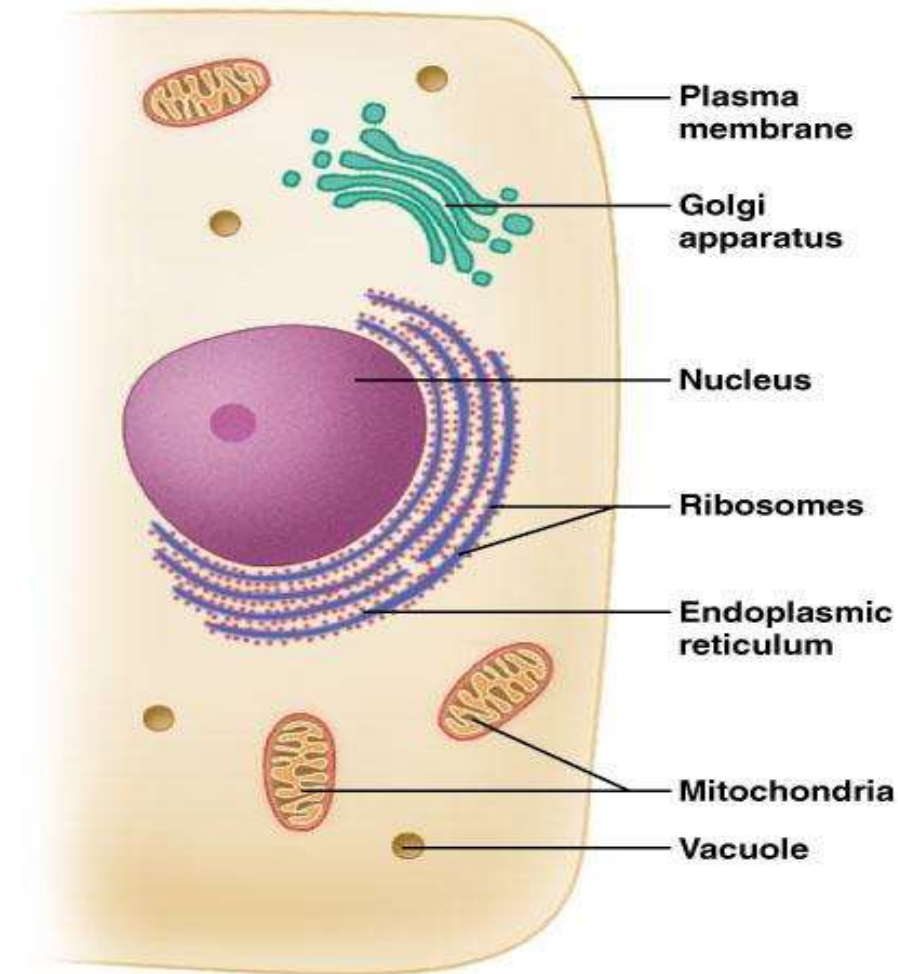


# UNIT 5

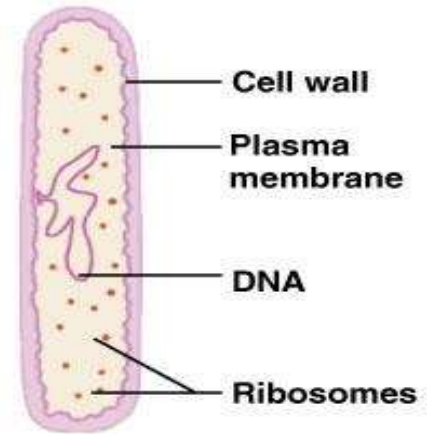
## ImmunoBiology

- Elements of the immune system,
- Types of the immune response
- Active and passive immunity
- Immunoinformatics
- epitope prediction tools

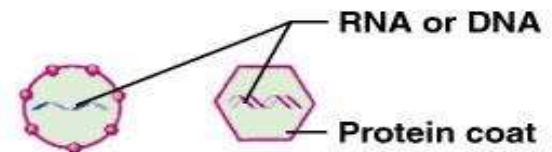
# Eukaryotic Cells, Bacteria, and Viruses



**(a) Eukaryotic cell**



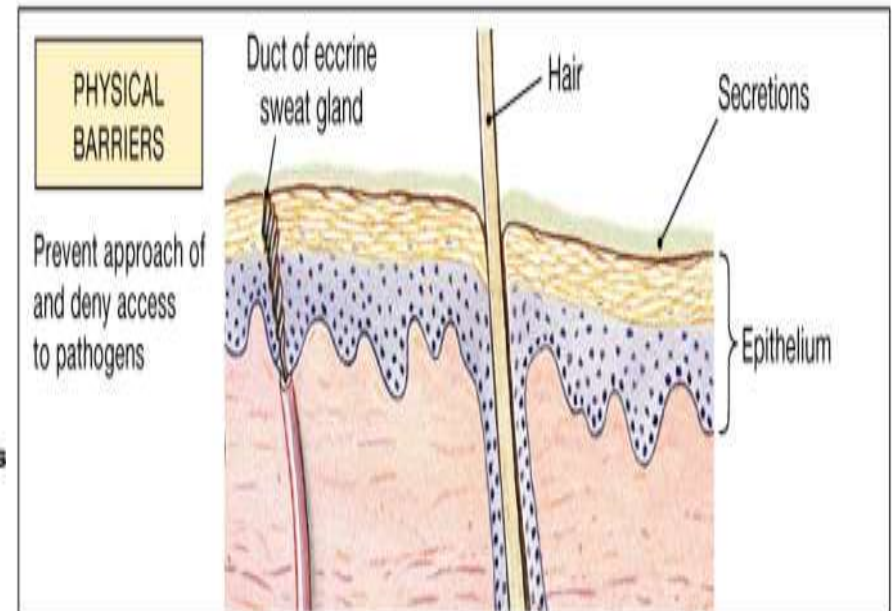
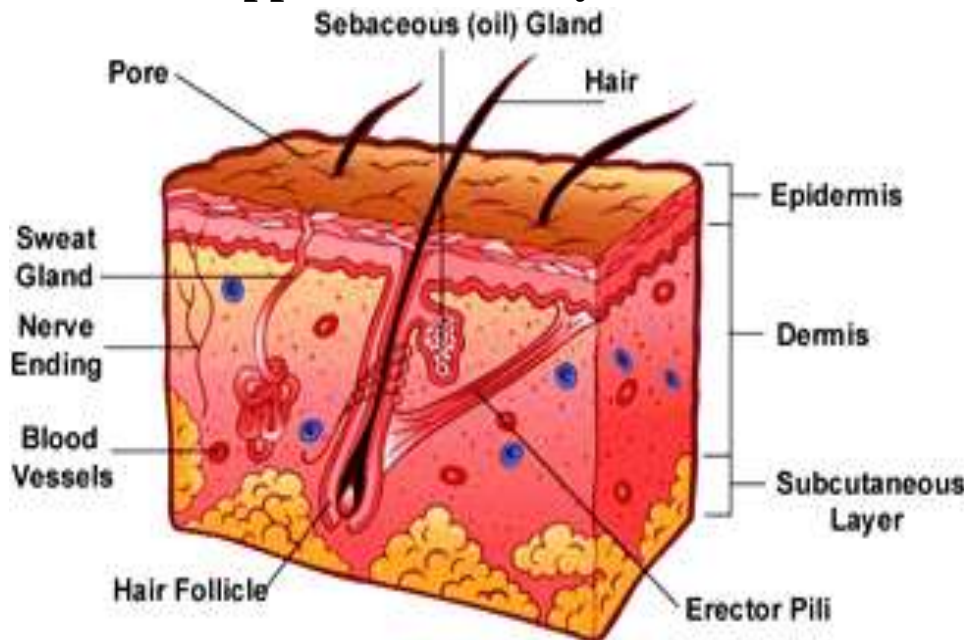
**(b) Bacterium**



**(c) Viruses**

# Body Defenses: Overview

- **Physical barriers: skin & epithelial linings & cilia**
- **Chemical: acids, mucous & lysozymes**
- **Immune defenses – internal**
  - **Innate, non-specific, immediate response (min/hrs)**
  - **Acquired – attack a specific pathogen (antigen)**
- **Steps in Immune defense**
  - **Detect invader/foreign cells**
  - **Communicate alarm & recruit immune cells**
  - **Suppress or destroy invader**

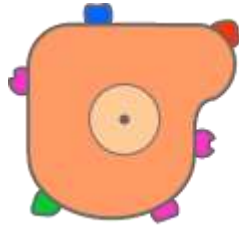


# Discrimination of self from non-self

“The success of the immune system depends on its ability to discriminate between foreign (nonself) and host (self) cells. Survival requires both the **ability** to mount a destructive immune response against **nonself** and the **inability** to mount a destructive response against **self**.”

# Markers of Self

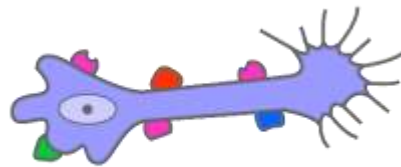
Epithelial  
cell



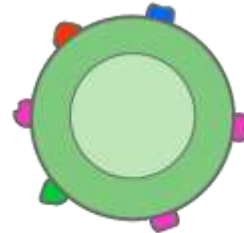
Muscle cell



Nerve  
cell



Leukocyte



At the heart of the immune response is the ability to **distinguish between “self” and “non-self.”**

- Every cell in your body carries the same set of distinctive surface proteins that distinguish you as “self.”
- Normally your immune cells do not attack your own body tissues, which all carry the same pattern of self-markers; rather, your immune system coexists peaceably with your other body cells in a state known as self-tolerance.

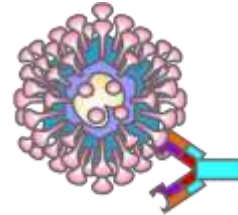
This set of unique markers on human cells is called the **major histocompatibility complex (MHC) proteins.**

# Markers of Non-self

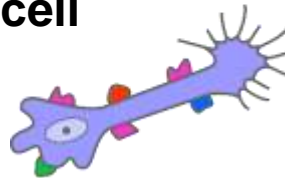
Bacteria



SARS virus



Non-self nerve cell



Non-self leukocyte



Epitope

Antigen

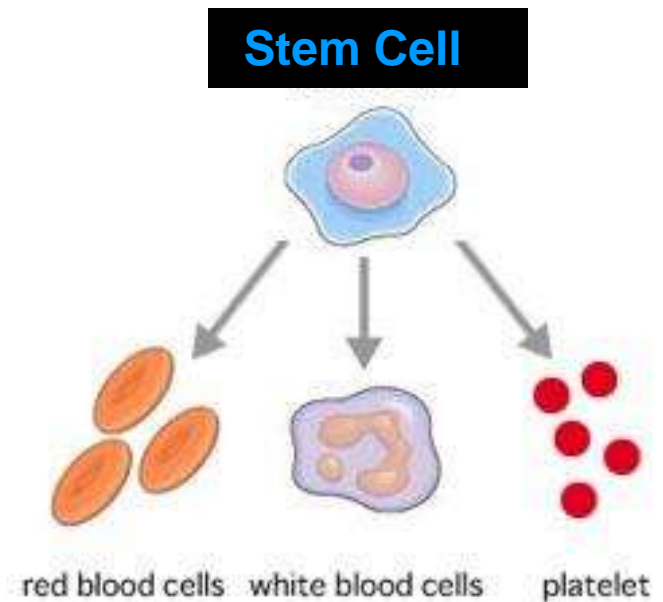
**Antigen** = any non-self substance

- Virus
- Bacteria
- Non-self cell (foreign cell)

**Epitope** = The distinctive markers on antigens that trigger an immune response

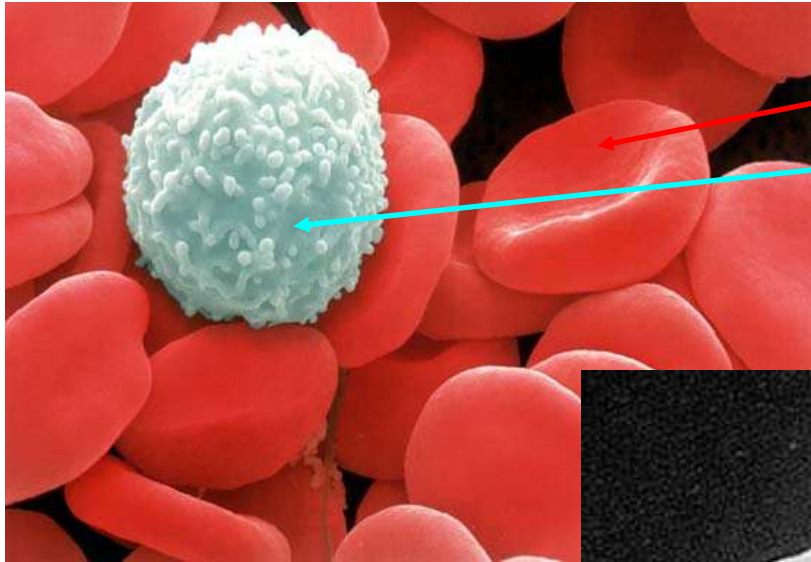
# Blood

- Blood is 55% liquid (plasma) and 45% cellular
- Cellular component of blood:
  - Red blood cells = carry oxygen
  - White blood cells = immune system
  - Platelets = clot blood
- All blood cells arise from a pluri-potent stem cell found in bone marrow





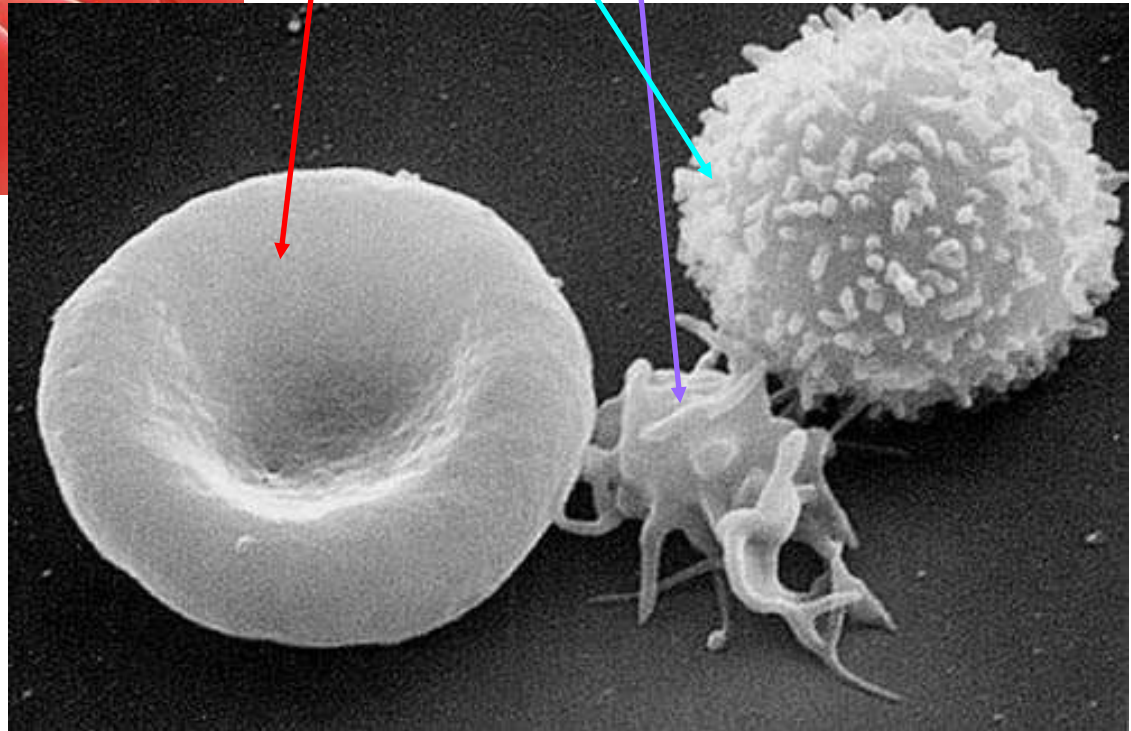
# Blood cells



**Red Blood cells**

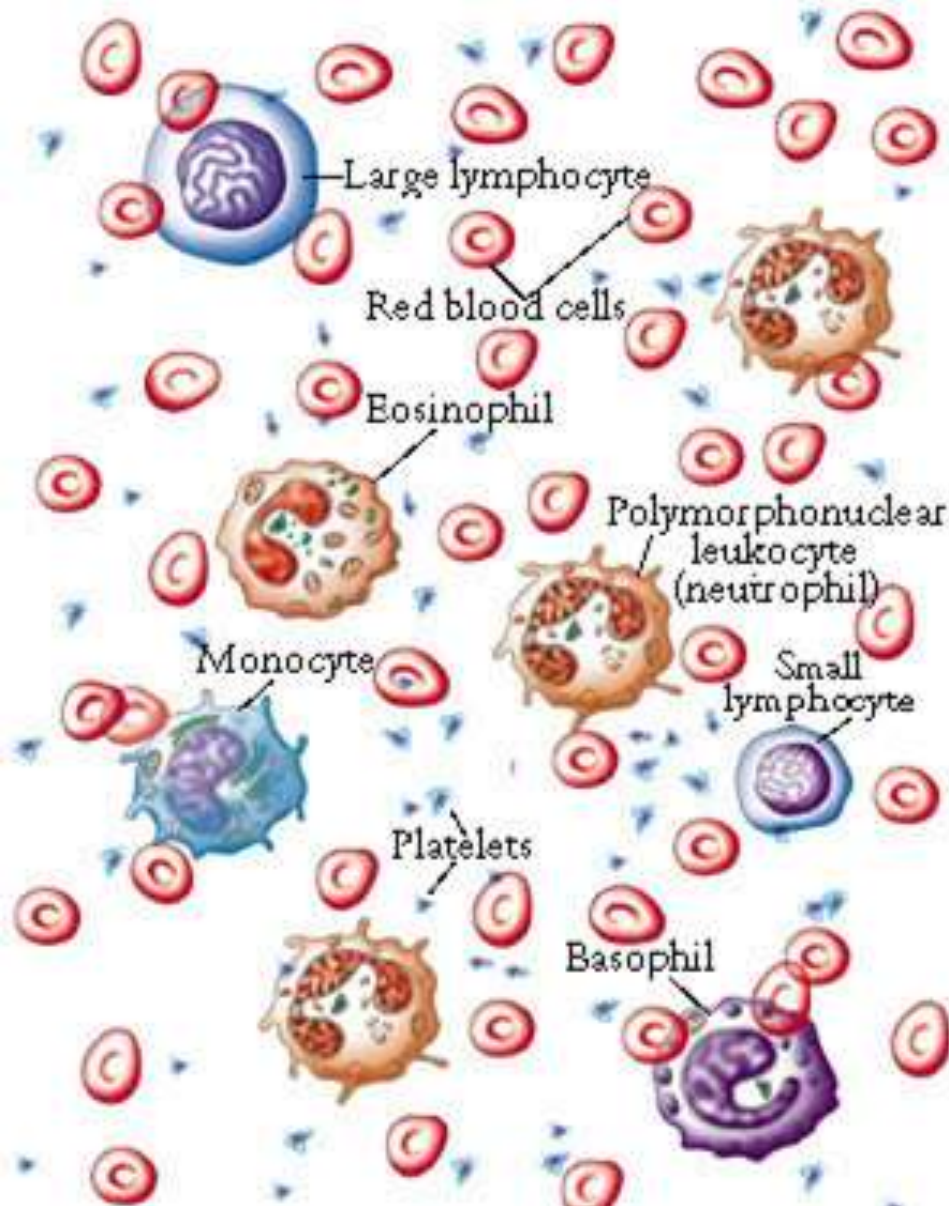
**White blood cells (immune cells)**

**Platelets**





# Leukocytes in the Blood



Red Blood Cells		$5.0 \times 10^6/\text{mm}^3$	
Platelets		$2.5 \times 10^5/\text{mm}^3$	
Leukocytes		$7.3 \times 10^3/\text{mm}^3$	
1	Neutrophil		50-70%
2	Lymphocyte		20-40%
3	Monocyte		1-6%
4	Eosinophil		1-3%
5	Basophil		<1%

# TYPES OF WHITE BLOOD CELLS:

## Leukocytes (white blood cells)

*agranular*

*granular*



Lymphocytes  
20-25%



Monocyte  
3-8%



Basophil  
0.5-1%



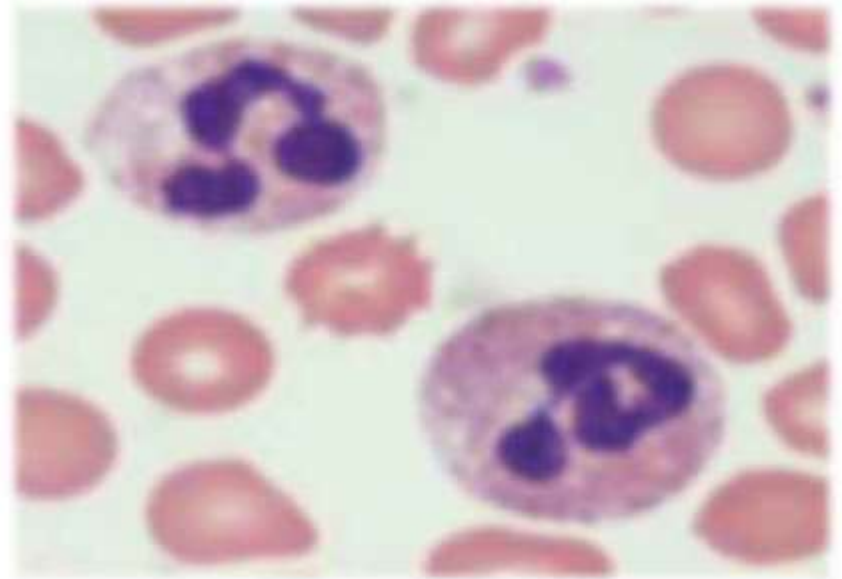
Neutrophil  
60-70%



Eosinophil  
2-4%

# Neutrophil

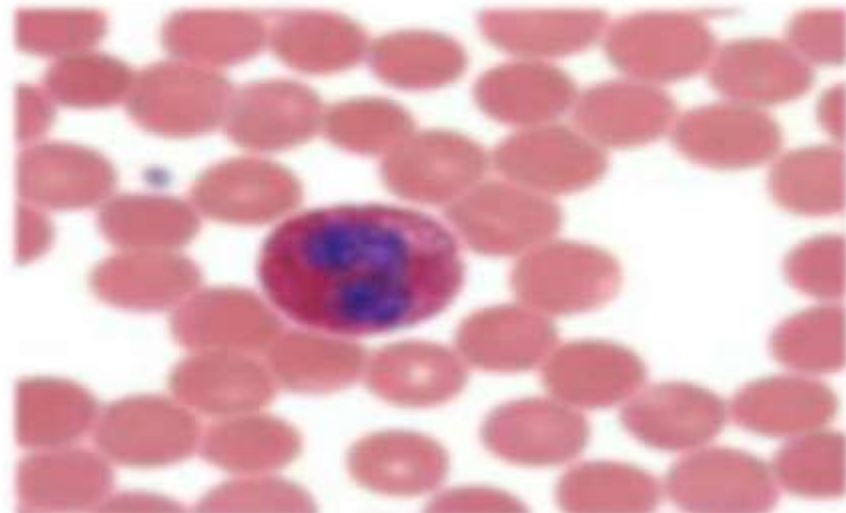
Neutrophil • 60% of WBC • Lifespan is 10 hours in blood • Seek out and destroy ingested bacteria in connective tissues • 100 billion manufactured daily





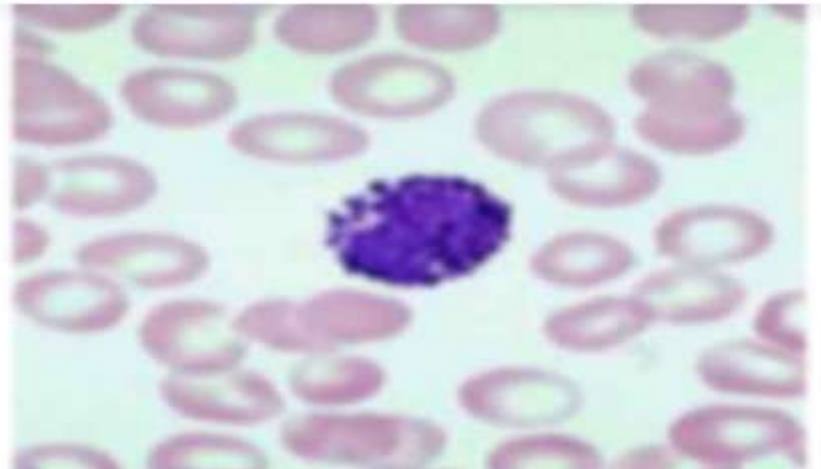
# Eosinophil

- 1-3 % of WBC
- Help control allergic reactions
- Release an enzyme histamine – a chemical released during allergic reactions



# Basophil

- Less than 1% of WBC
- Involved in allergic and inflammatory reactions
- Contains large amounts of histamines which may be released in injured tissue in order to increase inflammation
- Contains heparin an anti-clotting chemical



# Lymphocytes

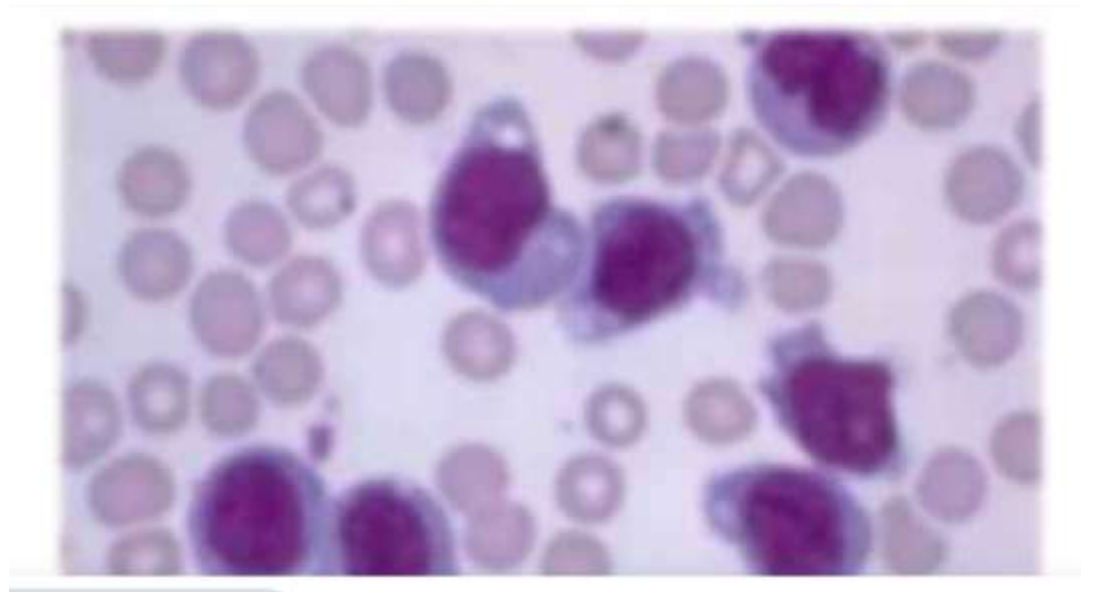
- 30% of all leukocytes
- Made from stem cells, but are released from lymph nodes, thymus, spleen and bone marrow
- Produce antibodies and destroy foreign cells found in infectious mononucleosis



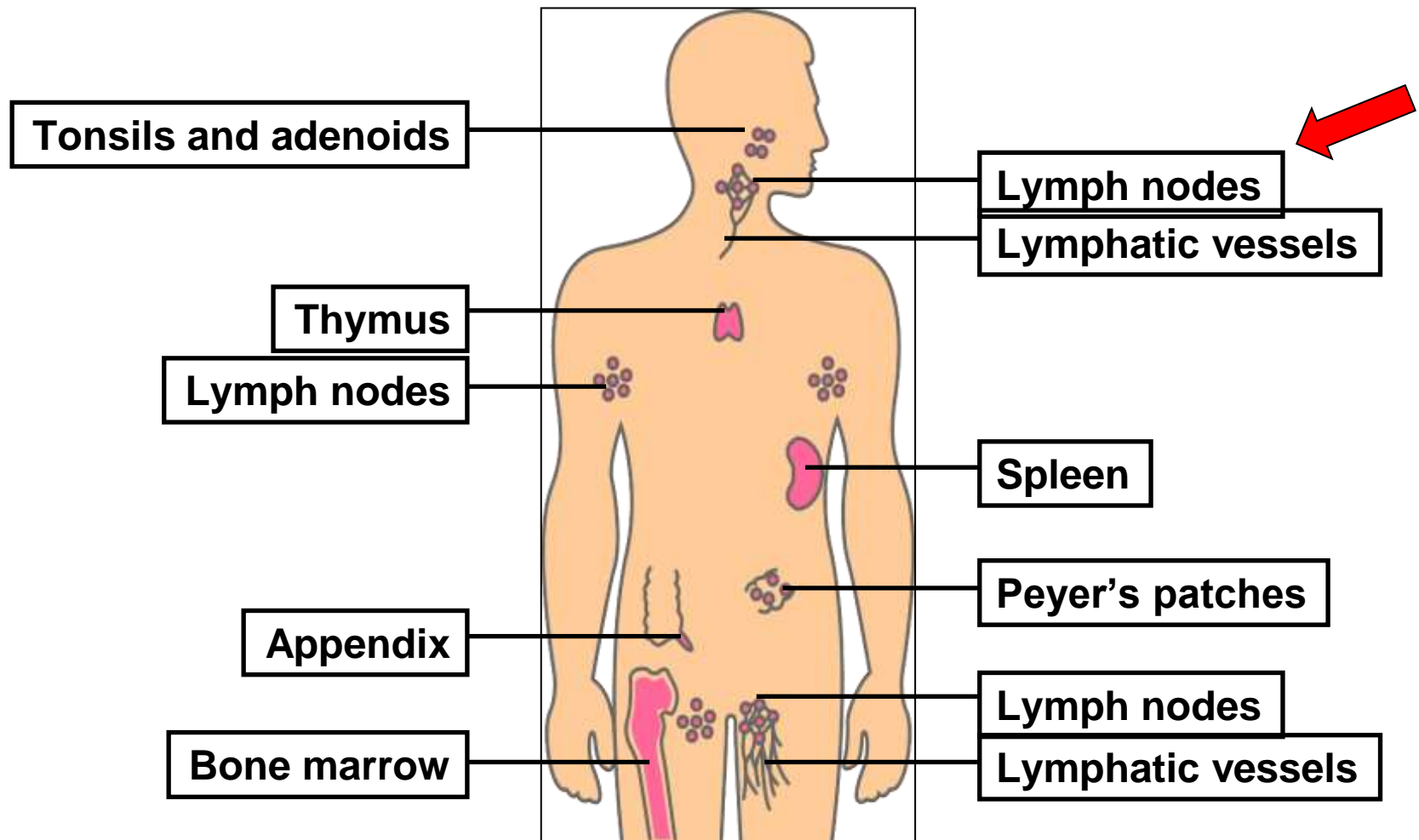


# Monocyte

- 6% of WBC's
- Enter connective tissue
- Eat bacteria, dead cells and other littering tissue



# Organs of the Immune System



Bone marrow, the soft tissue in the hollow center of bones, is the ultimate source of all blood cells, including the immune cells.

# Lymphatic System

The organs of your immune system are connected with one another and with other organs of the body by a network of lymphatic vessels.

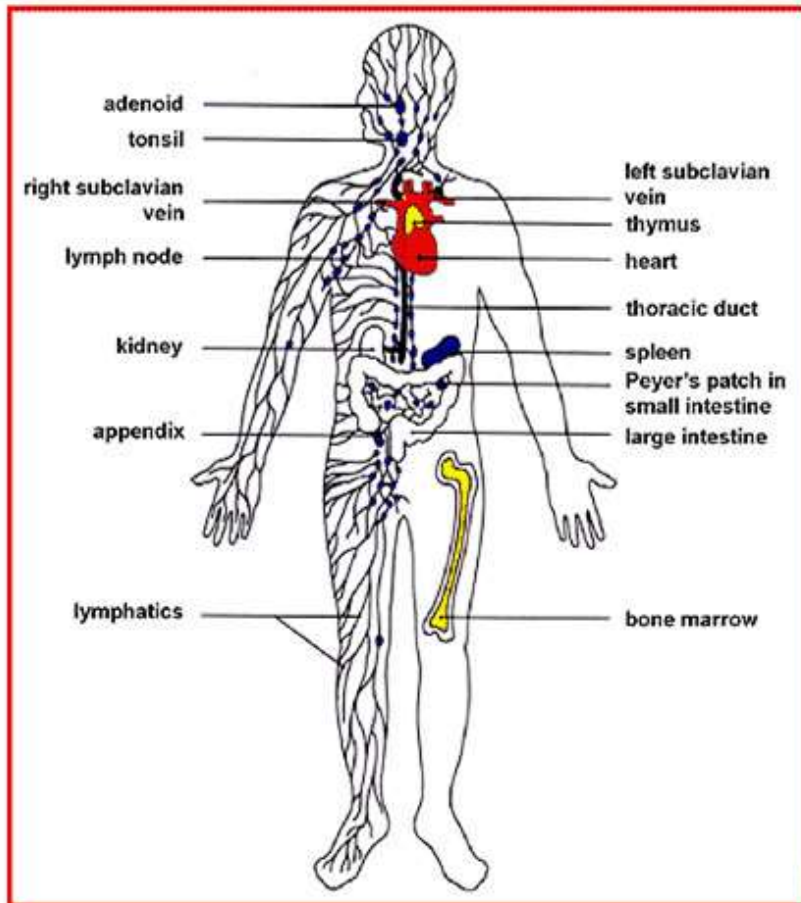
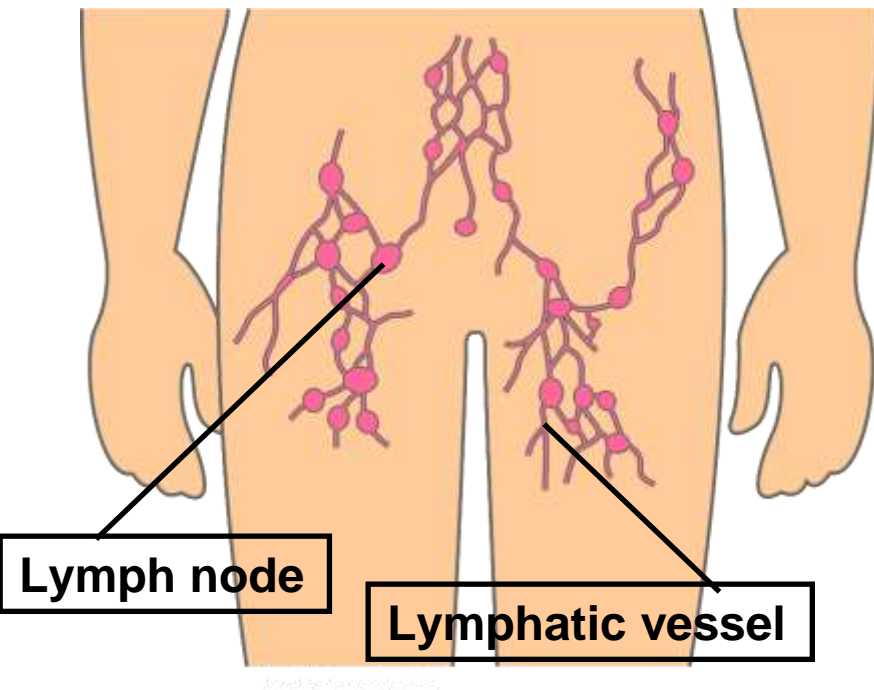


Figure 1. The immune system.

# Lymphatic System

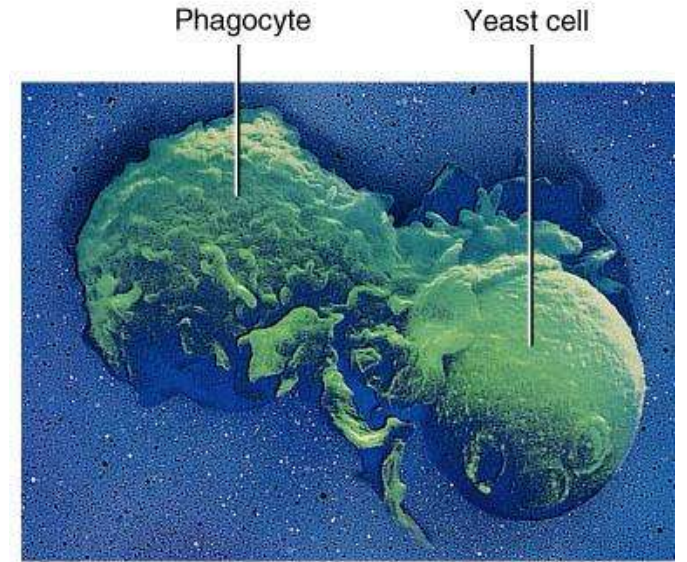
The organs of your immune system are connected with one another and with other organs of the body by a network of lymphatic vessels.



1. Lymphatic **vessels** closely parallels the body's veins and arteries
  - Lymphatic vessels carry *lymph*, a clear fluid that bathes the body's tissues
  - Cells/fluids are exchanged between blood and lymphatic vessels, **enabling the lymphatic system to monitor the body for invading microbes.**
2. Lymph **nodes** contain high levels of immune cells

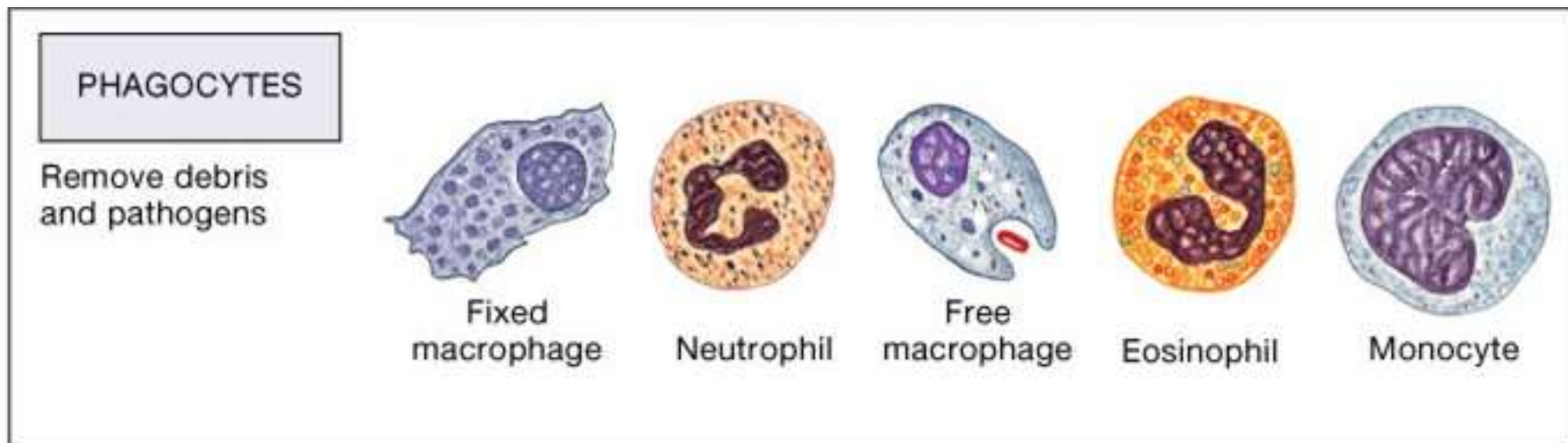
# Nonspecific Defenses, Phagocytes

- Remove cellular debris and respond to invasion by foreign pathogens
  - Monocyte-macrophage system - Fixed and free
  - Microphages – Neutrophils and eosinophils
  - Move by diapedesis
  - Exhibit chemotaxis



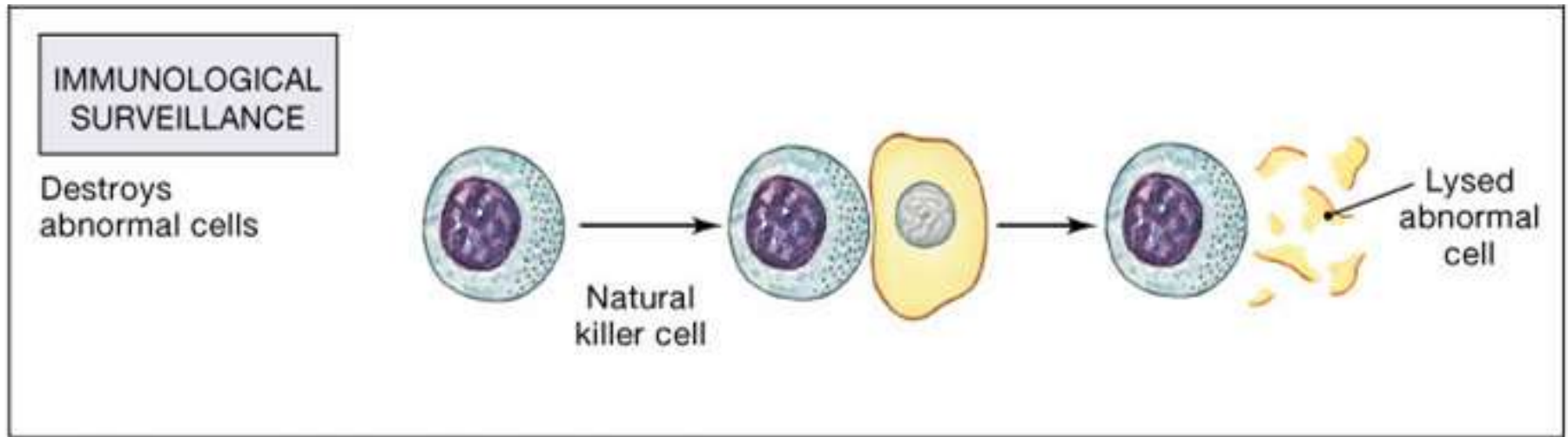
**SEM** about 2500x

(b) Phagocyte engulfing a yeast cell



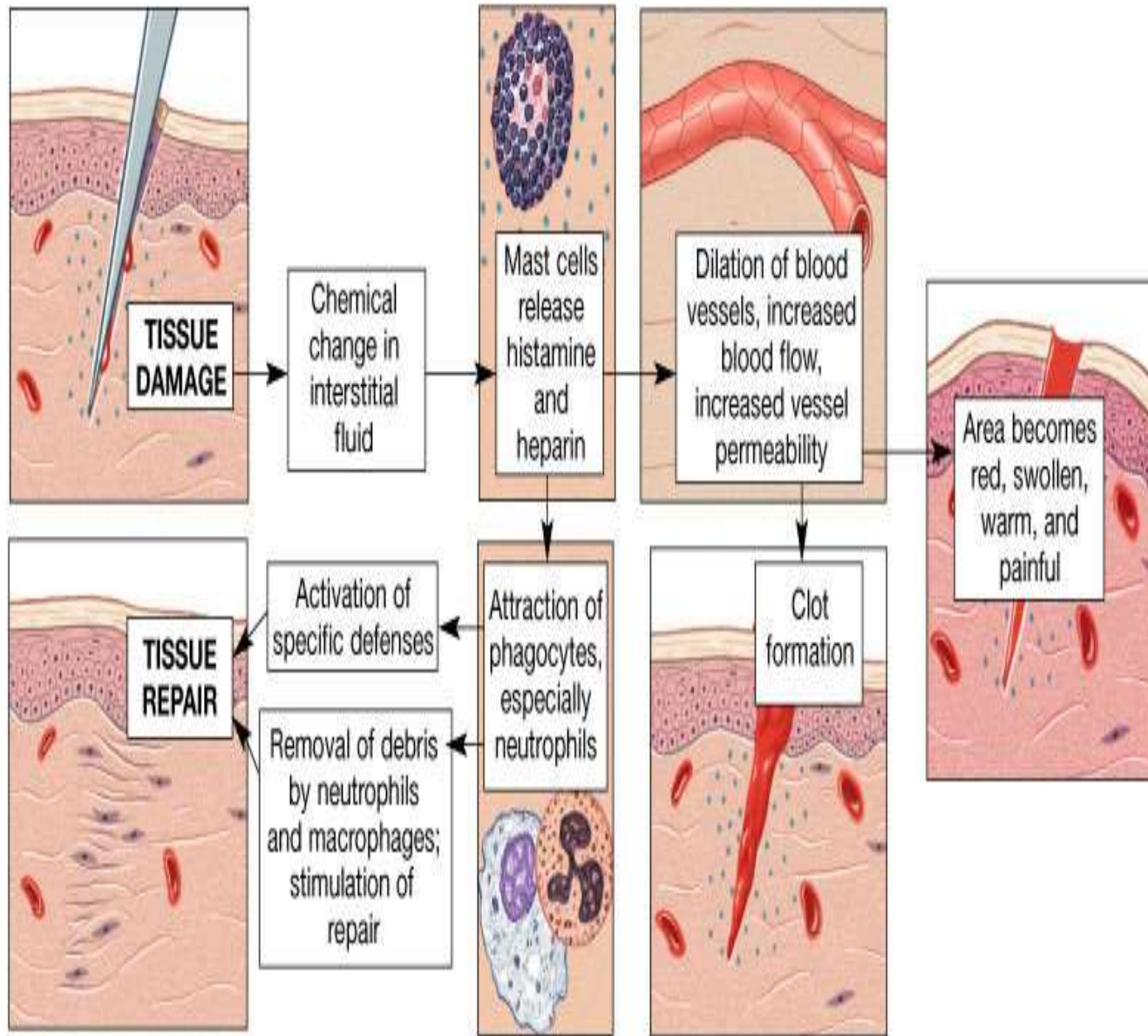
# Nonspecific Defenses, Immunological surveillance

- Constant monitoring of normal tissue by NK cells
- NK cells
  - Recognize cell surface markers on foreign cells
  - Destroy cells with foreign antigens

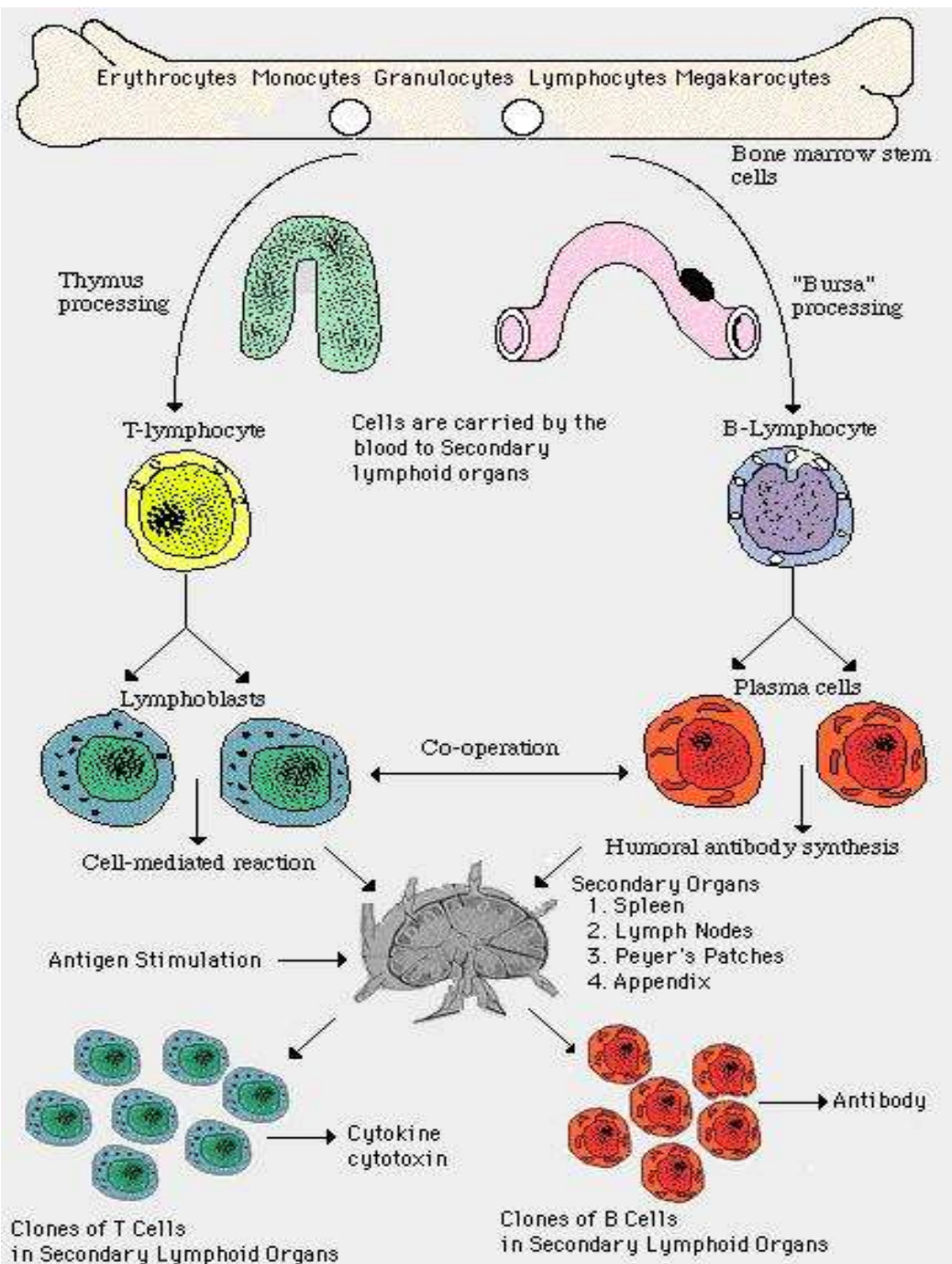




# Inflammation







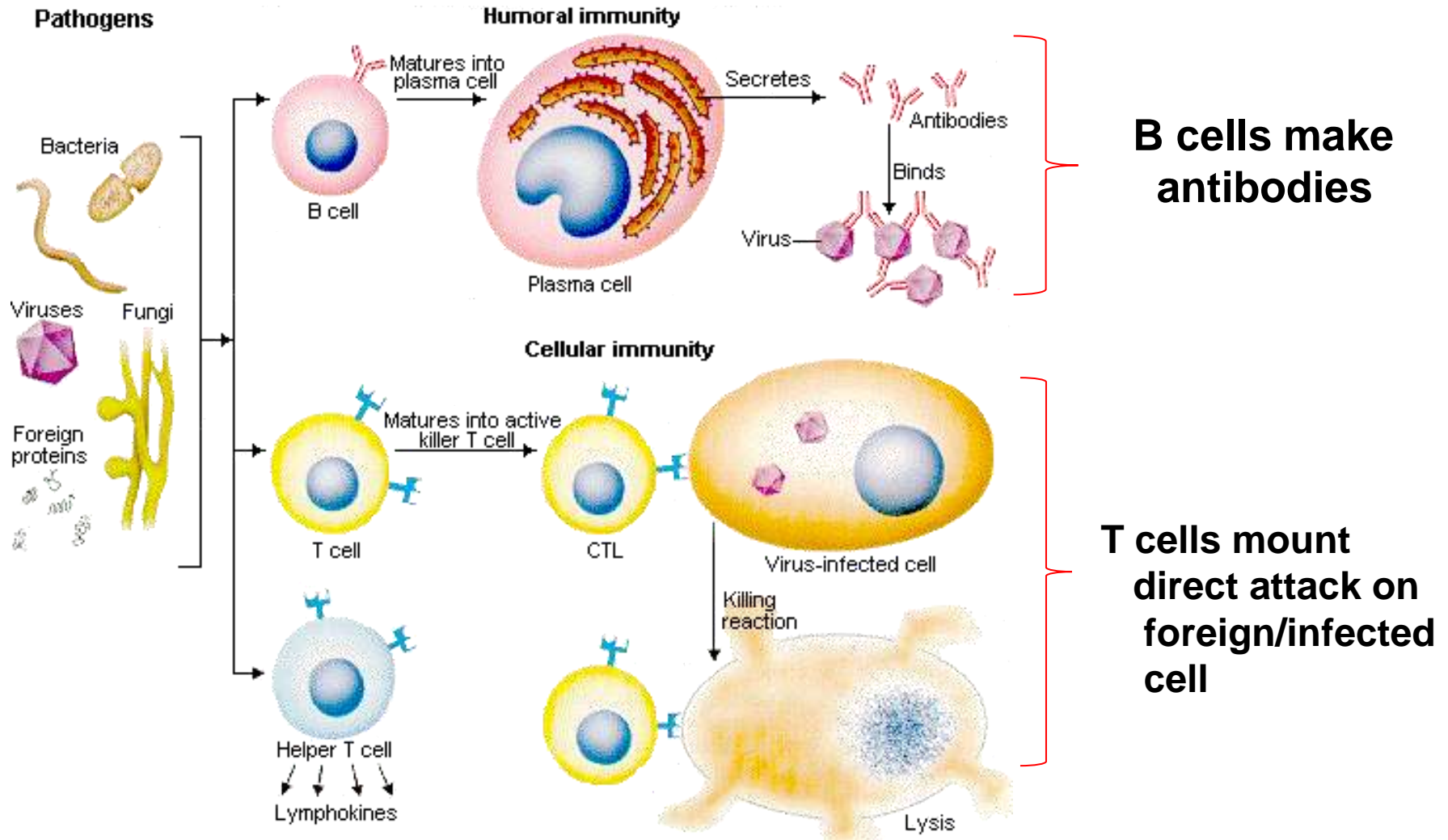
# Adaptive or Acquired Immunity

- Acquired after birth
- Seen only in vertebrates
- Characteristic features are:
  - Diversity
  - Specificity
  - Self vs non-self
  - Memory

# Immune Response System

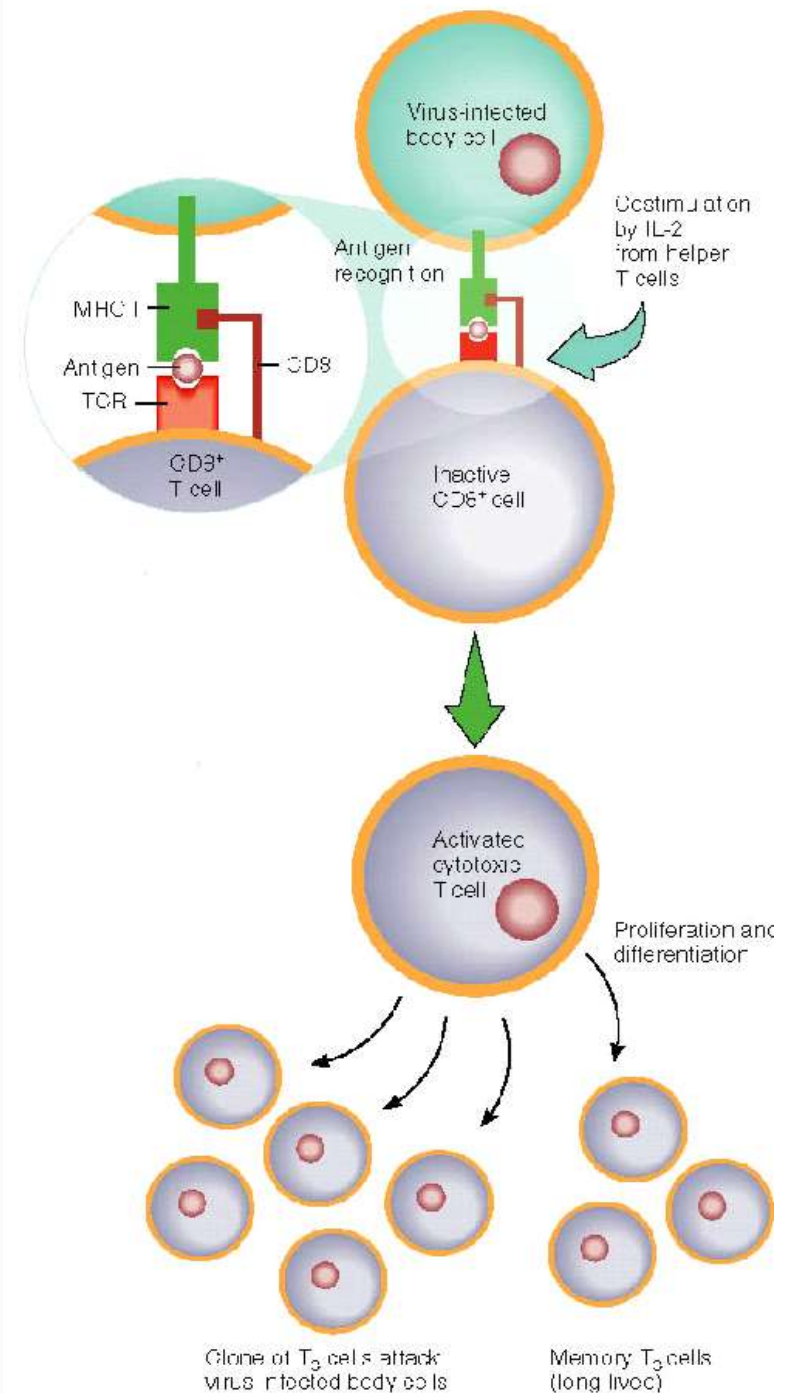
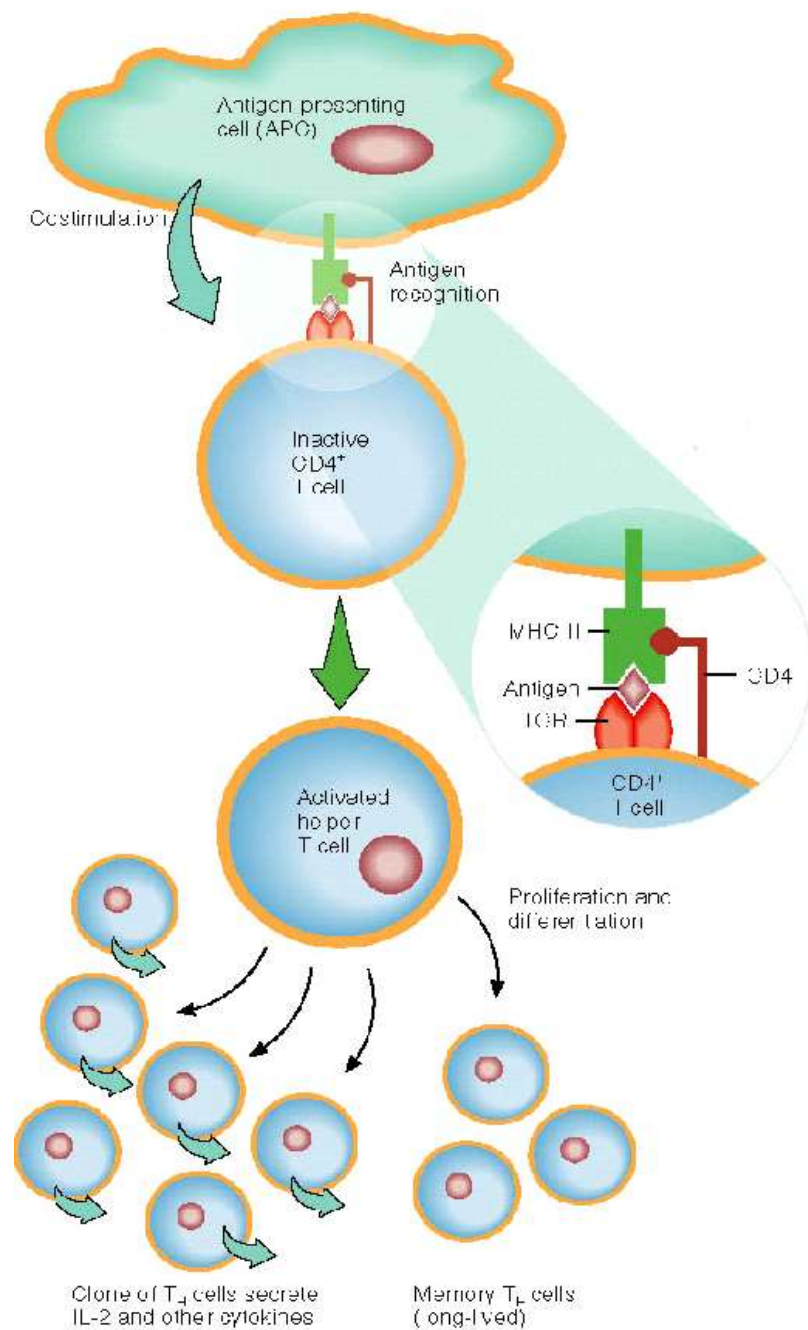
Made up of two cellular systems (lymphocytes)

1. Humoral immunity - **B cells**
2. Cell-mediated immunity - **T cells**



# **Cell mediated immunity**

- **T cells must be activated**
- **Must have both surface antigen recognition and costimulation to activate**
- **T cell receptors recognise and bind to specific antigen presented with MHC complexes**
  - **T cell only activated if binds to antigen and receives costimulation**
    - **Co-stimulation provided by cytokines or membrane proteins**
  - **Need for co-stimulation prevents immune responses occurring accidentally**
    - **Recognition (binding to receptor) without costimulation results in anergy (prolonged state of inactivity) in both B and T cells**
  - **Once T cell co-stimulated it is activated**
    - **Proliferates**
    - **Differentiates (forms more highly specialised cells)**
  - **Activation, proliferation and differentiation occurs in secondary lymphatic organs and tissues**

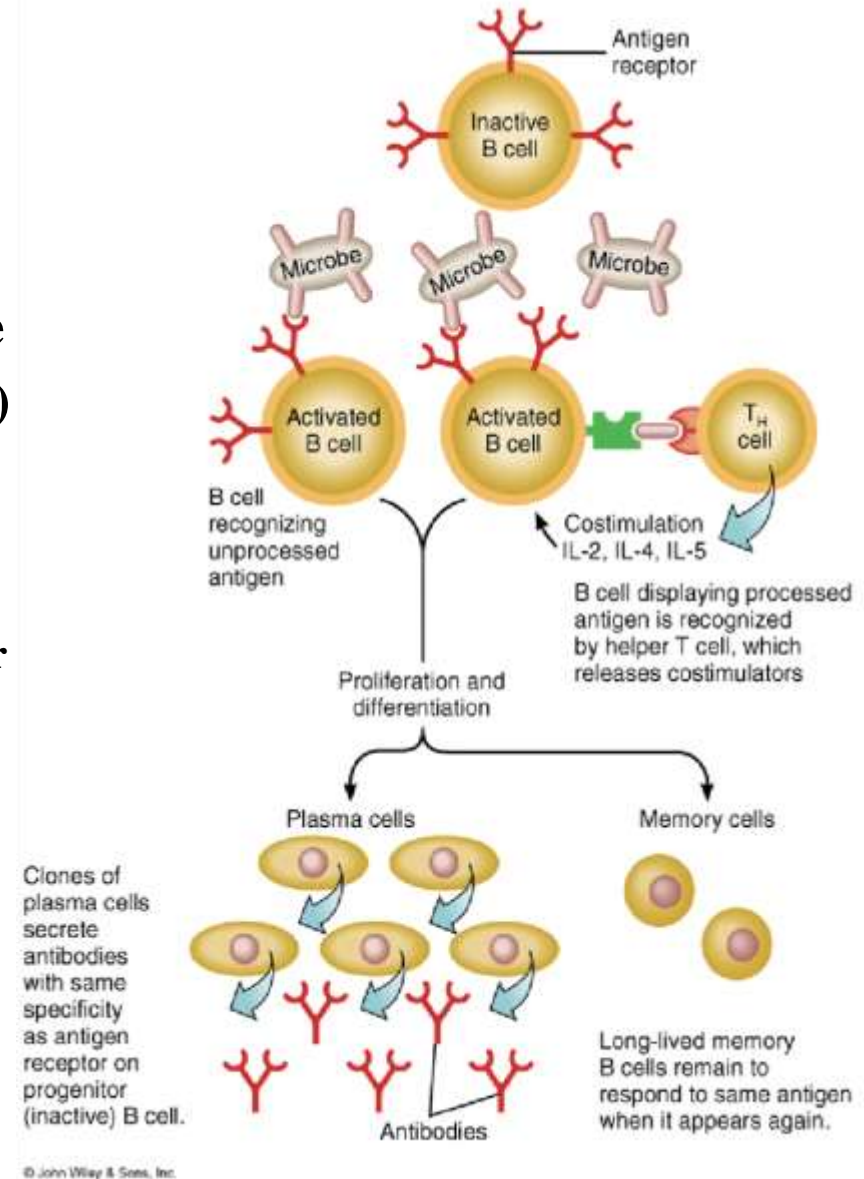






# Humoral (Antibody-mediated) immunity

- Mediated by B cells
- Antigen can activate B cell in two ways:
  - direct binding
    - provokes less vigorous response
  - B cells process antigen (act as APC) and display processed antigen with MHC proteins
    - $T_H$  cells recognise processed antigen
      - $T_H$  cells provide co-stimulation for B cell
- Activated B cell
  - proliferates and differentiates
    - plasma cells
      - secrete antibodies with same antigen binding properties as receptors
    - memory B cells



# Innate/Acquired-Adaptive Immunity

- Innate immunity rely on a global distinction of self/non-self recognition
  - Instantaneous
  - Non specific
  - Non adaptative
  - Memory less
- Adaptative immunity involve the cell-mediated and hummoral response.
  - Time limited to the eradication of the antigen
  - Specific to the given antigen
  - Adaptativ
  - Memory full



# Immunologic Memory

- First Response (4-5 days)
  - First adaptive response against a given antigen
- Secondary Response (1 day)
  - Shorter lag time
  - More rapid buildup
  - Higher overall level of response
  - Better fit to the invading antigen
  - Utilizes IgG instead of IgM

# Active and Passive Immunity

- Active immunity and passive immunity are two types of acquired immunity.

Active Immunity	Passive Immunity
Active immunity is usually long-lasting, sometimes life-long. It is produced by the antibodies of the host in response to direct contact with an antigen	Passive immunity lasts only for a few weeks or months. It is produced by the introduction of antibodies from outside into the host
It produces an immunological memory	It does not produce immunological memory
When the antigens enter the body, antibodies and other specialised lymphocytes are produced	Antibodies are introduced from an external source. For instance, a mother introduces antibodies to a fetus through the placenta and to an infant via mother's milk.
There are no side-effects	It may cause reactions
Immunity does not occur immediately	Immunity develops immediately

# Immunity: Active and Passive

Active Immunity

VS

Passive Immunity

Natural



Infection

Artificial



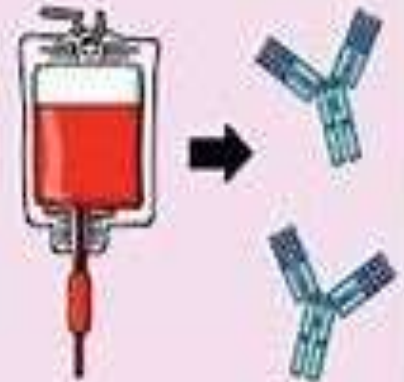
Vaccination

Natural



Maternal  
antibodies

Artificial



Monoclonal  
antibodies

# VACCINES

- Vaccination is the administration of antigenic agents applied to stimulate the immune system of an individual and to develop adaptive immunity to a disease.
- Vaccines can ameliorate, or often even prevent, the effects of infection. Vaccination is generally considered to be the most effective method of preventing infectious diseases
- A multi-stage tuberculosis vaccine and Covid has recently been developed to confer protection after the exposure to the pathogen . There are numerous vaccine examples, including experimental ones against AIDS, cancer and Alzheimer's disease.
- The core mechanism behind all the vaccinations is the ability of the vaccine to initiate an immune response in a quicker fashion than the pathogen itself.
- The purpose of every vaccination is to present a particular antigen or set of antigens to the immune system in order to evoke a relevant immune response.
- The main active component of a vaccine may be inactive, but still intact (attenuated bacteria or viruses), or purified components of the pathogen that are known to induce immune reaction.

# Types of Vaccines

## 1. Inactivated vaccines

- This type of vaccine consists of virus particles grown in cell culture and inactivated by applying high temperature or chemicals such as formaldehyde. Booster shots required Eg., Hepatitis A, Polio, Flu

## 2. Live attenuated vaccines

- The attenuated vaccines contain live virus particles with low levels of virulence. They have retained their ability to slowly reproduce, and thus they remain a continuous source of antigen for a certain period after the first vaccination, reducing the need of booster shots. Eg., MMR combined vaccine, Small pox, Rotavirus

## 3. Subunit vaccines

- Subunit vaccines use only the antigenic components that best stimulate the immune system, instead of dealing with the entire micro-organism. Eg., Hepatitis B, HPV (Human papillomavirus), Pertussis

#### 4. Toxoid vaccines

- The toxoid vaccines are typical solution for bacteria that secrete harmful metabolites or toxins. It is common to use them when the main reason for discomfort or sickness is a bacterial toxin. Eg., diphtheria and tetanus.

#### 5. DNA vaccines

- DNA vaccination is a very new approach for induction of humoral and cellular immune responses to protein antigens by administering genetically engineered DNA. DNA vaccines are still in the experimental stage, and have been tested in numerous viral, bacterial and parasitic models of disease

#### 6. Peptide vaccines

- The improved knowledge of antigen recognition at molecular level has contributed to the development of rationally designed peptide vaccines. chemical approach to synthesize the identified B-cell and T-cell epitopes that are immunodominant and can induce specific immune responses. The peptide vaccines against various cancers have been developed, and entered phase I and phase II of clinical trials



# Vaccines yet to developed!



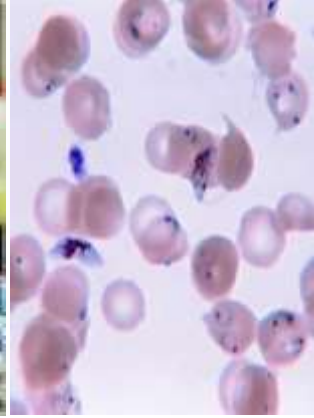
**HIV/AIDS Patient**



**Aging**



**Malaria-parasitic disease**



**Smoking causes cancer**



**Diabetes**

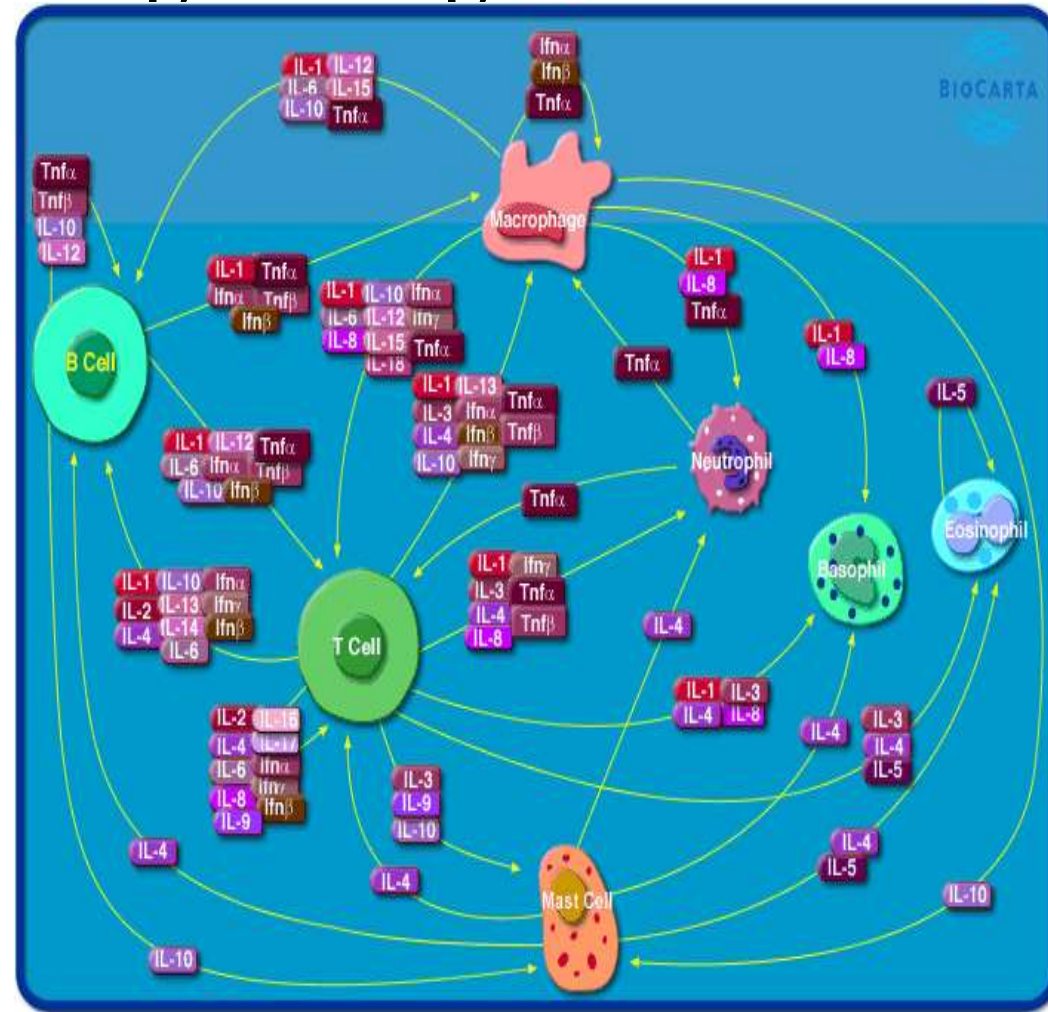


**Rheumatoid arthritis**



# Immune Engineering

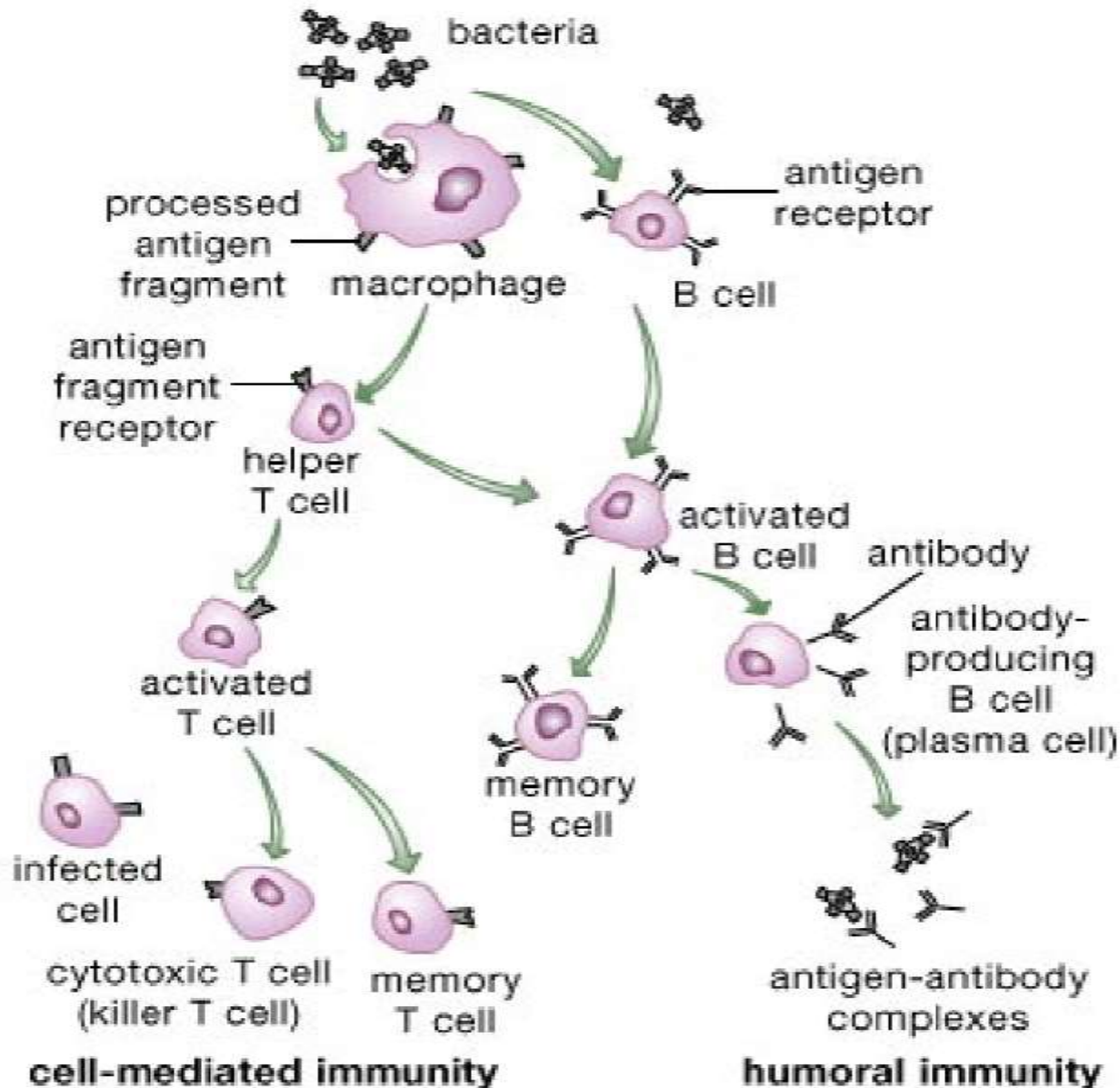
- **The complexity of the immune system can be compared to that of the brain.**
- **There is a vast number of cells, molecules, and organs that compose the immune system, and these have to act in concert, and together with other vital systems, so as to promote and maintain life.**
- **Neither can the immune system act in isolation to maintain life, nor can a higher organism live without an immune system.**



- **Artificial immune systems (AIS) compose a new computational intelligence approach inspired by theoretical and experimental immunology with applications to problem solving.**
- **Like all new approach, the field still lacks a more formal description and better theoretical foundations.**
- **The application of mathematical analysis and modeling to immunology may result in outcomes such as a deeper and more quantitative description of how the immune system works, a more critical analysis of hypothesis, it can assist in the prediction of behaviors and the design of experiments.**

- Immuno-informatics
- Epitope prediction tools

# Summary of Acquired Immunity



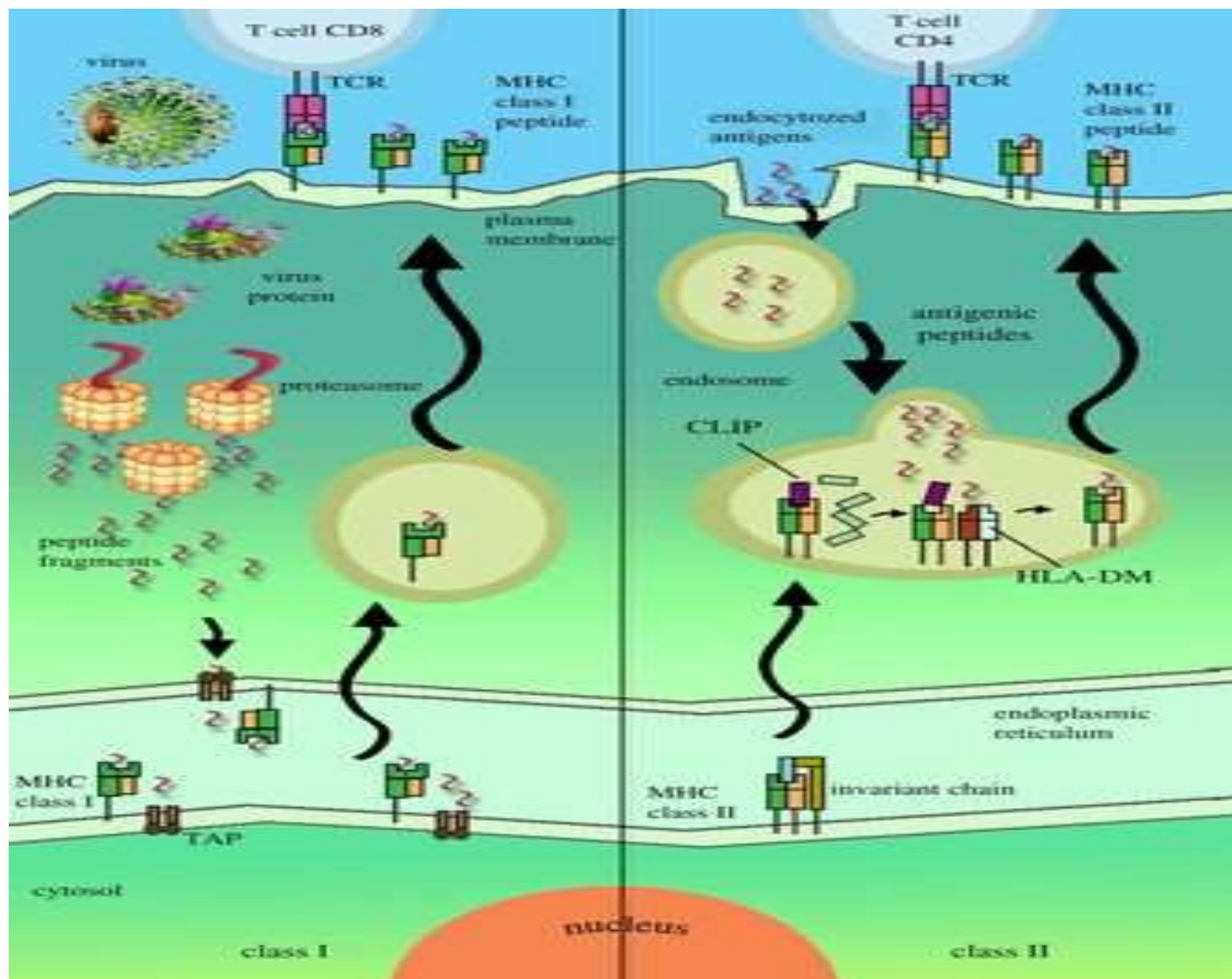
# Combinatorial Science

- Immunoinformatics also known as computational immunology
- Astounding diversity of immune system components
- Complexity of the regulatory pathways
- Complexity of the network-type interactions

# T-cell epitopes

- T-cell epitopes are presented on the surface of an antigen-presenting cell (APC), where they are bound to major histocompatibility (MHC) molecules in order to induce immune response
- MHC class I molecules usually present peptides between 8 and 11 amino acids in length, whereas the peptides binding to MHC class II may have length from 12 to 25 amino acids





# T cell Epitope Prediction Methods

- Binding affinity predictions for a range of MHC molecules is considered to be the most popular epitope prediction methods.
- Molecular binding between antigenic peptides and MHC molecules is necessary for recognition by cytotoxic T-cells.
- The peptides with ability to bind MHC molecules could be predicted by different methodologies including Hidden Markov model (HMM), Artificial neural networks (ANNs), Support Vector Machine (SVM), and Quantitative matrices.

**Table 6.** A list of reliable tools and databases for T-cell epitope prediction.

Database name	URL Link	Application
Allele frequencies	<a href="http://www.allelefrequencies.net">http://www.allelefrequencies.net</a>	HLA frequencies in worldwide population and polymorphism frequencies in immunologically
MHCPred	<a href="http://www.ddg-pharmfac.net/mhcpred/MHCPred/">http://www.ddg-pharmfac.net/mhcpred/MHCPred/</a>	Quantitative prediction of peptide-MHC binding
MMBPred	<a href="http://www.imtech.res.in/raghava/mmbpred/">http://www.imtech.res.in/raghava/mmbpred/</a>	Prediction of atypical MHC class I binders as well as mutations that allow high-affinity binding
NetCTL	<a href="http://www.cbs.dtu.dk/services/NetCTL">http://www.cbs.dtu.dk/services/NetCTL</a>	Prediction of CTL/HLA supertype epitopes
NetMHC	<a href="http://www.cbs.dtu.dk/services/NetMHC">http://www.cbs.dtu.dk/services/NetMHC</a>	Prediction of the MHC binding propensity of peptides
NetChop	<a href="http://www.cbs.dtu.dk/services/NetChop">http://www.cbs.dtu.dk/services/NetChop</a>	Prediction of proteasome/immunoproteasome cleavage
TAPPred	<a href="http://www.imtech.res.in/raghava/tappred/">http://www.imtech.res.in/raghava/tappred/</a>	Prediction of binding affinity of TAP protein
Pcleavage	<a href="http://www.imtech.res.in/raghava/pcleavage/">http://www.imtech.res.in/raghava/pcleavage/</a>	Prediction of proteasome/immunoproteasome cleavage
Propred	<a href="http://www.imtech.res.in/raghava/propred1/">http://www.imtech.res.in/raghava/propred1/</a>	Prediction of class II binding regions in antigenic protein sequence
ElliPro	<a href="http://www.tools.immuneepitope.org/tools/ElliPro">http://www.tools.immuneepitope.org/tools/ElliPro</a>	Prediction of linear and conformational antibody epitopes
EpiToolKit	<a href="http://www.epitoolkit.org">http://www.epitoolkit.org</a>	Prediction of MHC classes I/II ligands using several method
EpiVax	<a href="http://www.epivax.com">http://www.epivax.com</a>	Prediction of classes I/II conserved and promiscuous epitopes
MAPPP	<a href="http://www.mpiib-berlin.mpg.de/MAPPP/cleavage.html">www.mpiib-berlin.mpg.de/MAPPP/cleavage.html</a>	Prediction of proteasome cleavage sites
SYFPEITHI	<a href="http://www.syfpeithi.de">http://www.syfpeithi.de</a>	A database for MHC anchor motifs and binding specificity
IMGT <sup>®</sup>	<a href="http://www.imgt.org">http://www.imgt.org</a>	A high-quality integrated resource of IG, TR, MHC, and related proteins
IMGT/HLA	<a href="http://www.ebi.ac.uk/imgt/hla/allele.html">http://www.ebi.ac.uk/imgt/hla/allele.html</a>	A database with 5518 HLA class I and 1612 HLA class II alleles



# NetMHCpan

- Predicts **binding** of **peptides** to any **known MHC** molecule using **artificial neural networks** (ANN).
- Trained on more than **115,000 quantitative binding data** covering more than 120 different MHC molecules.
- MHC class I: humans, non-human primates (chimpanzee, rhesus macaque, gorilla), mice, **pigs**, and cattle.
- Includes the newest MHC allele releases from the IMGT/HLA & IPD-MHC databases.

# [https://services.healthtech.dtu.dk/service.php? NetMHCpan-4.1](https://services.healthtech.dtu.dk/service.php?NetMHCpan-4.1)

## NetMHCpan

The screenshot displays the NetMHCpan-4.1 web interface. At the top, there is a navigation bar with a logo on the left and a series of colored tabs (yellow, green, orange, red, purple, blue, brown) for different services. Below the navigation bar, a pink bar contains the text "SUBMISSION", "Output format", "API/CSV download", and "Evaluation Page". The main content area is titled "SUBMISSION" and includes a dropdown menu for "Type of input: Peptide". Below this, a text box is labeled "Paste a single sequence or several sequences in FASTA format into the field below:". To the right of this text box, an arrow points to the text "Enter your protein(s) sequence(s)". Below the text box, there is a link "or submit a file in FASTA format directly from your local disk:" followed by a "Browse..." button. Further down, there is a "Peptide length" dropdown set to "Over peptides". Below that, a "Select species" dropdown is set to "Pig (SLA)". Underneath, a "Select SLA (max 20 per submission)" dropdown is shown with a list of SLA alleles: SLA-1\*0101, SLA-1\*0201, and SLA-1\*0202. To the right of this list, an arrow points to the text "45 SLA alleles".

ENTER YOUR PROTEIN(S) SEQUENCE(S)

45 SLA alleles

# NetMHCpan Prediction Results

NetMHCpan Server - prediction results

Technical University of Denmark

# NetMHCpan version 2.4

# Input is in FSA format

# Peptide length 9

SLA-1\*0201 : Estimated prediction accuracy 0.416 (using nearest neighbor HLA-A\*80:01)

# Threshold for Strong binding peptides 50.000

# Threshold for weak binding peptides 500.000

pos	HLA	peptide	Identity	1-log50k(aff)	Affinity(nM)	%Rank	BindLevel
131	SLA-1*0201	ITYLNNMGY	gi_397586_emb_C	0.604	72.62	0.05	<= WB
440	SLA-1*0201	GILSYTSLY	gi_397586_emb_C	0.502	218.19	0.20	<= WB
458	SLA-1*0201	KVCNMLIAY	gi_397586_emb_C	0.486	259.13	0.25	<= WB
39	SLA-1*0201	YGAYMLFMY	gi_397586_emb_C	0.464	331.85	0.40	<= WB
569	SLA-1*0201	NVPDKHGLY	gi_397586_emb_C	0.452	375.69	0.40	<= WB
660	SLA-1*0201	GQYNLKLIV	gi_397586_emb_C	0.445	403.38	0.40	<= WB
687	SLA-1*0201	YITCPDSL Y	gi_397586_emb_C	0.423	511.67	0.50	
998	SLA-1*0201	IVHRQCYKY	gi_397586_emb_C	0.420	532.53	0.80	
98	SLA-1*0201	LTQRPVMGY	gi_397586_emb_C	0.418	542.64	0.80	
22	SLA-1*0201	RTNAPLLFM	gi_397586_emb_C	0.417	546.54	0.80	
183	SLA-1*0201	VTPEDLVSY	gi_397586_emb_C	0.410	591.08	0.80	
162	SLA-1*0201	TTWILQLHY	gi_397586_emb_C	0.409	597.13	0.80	



# B-cell epitope prediction

- Mapping B-cell epitopes within the protein sequence of antigens plays a crucial role in vaccine design, immunodiagnostic design and immunogen design for antibody production efforts
- B-cell epitopes are determinant regions of the antigen with the ability to bind immunoglobulins and B-cell receptors (BCRs).
- Identification of B-cell epitopes capable of invoking strong immune responses plays a pivotal role in effective vaccine designing efforts

**Table 7.** A list of reliable tools and databases for B-cell epitope prediction.

Database name	Web address	Application
ABCPred	<a href="http://www.imtech.res.in/raghava/abcpred">http://www.imtech.res.in/raghava/abcpred</a>	Prediction of linear B-cell epitopes based on artificial neural net
BCIPEP	<a href="http://www.imtech.res.in/raghava/bcipep">http://www.imtech.res.in/raghava/bcipep</a>	B-cell epitope database
Bepipred	<a href="http://www.cbs.dtu.dk/services/BepiPred">http://www.cbs.dtu.dk/services/BepiPred</a>	Prediction of B-cell epitopes using HMM method
IMGT ®	<a href="http://www.imgt.org">http://www.imgt.org</a>	A high-quality integrated resource of IG, TR, MHC, and related proteins
Bcepred	<a href="http://www.imtech.res.in/raghava/bcepred/">http://www.imtech.res.in/raghava/bcepred/</a>	Prediction of epitopes with 58.7% accuracy
BEPITOPE	<a href="mailto:jlpelequer@cea.fr">jlpelequer@cea.fr</a>	Prediction of linear epitopes location and pattern
DiscoTope	<a href="http://www.cbs.dtu.dk/services/DiscoTope/">http://www.cbs.dtu.dk/services/DiscoTope/</a>	Prediction of conformational B-cell epitopes based on 3D structure
COBEpro	<a href="http://scratch.proteomics.ics.uci.edu">http://scratch.proteomics.ics.uci.edu</a>	A two-stage system to predict linear B-cell epitopes, associated with the SCRATCH database
CEP	<a href="http://bioinfo.ernet.in/cep.htm">http://bioinfo.ernet.in/cep.htm</a>	Prediction of B-cell epitopes
AgAbDb	<a href="http://www.115.111.37.206:8080/agabdb2/home.jsp">http://www.115.111.37.206:8080/agabdb2/home.jsp</a>	Antigen-antibody interaction database
MIMOP	Request from <a href="mailto:franck.molina@cpbs.univ-montp1.fr">franck.molina@cpbs.univ-montp1.fr</a>	prediction of 3D epitopic region from the mimotope peptide sequences
MIMOX	<a href="http://www.immunet.cn/mimox/">http://www.immunet.cn/mimox/</a>	Epitope mapping based on phage display method
Pepitope	<a href="http://www.pepitope.tau.ac.il/">http://www.pepitope.tau.ac.il/</a>	Epitope mapping based on affinity-selected peptides
3DEX	<a href="http://www.schreiber-abc.com/3dex/">http://www.schreiber-abc.com/3dex/</a>	Conformational epitopes mapping in 3D protein structures
IEDB	<a href="http://www.immuneepitope.org">http://www.immuneepitope.org</a>	Epitope prediction database
AntiJen	<a href="http://www.jenner.ac.uk/antijen/">http://www.jenner.ac.uk/antijen/</a>	Quantitative binding data for B-cell epitopes
CED	<a href="http://immunet.cn/ced">http://immunet.cn/ced</a>	Conformational Epitope Database

# BepiPred

- BepiPred-2.0 is based on a random forest algorithm trained on epitopes annotated from antibody-antigen protein structures.
- This new method was found to outperform other available tools for sequence-based epitope prediction both on epitope data derived from solved 3D structures, and on a large collection of linear epitopes downloaded from the IEDB database

<https://services.healthtech.dtu.dk/service.php?BepiPred-2.0>

## BepiPred - 2.0

### Prediction of potential linear B-cell epitopes

Paste or upload protein sequence(s) as fasta format to predict potential B-cell epitopes. Prediction can take a few minutes per sequence.

Submission

Instructions

Datasets

Abstract

Downloads

## Submit data

*At most 50 sequences and 300,000 amino acids per submission; each sequence not less than 10 and not more than 6000 amino acids.*

Enter protein sequence(s) in fasta format...

For example file [Click here](#)

Format directly from your local disk:  No file chosen

## Sequence Markup Types

- › **Epitopes:** Positions above epitope threshold
- › **Predictions:** The protein sequence displayed with orange gradient, illustrating BepiPred-2.0 predictions

Gradients



Low

High

Epitope Threshold ?

0

0.5

1



Name	Sequence Markup
5CON_A	<p>Epitopes : .....EEEEEEEEEEEEEEEEEEEEEEEE.....EEEE.....EEEEEEEE.....</p> <p>Predictions: CDAFVGTWKLVSSENFDDYMKEVGVGFATRKVAGMAKPNMIISVNGDLVTIRSESTFKNTEISFKLGVEFDEITADDRKVKSIITLDGGA</p> <p>1-----10-----20-----30-----40-----50-----60-----70-----80-----9</p>

# Discussion

- Immunoinformatics accelerate the knowledge acquisition in clinical immunology
- Bioinformatics has broad applicability to immunology
- Development of *in-silico* models of entire systems – towards a virtual immune system



**THANK YOU**