



# On the Effectiveness of Distributed Worm Monitoring

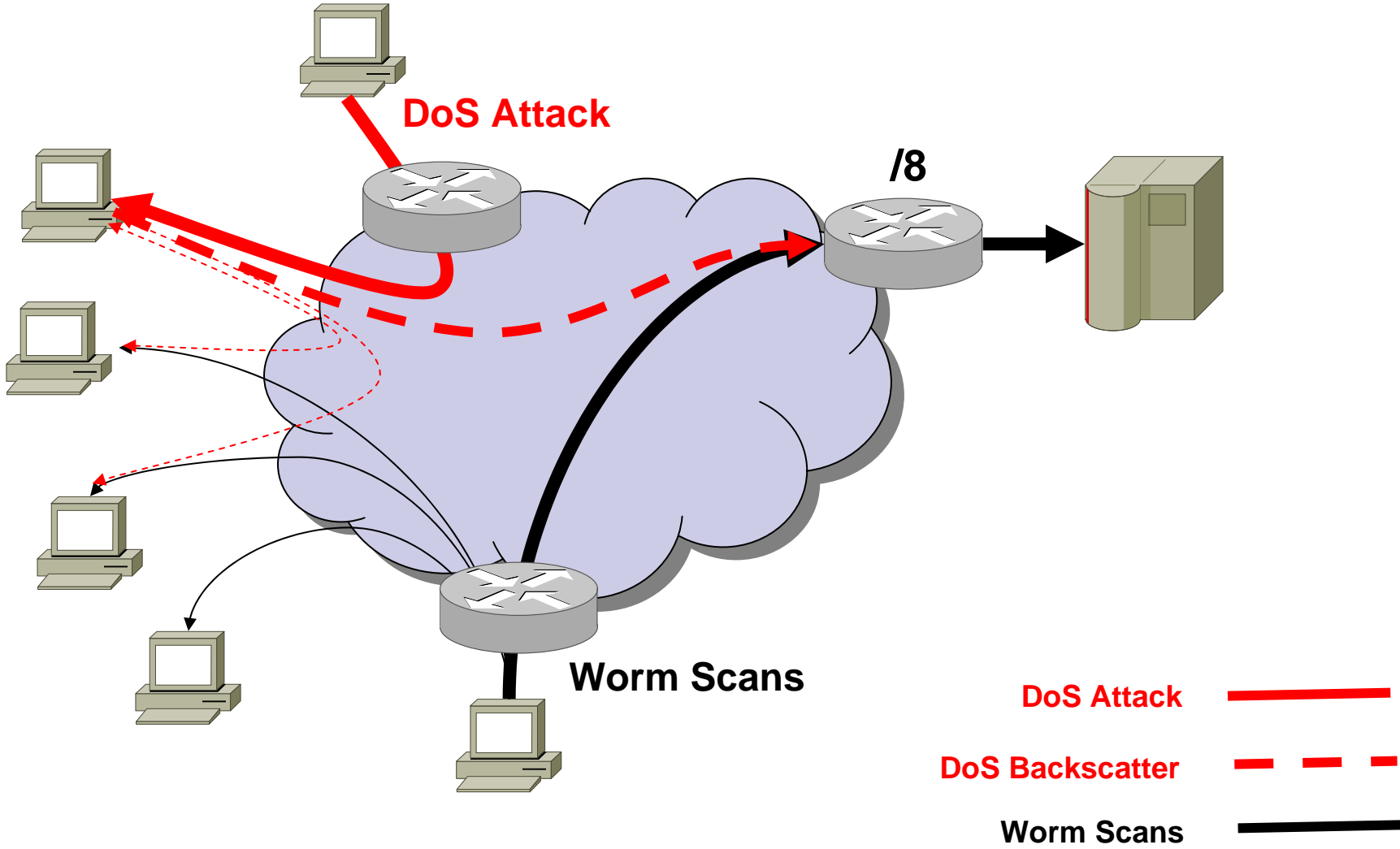
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# Monitoring Internet Threats

- Threat monitoring techniques:
  - Intrusion detection systems monitoring active networks
  - **Monitoring routable unused IP space** [ Moore et al, 2002 ]
- Monitoring unused address space is attractive
  - No legitimate traffic
  - Forensic analysis and early warning
- CAIDA deployed the first /8 telescope

# Single Monitor Case

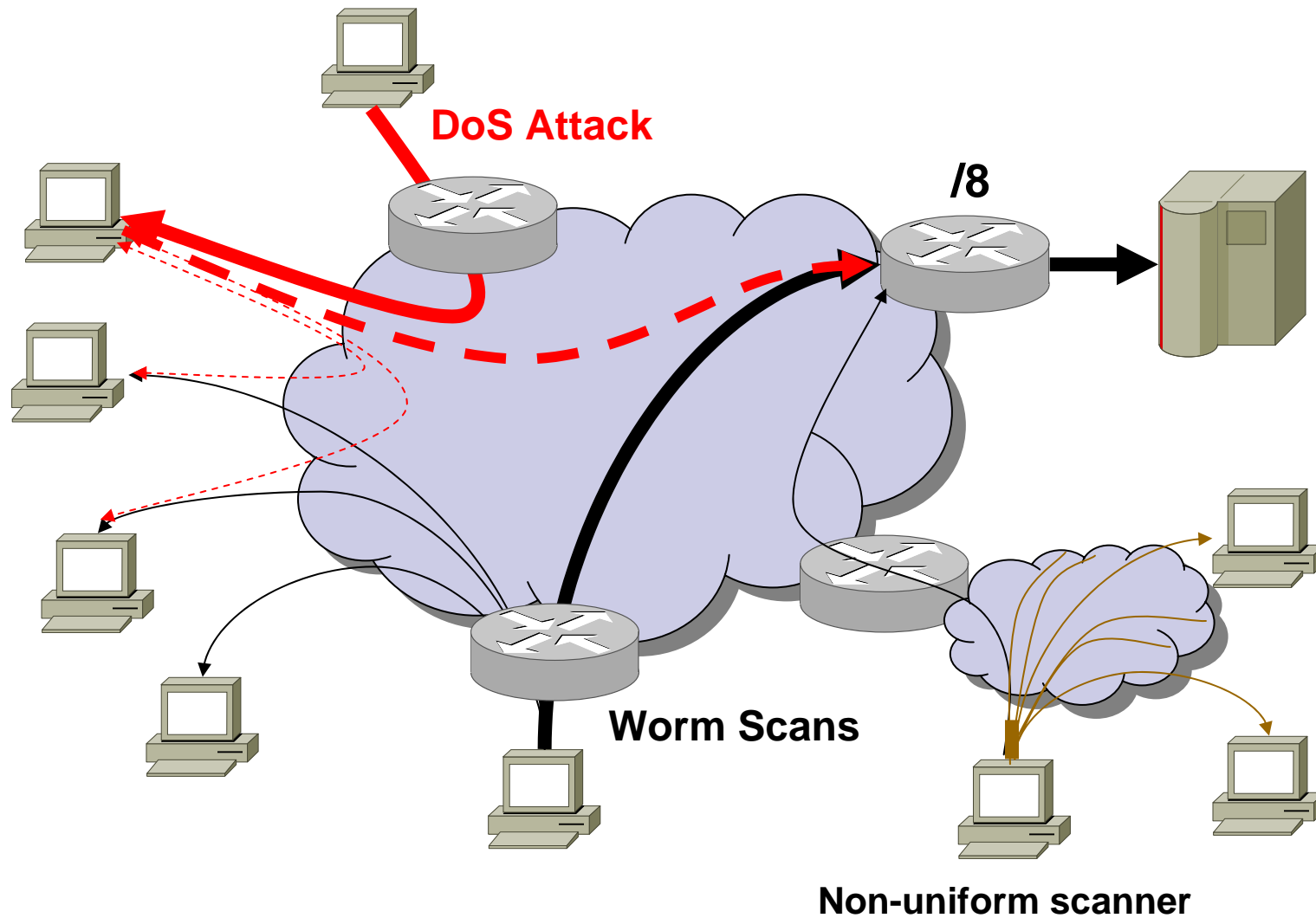




# Size Matters!

- Size of the monitor is an important factor in providing an accurate view of a worm breakout [Moore et al, 2002]
- But there are several other factors yet to be explored

# Single monitor view is too limited





# Goals

- Provide a model to evaluate the performance of distributed monitoring systems in terms of:
  - Number of monitors?
  - Sizes of monitors and the overall IP space requirements?
- Provide guidelines for better design and monitor deployment practices.



# Outline

- Problem and Motivation
- A Worm Propagation Model
  - Population Distribution
  - Extended worm model
- Distributed Worm Monitoring
  - Distributed Telescope Model
  - Design parameters
- Summary

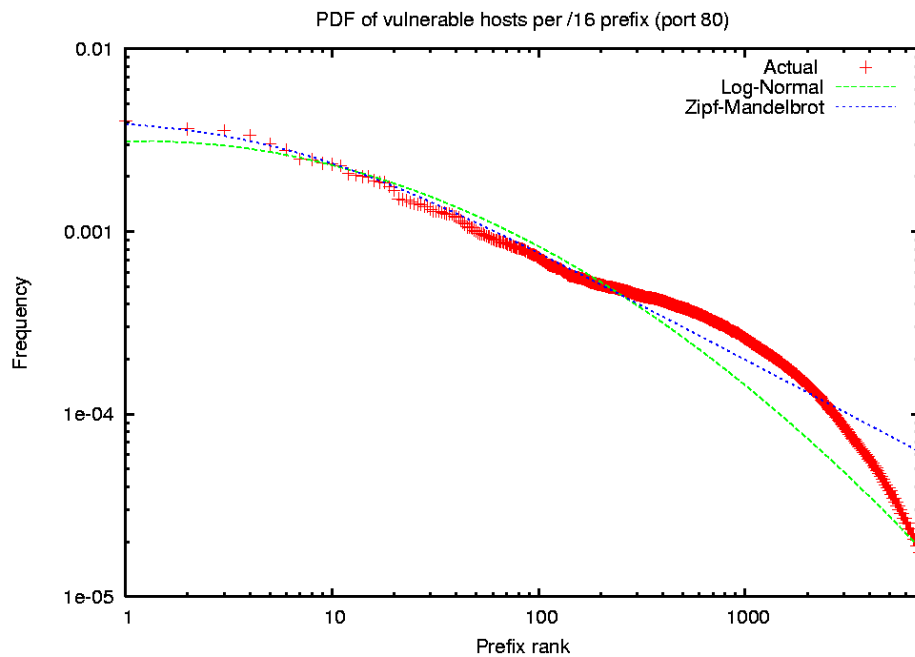


# Why another worm model?

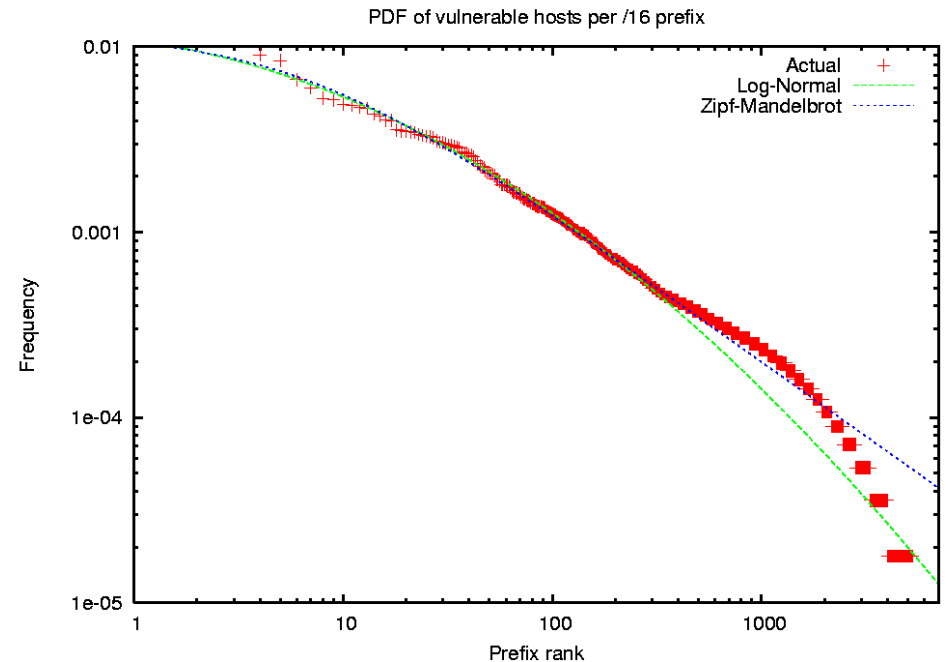
- Previous worm models assumed that the vulnerable population is **uniformly** distributed over the whole IP space.
- Sources of non-uniformity in population distribution
  - Un-allocated address space
  - Highly-clustered allocated space
  - Usage of the allocated space



# Population distribution



DShield dataset



CAIDA's dataset (Witty Worm)

**The distribution of Vulnerable population over the IP space is far from uniform  
Best fits a Log-normal distribution**



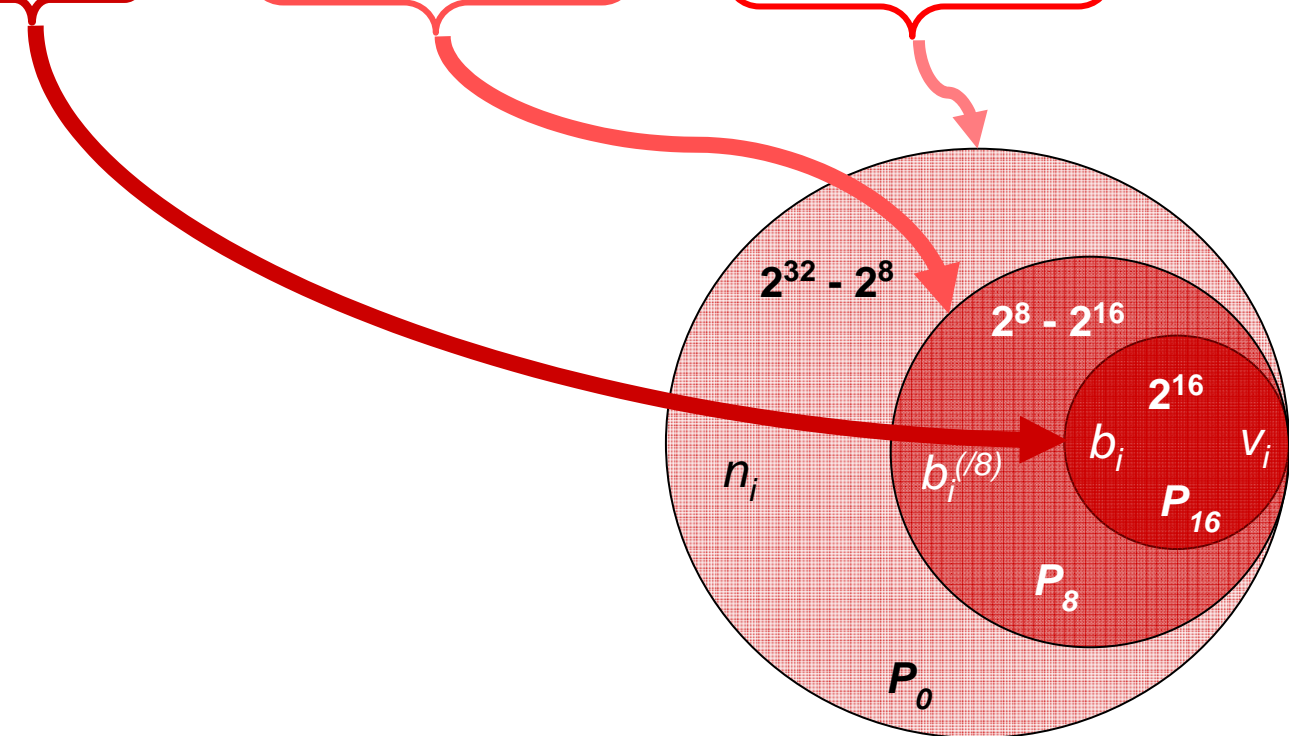
# Extended Worm Propagation Model

- Worm propagation models **must** incorporate population density distribution.
- Especially Non-uniform scanning worms:
  - Probability of scanning a host depends on its **location** relative to the infected scanner

# Non-uniform worm propagation model

- Expected number of scans per /16 subnet

$$k_i^j = \underbrace{p_{16} s b_i^j}_{\text{P}_{16}} + \underbrace{p_8 s b_i^{(j/8)} \frac{2^{16}}{2^{24}}}_{\text{P}_8} + \underbrace{p_0 s n_i \frac{2^{16}}{2^{32}}}_{\text{P}_0}$$



# Non-uniform worm propagation model

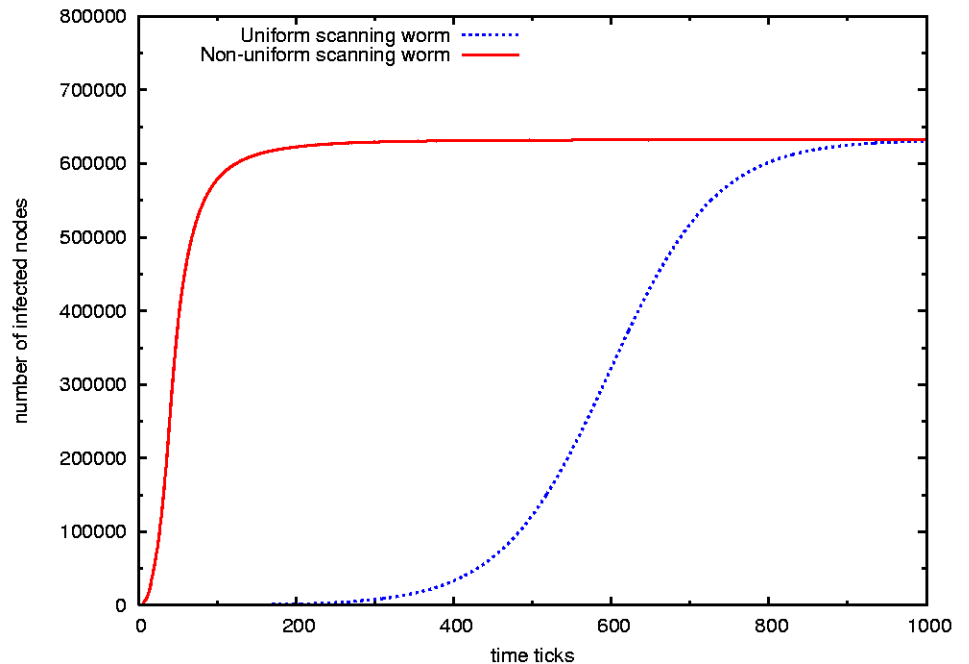
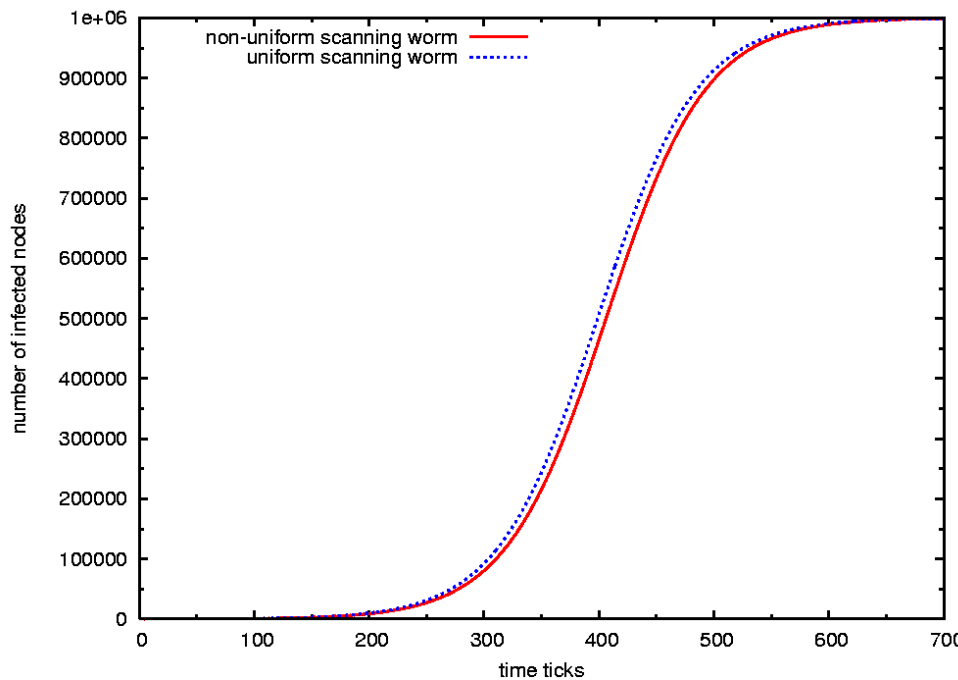
- The expected number of infected hosts per /16 subnet (AAWP Model [*Chen et al, 2003*])

$$b_{i+1}^j = b_i^j + \underbrace{(v_i - b_i^j)}_{\substack{\text{Vulnerable non-} \\ \text{infected} \\ \text{hosts}}} \left[ 1 - \left( 1 - \frac{1}{2^{16}} \right)^{k_i^j} \right]$$

- The expected total infection

$$n_{i+1} = \sum_{j=1}^{2^{16}} b_i^j$$

# Impact of population distribution



Number of Infected hosts vs time, for a Nimda-like worm  
 $s = 100$  scans/time tick,  $P_{16} = 0.5$ ,  $P_8 = 0.25$ ,  $P_0 = 0.25$

$N = 10^6$  hosts uniformly distributed  
Over the IP space

$N = 620,000$  hosts extracted from  
DShield data set



# Outline

- Problem and Motivation
- Better Worm Model
  - Population Distribution
  - Extended worm model
- Distributed Worm Monitoring
  - Distributed monitoring system model
  - Design parameters
- Summary



# Using the Model---

## Distributed Monitoring:

- What do we want to evaluate?
  - System detection time: the time it takes the monitoring system to detect (with particular confidence) a new scanner.

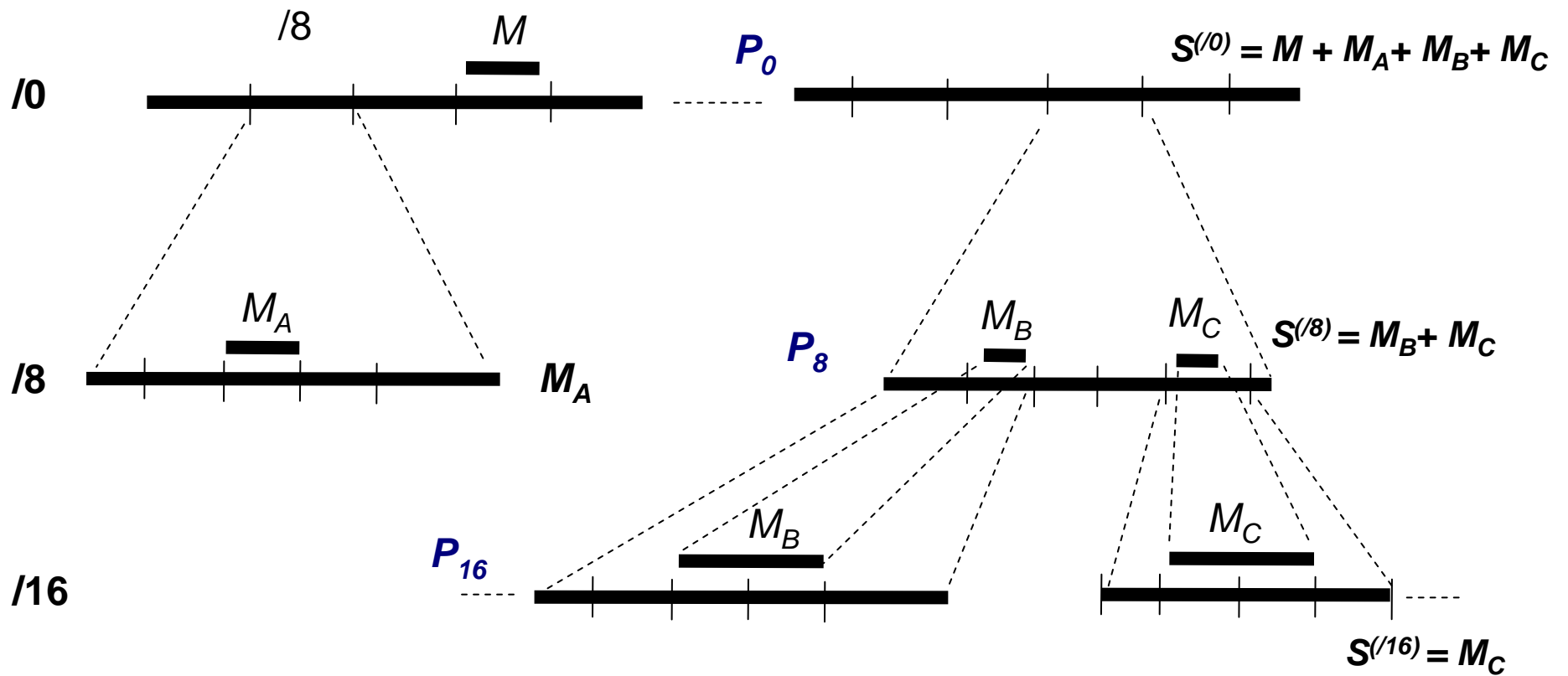


# Assumptions

- Single scan detection
- Information sharing and aggregation infrastructure among all monitors.



# Monitors Logical Hierarchy



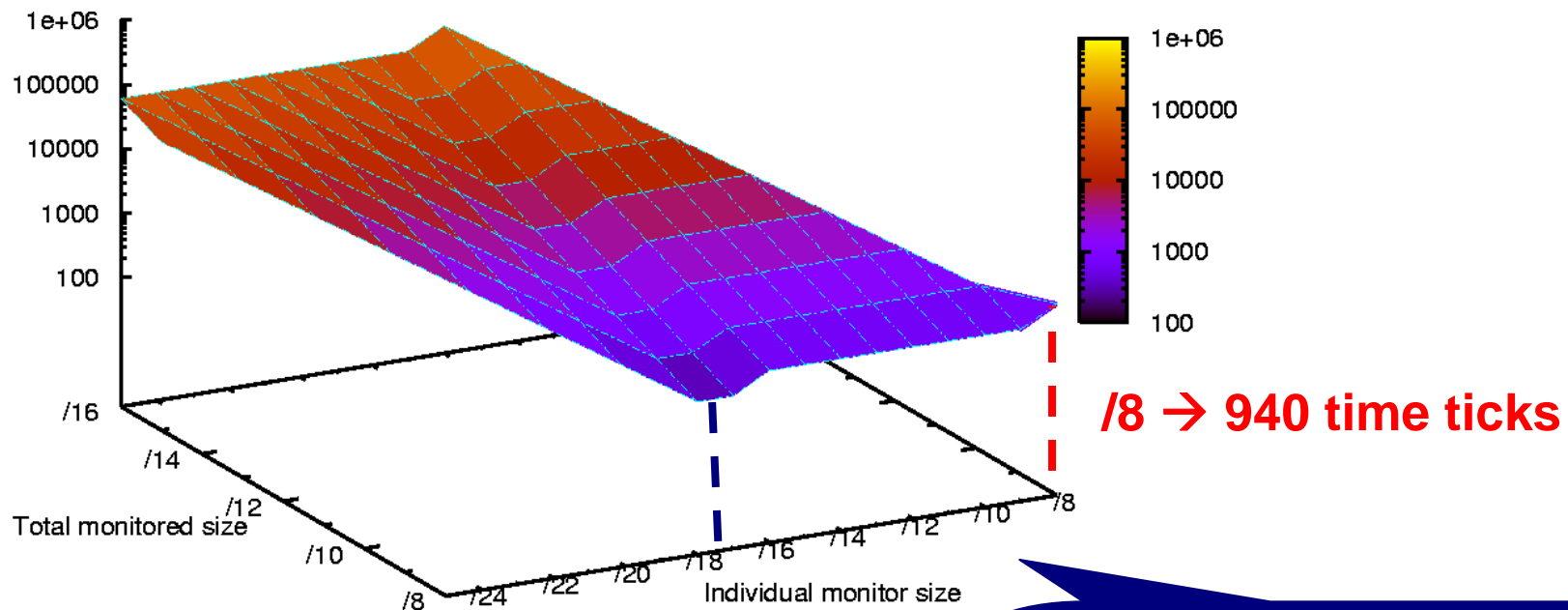


# Evaluation

- Nimda-like scanner
- Three Monitor deployment scenarios:
  - Random monitor deployment
  - Full knowledge of population distribution
  - Partial population knowledge

# Evaluation (Random monitor placement)

Detection Time (ticks)



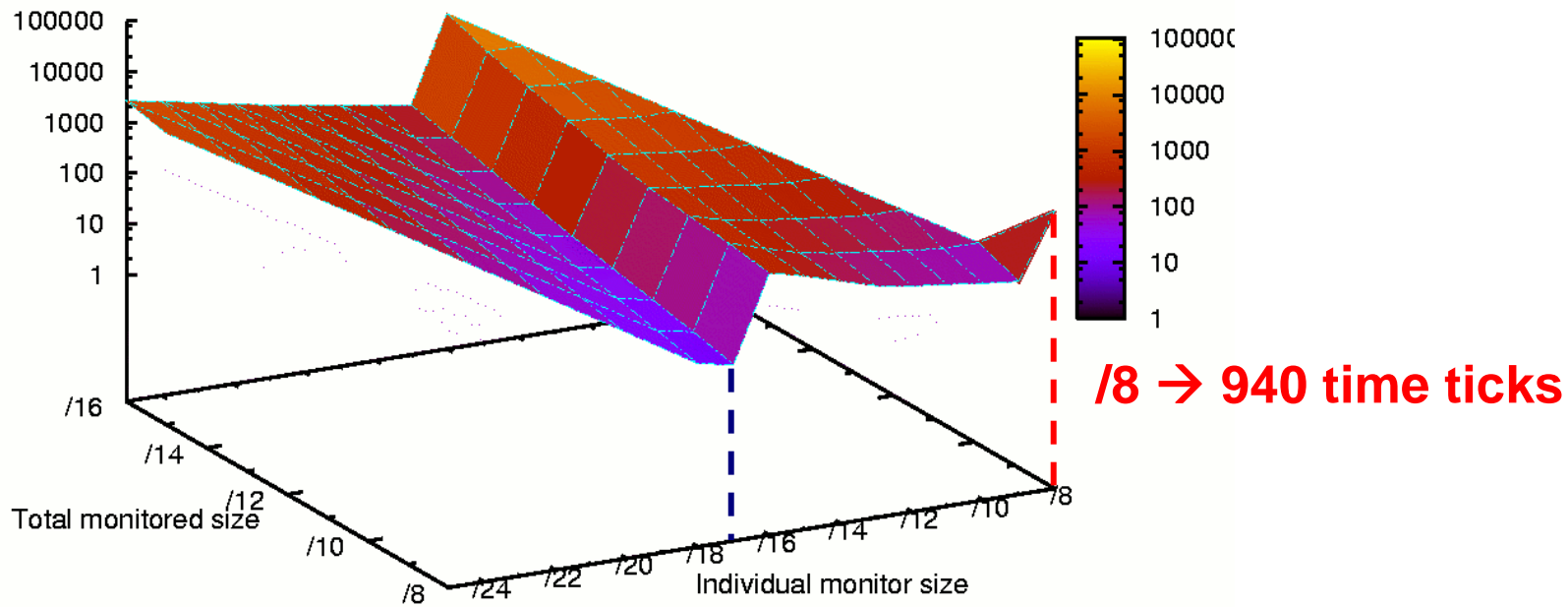
**512 /17 → 230 time ticks**

**with only 40 hosts per /16,  
7100 more scans will  
cause infecting 2 victims  
before being detected**

Random Monitor placement  
 $P_r = 0.999$ ,  $s = 10$  scans/time tick  
Nimda-like scanning

# Evaluation ( Full vulnerable distribution knowledge)

Detection Time (ticks)

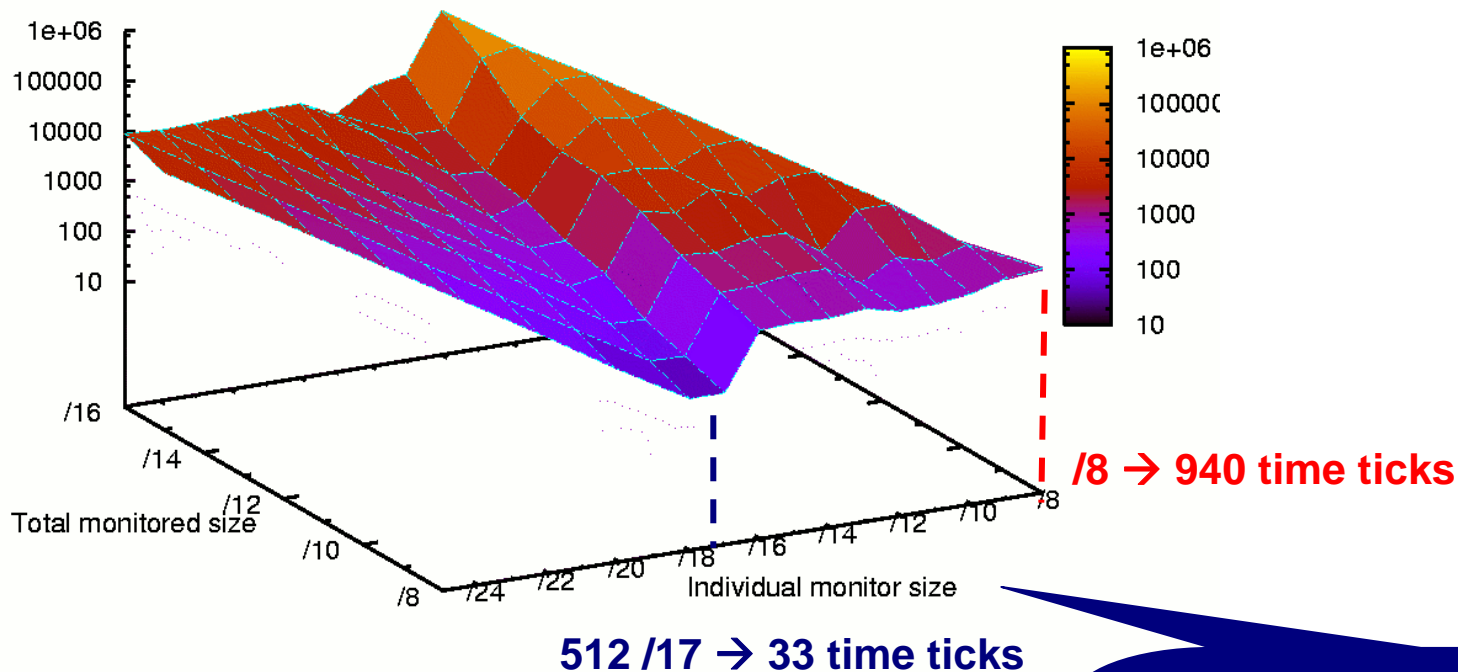


**512 /17 → 9 time ticks**

Monitors deployed in top populated prefixes

# Evaluation (Partial Knowledge )

Detection Time (ticks)



Monitors deployed **randomly** over the 5000 most populated /16 prefixes (contain 90% of the vulnerable population)

**Example: 512 monitors with 2048 IP addresses/monitor → 160 time ticks**



# Practical Considerations

- Monitors will be deployed at different administrative domains.
- How many domains are needed to deploy these 512 monitors?
- Mapping the monitors to AS space, only **130 AS's** among the top address space owners are required to achieve detection time of 160 time ticks



# Summary

- Population distribution has a profound impact on worm propagation speed.
- Distributed Monitoring provides an improved detection time (three times faster than a single monitor of equivalent size).
- Even partial knowledge of the population distribution can improve detection time by roughly 30 times.
- Effective distributed monitoring is possible with cooperation among top address space owners.



# Questions?