

*An Immunity-Based Technique to  
Characterize Intrusions in Computer  
Networks*

By D. Dasgupta and F. Gonzalez,  
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# *Overview*

- ❖ Artificial Immune Systems (AIS)
- ❖ Paper credentials
- ❖ New and Significant
- ❖ Results
- ❖ Other Methods
- ❖ Intro and Definitions
- ❖ Positive Characterization
- ❖ Negative Characterization
- ❖ Comparisons
- ❖ Criticism



# *Artificial Immune Systems*

- ❖ Ideas first published in 1994
- ❖ Dedicated conference since 2001 - International Conference on Artificial Immune Systems (ICARIS)
- ❖ 805 entries found by Google Scholar
- ❖ DOD Office of Force Transformation described Network Immune Systems as hot trend. One of the companies listed, Okena, bought by Cisco.

# *Paper Credentials*

- ❖ Published in IEEE Transactions on Evolutionary Computation - peer-reviewed quarterly journal of IEEE Computational Intelligence Society
- ❖ Google Scholar: Cited by 35
- ❖ Prof. Dasgupta:
  - 1998 editor of Springer book on AIS
  - AIS track organizer for Genetic and Evolutionary Computation Conference (GECCO) 2004/2005
  - Chair of IEEE Working Group on AIS

# *New and Significant*

- ❖ Approach inspired by immune system
  - Extends previous two-class self/non-self approach to multi-class approach
  - No reliance on structured representation of data
  - Need only positive data to build normal profile of system
  - Applied to anomaly detection for network security but general approach can be applied to different anomaly detection problems

# *Results*

- ❖ Negative Characterization (NC) approach compares very favorably to Positive Characterization (PC) approach
  - Only 10% of space needed
  - Results “closely” match PC results

# *Introduction*

- ❖ Immune system: Distinguish self from non-self
- ❖ Computer system: Legitimate users and data from unauthorized users and viruses
- ❖ 1994 Stephanie Forrest at U of NM used negative-selection algorithm (detect only non-self)

# *Statistical Methods*

- ❖ Simple model: individually different state variables.
  - Normalcy depends on time
  - Normalcy depends on correlations among parameters
- ❖ Temporal and multi-var correlations
  - Require too much training time
  - Infeasible due to large data set size



# *Data Mining*

- ❖ Some success in capturing multi-variable correlations but
  - Data must have some degree of structure
  - Need information on which connections are normal versus which connections are attacks for some algorithms

# *Definitions*

## ❖ System State Space

- State  $S$  consists of vectors of features corresponding to all possible states of system.

$$x^i = (x_1^i, x_2^i, \dots, x_n^i) \quad [0.0, 1.0]^n$$

## ❖ Normal Subspace:

$$\textit{Self} \quad S$$

$$\textit{Non\_Self} = S - \textit{Self}$$

# *More Definitions*

❖ Characteristic function:

$$\mu_{self} : [0.0, 1.0]^n \quad [0.0, 1.0]$$

$$m_{self,t}(\vec{x})$$

1 if  $> t$

0 if  $\leq t$

Degrees of normalcy: 1 normal - 0 abnormal

# *Anomaly Detection Problem*

- ❖ Given a set of normal samples build good estimate of  $\mu_{self}$
- ❖ This function decides whether or not observed state is anomalous

# *Positive Characterization (PC) Approach*

- ❖ Use positive sample set for representation of *Self* space as *Self* set
- ❖ Characteristic function is distance from that element to nearest neighbor in *Self* set
  - Euclidean distance but also used:

$$D(\vec{x}, \vec{y}) = \max(|x_1 - y_1|, \dots, |x_n - y_n|)$$

# *PC Implementation*

- ❖ Use Bentley's KD-tree
  - Represents k-dimensional points
  - Generalization of one-dim binary search tree
  - Amortized cost of nearest-neighbor query is  $O(\log N)$

# *Data Sets for Experiments*

- ❖ Data sets on intrusion detection from MIT Lincoln Labs
  - Normal and abnormal data from test lab with simulated attacks
  - Standard corpora for IDS
  - Network traffic (tcpdump), audit data (Solaris BSM) and file system data
  - Five weeks of data with ~300k per day compressed (3 training/labeled, 2 test)

# *Experiment Setup*

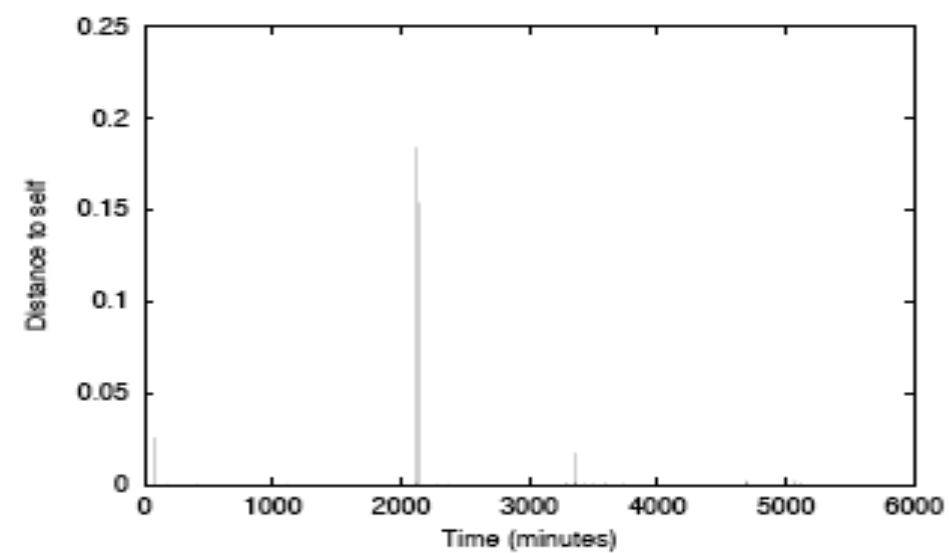
- ❖ Train using first week - attack free data
- ❖ Use second week for test - only network attacks
- ❖ Parameters used:
  - Number of bytes per second
  - Number of packets per second
  - Number of Internet Control Message Protocol (ICMP) packets per second
- ❖ Sample each minute. Built set  $S$  of normal descriptors from time series  $R$  in overlapping sliding windows with window size  $w$ :

$$S = \{(r_1, \dots, r_w), (r_2, \dots, r_{w+1}), \dots, (r_{n-w+1}, \dots, r_n)\}$$

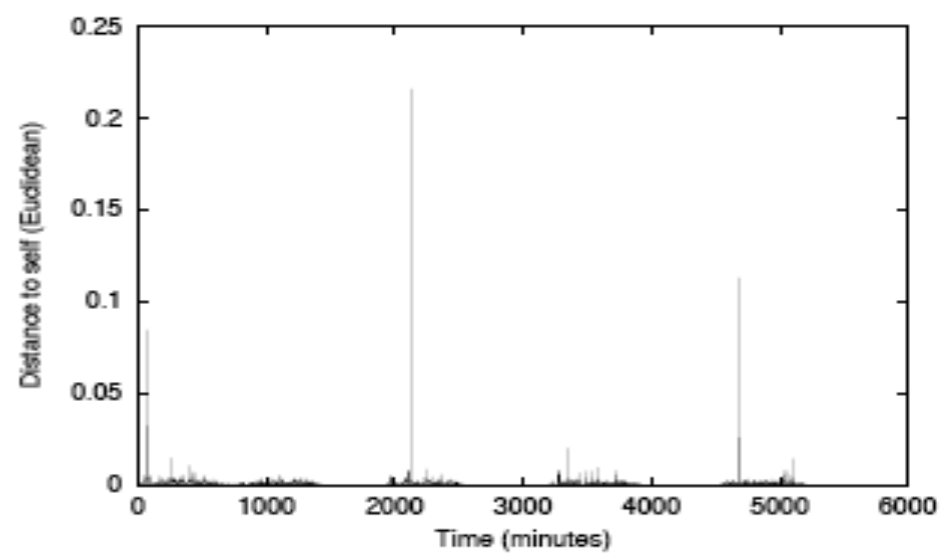


# *Experiment 1*

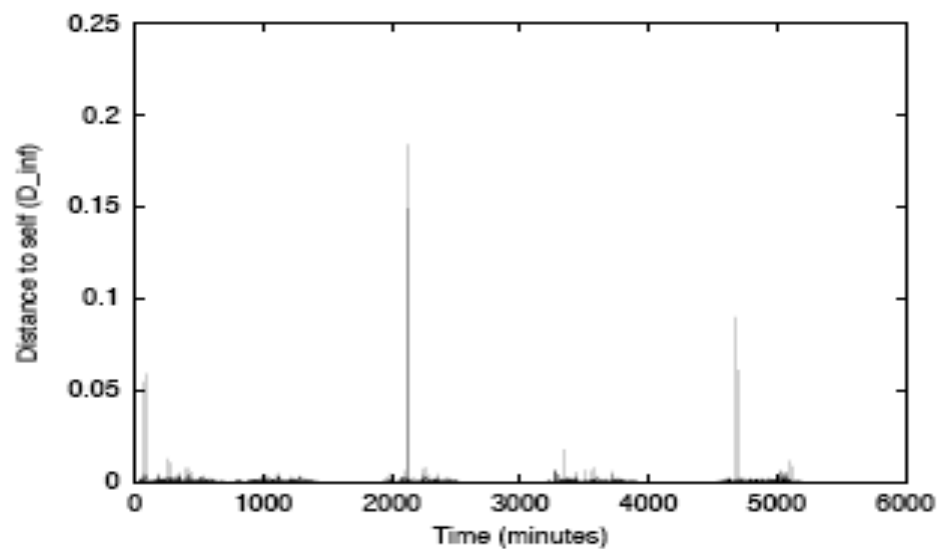
- ❖ Considers variables independently
  - Not enough to detect all five attacks
  - Higher window size increases sensitivity
  - Higher window size allows for detection of temporal patterns
  - Change of metric does not modify number/type of deviations detected



(a)  $ws = 1$



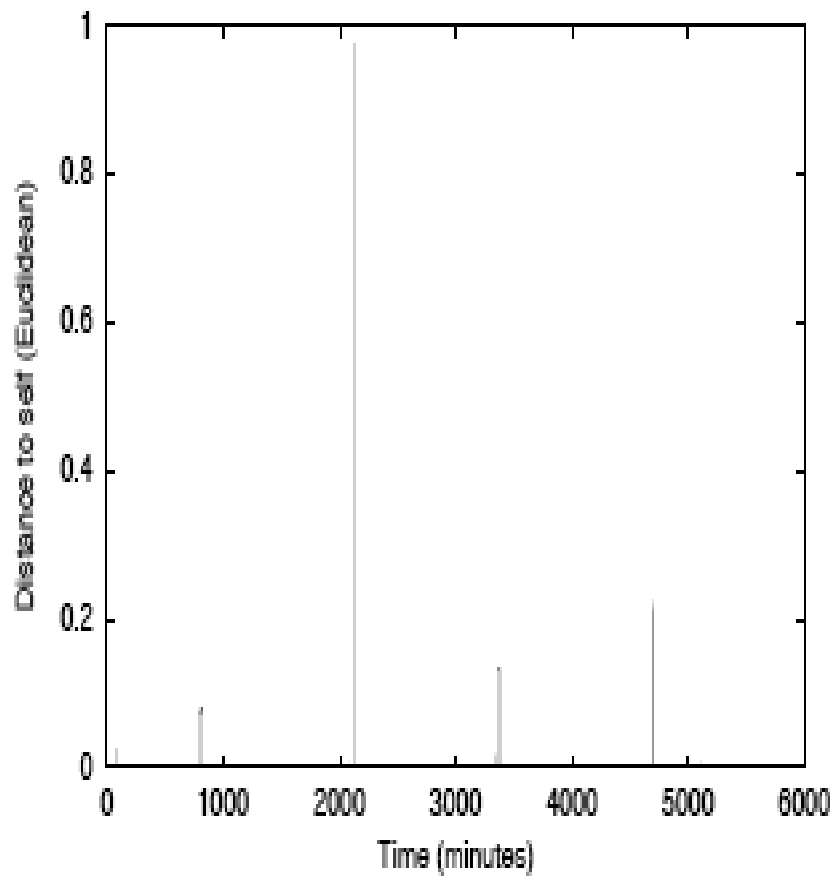
(b)  $ws = 3$ , Euclidean



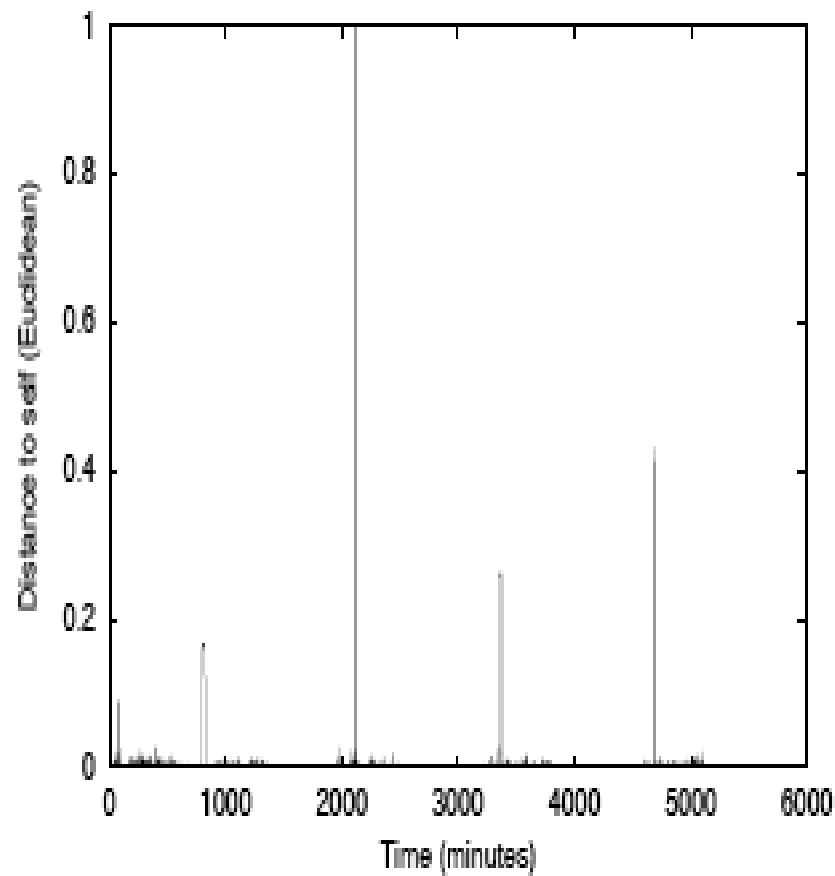
(c)  $ws = 3$ ,  $D_\alpha$

## *Experiment 2*

- ❖ Uses all three network params as features
- ❖ Use window size 1 and 3
- ❖ Detects all five attacks
- ❖ Increasing window size increases sensitivity



(a)  $ws = 1$



(b)  $ws = 3$

# *Positive Characterization*

- ❖ Worked well on performed experiments
- ❖ Problem:
  - Must store all samples that constitute normal profile
  - Infeasible for large data generated by network traffic!

# *Human Immune System*

- ❖ Self/Non-Self distinction with T-Cells
  - T-Cell receptors made with pseudo-random genetic rearrangement process.
  - Negative selection: Censoring process in thymus. Destroy T-cells that react against self-proteins
  - Circulate throughout body to detect non-self proteins (antigens).

# *Original Negative Selection Algorithm*

- ❖ Forrest et al. 1994
- ❖ Self:  $S$  - collection of strings of length  $l$  over finite alphabet (e.g. normal activity segmented into equal-sized substrings)
- ❖ Set of detectors  $R$  - fail to match any string in  $S$
- ❖ Monitor  $S$  for matches of  $R$  - change!
- ❖ Used binary representation (either Self or Non-Self),  $O(N)$  runtime ( $N$  = number of strings in  $S$ )

# *Extended Negative Selection*

- ❖ Use real-valued representation for detector rules

R\_1: If Cond\_1 then non-self (or level of variability)

...

R\_m: if Cond\_m then non-self

*Cond\_i: consists of ranges between a lower and upper value for each feature of a feature vector. The condition is met if all features fall within their respective ranges.*

*Levels: Different levels of variability  $v$  within normal space. Match highest level rule.*



# *Detector Rules*

- ❖ Conditions define hypercube in descriptor space
- ❖ Set of rules try to cover non-self space with hypercubes
- ❖ If a feature vector matches a condition it must be part of non-self space or that variable level (pick highest matching rule)
- ❖ How to create rules? Genetic algorithm



Self

Level 1

Level 2

# *Criteria for GA evolution*

- ❖ Multi-objective optimization problem
  - Don't cover positive examples
  - Large area (try to minimize number of rules)
  - Minimize overlap with other rules (use sequential niching algorithm)

# *GA Input*

- ❖ Feature vectors set  $S$  of normal behavior vectors (training set)
- ❖  $v$ : Level of variability
- ❖  $\text{maxRules}$ : max rules in solution set
- ❖  $\text{minFitness}$ : min value for rule to be included in solution set
- ❖  $\text{maxAttempts}$ : max number of attempts to evolve a rule with fitness  $> \text{minFitness}$

# *GA Representation*

- ❖ Each chromosome = condition part of a rule
- ❖ Used fixed eight bits for each element [low,high]
- ❖ String of bits mapped to interval [0.0,1.0]

# *Fitness Evaluation*

- ❖ Rule R matched for feature vector  $x^j$  if hypersphere centered at  $x^j$  and radius  $v$  intercepts hyperrectangle defined by points  $(low_1, high_1) \dots (low_n, high_n)$  of rule.
- ❖ Volume of R:

$$volume(R) = \prod_{i=1}^n (high_i - low_i)$$

# *Raw Fitness and Sequential Niching*

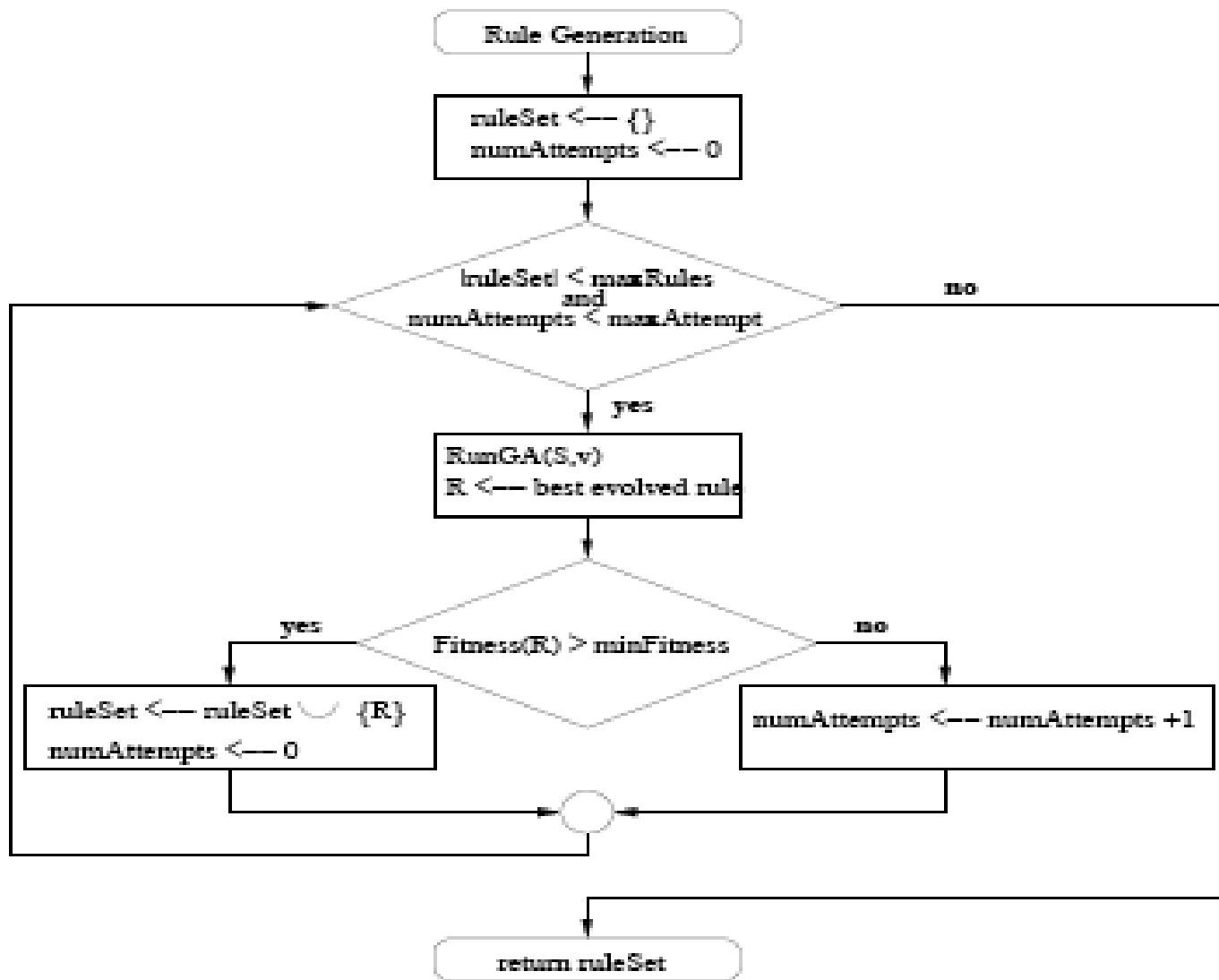
```
raw_fitnessR = volume(R) - C · num_elements(R)
```

```
fitnessR ← raw_fitnessR
```

```
for each Rj ∈ ruleSet do
```

```
    fitnessR ← raw_fitnessR - volume(R ∩ Rj)
```

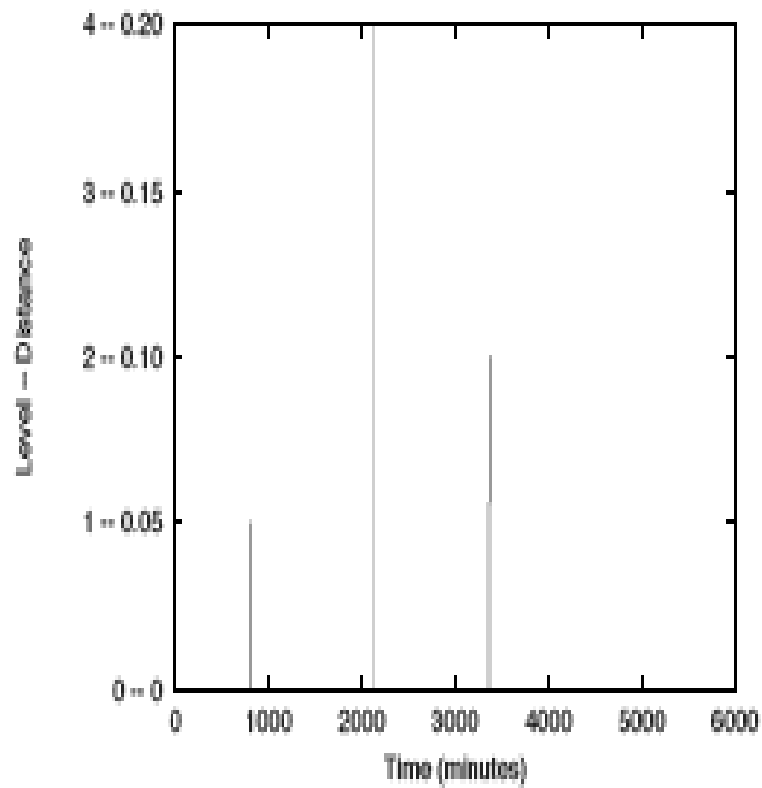
```
end-For
```



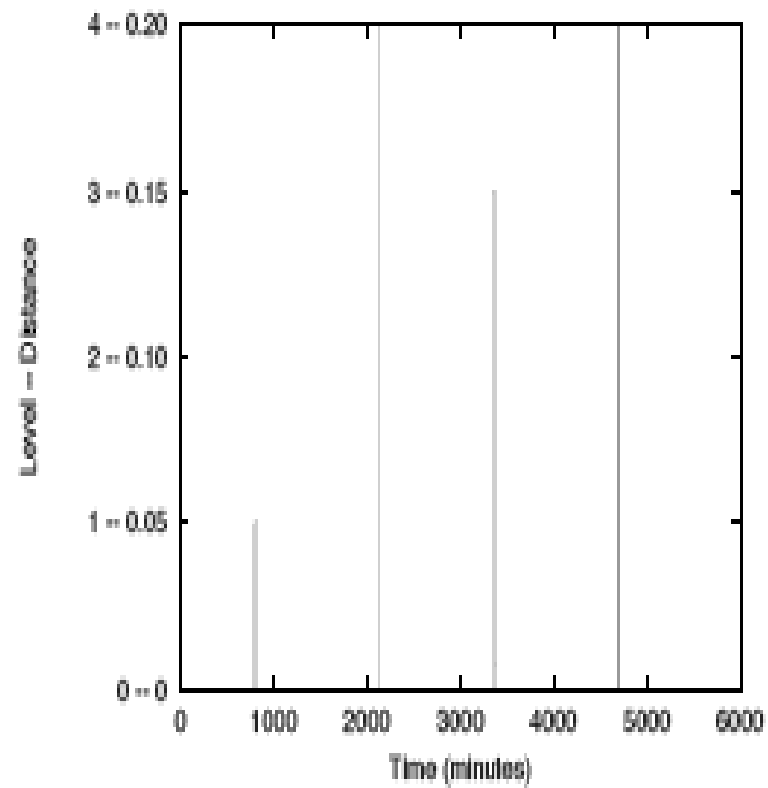


# *Experiments*

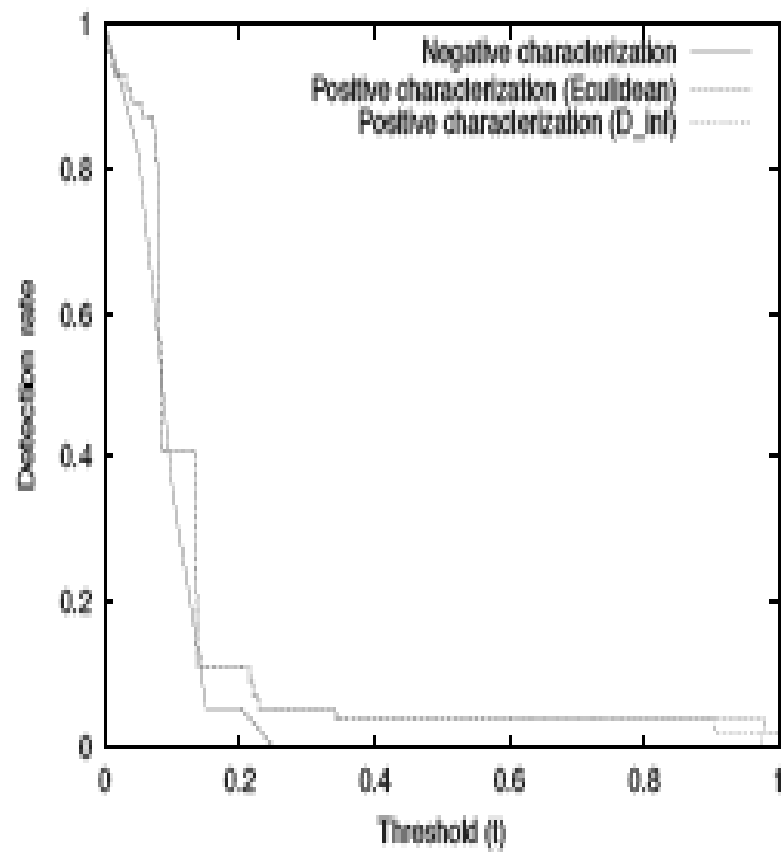
- ❖ GA params:
  - Population size 100
  - Number of generations 1500
  - Mutation rate 0.2
  - Crossover rate 1.0
  - Sensitivity 1.0
- ❖ Variability: 0.05, 0.1, 0.15 and 0.2
- ❖ Run each process 10 times against test, report average results



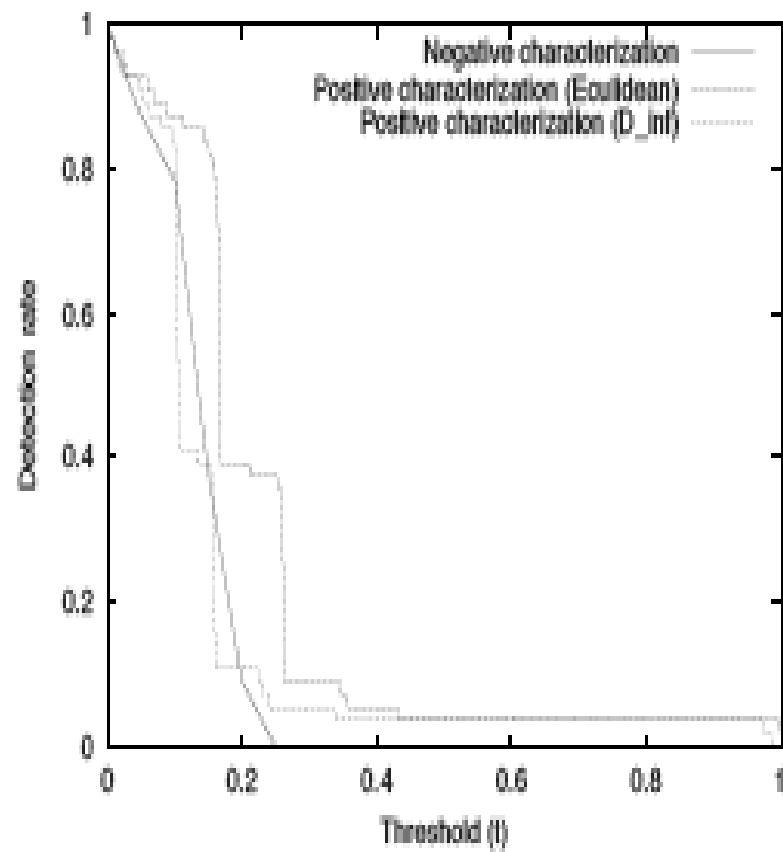
(a)  $ws = 1$



(b)  $ws = 3$



(a)  $ws = 1$



(b)  $ws = 3$

Table 4: Best true positive rates for the different techniques with a maximum false alarm rate of 1%.

Detection Technique	Window size 1	Window size 3
Positive Characterization (Euclidean)	92.8%	96.4%
Positive Characterization ( $D_\infty$ )	92.8%	92.8%
Negative Characterization	82.1%	87.5%

# *Tradeoffs*

- ❖ NC more efficient in space than PC
- ❖ Tradeoff between compactness and accuracy (but similar)
  - NC uses only 10% of space
  - Detects 4 vs. PC 5 attacks

# *Future Work*

- ❖ Different strategies for covering non-self space (e.g. hyperspheres)
- ❖ Apply to different IDS data
- ❖ Development of new algorithms to generate non-self covering rules

# *Criticism*

- ❖ How does difference between PC/NC scale for number of attacks?
- ❖ How does approach scale to larger data sets?
- ❖ How does approach compare to best alternative approach?

# *References*

- ❖ D. Dasgupta and F. Gonzalez, “An Immunity-Based Technique to Characterize Intrusions in Computer Networks”, *IEEE Transactions on Evolutionary Computing*, 6(3), pages 1081-1088, June 2002
- ❖ MIT Lincoln Lab Intrusion Detection Evaluation, <http://www.ll.mit.edu/IST/ideval/>