ Rejection of unknown behavior
 - ▶ Probabilistic fit on output of classifier (reject if <0.5)



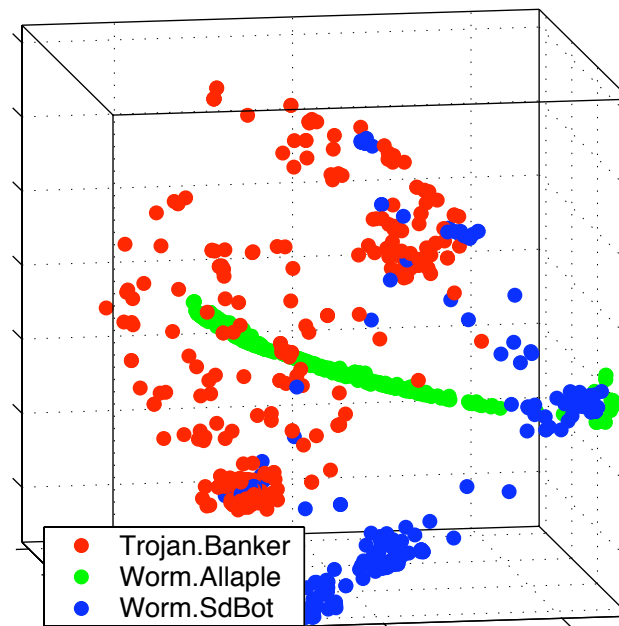
Known families



Unknown families

- ▶ Reliable rejection of unknown behavior, yet accuracy decreases from 88% to 73%

- ▶ Embedding to high-dimensional vector space
 - ▶ Each operation spans several dimensions
 - ▶ > 1,000,000 and more dimensions



- ▶ Visualization using projections (e.g. with PCA)