

# **Artificial Immune Systems and Data Mining: Bridging the Gap with Scalability and Improved Learning**

**Olfa Nasraoui, Fabio González  
Cesar Cardona, Dipankar Dasgupta  
The University of Memphis**

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# Inspired by Nature...

- living organisms exhibit extremely sophisticated learning and processing abilities that allow them to survive and proliferate
- nature has always served as inspiration for several scientific and technological developments, exp: Neural Networks, Evolutionary Computation
- immune system: parallel and distributed adaptive system w/ tremendous potential in many intelligent computing applications.

# What is the Immune System?

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- **Protects** our bodies from foreign pathogens (viruses/bacteria)
- **Innate** Immune System (initial, limited, ex: skin, tears, ...etc)
- **Acquired** Immune System (**Learns** how to respond to NEW threats adaptively)
- **Primary** immune response
  - First response to invading pathogens
- **Secondary** immune response
  - Encountering similar pathogen a second time
  - **Remember** past encounters
  - Faster and stronger response than primary response

# Points of Strength of The Immune System

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- **Recognition** (Anomaly detection, Noise tolerance)
- **Robustness** (Noise tolerance)
- **Feature extraction**
- **Diversity** (can face an entire repertoire of foreign invaders)
- **Reinforcement learning**
- **Memory** (remembers past encounters: basis for vaccine)
- **Distributed** Detection (no single central system)
- **Multi-layered** (defense mechanisms at multiple levels)
- **Adaptive** (Self-regulated)

# Major Players: B-Cells

- Through a process of recognition and stimulation, B-Cells will clone and mutate to produce a diverse set of antibodies adapted to different antigens
- B-Cells secrete antibodies w/ paratopes that can bind to specific antigens (epitopes) and destroy their host invading agent through a KILL, SUICIDE, or INGEST signal.
- ❖ B-Cells antibody paratopes also can bind to antibody idiotypes on other B-Cells, hence sending a STIMULATE or SUPPRESS signal → hence the **Network** → **Memory**

# Requirements for Clustering Data Streams (Barbara, 02)

## ■ Compactness of representation

- Network of B-cells: each cell can recognize several antigens
- B-cells compressed into clusters/sub-networks

## ■ Fast incremental processing of new data points

- New antigen influences only activated sub-network
- Activated cells updated incrementally
- Proposed approach learns in **1 pass**.

## ■ Clear and fast identification of “outliers”

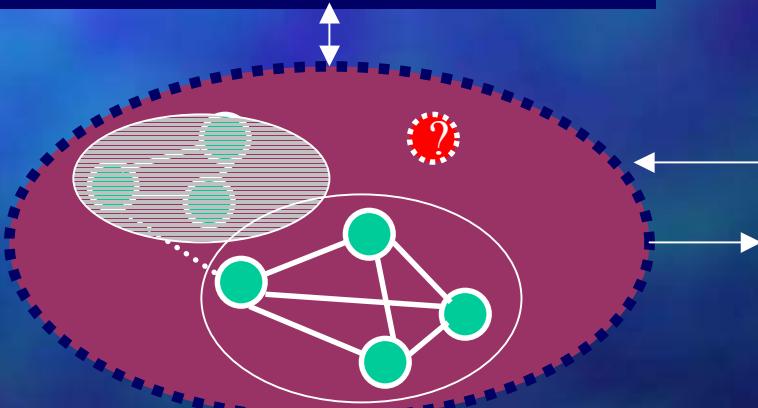
- New antigen that does not activate any subnetwork is a potential outlier → create new B-cell to recognize it
- This new B-cell could grow into a subnetwork (if it is stimulated by a new trend) or die/move to disk (if outlier)

# General Architecture

Evolving data →

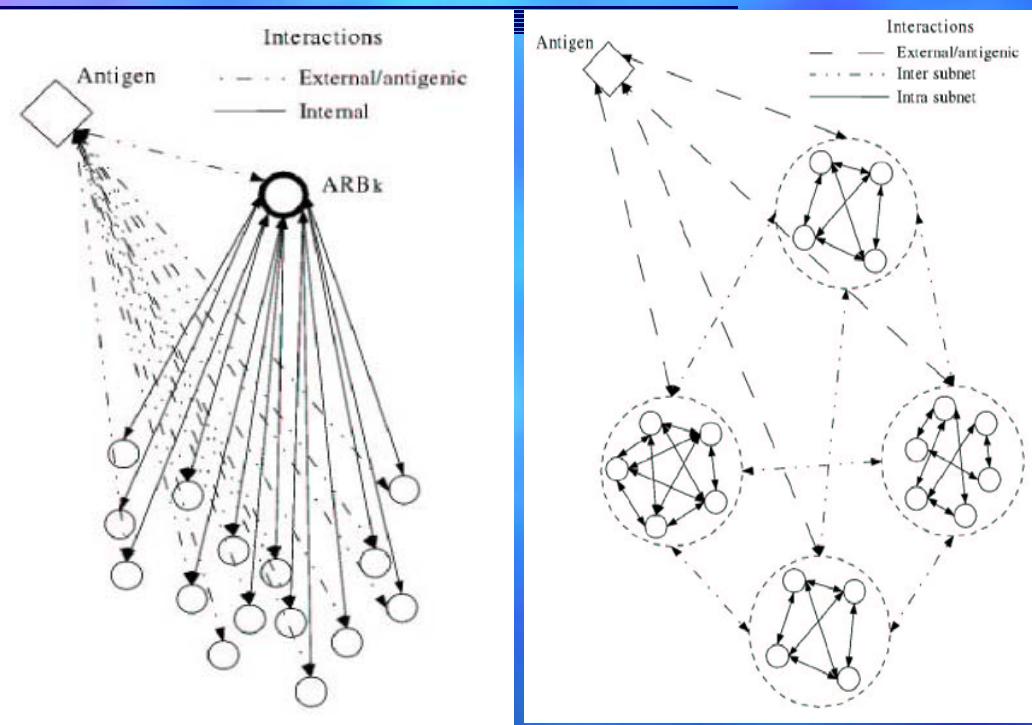
1-Pass Adaptive  
Immune  
Learning

Evolving Immune  
Network  
(compressed into  
subnetworks)

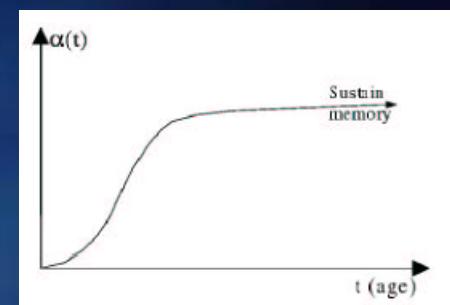
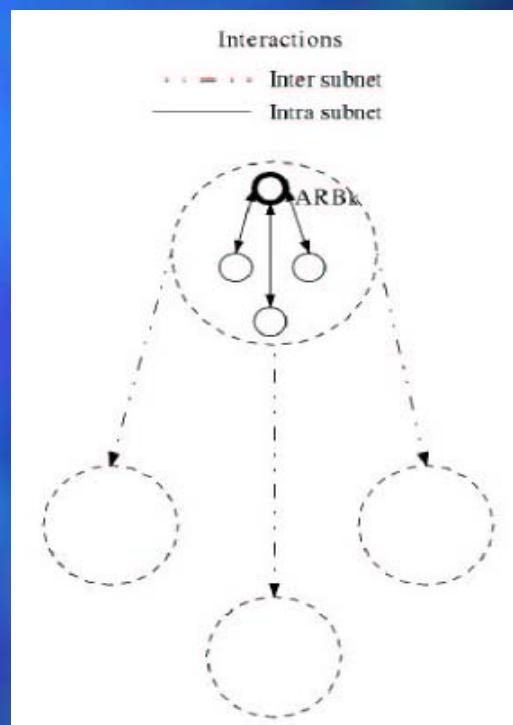


Immune network  
information system  
*Stimulation (competition  
& memory)*  
*Age (old vs. new)*  
*Outliers (based on  
activation)*

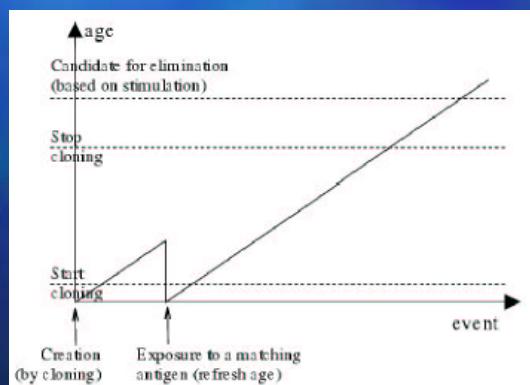
# Internal and External Immune Interactions: Before & After



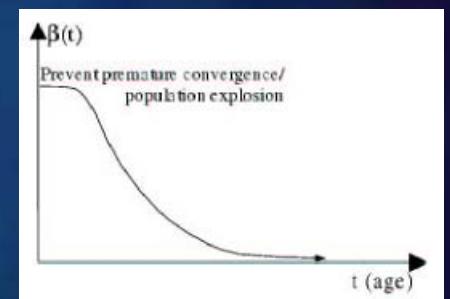
# Internal Immune Interactions



## Internal Stimulation

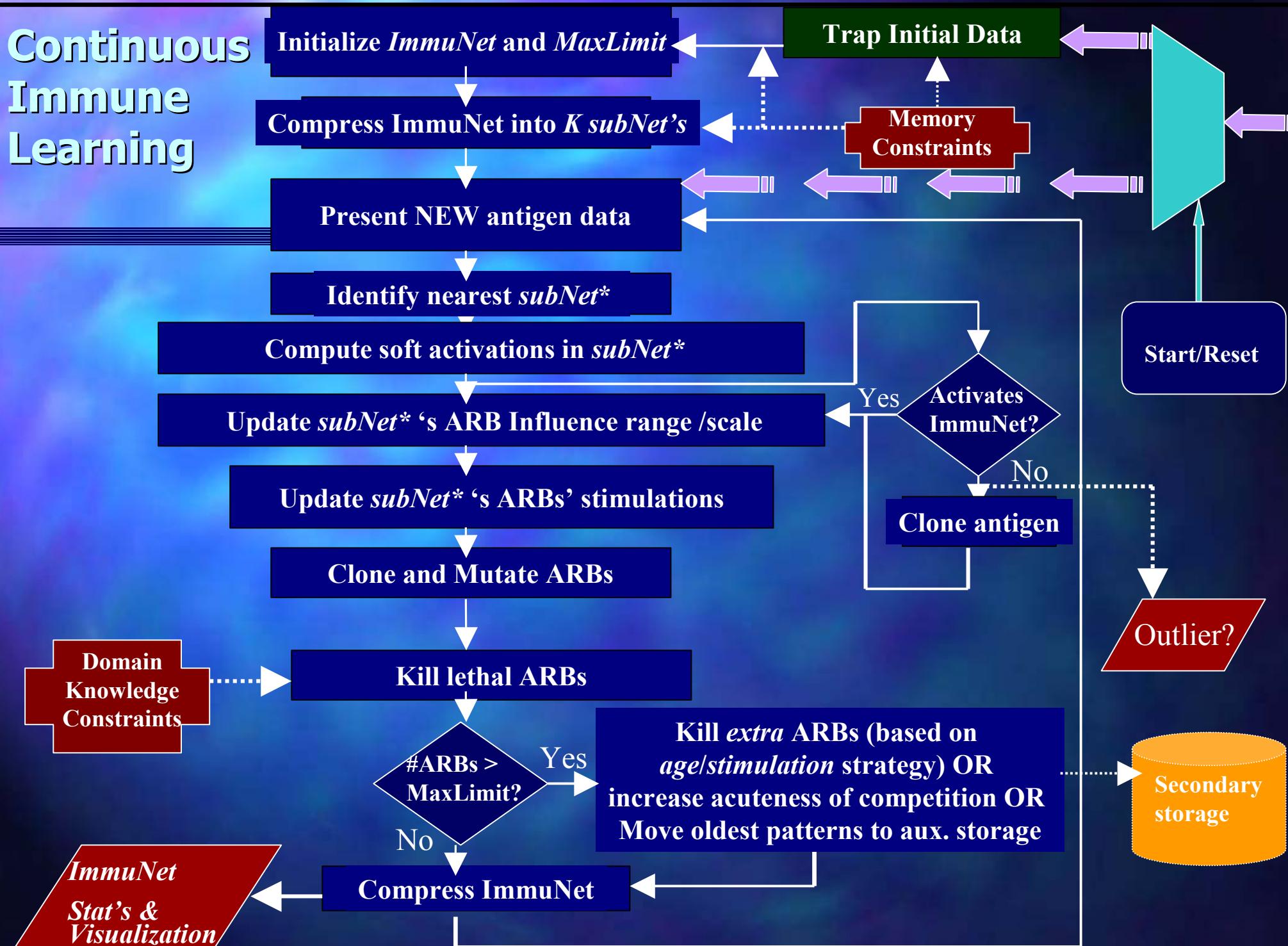


## Lifeline of B-cell



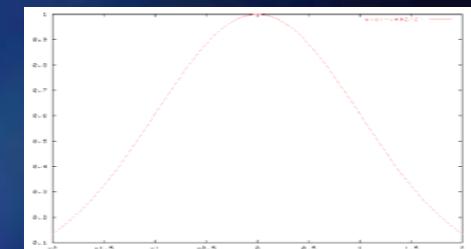
## External Stimulation

# Continuous Immune Learning



# Model for Artificial Immune Cell

- Antigens represent data and the B-Cells represent clusters or patterns to be learned/extracted
- ARB/B-cell object:
  - Represents not just a single item, but a fuzzy set
  - Better Approximate Reasoning abilities
  - Each ARB is allowed to have its own **zone of influence** with size/scale:  $\sigma_i$
  - ARBs dynamically adapt their influence zones/hence stimulation level in a strife for survival.
  - Membership function dynamically **adapts** to data
  - **Outliers** are easily detected through weak activations
  - No more dependence on hard threshold-cuts to establish network
  - Can include most probabilistic and possibilistic models of uncertainty
  - Flexible for different attributes types (numerical, categorical, ...etc)



# Immune Based Learning of Web profiles

- The Web server plays the role of the human body, and the incoming requests play the role of antigens that need to be detected
- The input data is similar to web log data (a record of all files/URLs accessed by users on a Web site)
- The data is pre-processed to produce session lists:
  - A session list  $S_i$  for user # $i$  is a list of *URLs visited by same user*
  - In discovery mode, a session is fed to the learning system as soon as it is available
- B-cell $_i$ :  $i^{\text{th}}$  candidate profile:
  - List of URLs
  - Historic Evidence/Support: List of supporting cumulative conditional probabilities ( $\text{URL}_k, \text{prob}(\text{URL}_k)$ ) with  $\text{prob}(\text{URL}_k) = \text{prob}(\text{URL}_k \mid \text{B-cell}_i)$
  - Each profile has its own influence zone defined by  $\sigma_i$