

Malware family classification via efficient Huffman features

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Malware family classification via efficient Huffman features

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MDFRWS Background

- Malware is constantly on the rise.
- Classification is important to group according to similar traits, behaviours etc.
- Malware feature extraction approaches:
 - can be labour-intensive so may not scale well.
 - require knowledge of malware's internal binary structure.
- Feature selection crucial to classification performance.
 - Malware analysts are not data scientists
- Need for an "automated" solution.





IDFRWS Compression

- Think 7-zip, Bzip, LZMA etc.
- Encodes data to a reduced representation (fewer bits)
 - reduced storage space or bandwidth for transmission
- For our purposes, we can represent a full binary file with less data.
- Compression can be applied to data in any format wide range of possible domains.





MDFRWS Related Work

Distance Metrics

- Normalized Compression Distance (NCD) (Cilibrasi & Vitanyi, 2005).
- LZJD and SHWel (Raff & Nicholas, 2017)

Feature space

- Dictionaries of substrings can be engineered as feature vectors (Sculley & Brodley, 2006).
- Applied to text classification problems (Paskov et al., 2013).

Limitations:

- NCD computationally inefficient
- LZ variants feature space infeasibly large

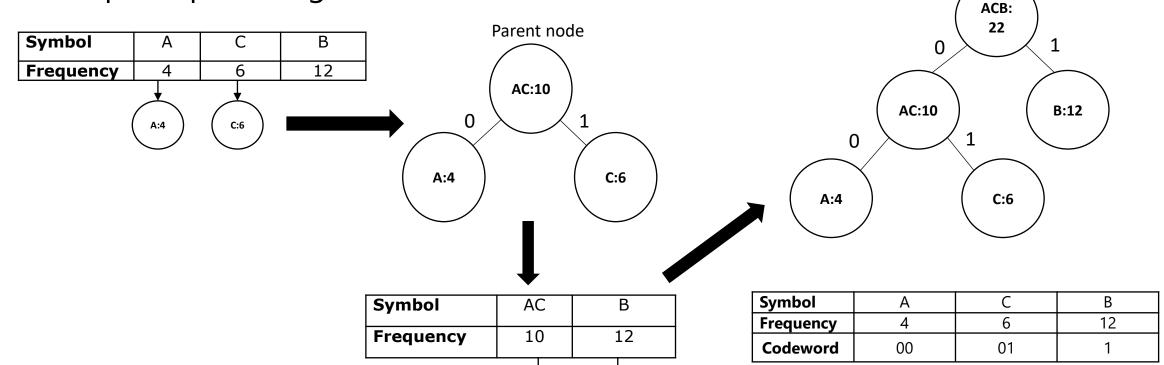




MDFRWS Huffman Coding

 Prefix codes generated from a set of symbol (character) frequencies.

■ Example: input string: AAAABBBBBBBBBBBBCCCCCC



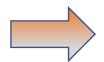
12



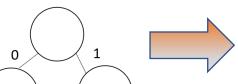


IDFRWS eHf algorithm





 $h \{ [\nu, [\kappa, c = \{\emptyset\}]] \}$



sorted $h\{[\nu, [\kappa, c]]\}$

Malware sample

 $h = heap; v = frequency; \kappa = symbol, c = codeword$

Symbol converted to decimal:

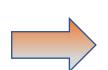
- A = 65
- = 66
- C = 67

Initial heap

 $[4, [65, \emptyset]]$

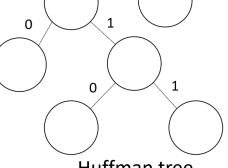
 $[6, [67, \emptyset]]$

 $[12, [66, \emptyset]]$



Heap after processing

[22, [66, '1'],[65,'00'], [67, '01']]



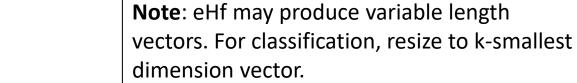
Huffman tree



foreach i in h:

$$eHf_i = \kappa_i + \nu_i + int(c_i)$$





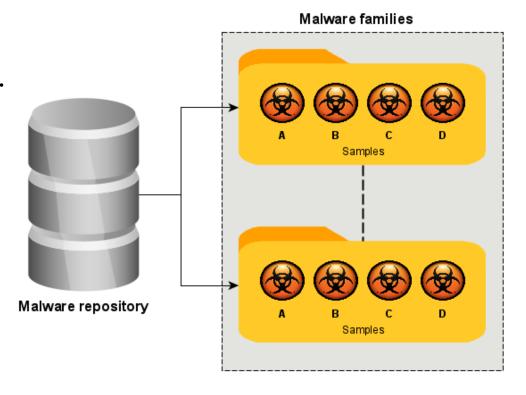
eHf feature vector = [69, 74, 79]





MDFRWS Data

- Win32 portable executable files.
- VirusTotal academic share (~2018 present).
- AV Class Labeller (Sebastian et al., 2016).
- Initial training dataset 8,232 from 12 families.
- Extended to 14,694 from 23 families.

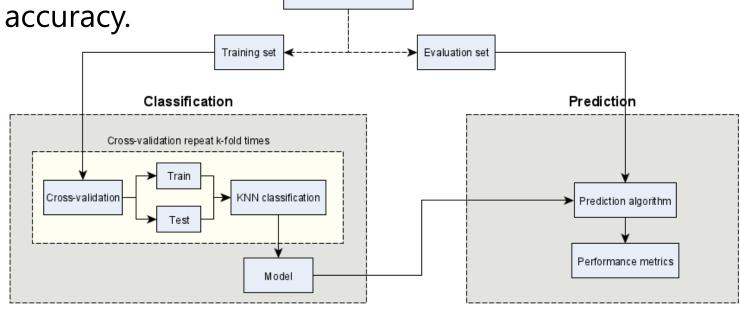






MDFRWS Classification

- Data and label sets split 90:10 for training and evaluation.
- k-nearest neighbour algorithm (k = 3, Minkowski).
- Parameter tuning using GridSearchCV.
- 5-fold stratified cross-validation.
- Metrics: precision, recall and accuracy.



Huffman features data



MDFRWS Results

Family	Precision	Recall	Accuracy	
Agent.BDMJ	0.989	1.000	0.994	
Autoit	0.961	0.976	0.969	
Berbew	0.993	0.986	0.990	
Dinwod	0.994	0.983	0.988	
Dorkbot	0.977	0.988	0.982	
Dridex	1.000	1.000	1.000	
Oberal	0.976	1.000	0.988	
Scar	0.857	0.854	0.855	
Sfone	0.987	0.996	0.991	
Socks	0.991	0.980	0.986	
Sytro	0.994	0.999	0.997	
Vilsel	0.985	0.971	0.978	
Weighted avg.	0.982	0.982	0.982	

- Initial dataset *8,232* samples.
- 11 out of 12 class prediction true positive rates (TPR) of 97% or above.
- Scar family poorest performer some mislabelling discovered on VT.
- 10% evaluation testing returned ~97% precision, recall and accuracy.





MDFRWS Comparison with NCD

Dist. metric	Runtime (secs)	Prec.	Recall	Acc.
eHf-Jaccard	1.42×10^{-3}	0.969	0.968	0.968
eHf-Minkowski	1.02×10^{-3}	0.972	0.973	0.972
eHf-Euclidean	1.06×10^{-3}	0.970	0.971	0.969
NCD	1.2	0.782	0.774	0.772

■ Compare eHf + standard distance metrics with NCD



IDFRWS Comparison with LZJD¹

	Run-time ef	Size	
	feat. gen.	training	feat. dims.
LZJD-sh	5.15×10^{-1}	9.92×10^{-2}	1024
LZJD	1.21×10^{-1}	1.86×10^{-1}	1024
eHf	5.0×10^{-2}	2.71×10^{-2}	229

Comparison of time complexities on larger dataset.





IDFRWS Comparison with LZJD²

	Training		Validation			
	precision	recall	accuracy	precision	recall	accuracy
LZJD-sh	0.977	0.973	0.974	0.882	0.878	0.873
LZJD	0.951	0.950	0.950	0.745	0.752	0.746
eHf	0.972	0.973	0.972	0.974	0.974	0.974

Comparison of classification performance.





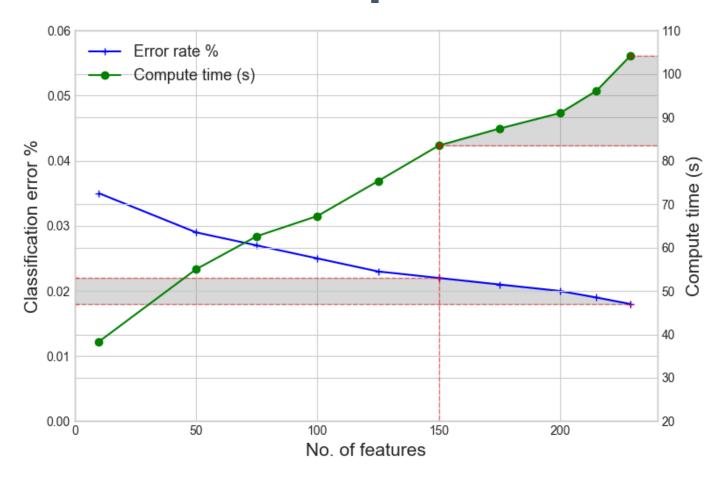
IDFRWS Code reordering obfuscation

- Import tables extracted from all samples in the VT dataset.
- KNN classifier trained as previously using the eHf import table vectors.
- SCYTHE tool used to reorder ~3k sub-sample import tables.
- KNN model: precision, recall and accuracy of 99.8%, 99.7% and 99.7% respectively.
- Ordering of the input sequence is not a consideration in the generation of the codewords as data is stored according to frequency of symbols.





MDFRWS Feature optimization



- Features can be optimized.
- Reduction from 229 dimensions to 150 = 20% less compute time.
- Error rate increase of 0.4%.
- Dependent on data.





IDFRWS Summary of contributions

- Novel method of representing binary features.
- Negates the need for invasive analysis techniques.
- Does not require intricate knowledge of binary structures.
- (Quite) Fast and scalable.
- FOSS
- Potential to apply to other domains.
- Outputs can be "plugged in" to other ML algorithms.





MDFRWS Study limitations

- eHf developed in January 2021.
 - Limited to malware executables and import table dumps.
 - Only shows results from KNN classifier.
 - Other forms of obfuscation not tested.





MDFRWS Future work

- Test at scale (Sorel 20M dataset)¹
- Improve processing speeds.
- Other forensic scenarios.
- Further obfuscation testing.





MDFRWS Thank you!





