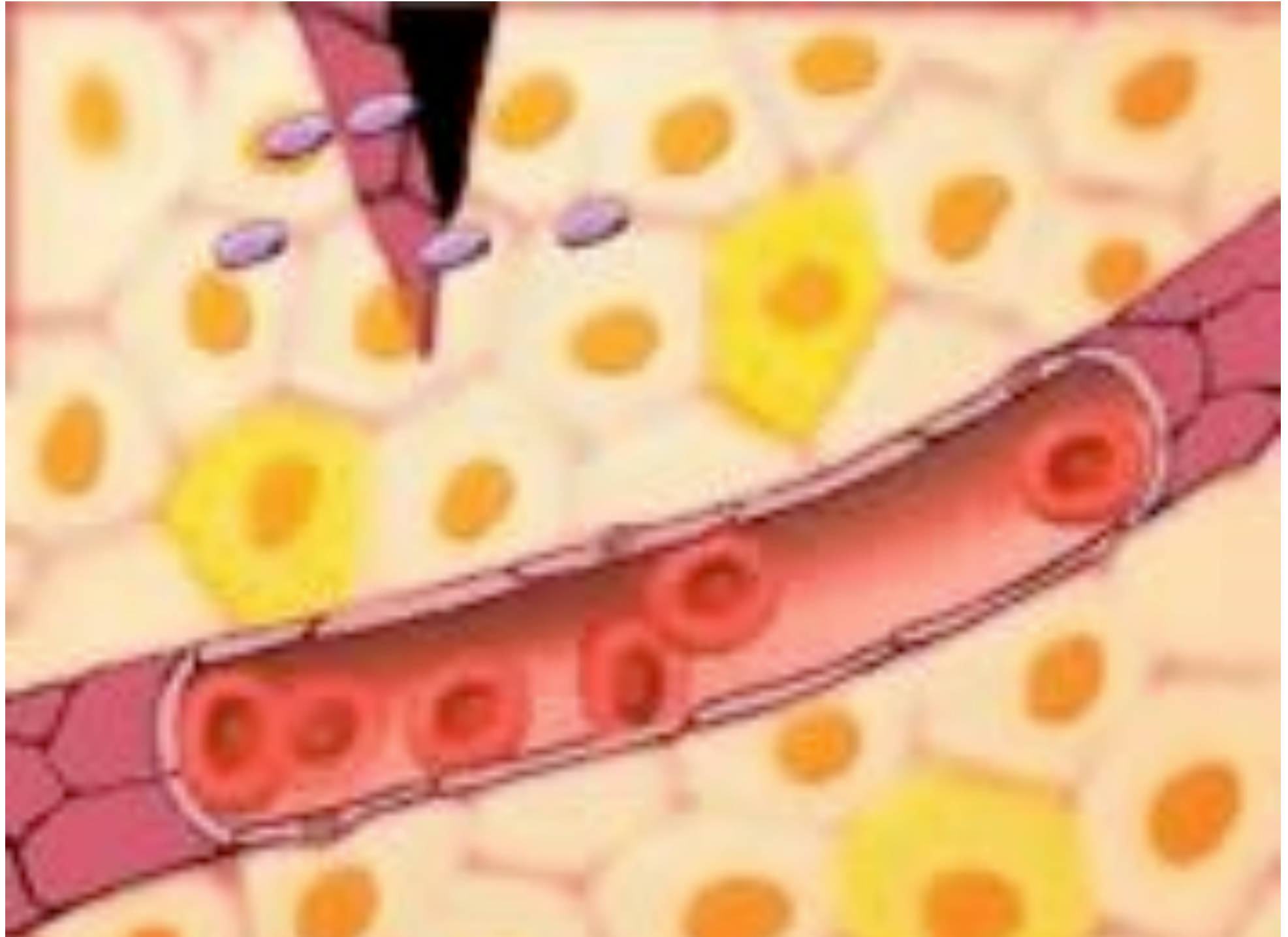


INFLAMMATION RESPONSE!



THE INNATE IMMUNITY AND INFLAMMATION RESPONSE BIOLOGY THE ACUTE PHASE PROTEINS (APP)

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response are microbial, such as viruses and bacteria
which are the major
extracellular DAMPs or Danger-Associated Molecular
Patterns
and contain
DAMPs or Pathogen Associated Molecular Patterns...

Viruses infecting a cell,
multiplication and release

Streptococcus pneumoniae

Growth of pathogenic
bacteria shown in time-laps

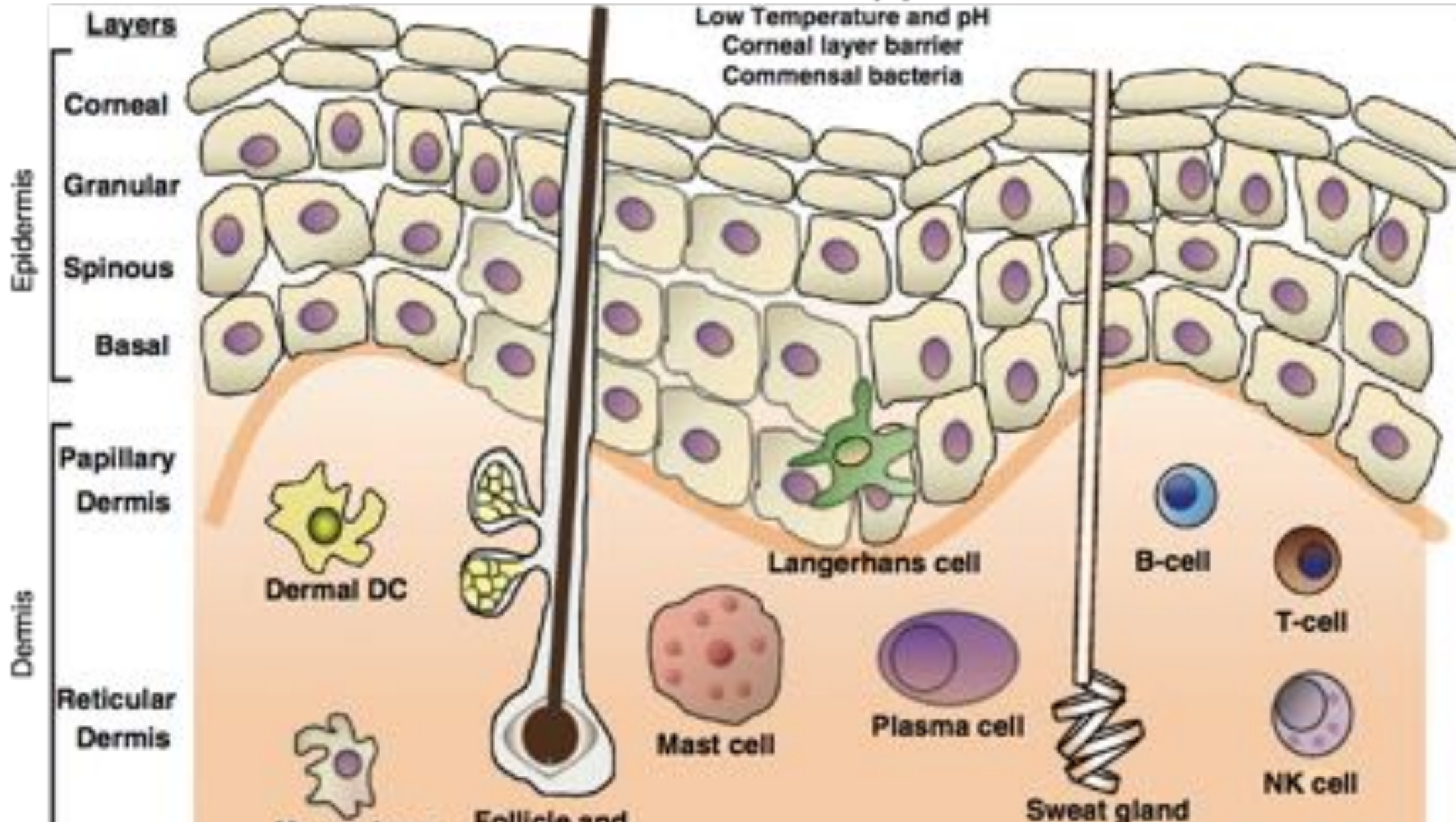
Speed = x 540

impenetrable escape from the potent immunologic tissue barriers!

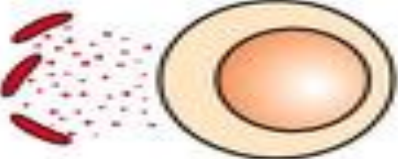
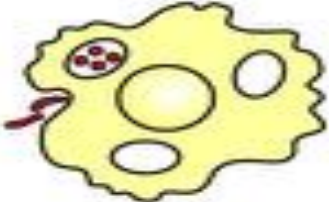
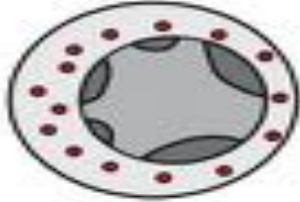
(a)

Skin surface host defenses:

- Antimicrobial peptides
- Low Temperature and pH
- Corneal layer barrier
- Commensal bacteria



DAMAGE WITH THREE MAIN DIRECT MECHANISMS.....

Pathogenic mechanism	Direct mechanisms of tissue damage by pathogens		
	Exotoxin production	Endotoxin	Direct cytopathic effect
			
<p><i>Streptococcus pyogenes</i> <i>Staphylococcus aureus</i> <i>Corynebacterium diphtheriae</i> <i>Clostridium tetani</i> <i>Vibrio cholerae</i></p>	<p><i>Escherichia coli</i> <i>Haemophilus influenzae</i> <i>Salmonella typhi</i> <i>Shigella</i> <i>Pseudomonas aeruginosa</i> <i>Yersinia pestis</i></p>	<p>Variola Varicella-zoster Hepatitis B virus Polio virus Measles virus Influenza virus Herpes simplex virus Human herpes virus 8 (HHV8)</p>	
<p>Tonsillitis, scarlet fever Boils, toxic shock syndrome, food poisoning Diphtheria Tetanus</p>	<p>Gram-negative sepsis Meningitis, pneumonia Typhoid Bacillary dysentery Wound infection</p>	<p>Smallpox Chickenpox, shingles Hepatitis Poliomyelitis Measles, subacute sclerosing</p>	

THEY CAN ENTER OUR CELLS, USING NOT ONLY SPECIFIC RECEPTORS

Trends in Biotechnology June 2012, Vol. 30, No. 6

Pathogenic microbes and their membrane receptor targets

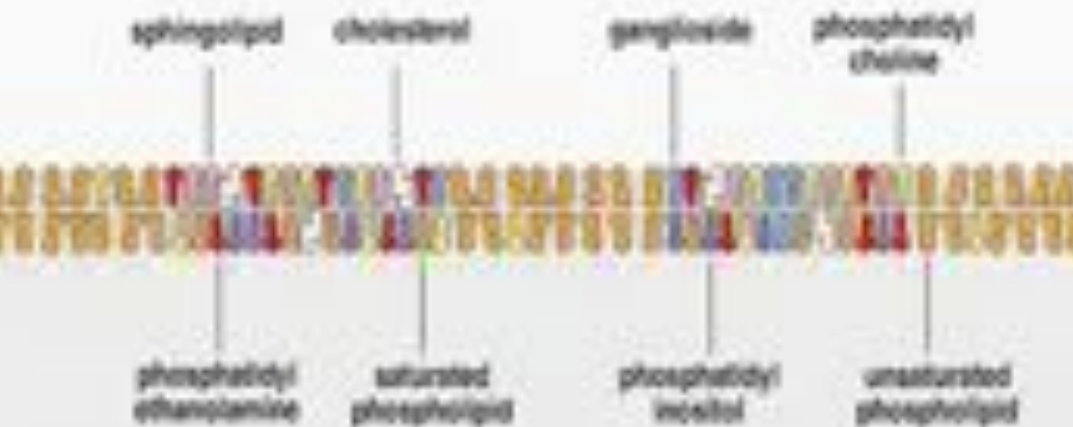
	Species	Virulence factor	Cell receptor*
virus	<i>E. coli</i>	Heat-labile enterotoxin, endotoxin	Ganglioside
	<i>V. cholera</i>	Cholera toxin	Ganglioside
	<i>Streptococcus, Staphylococcus</i>	Lipoteichoic acid, hemolysin	Phospholipid
	Influenza	Hemagglutinin, neuraminidase	Ganglioside
	HIV	GP120 protein	Galactosyl ceramide
enveloped virus	Paramyxovirus	Attachment protein G	EphrinB2 protein
	Polyomavirus, rhinovirus	Capsid coat protein	Ganglioside, ceramide, ICAM-1 and LDLR proteins
	Adenovirus	Capsid protein knob domain	CAR and LDLR proteins

*ICAM-1, intercellular adhesion molecule 1; CAR, coxsackie virus and adenovirus receptor; LDLR, low-density lipoprotein receptor.

Understanding host-pathogen interactions
using membrane-based nanostructures

Santos Manes, Gustavo del Real & Carlos Martinez-A

Nature Reviews Immunology 3, 557-568 (2003)



I rafts lipidici sono delle strutture di membrane eterogenee, insolubili in detergenti non ionici come Triton X-100 ed arricchite in colesterolo, glicosfingolipidi come GM1 o GM3 e proteine come le caveoline, flotilline

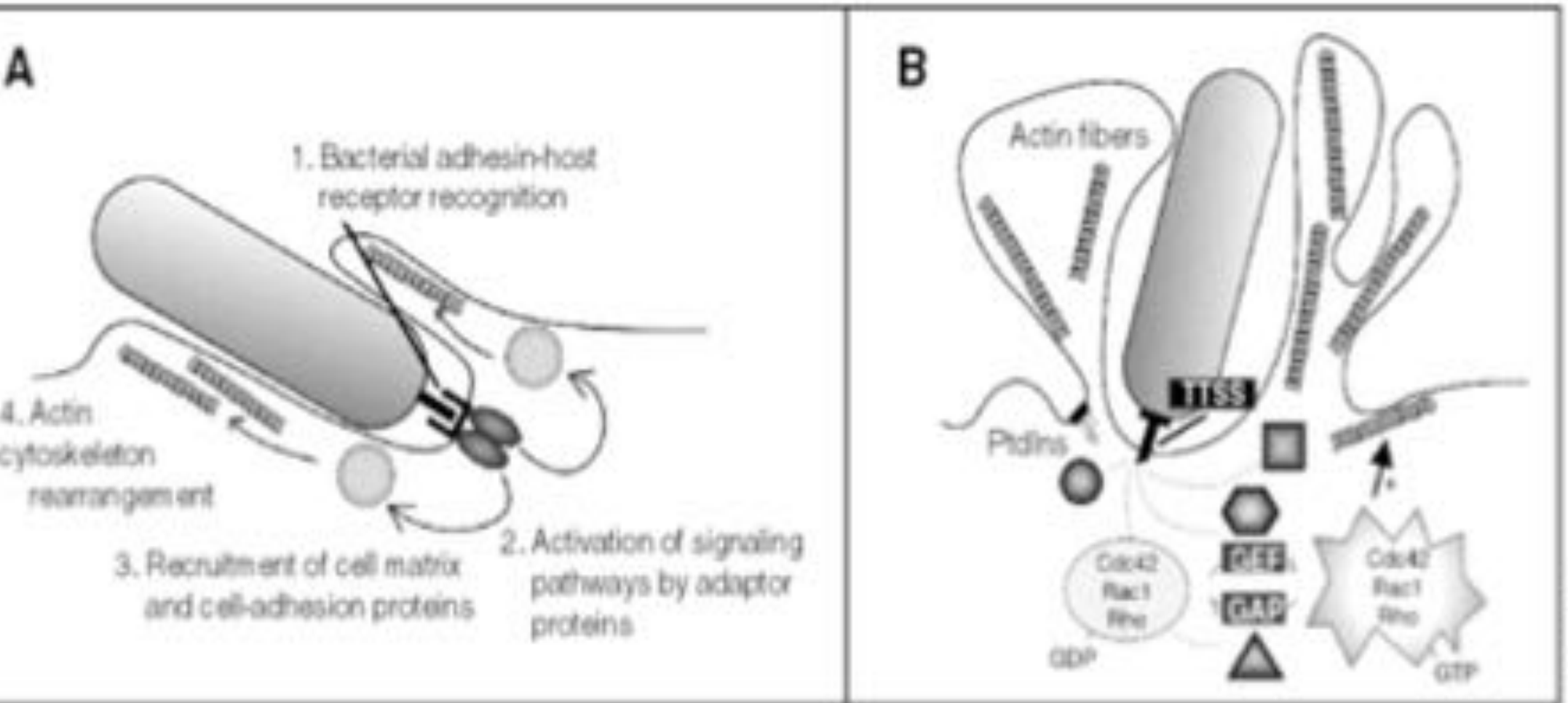
Viruses	
Adenovirus 40	Entry/trafficking
Adenovirus 11	Entry/trafficking
Adenovirus 1	Entry
Human sarcoma and leukosis virus	Entry
Human milk forest virus	Entry/budding
Sendai virus	Entry/budding
Human T-cell leukaemia virus type 1	Entry/budding
HIV-1	Entry/budding/transcytosis
Ebola and Marburg viruses	Entry/budding

Bacteria

<i>Campylobacter jejuni</i>	Intracellular survival
<i>Legionella pneumophila</i>	Intracellular survival
<i>Pseudomonas aeruginosa</i>	Host response, signalling
<i>Brucella</i> spp.	Entry/intracellular survival
FimH and Dr+ <i>Escherichia coli</i>	Entry/intracellular survival
<i>Salmonella typhimurium</i>	Entry/intracellular survival
<i>Shigella flexneri</i>	Entry/intracellular survival
<i>Chlamydia</i> spp.	Entry/intracellular survival
<i>Mycobacterium</i> spp.	Entry/intracellular survival
<i>Vibrio cholerae</i> (cytolysin)	Toxin binding/oligomerization
<i>Aeromonas hydrophila</i> (aerolysin)	Toxin binding/oligomerization
<i>Clostridium</i> spp.	Toxin binding/oligomerization
<i>Streptococcus pyogenes</i> (streptolysin O)	Toxin oligomerization
<i>Bacillus anthracis</i> (anthrax toxin)	Toxin oligomerization
<i>Bacillus thuringiensis</i> (Cry1A toxin)	Toxin binding/oligomerization

ZIPPER MECHANISM

TRIGGER MECHANISM



bacteria to enter the cells, move and escape the cytoplasm and modulate the functions using proteins that mimic the functions of the structural and signaling proteins (such as small G proteins Rho, Rac and cdc42) and their effectors (such as Wasp,

Microbial pathogenesis and cytoskeletal function

Samantha Gruenheid and Brett Finlay

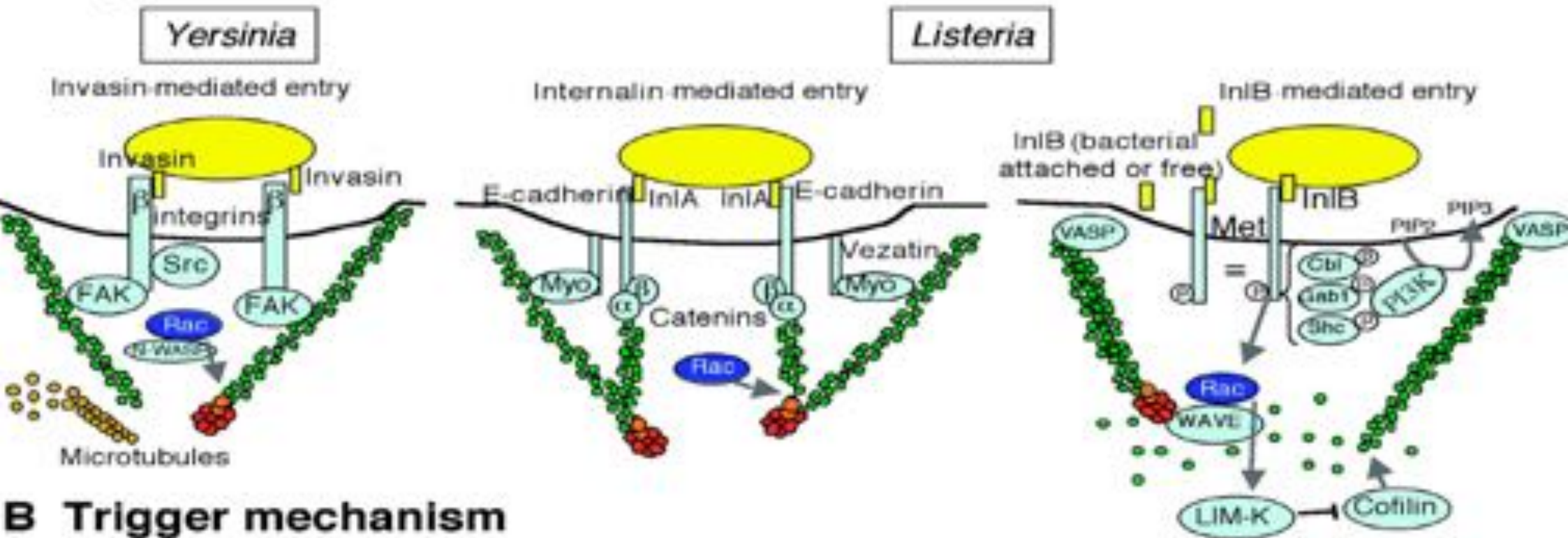
Nature 422, 775-781 (17 July 2003)

Mechanisms used by bacteria to enter cells.

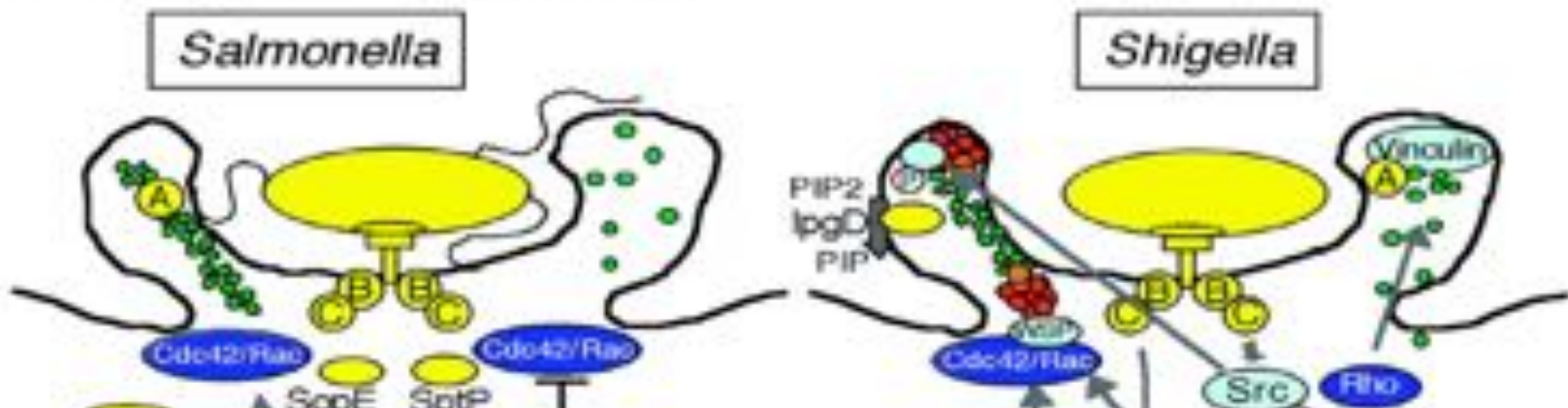
The zipper mechanism used by *Yersinia* and *Listeria*.

The trigger mechanism used by *Salmonella* and *Shigella*.

A Zipper mechanism

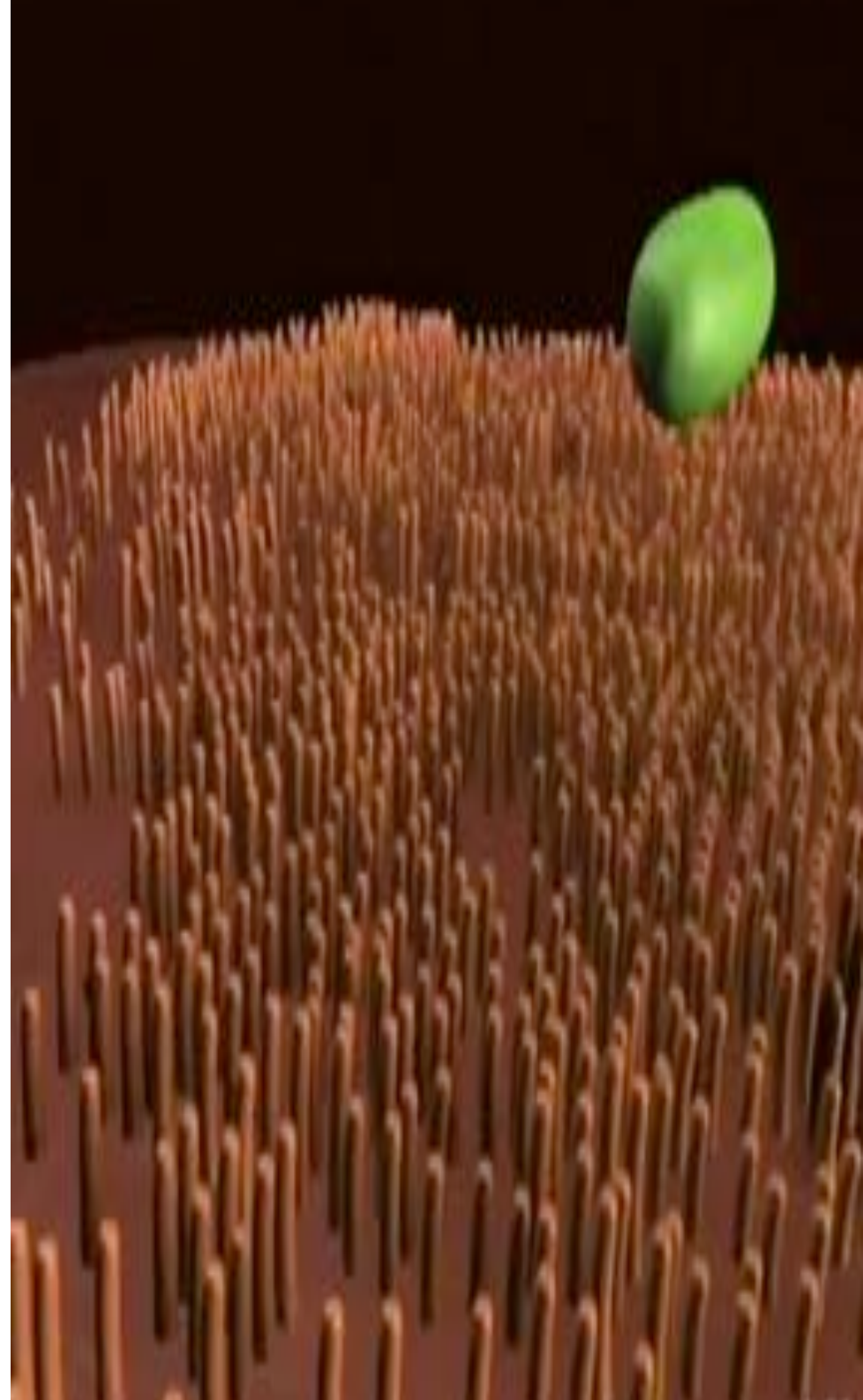


B Trigger mechanism



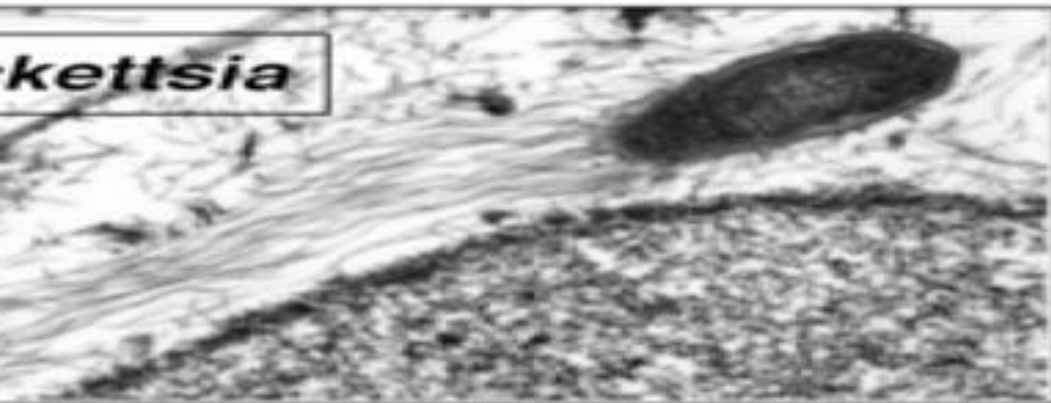
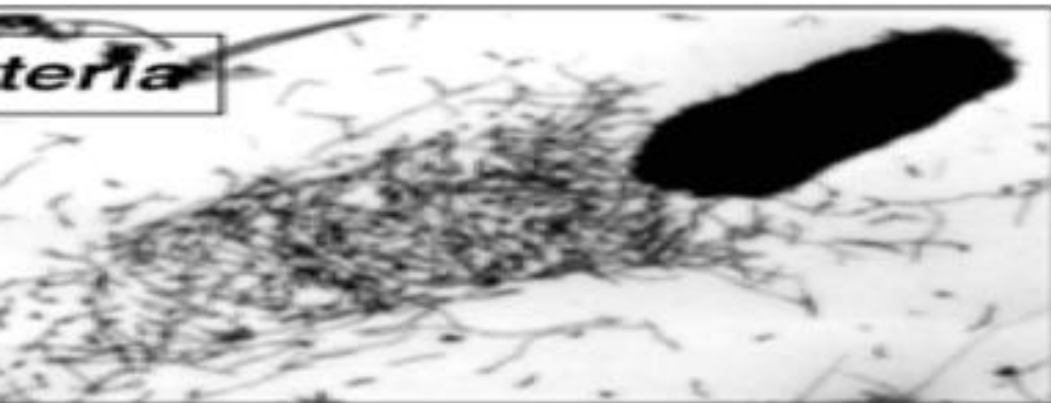
	Type III secretion apparatus
	Actin
	Arp2/3 complex
	Rho GTPases

The
invasion
and cell
migration
of
Salmonella!

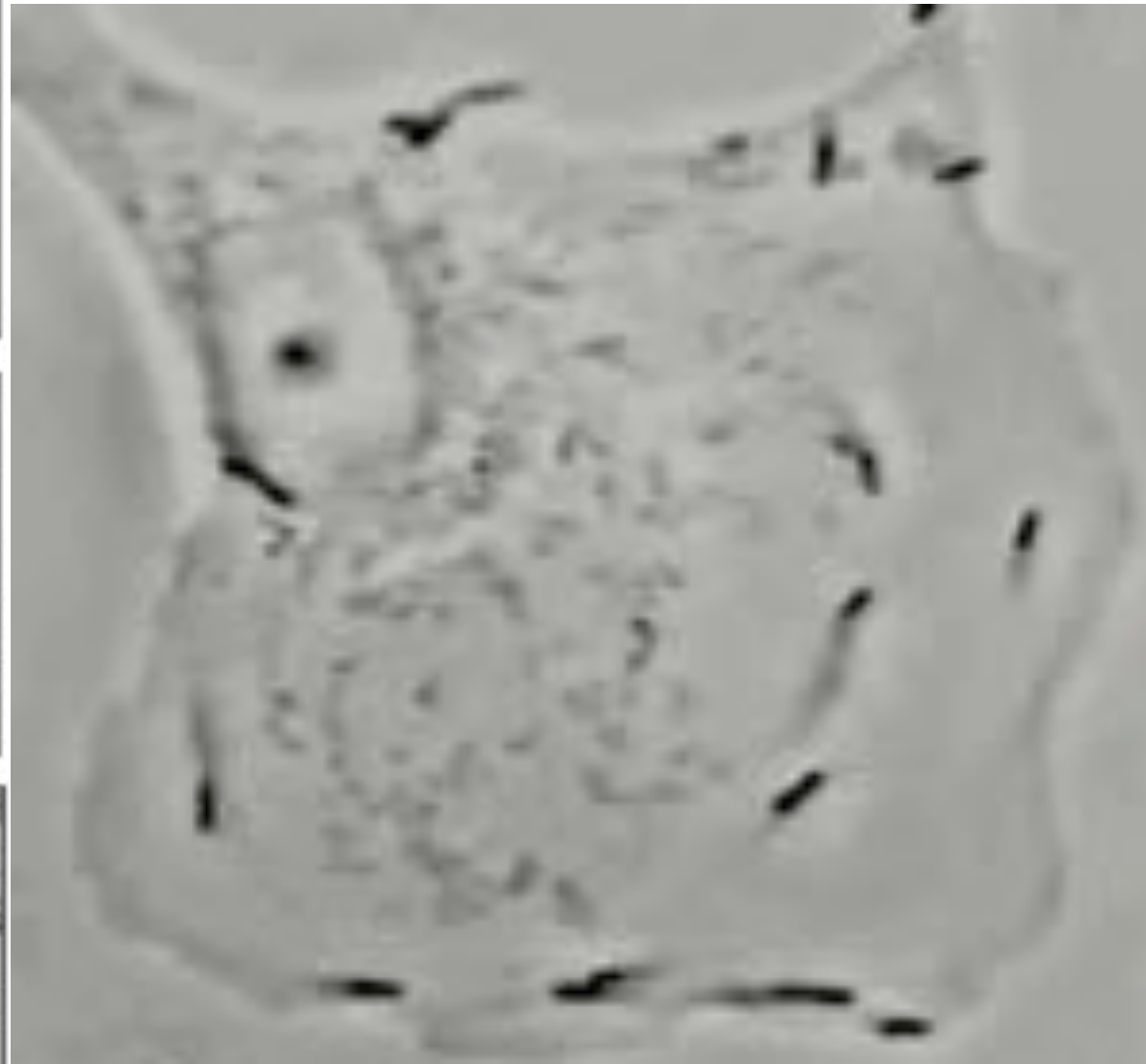


-based motility of *Listeria*,
Yersinia, and *Shigella*.

Electron micrographs of actin tails
decorated with fragment S1 of myosin!



The invasion and cell
migration of **LISTERIA**



Inducers → Sensors → Mediators → Effectors

**Many endogenous inducers activate or are "alarmins"
or
DAMP, danger-associated molecular patterns!**

Currently known alarmins include **defensins, cathelicidins, eosinophil-derived**

neurotoxin, lactoferrin, some high-mobility group (HMG) proteins, granulysin,

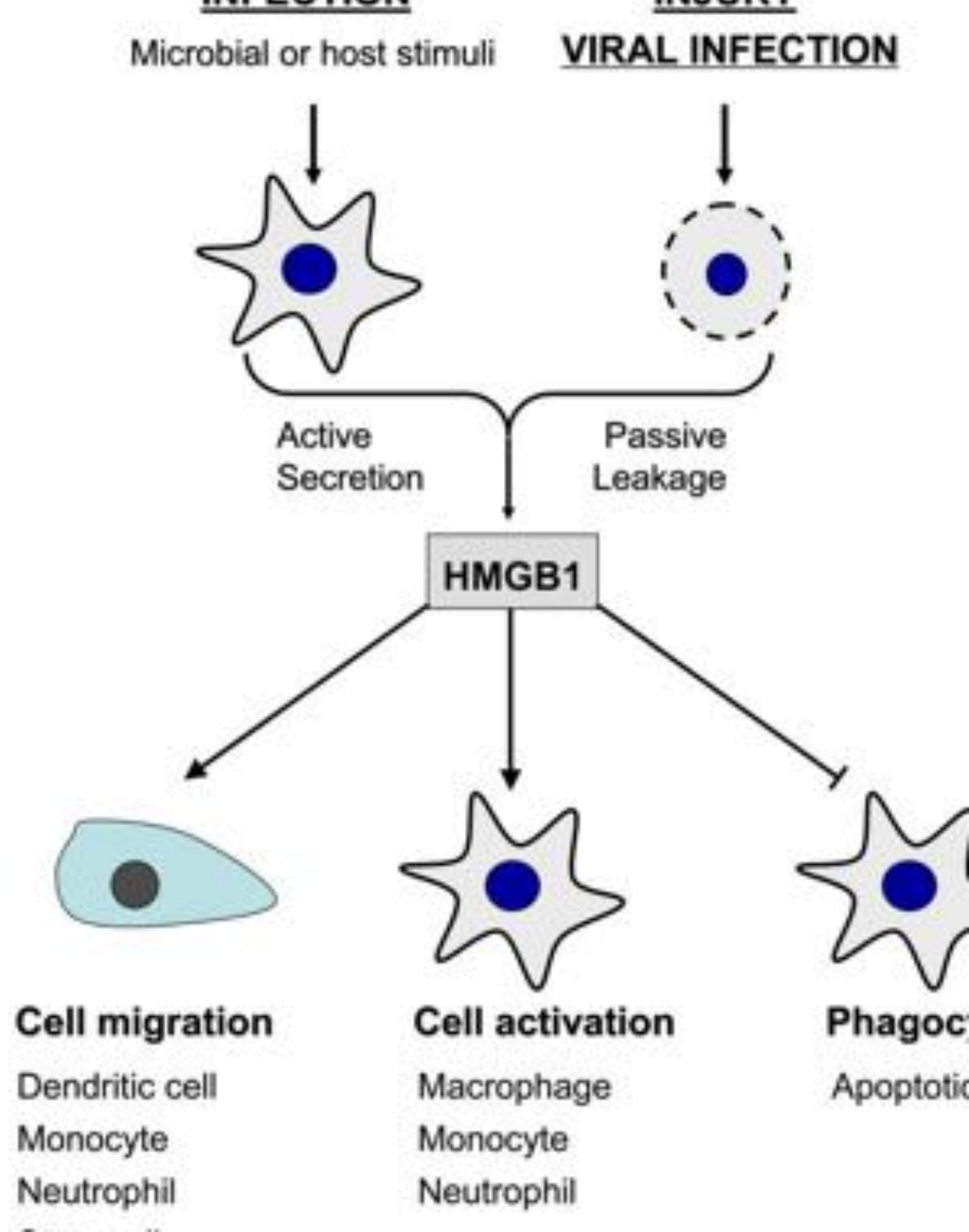
probably also **ATP** and **histamine**, while endogenous mediators that may eventually

be alarmins include some members of the **S100 family proteins, heat-shock**

proteins, and certain degraded products of extracellular matrix (e.g. **hyaluronan and**

chondroitin sulfate).

HMGB1 is actively secreted by immune cells in response to various microbial products (e.g., CpG-DNA) or endogenous stimuli (TNF, IFN- γ , or hydrogen peroxide), and passively released by damaged or virus-infected cells. Extracellular HMGB1 sustains an inflammatory response by stimulating migration of immune cells, facilitating recognition of bacterial products, activating various innate immune cells, and suppressing apoptosis of apoptotic cells. HMGB1 can function as an alarm signal to recruit, alert and activate various innate immune



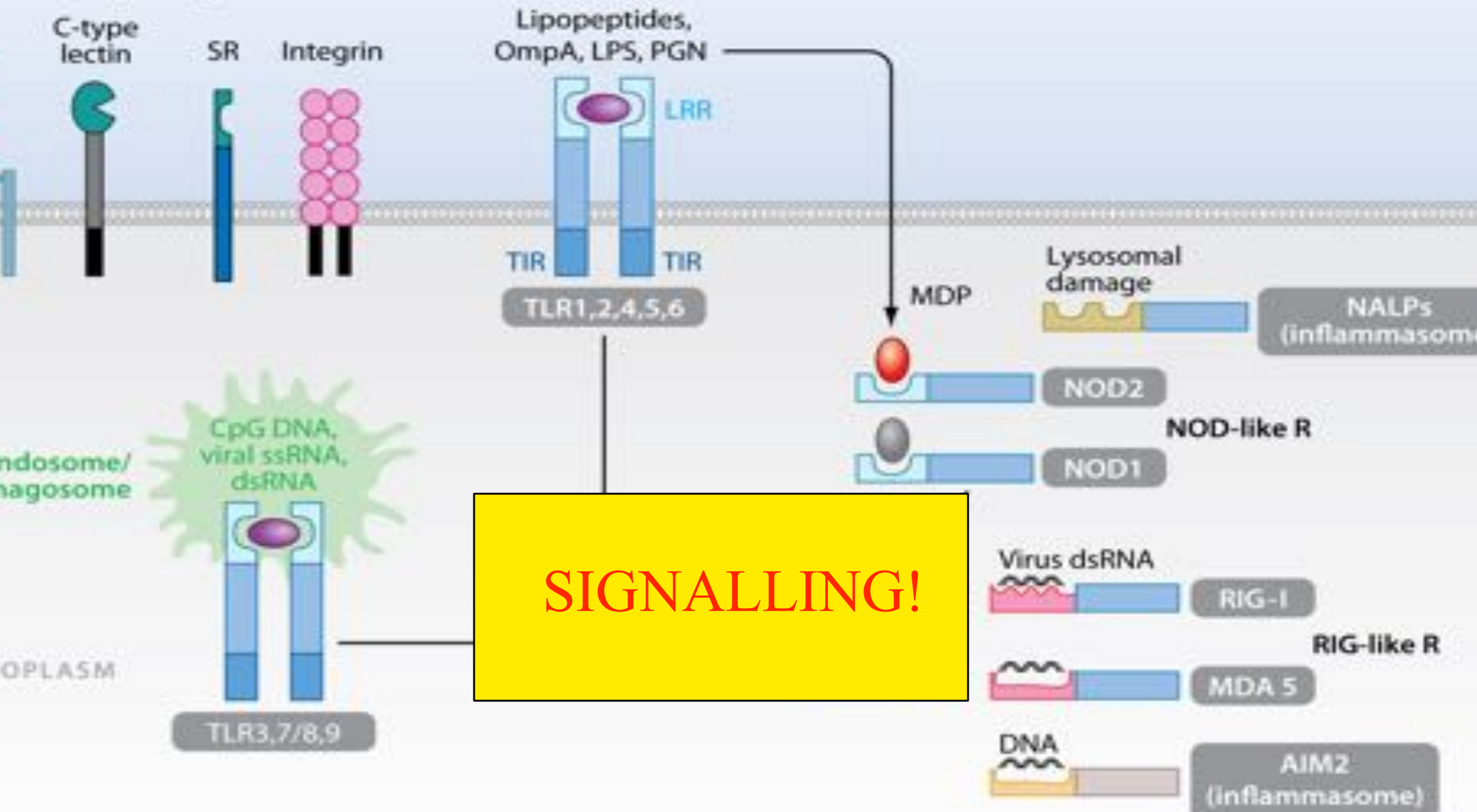
can activate inflammatory and the immune response through **RECEPTORS!**

Our body feels the damage (mainly from biological, chemical and physical stimuli) through **RECEPTORS!**

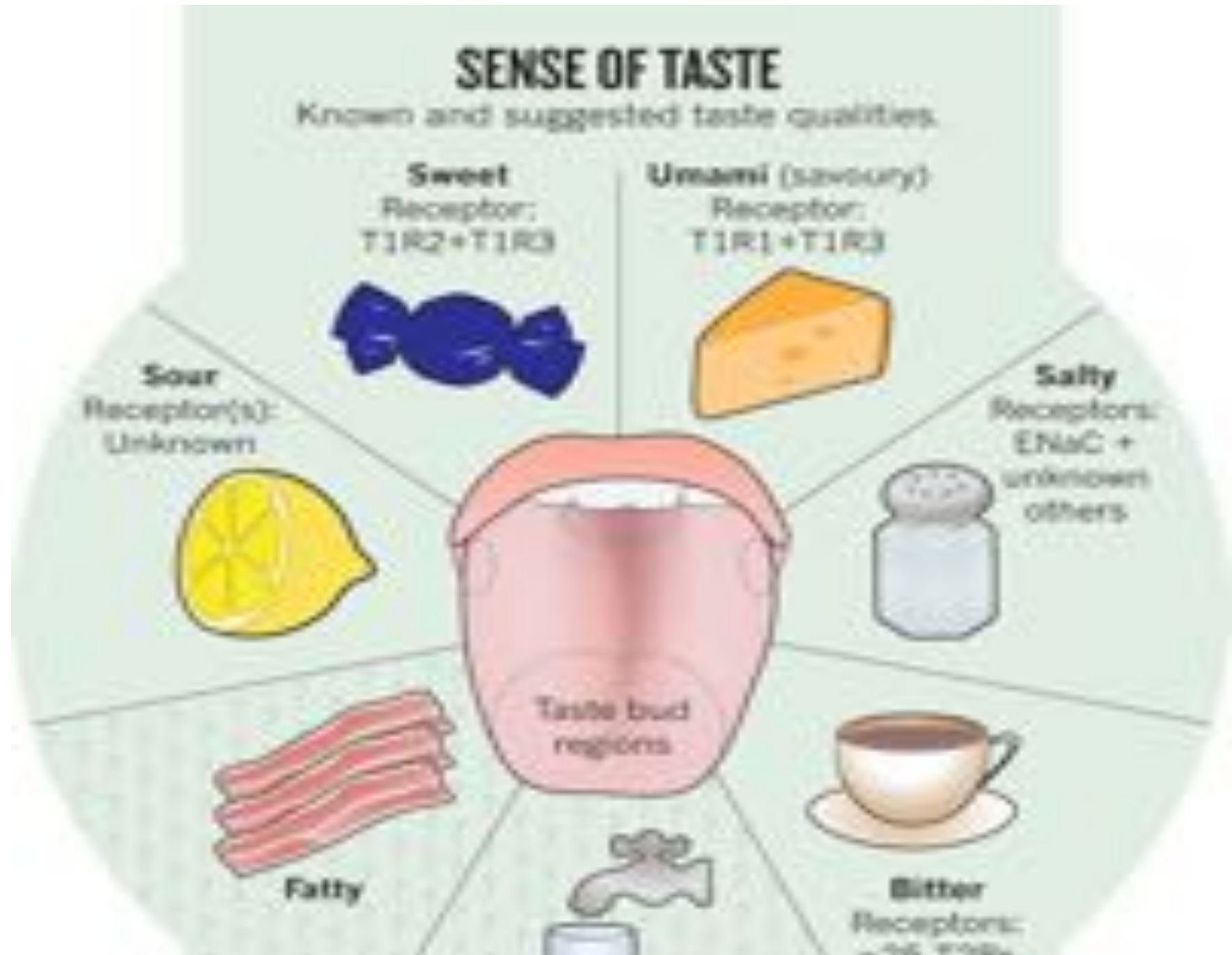
The **MAIN RECEPTORS** of the damage (by stimuli biologicals, chemicals, physicals etc) are:

- **MEMBRANE RECEPTORS**

DAMAGE RECEPTORS OF NATURAL IMMUNITY AND INFLAMMATION!



INFLAMMATION!



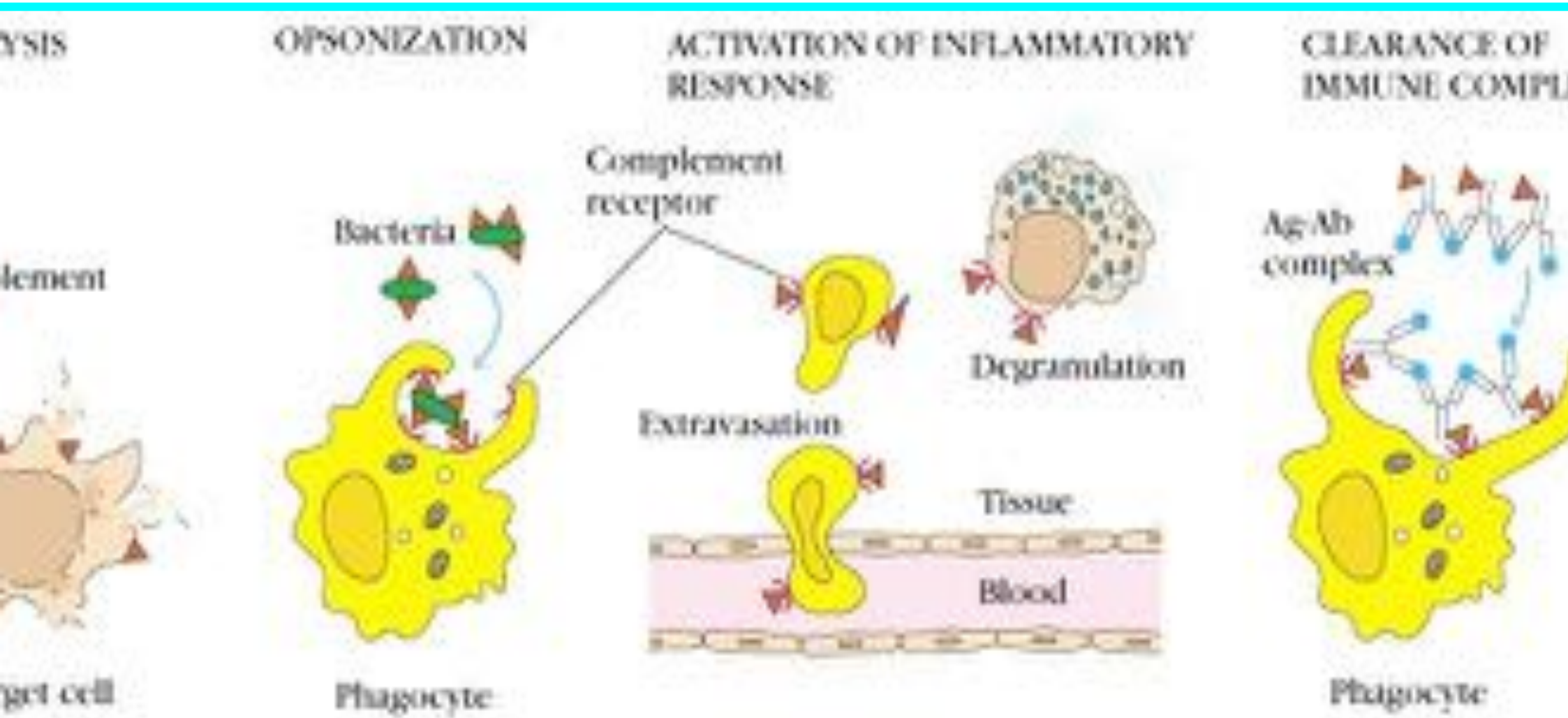
Respiratory innate immunity.

Taste receptors (T2Rs) are emerging as novel regulators of innate immunity in the respiratory tract. They are expressed in respiratory ciliated cells and in solitary chemosensory cells (SCCs), which contain the T1R2 and T1R3 subunits comprising the human sweet taste receptor. Activation of these

T2Rs are expressed in ciliated cells and solitary chemosensory cells of the respiratory tract. Bitter chemicals released by microbes during upper respiratory tract infections activate T2Rs and induce epithelial secretion of antimicrobial peptides, such as β -defensins 1 and 2!

INFLAMMATION ACTIVATE AND REGULATE:

- THE COMPLEMENT SYSTEM!



D INFLAMMATION ACTIVATE AND REGULAT

- **THE PHAGOCYTOSIS!**



INDUCE INFLAMMATION, ACTIVATE AND REGULATE

- **THE 5 CARDINAL SIGNS OF ACUTE INFLAMMATION (rubor, tumor, calor, dolor and functio lesa)!**



ID INFLAMMATION ACTIVATE AND REGULAT

The acute phase RESPONSE!

**FEVER, ESR and
LEUCOCYTOSIS
ARE**

THE MARKERS

OF INFLAMMATION

**APR (Acute Phase
Reactions) are fully
characterized:**

a) by neuroendocrine changes, fever, lethargy and anorexia, increased secretion of corticotropin-releasing hormone, cortisol, nitric oxide and decreased secretion of growth insuline-like factor;

b) hematopoietic modifications such as anemia, leukocytosis, thrombocytosis;

**c) metabolic changes such
muscle loss and negative
nitrogen balance, impaired
gluconeogenesis, osteoporosis,
increased hepatic lipogenesis,
increased lipolysis in adipose
tissue, cachexia;**

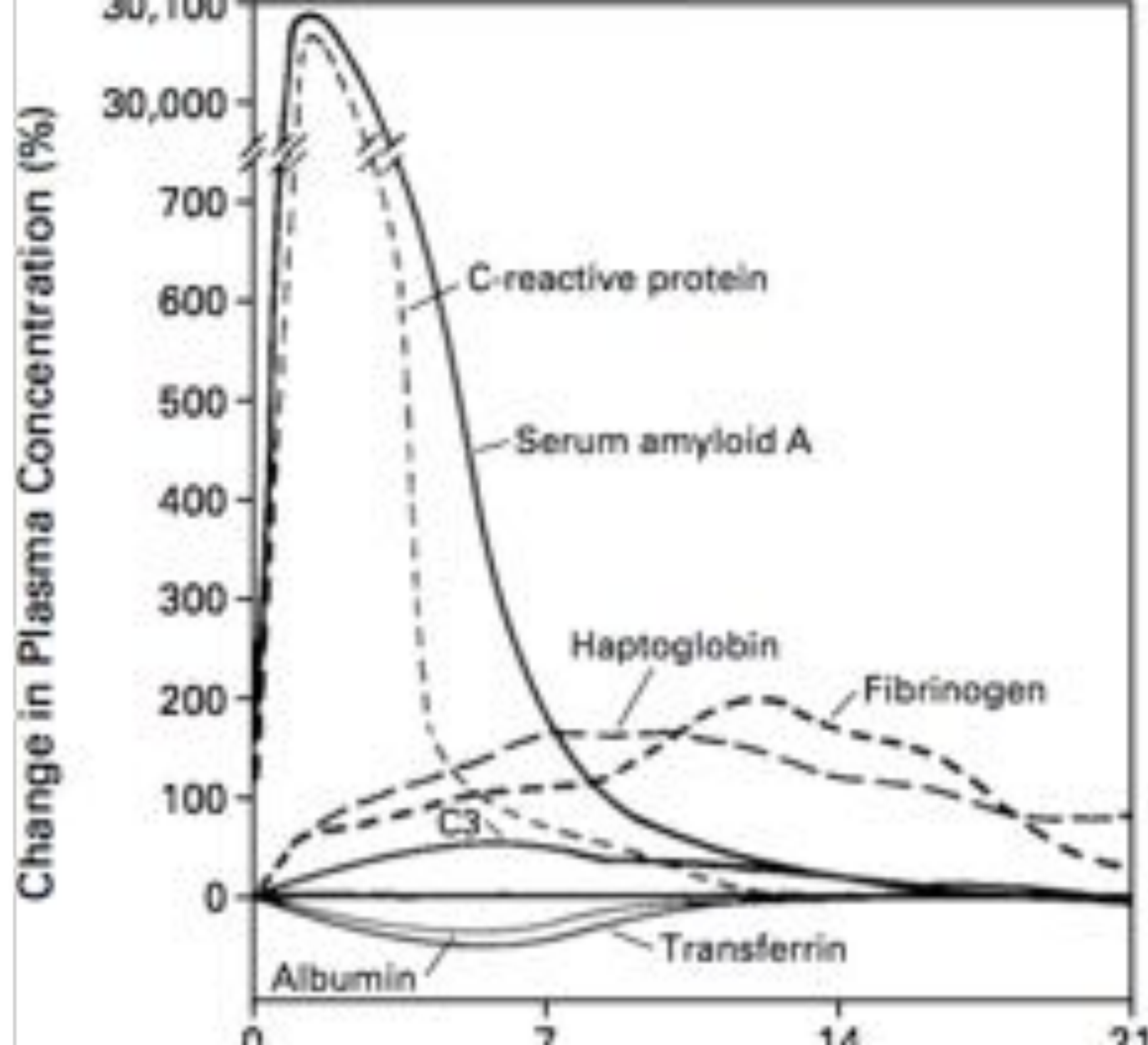
d) plasma modification of some metals such as calcium, iron and zinc, and vitamins, but mostly of some **proteins and lipoproteins!**

The acute phase proteins or APP!

The study of APP was born with the identification of the first APP, the C-reactive protein, or CRP, with the subsequent resurgence of research activities from the 1930s until today. The discovery of PCR was made by Tillet and Francis in 1930 with the publication of a paper titled "Serological Reactions In Pneumonia With Somatic Nonprotein Fraction Of Pneumococcus" in the Journal of Experimental Medicine:

Tillett WS, Francis T: Serological reactions in pneumonia with a non-protein somatic fraction of

er was observed, in addition to the CPR, the parallel increase in concentration of other plasma proteins including the Amyloid Protein A or SAA, the fibrinogen, the C3 complement component, the alpha-1-macroglobulin, the alpha-2-macroglobulin, the alpha-1-antitrypsin and more proteins!



Today we define as **acute phase protein (APP)** that protein whose plasma concentration increases (**positive APP**) or decreases (**negative APP**) by at least 25% during the acute phase reaction!

the positive and negative A.P.R.

Positive acute phase reactants (concentrations increase with acute inflammation)

Immune-related

Complement (C')

Mannose-binding lectin (MBL)

C-reactive protein (CRP)

Orosomucoid (alpha-1 acid glycoprotein)

Antiproteases (anti-enzymes)

Alpha-1 antitrypsin (A1-AT)

Alpha-2 macroglobulin (A2M)

Anti-oxidants

Ceruloplasmin

Coagulation factors

Fibrinogen

Factor VIII

Others

Haptoglobin

Serum amyloid A (SAA)

Plasma fibronectin

Lipopolysaccharide-binding protein (LBP)

Ferritin

Negative acute phase reactants (concentrations decrease with acute inflammation)

positive and negative APP!!!

Positive APP:

Short pentraxins;
Collectins;
Proteins of the complement system;
LPS binding protein or LBP;
Proteins of the coagulation system and fibrinolysis;
Antiproteases;

Negative APP:

- a) Albumin;
- b) Transferrin;
- c) Transthyretin;
- d) Alpha-fetoprotein;
- e) Thyroxin-binding globulin;
- f) Factor XII.

Negative APP:

albumin;
transferrin;
transthyretin;
alpha-fetoprotein;
tyroxin-binding globulin;
Factor XII.

The negative APP do diminish just because **hyper-production** of positive APP which are limiting an acid reserve or because **escaping** from the vessel the inflammatory exudate also for **anorexia** increased energy and **pro-catabolism** that arise in course of inflammation. Among the negative APP

Albumin is a single polypeptide which consists of 585 amino acids with a molecular weight of about 69 kDa. **The total pool of albumin is 4-5 g/kg of body weight and 40-45% is in the intravascular space and the other 60 % is in the interstitial space.**

Physiological functions of albumin are impressive:

Maintains 75-80 % of the plasma colloid osmotic pressure;

Stores and transports not only free fatty acids, calcium, certain steroid hormones, thyroxine, bilirubin, copper and tryptophan, but also drugs, such as penicillin, salicylic acid, and non-steroidal antiinflammatory drugs (NSAIDs);

Acts as an important source of sulfhydryl groups, which remove nitrogen and oxygen free radicals and other toxins; in this context, the antithrombotic and anticoagulant properties of albumin may be due to the uptake of the free radical nitric oxide (NO).

Concentration of albumin in the blood (serum albumin) varies between 3.5-5.0 g/dl and its decrease during inflammation can be significant even if it is non-specific, as the hypoalbuminemia may occur in various physiopathological conditions, such as **rheumatoid arthritis, cholecystitis, acute ulcerative colitis,**

Transferrin is the major β -globulin that transports iron (siderophilin). The transferrin contains 687 amino acids and has a calculated molecular weight of approximately 79 kDa. The transcription of the mRNA for the synthesis of transferrin in the liver is regulated by the concentration of iron in the hepatic plasma.

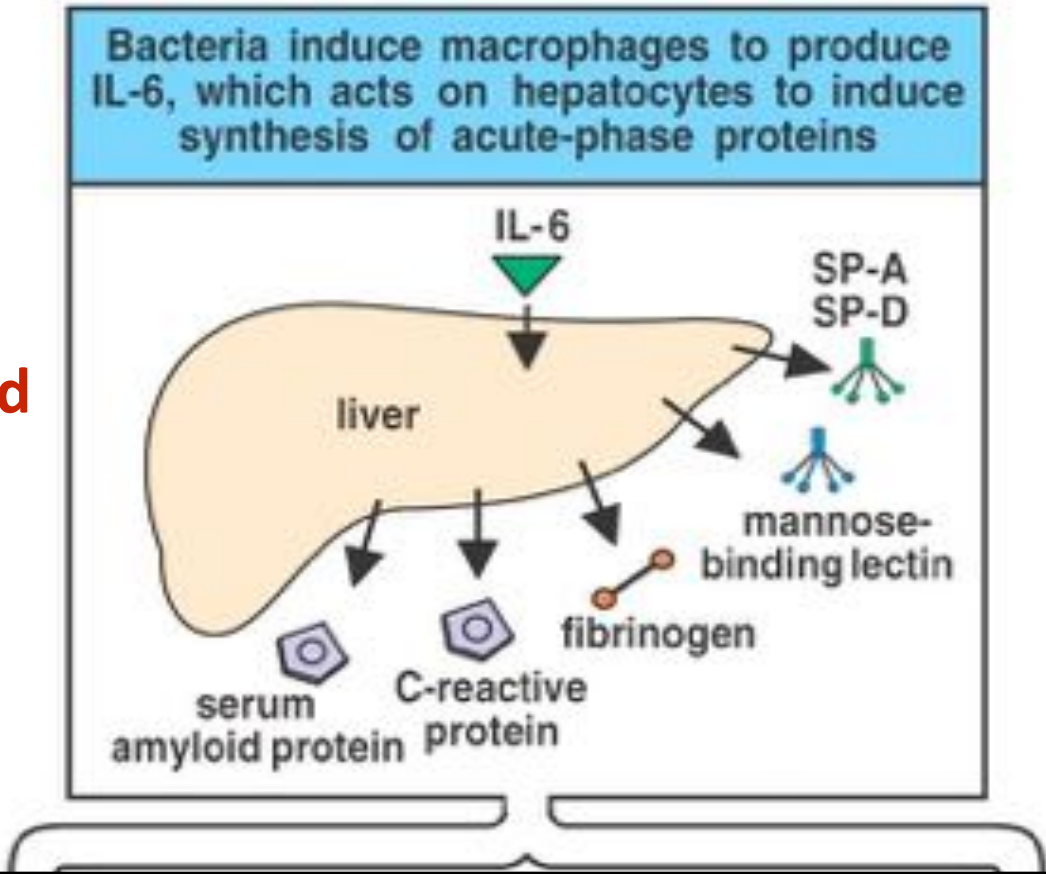
Normal transferrin levels in the blood are 200-360 mg/dL.

Transferrin levels increase during the use of birth control pills, during pregnancy and in cases of insufficient levels of iron and are increased by acute and chronic inflammatory diseases, especially for malnutrition, treatment with iron or with steroids, liver disease and nephrotic syndrome.

Electrophoretic variants of transferrin in serum are found occasionally.

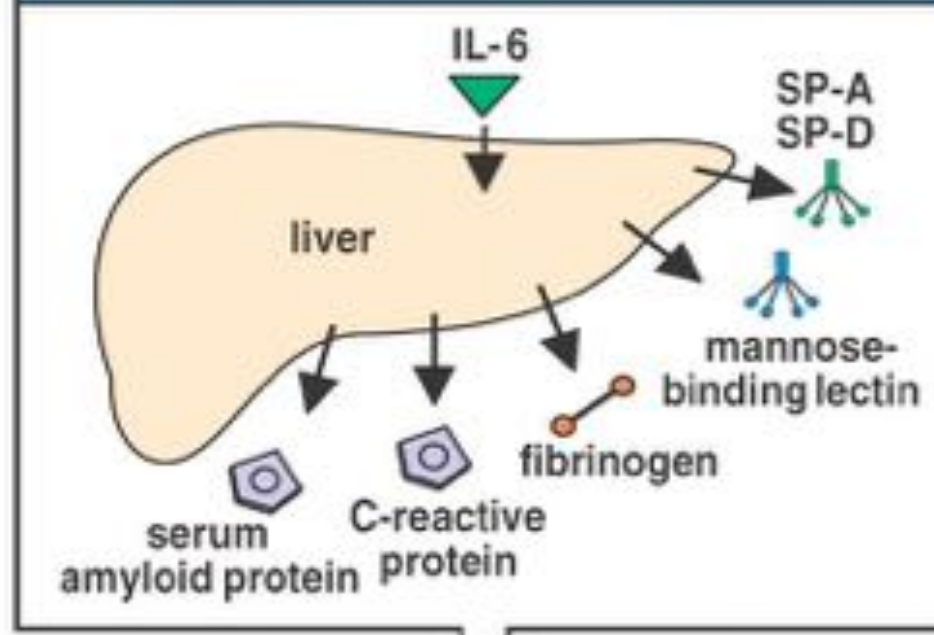
INDUCTION FROM THE LIVER:

short pentraxins;
lectins;
proteins of the complement system;
LPS binding protein or LBP;
proteins of the coagulation system and
fibrinolysis;
serine proteases;
transport proteins.



Inflammatory cytokines bind to specific receptors on hepatocytes and induce the activation of **Janus family kinases (JAK)**, the **STAT (signal transducers and Activators of transcription)**, and inflammatory transcription factors such as **NF- κ B**. Transcription factors most characteristic of acute phase response regulate the synthesis of APP belong to the family of leucine zipper, C/EBPA (C/EBP α), **C/EBP β** (C/EBP β), **C/EBP δ** (C/EBP δ) and **NF-IL6 (nuclear factor associated with IL-6)**, the transcription by binding to a site called **bZIP1**. **It is also been shown that the induction of acute phase proteins is correlated with a decreased synthesis of C/EBPA and an increase of**

Bacteria induce macrophages to produce IL-6, which acts on hepatocytes to induce synthesis of acute-phase proteins



Recently, it has been proposed a new classification of APP positive according to the cytokines that induce them from the LIVER!

APP positive were therefore divided into two classes:

APP Type-1 induced by IL-1 α and β , TNF- α and β , whose prototype is **AA**;

APP-type 2, induced by IL-6, but also by IL-11, Leukemia inhibitory f

Acute phase reactant pentraxins such as CRP and SAA or amyloid protein A, also indicate acute phase reactant protein serum amyloid or SAP;

Lectins, structurally related to C1q, such as mannose-binding lectin or MBL, proteins A and D of the pulmonary surfactant or SPA and SPD and ficolins, which include L-, M- and H-ficolins;

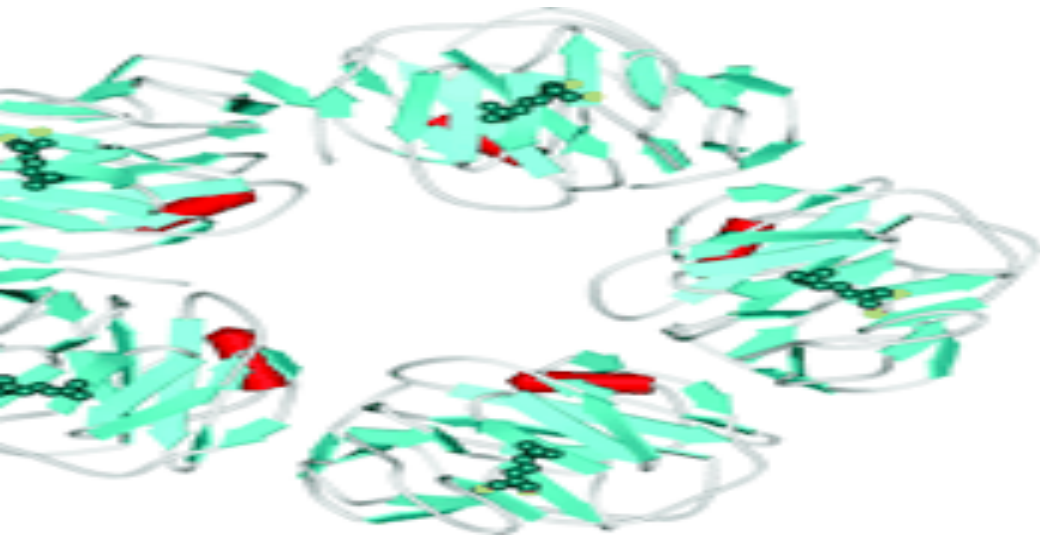
Proteins of the complement system, such as C3, C4, C9, Factor B, the inhibitor of C1q, C4b-binding protein;

S binding protein or LBP;

Proteins of the coagulation system and fibrinolysis, such as fibrinogen, plasminogen, tissue plasminogen activator, Protein S, vitronectin;

Protease inhibitors, such as α 1-antitrypsin (AAT) and the α 1 anti-chymotrypsin

- **SHORT PENTRAXINS:**



CPR

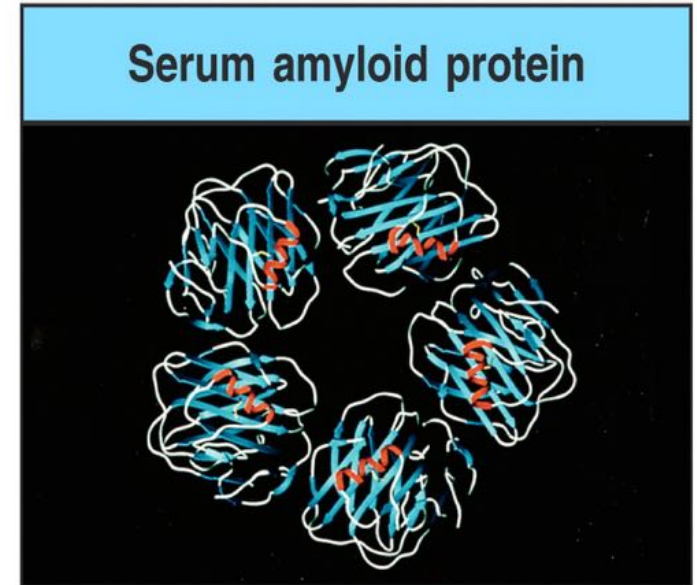
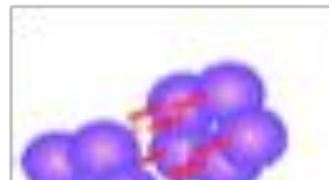


Figure 2-47 part 2 of 2 Immunobiology, 6/e. (© Garland Science 2005)

SAP

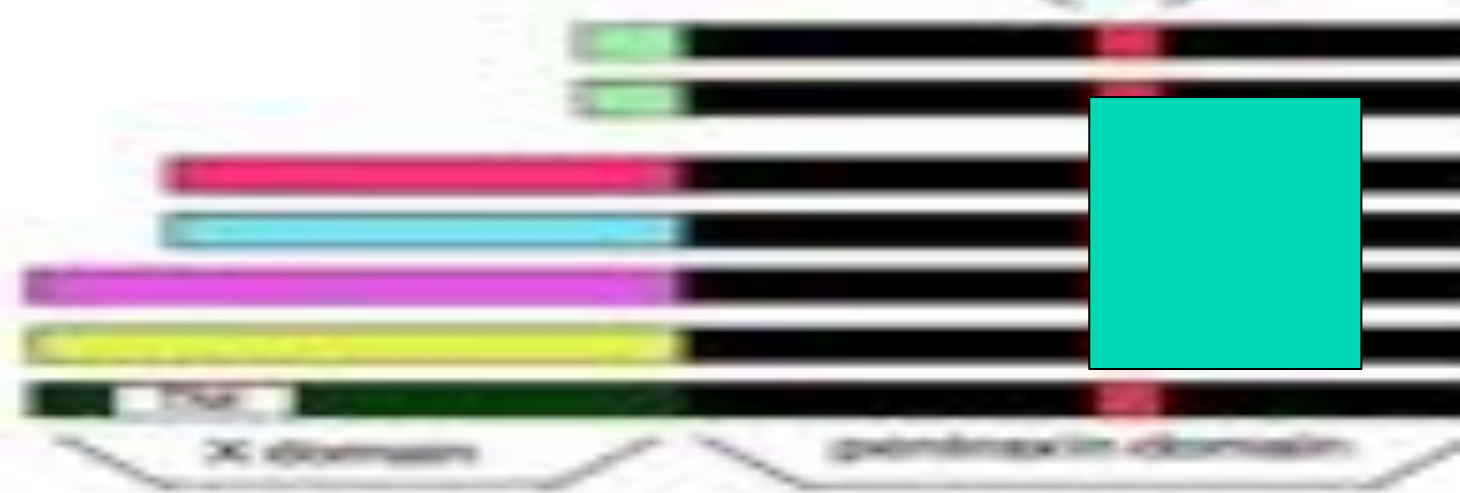
- **LONG PENTRAXINS: PTX3-PTX4**



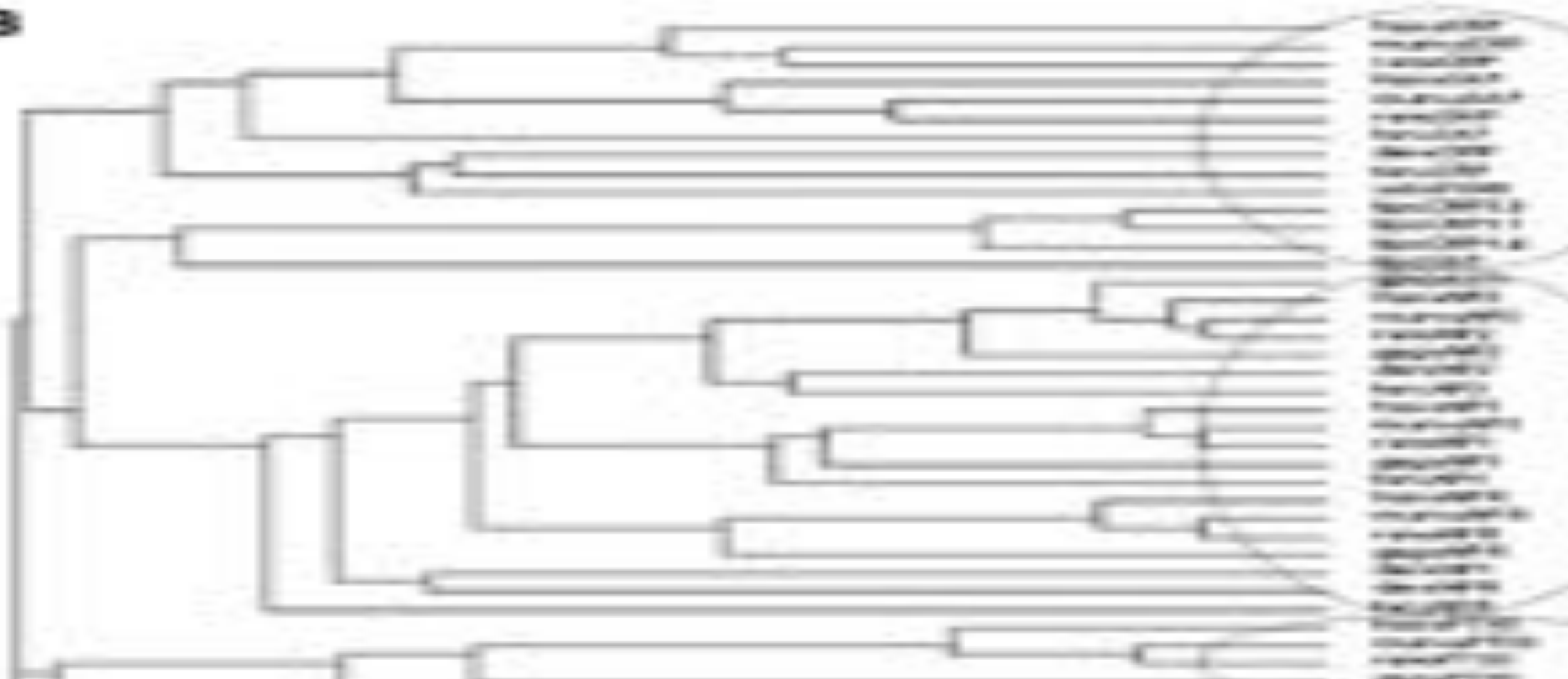
NCOR2
INTRACONS

NCOR1
INTRACONS

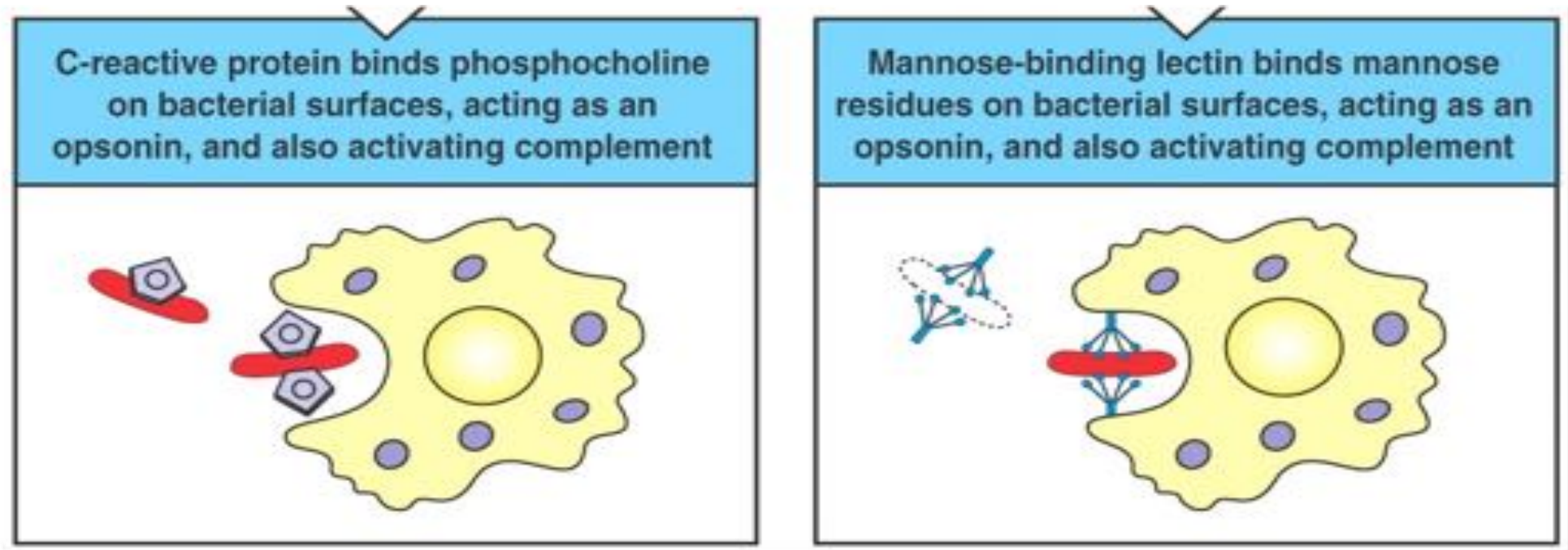
NCORP
NCORP
NPTD3
NPTD4
NPF1
NPF2
NPF3



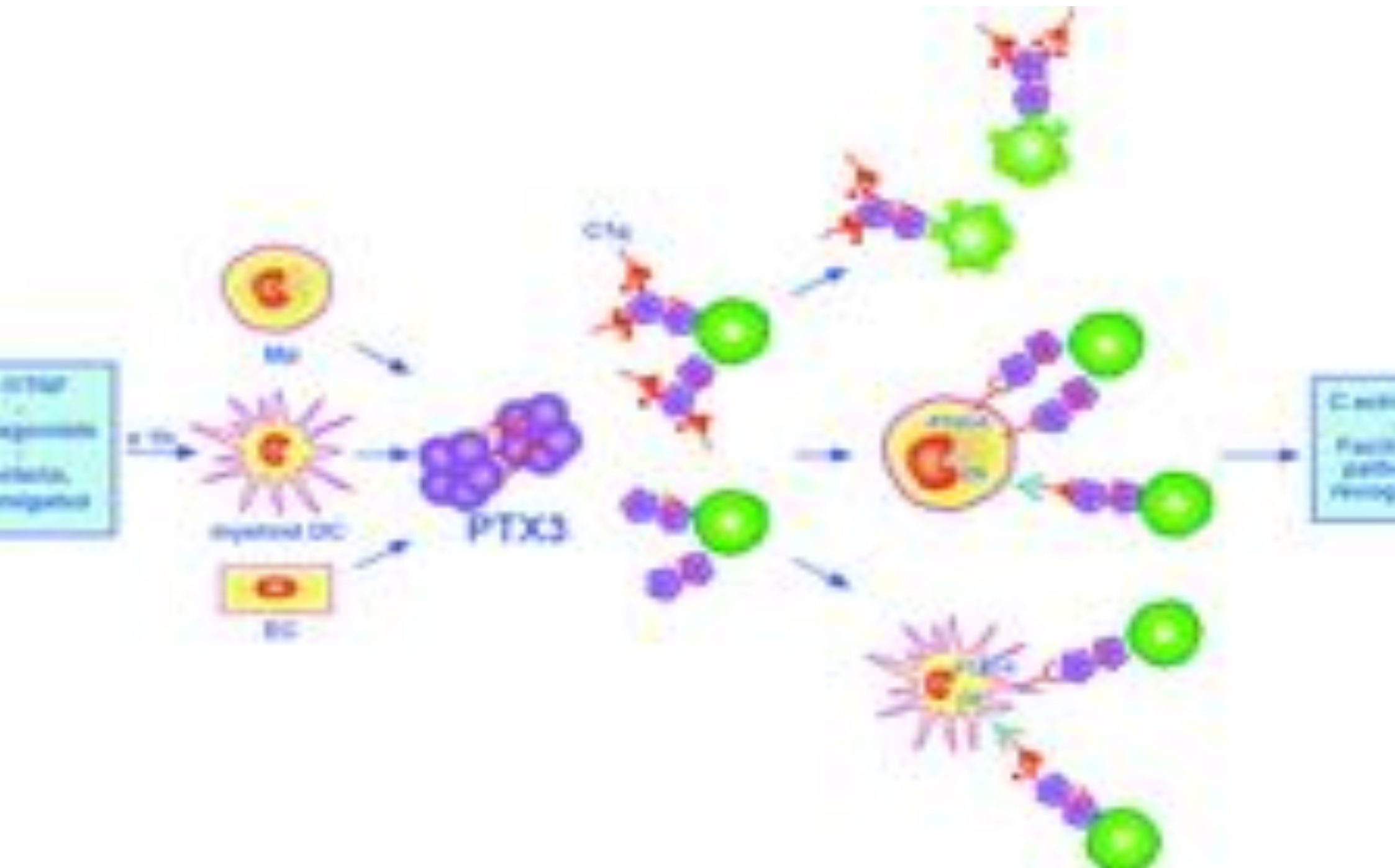
B



**of bacteria
and phosphoethanolamine
of apoptotic cells
and activate complement and phagocytosis!**

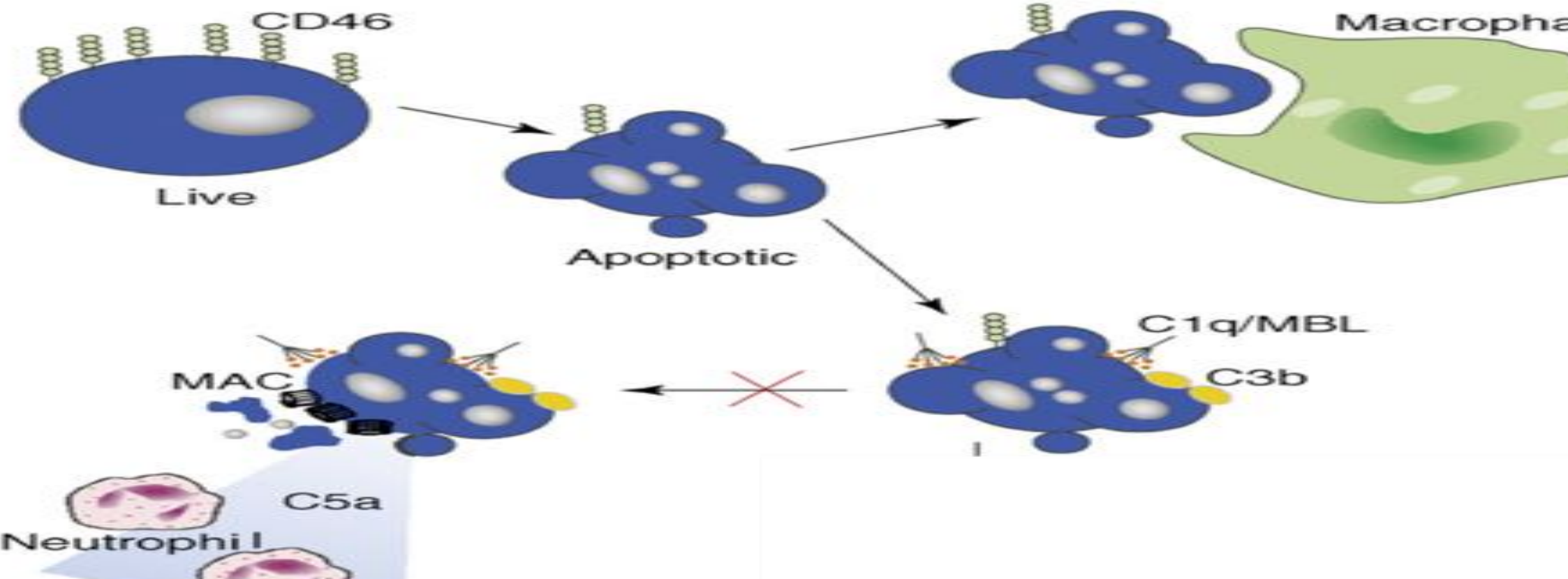


Role of the long pentraxin PTX3 in antimicrobial resistance

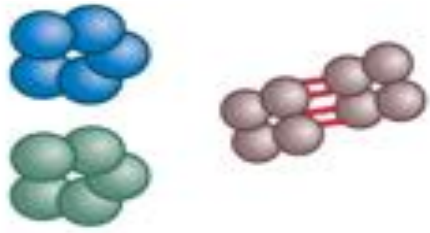


activated by
PENTRAXINS:

CLASSICAL COMPLEMENT ACTIVATION IN THE NATURAL IMMUNITY AND INFLAMMATION!

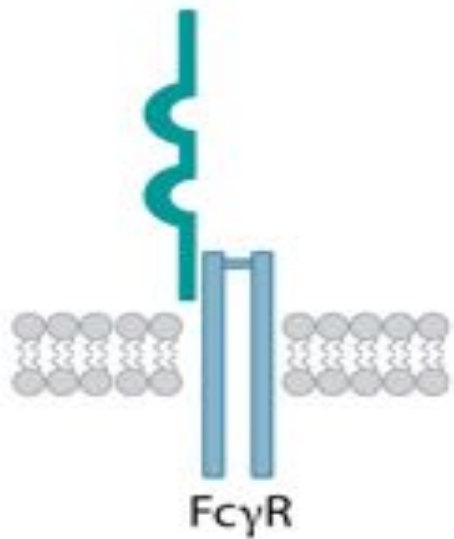


antibodies and activate phagocytosis!!

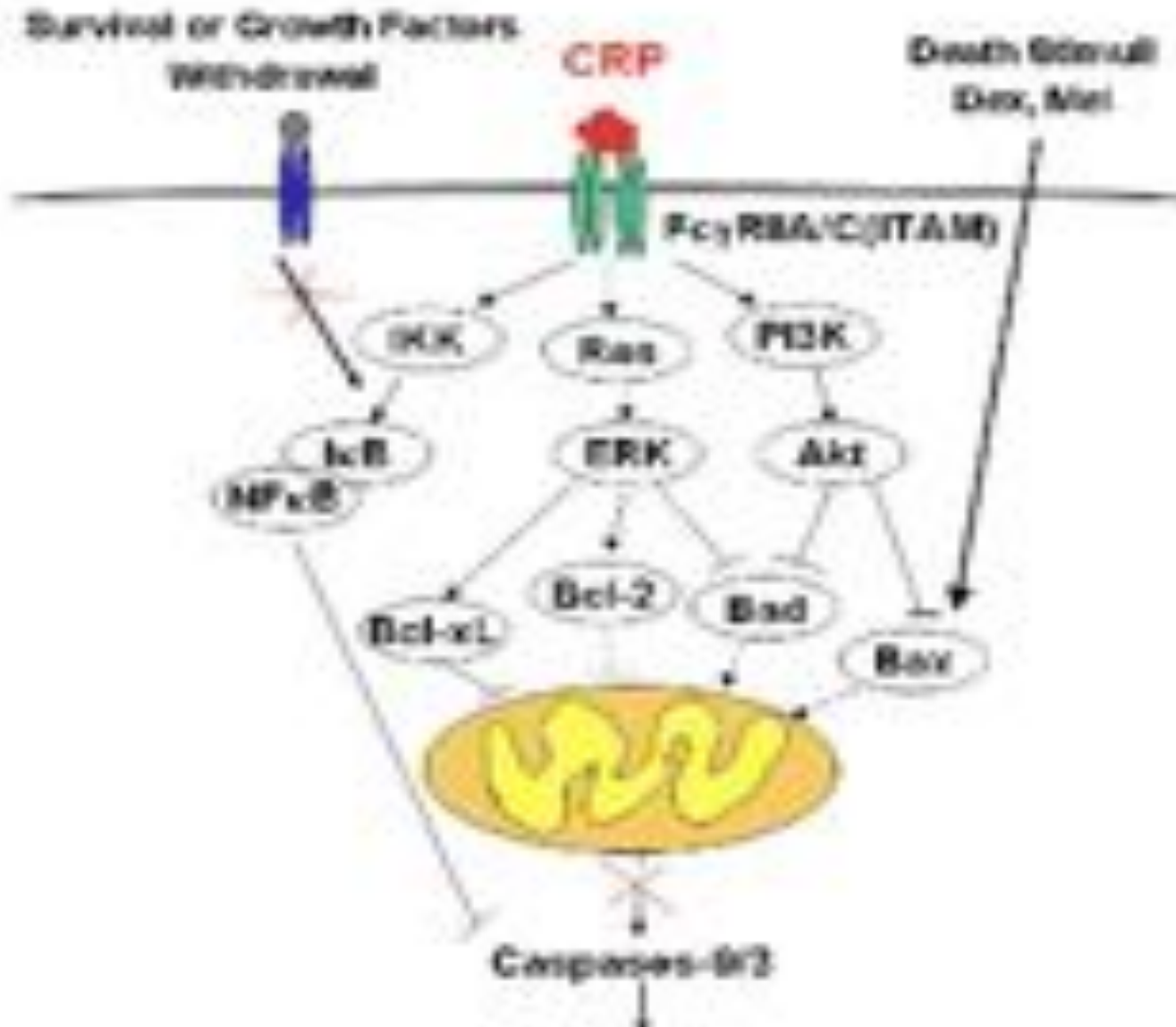


CRP/SAP

PTX3



FcγR

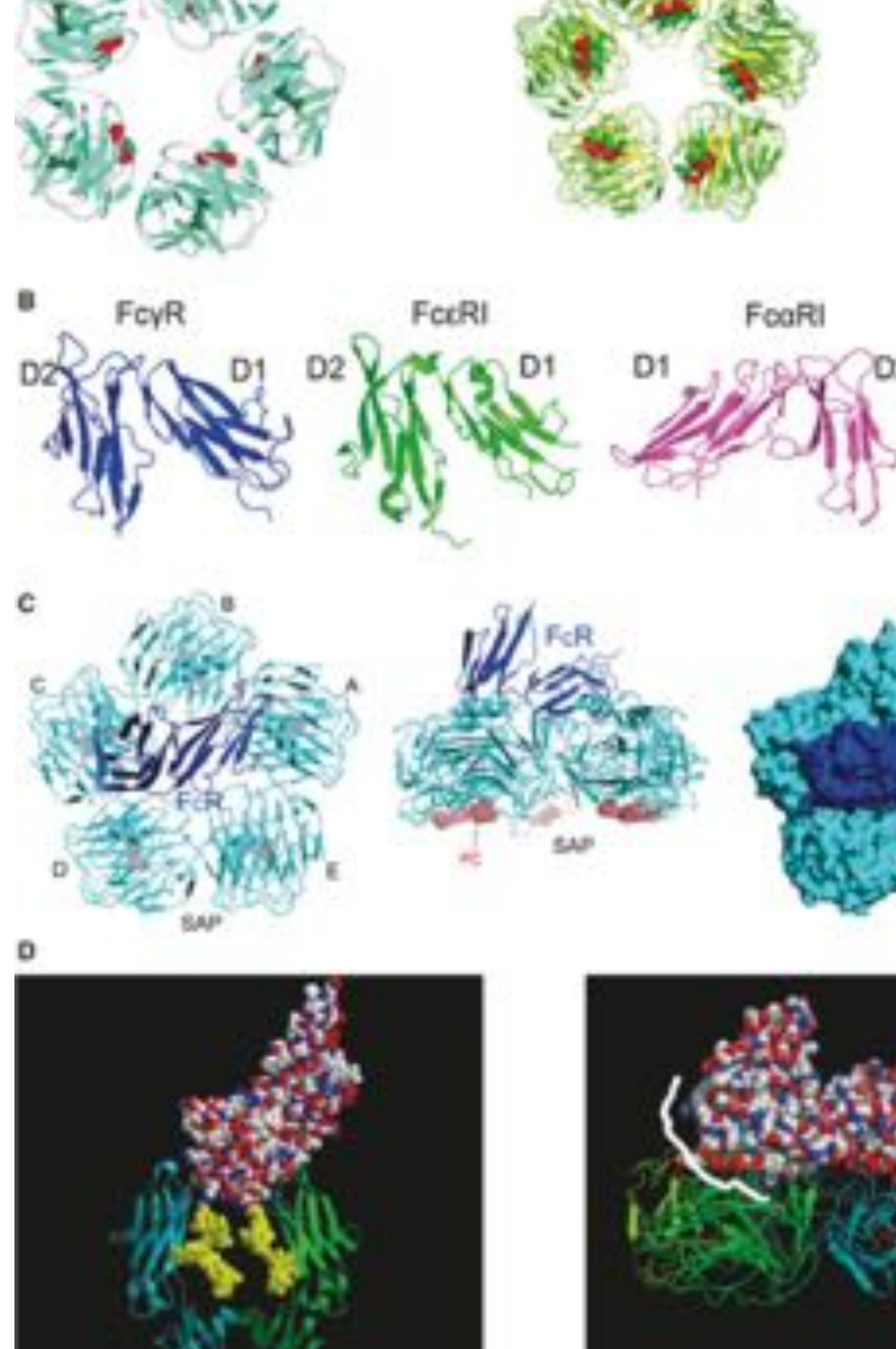


Crystal structure of phosphocholine and CRP (PDB entry 1B09, left) and structural superposition between CRP and SAP (right).

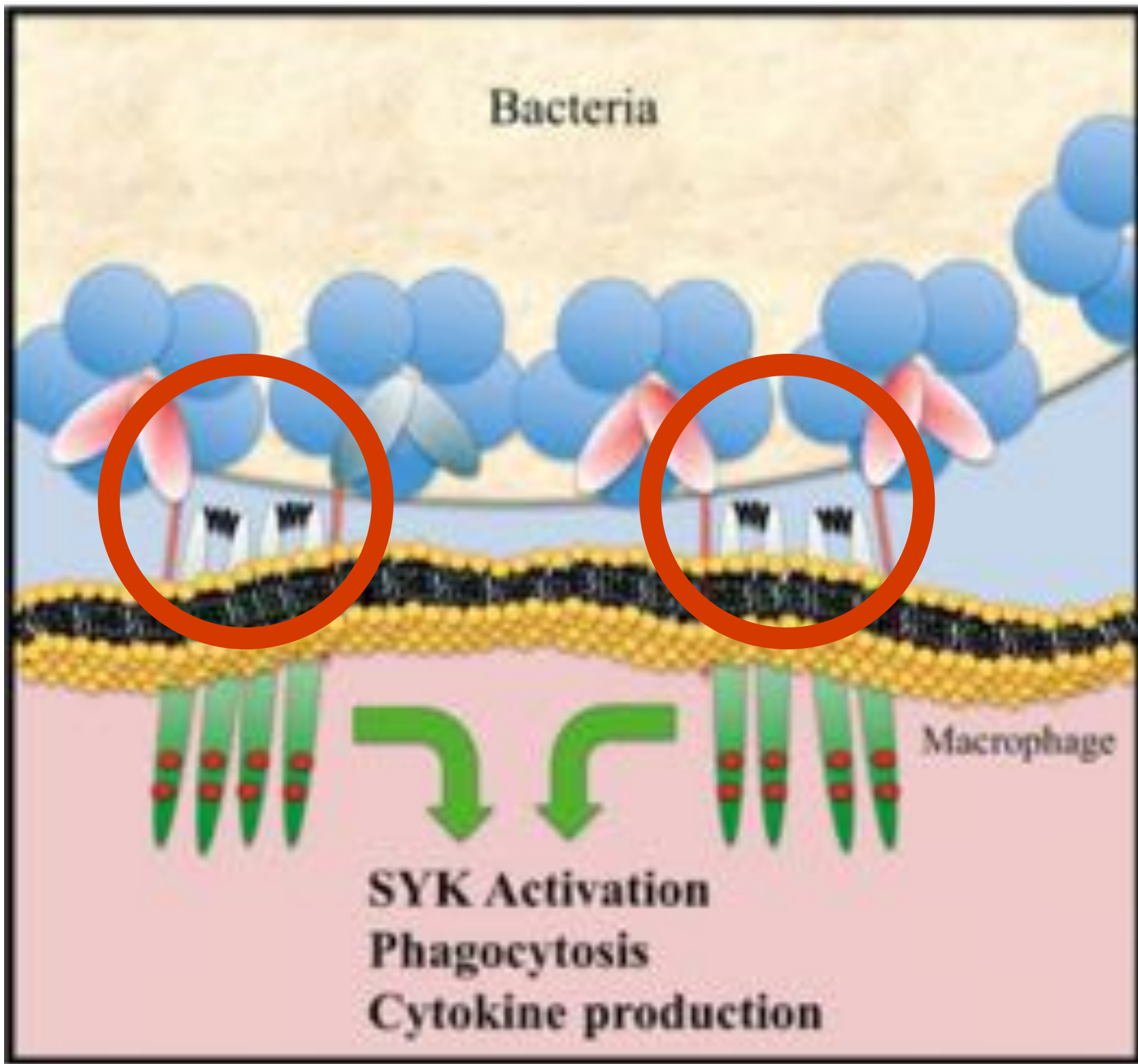
Structures of FcγRIIA, FcεRI (PDB entry 1F2Q), and FcαRI (PDB entry 1VZ).

Structural complex between human CRP (cyan) and FcγRIIA (blue) in two orthogonal views (left and middle panels) and in space filling model (right panel).

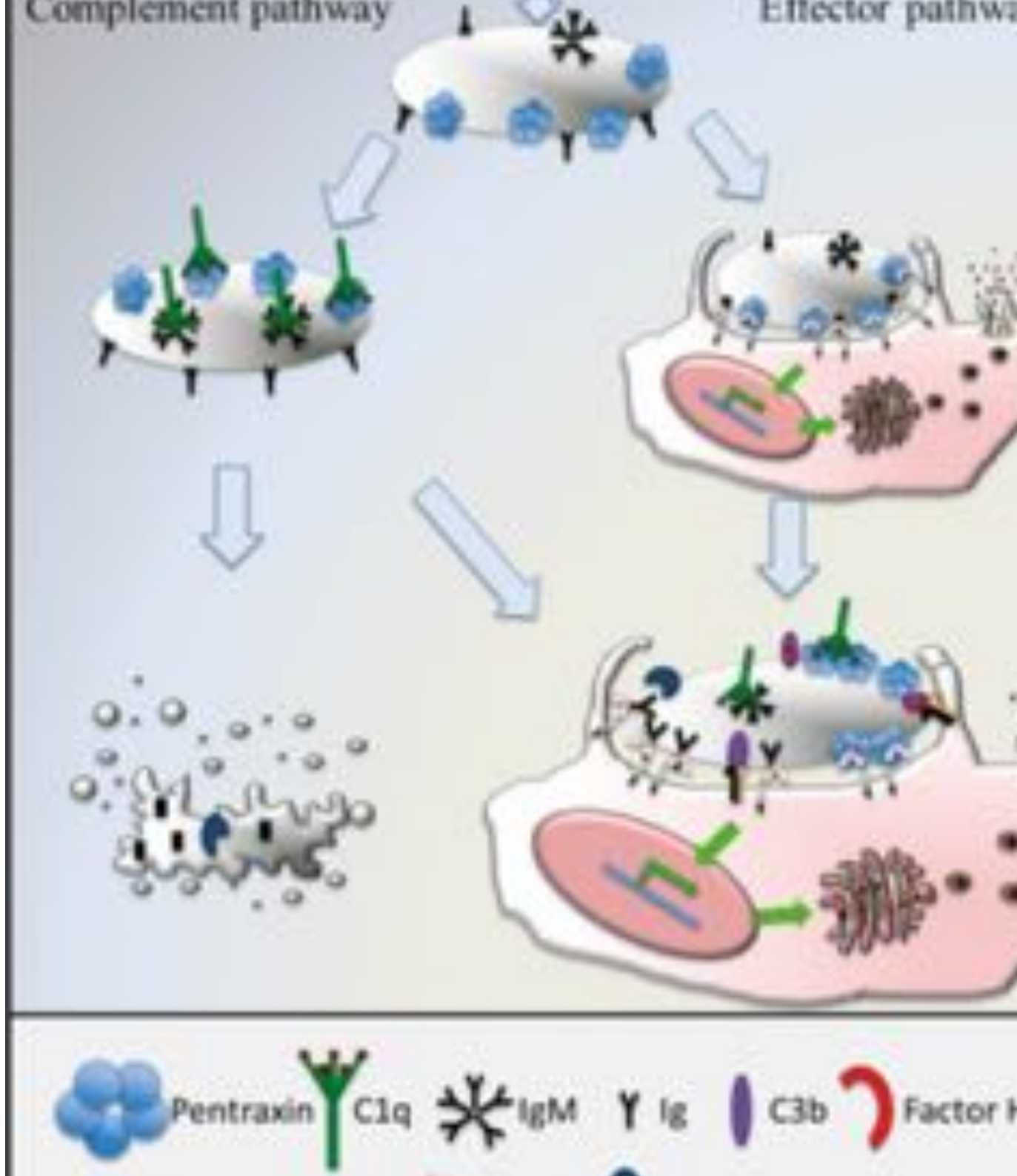
Binding mode of IgG-Fc on Fc receptor (left panel) partially overlap with that of SAP (right panel). The IgG-Fc surface region is highlighted in white on the SAP complex structure.



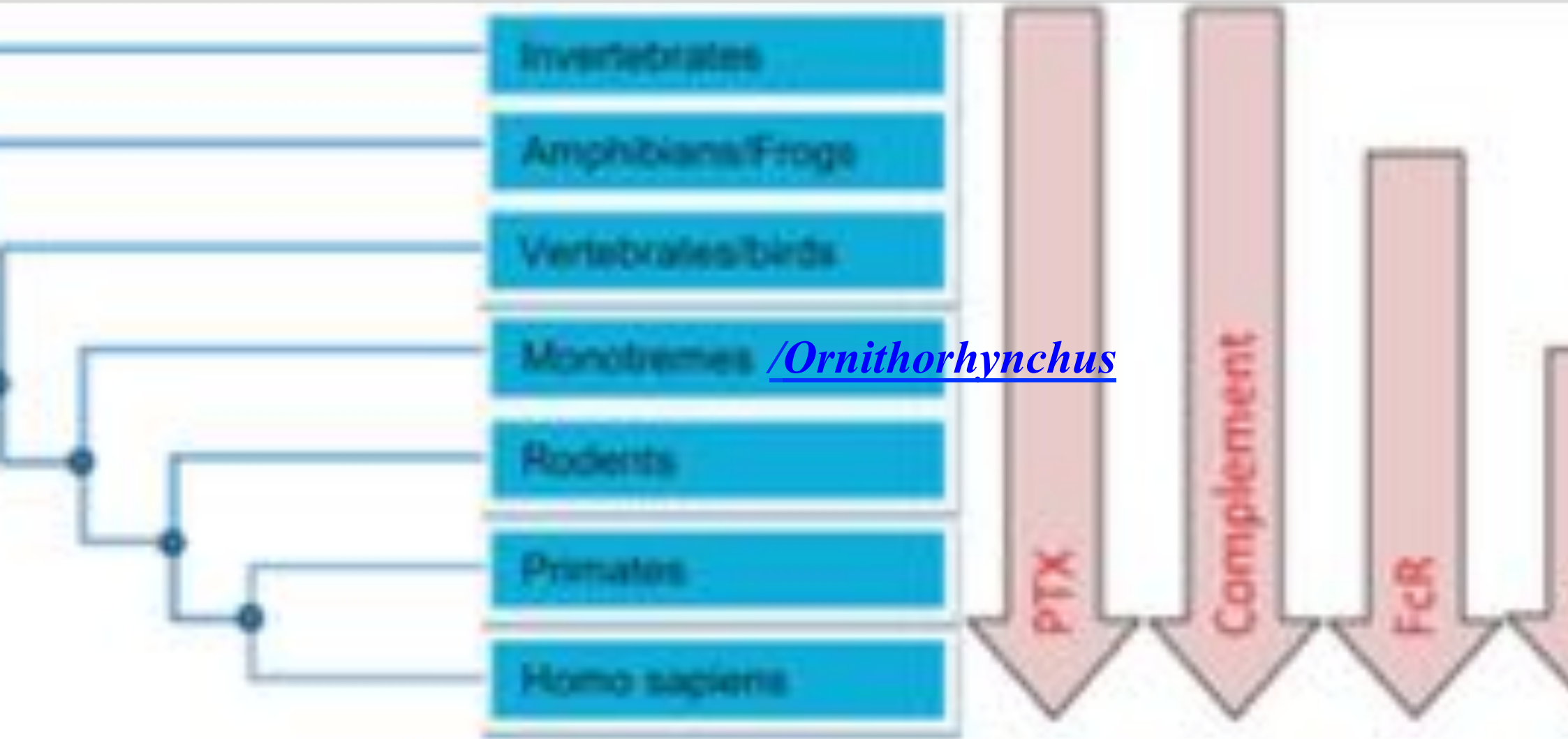
For IgG antibodies:



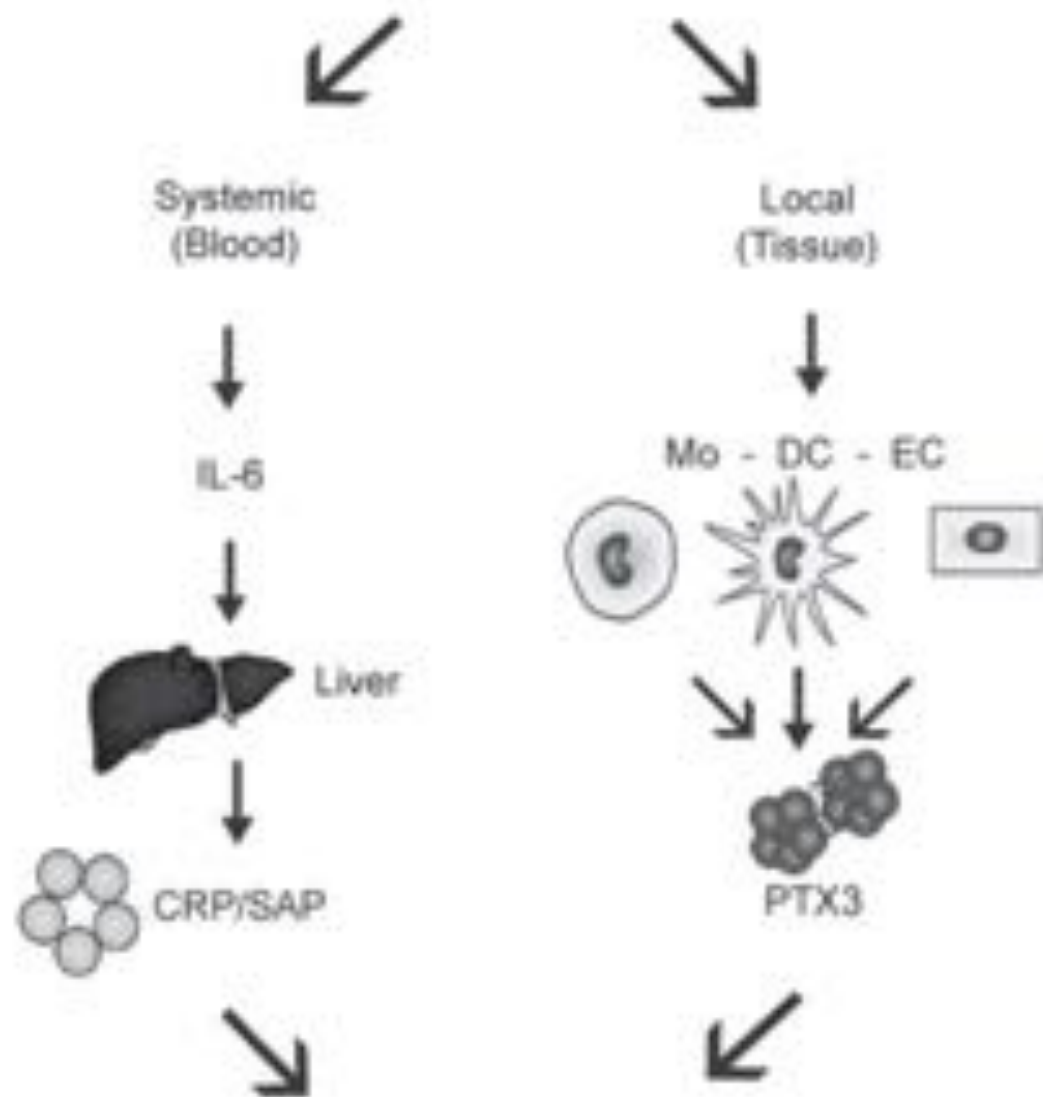
Comparison of Pentraxins and Antibodies in Complement and Fc receptor activation!



Pentraxins and Fc receptors



INFLAMMATION / MICROBIAL SENSING



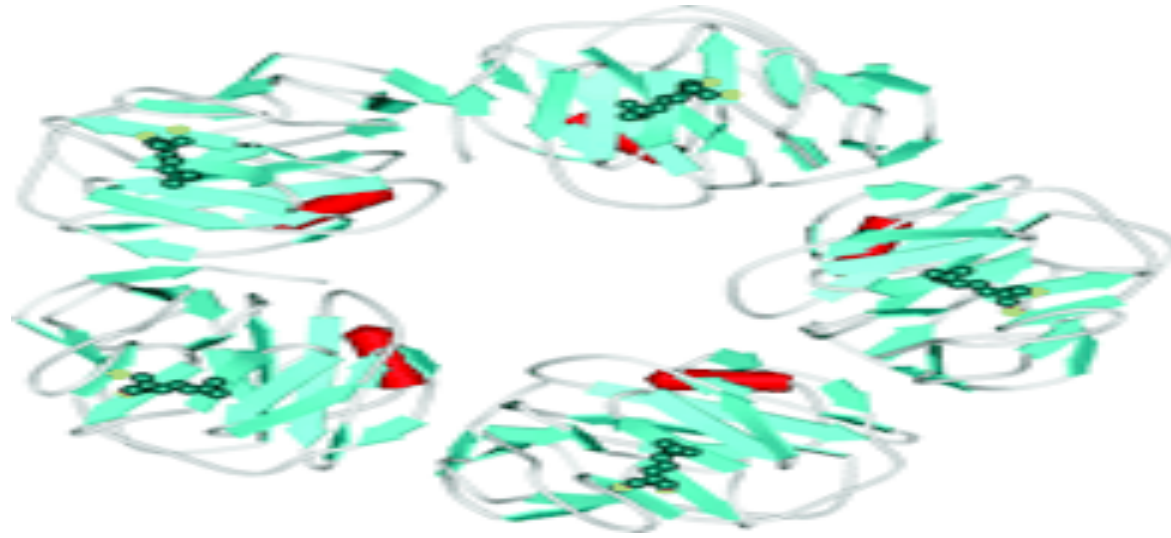
Pathogen
recognition

Complement
activation

Tissue
remodeling

Self / non self
discrimination

MAJOR POSITIVE APP!



Physiological concentration is less than 1 $\mu\text{g/ml}$ (100 ng/ml at birth, 170 ng/mL in children and from 470 to 1000 ng/mL in adults), but increases by 100-1000 times during inflammation.

Though for a long time CRP levels have been used as a quick test for the presumptive diagnosis of bacterial infection (high CRP) distinct from viral infection (low CRP), today a rise of the PCR can be observed in viral hepatitis, in bacterial acute flu-like syndromes, in TB, gout, in burns, in peritonitis, in rheumatic fever, rheumatoid arthritis, and a significant increase occurs in scarlet fever and Guillon-Barré syndrome. CRP is often used by hematologists to follow the progress or remission of autoimmune diseases and

[F](#), [Chen J](#), [Zheng R](#), [Liu H](#), [Li X](#), [Yang P](#), [Liu G](#), [Jia Y](#).

CONCLUSIONS:

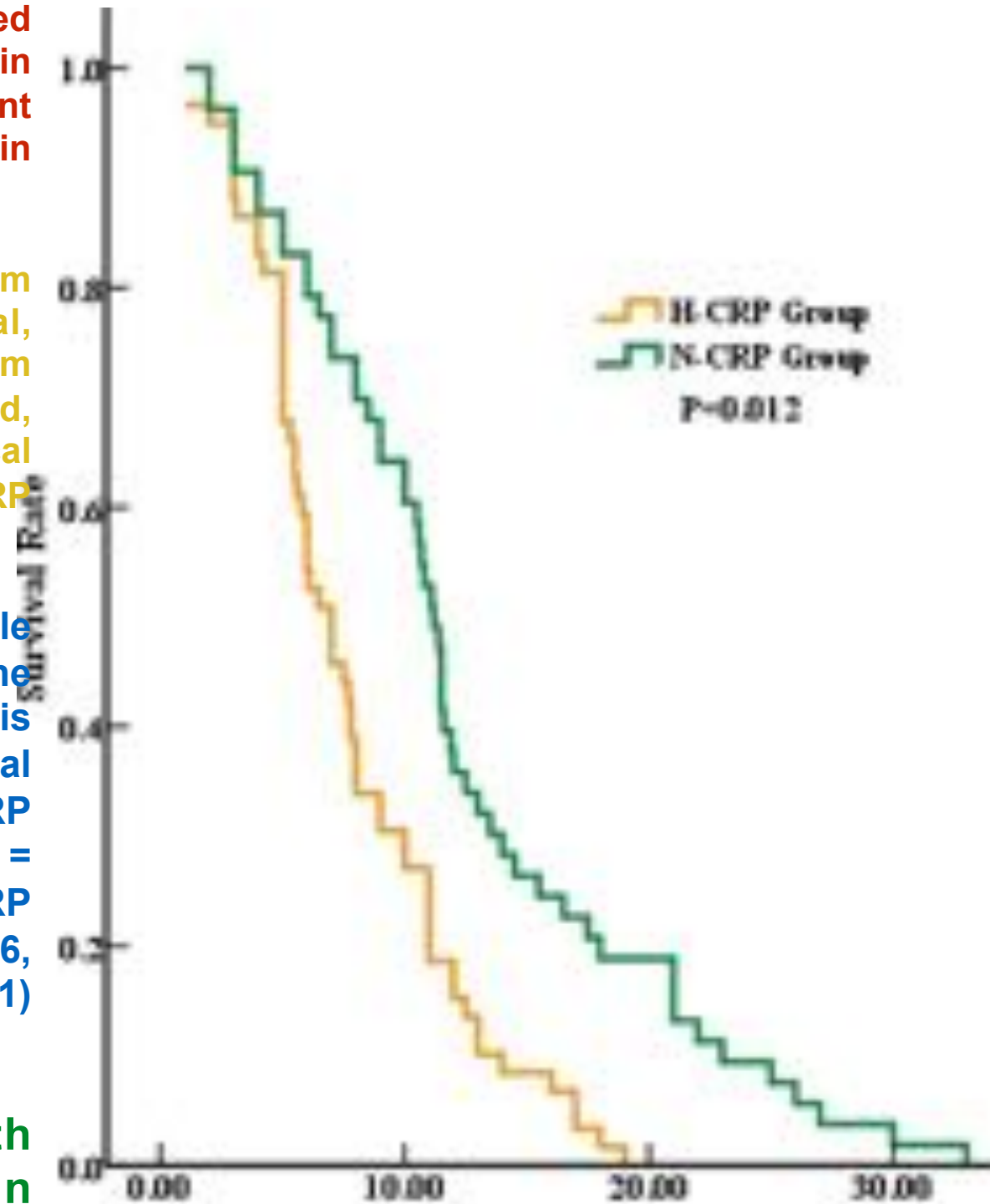
High serum C-reactive protein (CRP) was found to be associated with poor prognosis in kinds of solid tumors, however, its role in recurrent gastric cancer (RGC) is unknown. The present study aimed to explore the prognostic value of serum CRP in RGC patients.

A total of 100 RGC patients who underwent radical surgery from January 2005 to May 2008 were enrolled. The clinical, pathological and survival information were collected. The serum CRP was measured when the recurrence was confirmed. The association between serum CRP and clinicopathological features was analyzed. The prognostic value of serum CRP was investigated.

High CRP was elevated in 39 patients (H-CRP), while 61 patients were within the normal range (N-CRP). The H-CRP was associated with Lymph node metastasis (p = 0.004) and tumor size (p = 0.004). The median survival time to recurrence was significantly worse in the H-CRP group than N-CRP group (6.5 months vs. 11.5 months, p = 0.001). Multivariate analyses identified that elevated CRP (HR = 2.325, p < 0.001), time to recurrence (HR = 0.466, p = 0.001) and the follow-up treatment (HR = 2.650, p = 0.001) were independent prognostic factors.

CONCLUSIONS:

High serum CRP level was associated with poor pathological features, was an



POSITIVE APP.

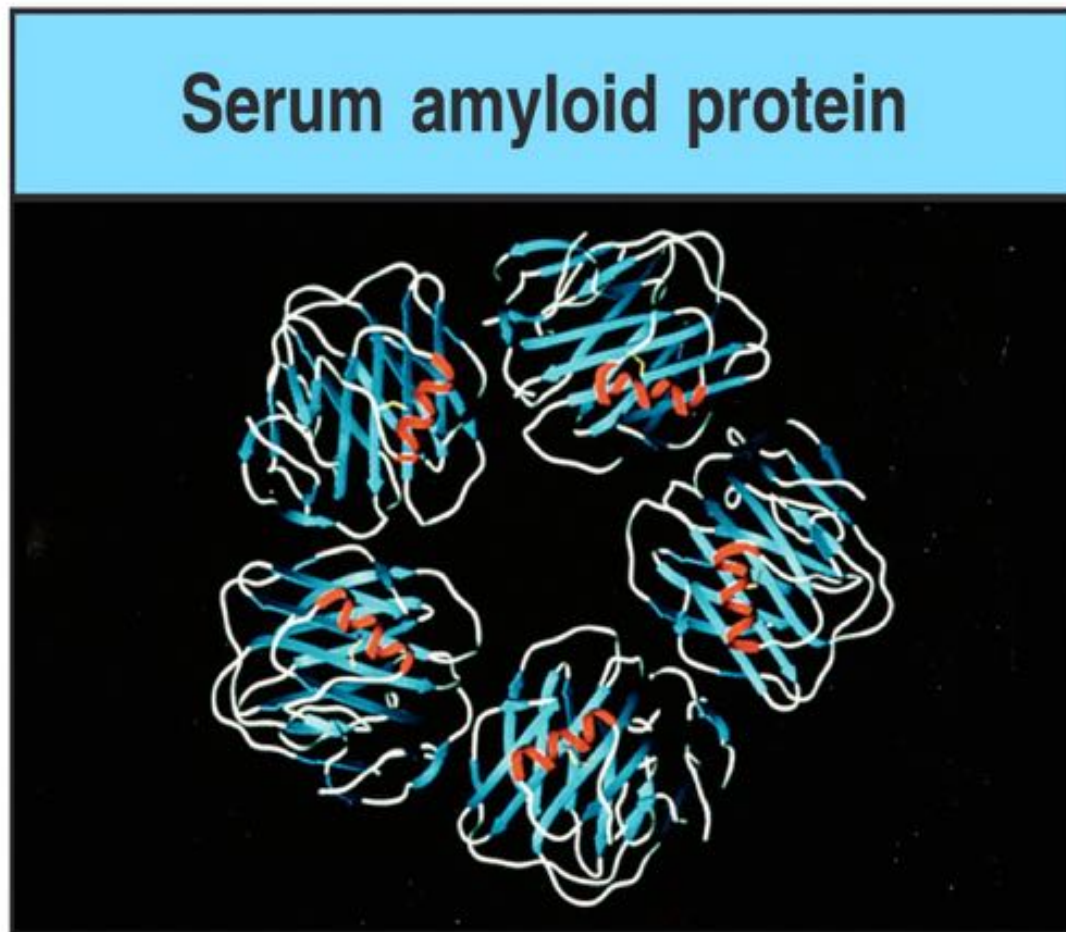


Figure 2-47 part 2 of 2 Immunobiology, 6/e. (© Garland Science 2005)

Authors have reported the existence of various types of SAA that can be classified into two types: CSAA (Chronic phase SAA) and ASAA (Acute phase SAA).

ASAA have:

biological functions, such as promoting the lysis of apoptotic cells; promoting cell proliferation and adhesion and chemotaxis of leukocytes; inducing ECM-degrading enzymes (collagenase, stromalysin, MMP2 and 3) and inflammatory cytokines (IL-6, TNF- α);

Clinical importance of determination of serum amyloid A

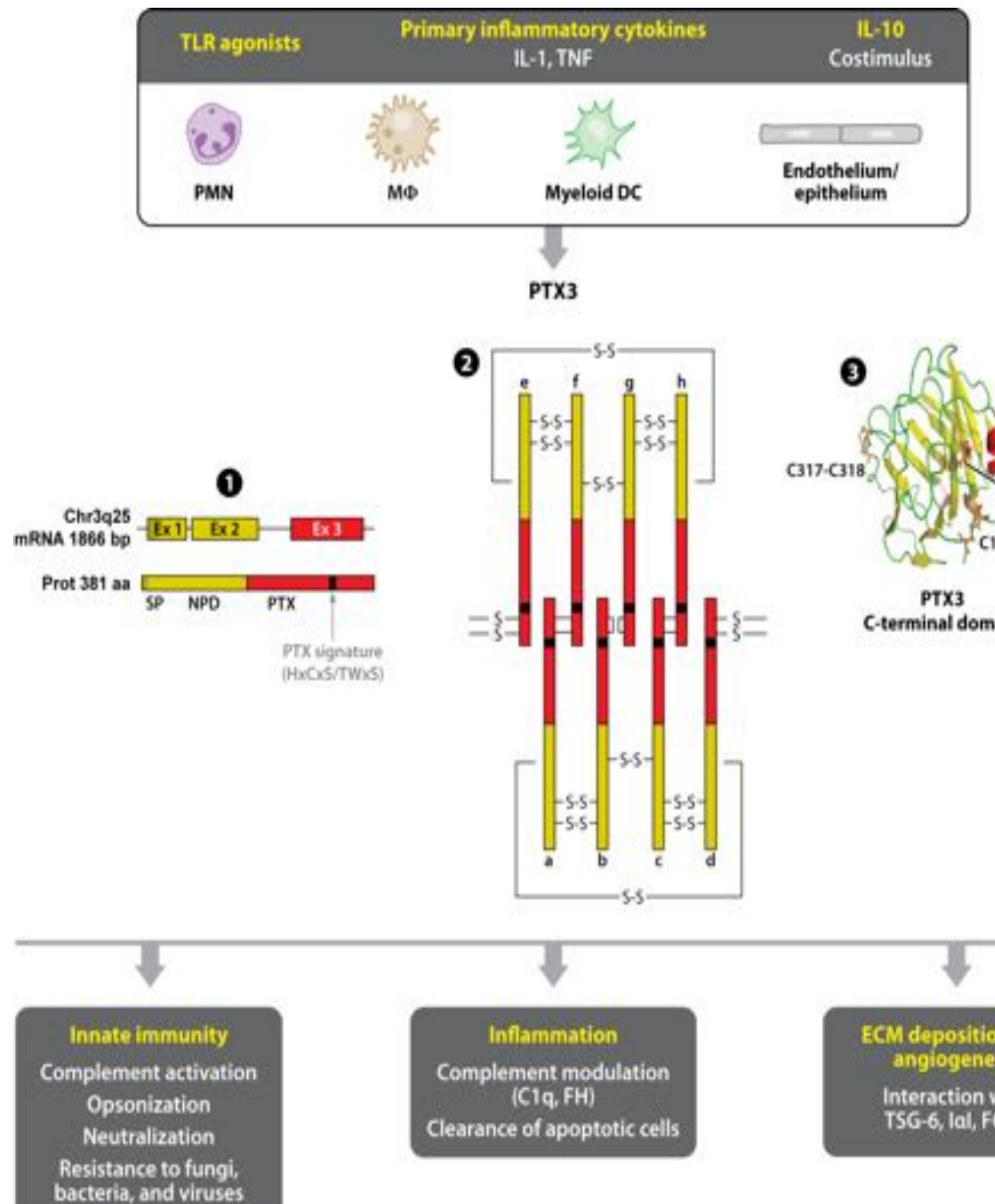
Serum amyloid A (SAA) is an acute phase first class protein discovered a quarter of a century ago. Its concentration depends on clinical findings of the patient, illness activity and the therapy applied.

SAA increases moderately to markedly (100-1000 mg/l) in bacterial and fungal infections, invasive malignant diseases, tissue injuries in the acute myocardial infarction and autoimmune diseases such as rheumatoid arthritis and vasculitis.

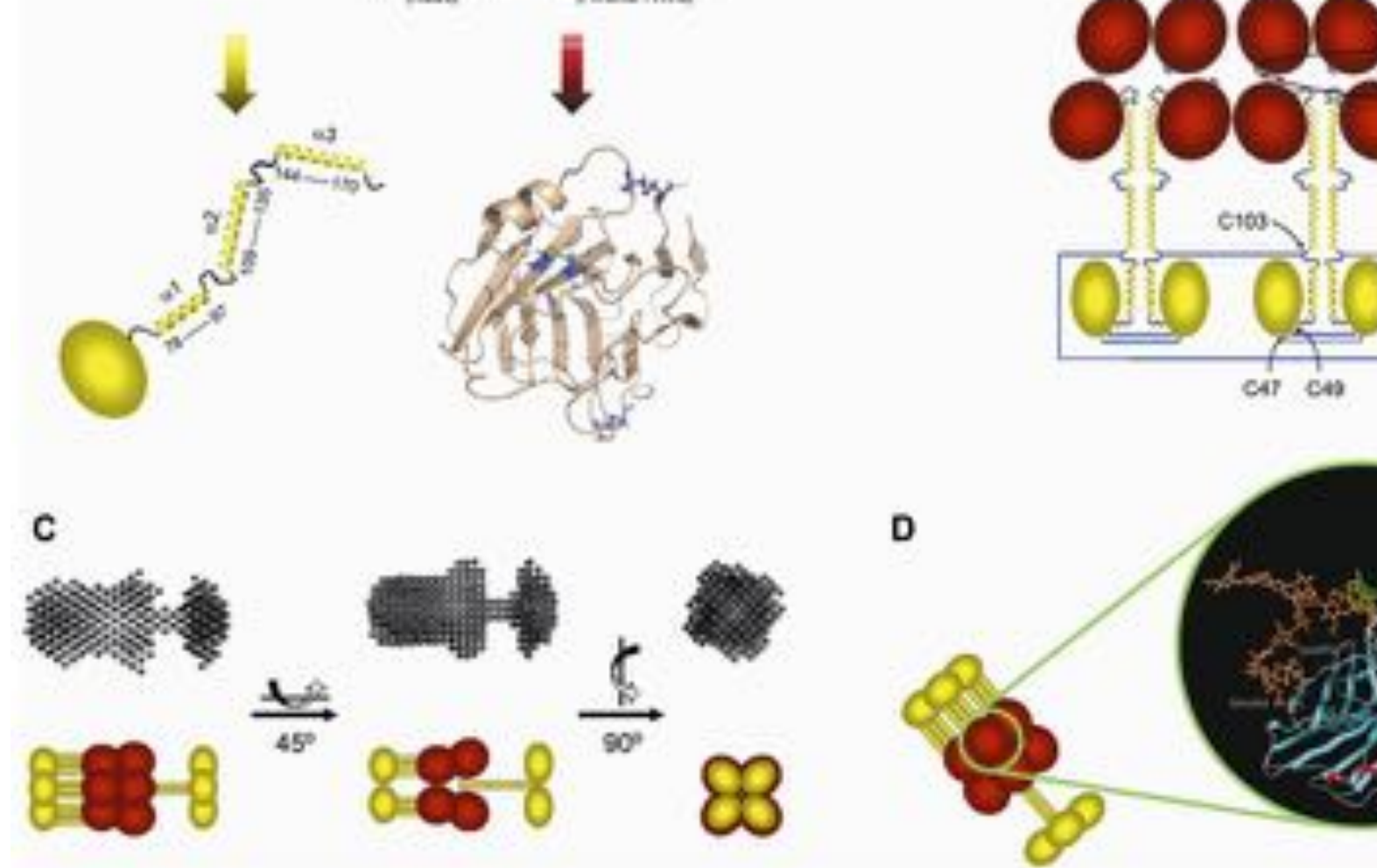
Mild elevation (10-100 mg/l) is often seen in viral infections, systemic lupus erythematosus and localized inflammation or tissue injuries in cystitis and cerebral infarction.

SAA as sensitive, non-invasive parameter is used in organ transplantation where early and correct diagnosis is needed as well as where prompt therapy is required. Simultaneous determination of C-reactive protein (CRP) and SAA may point to acute kidney allograft rejection.

Cellular sources and inducers of the long pentraxin PTX3!



The
 Pentraxins
 INCLUDING PTX3/
 PTX4, after
 their
 intrahepatic
 production
 can be
 glycosylated!

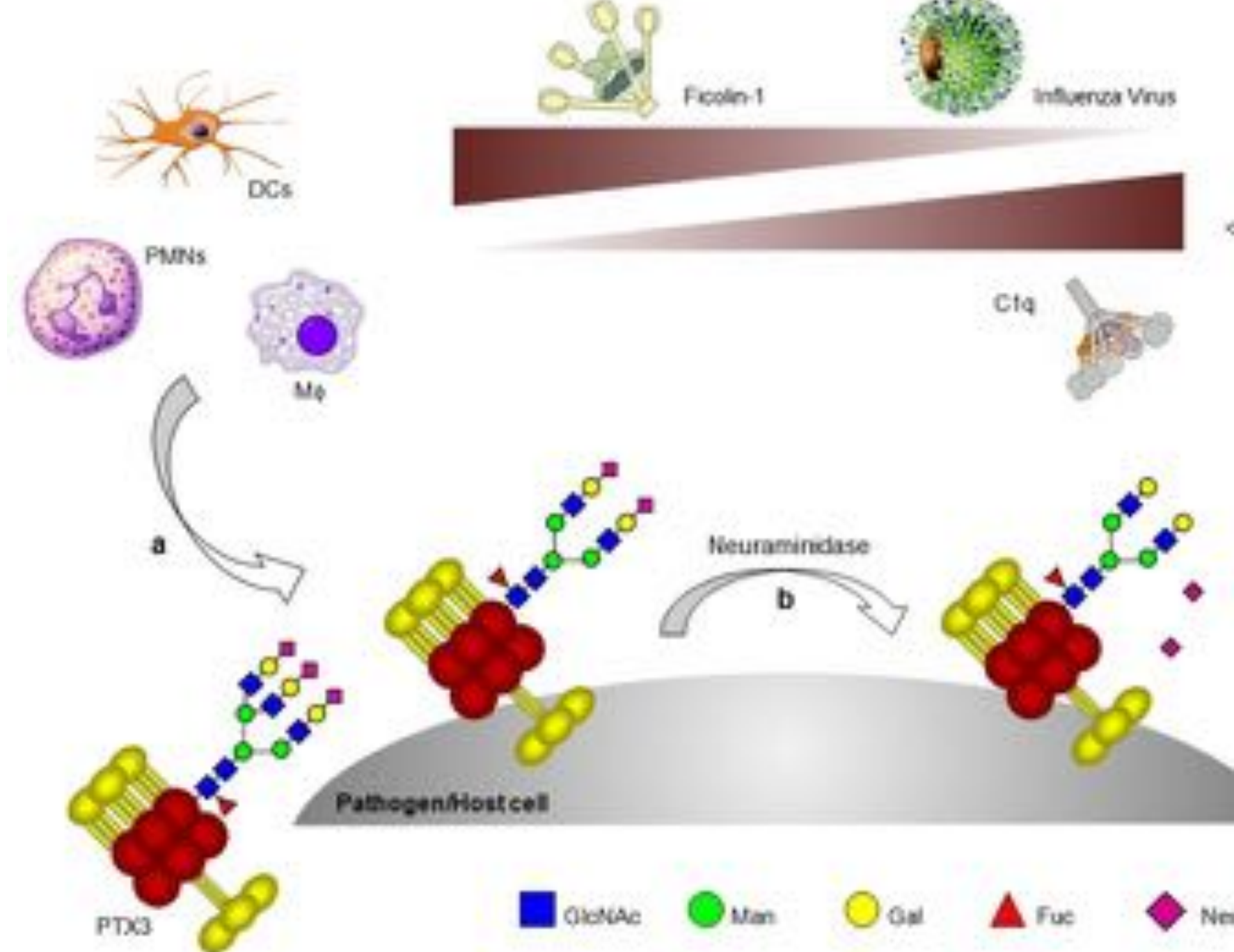


(A) Schematic representation of the PTX3 protomer subunit showing the N-terminal in yellow, followed by the globular pentraxin domain in red. Positions residues, the N-glycosylation site at Asn220 and the pentraxin signature indicated.

(B) Disulfide bond organization of the PTX3 octamer.

(C) Schematic model of PTX3 based on the two different structural arrangements for the N-terminal domain. The α -helical segments of the N-terminal are depicted as yellow rods. The C-terminal pentraxin domains are in red.

nt
munol.
2; 3: 407. 7



number of both somatic and immune cell types produce PTX3 at sites of inflammation. The glycosylation status of PTX3 (e.g., branching and sialylation) might depend on cellular source and inducing stimuli (a). In addition, the protein oligosacch

PTX3 is related to the CPR, whose levels in case of infection rise dramatically in a matter of hours. Compared to PCR, PTX3 is the fact of being produced from any tissue in response to inflammation. In this way its concentration increases much more quickly, allowing an early diagnosis. For example, in the case of infarction was found that when a patient arrives in the ER, PTX3 levels are already very high, while those of CRP rise after a few hours. It can also give important information about prognosis.

Condition in which the evaluation of the levels of PTX3 could be very useful, it is preeclampsia. This is a very serious complication that may occur during pregnancy, with a sharp increase of blood pressure. In these cases the raising of PTX3 concentration occurs long before while the clinical manifestations due to altered vascularization of the placenta appear much later. Even in this case PTX3 could become important for an early diagnosis.

Absence of PTX3 corresponds to a condition of infertility, since this protein is a key component of the structure of cells (the cumulus oophorus) surrounding the oocyte. Without it, ovulation occurs. Without it, the egg cell remains virtually 'naked', deprived of the natural elements that have the function of guiding the sperm in the right direction and thus fertilization can not occur.

Long pentraxin PTX3: a paradigm for humoral pattern recognition molecules

Mantovani,^{1,2} Sonia Valentino,¹ Stefania Gentile,¹ Antonio Inforzato,¹ Cattazzi,¹ and Cecilia Garlanda¹

¹Clinical and Research Center, Rozzano, Milan, Italy. ²Department of Medical Biotechnology and Translational Medicine, University of Milan, Milan, Italy

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Pattern recognition molecules (PRMs) are components of the humoral arm of innate immunity; they recognize pathogen-associated molecular patterns (PAMP) and are functional ancestors of antibodies, promoting complement activation, opsonization, and agglutination. In addition, several PRMs have a regulatory function on inflammation. PTX3 is a family of evolutionarily conserved PRMs characterized by a cyclic multimeric structure. On the basis of their properties, pentraxins have been operationally divided into short and long families. C-reactive protein (CRP) and mannose-binding lectin (MBL) are prototypes of the short pentraxin family, while pentraxin 3 (PTX3) is a prototype of the long pentraxins. PTX3 is produced by somatic and immune cells in response to proinflammatory stimuli and pathogen receptor engagement, and it interacts with several ligands and exerts multifunctional properties. Unlike antibodies, its gene organization and regulation have been conserved in evolution, thus allowing its pathophysiological role to be evaluated in genetically modified animals. Here we will briefly review the general properties of CRP and MBL, the prototypes of short and long pentraxins, respectively, emphasizing in particular the functional role of PTX3 as a humoral PRM with antibody-like properties.

innate immunity; pentraxins; PTX3; pattern recognition molecules

Bacteria	
<i>Pseudomonas aeruginosa</i>	NT*
<i>Klebsiella pneumoniae</i>	NT
<i>Salmonella typhimurium</i>	-
Fungi and yeasts	
<i>Aspergillus fumigatus</i>	+
<i>Saccharomyces cerevisiae</i> (zymosan)	+
<i>Paracoccidioides brasiliensis</i>	NT
Viruses	
Influenza virus	-
Human cytomegalovirus (HCMV)	NT
Membrane moieties	
Phosphocholine (PC)	+
Phosphoethanolamine (PE)	-
LPS	-
Outer membrane protein A from <i>Klebsiella pneumoniae</i> (KpOmpA)	NT
Complement components	
C1q	+
Factor H	+
C4BP	+
M-, L-ficolin	+
MBL	-
Extracellular matrix proteins	
TNF-stimulated gene-6 (TSG-6)	NT
Inter- α -trypsin-inhibitor (I α I)	-
Hyaluronan	NT
Laminin	+
Collagen IV	NT
Fibronectin	+
Growth factors	
EGF	-

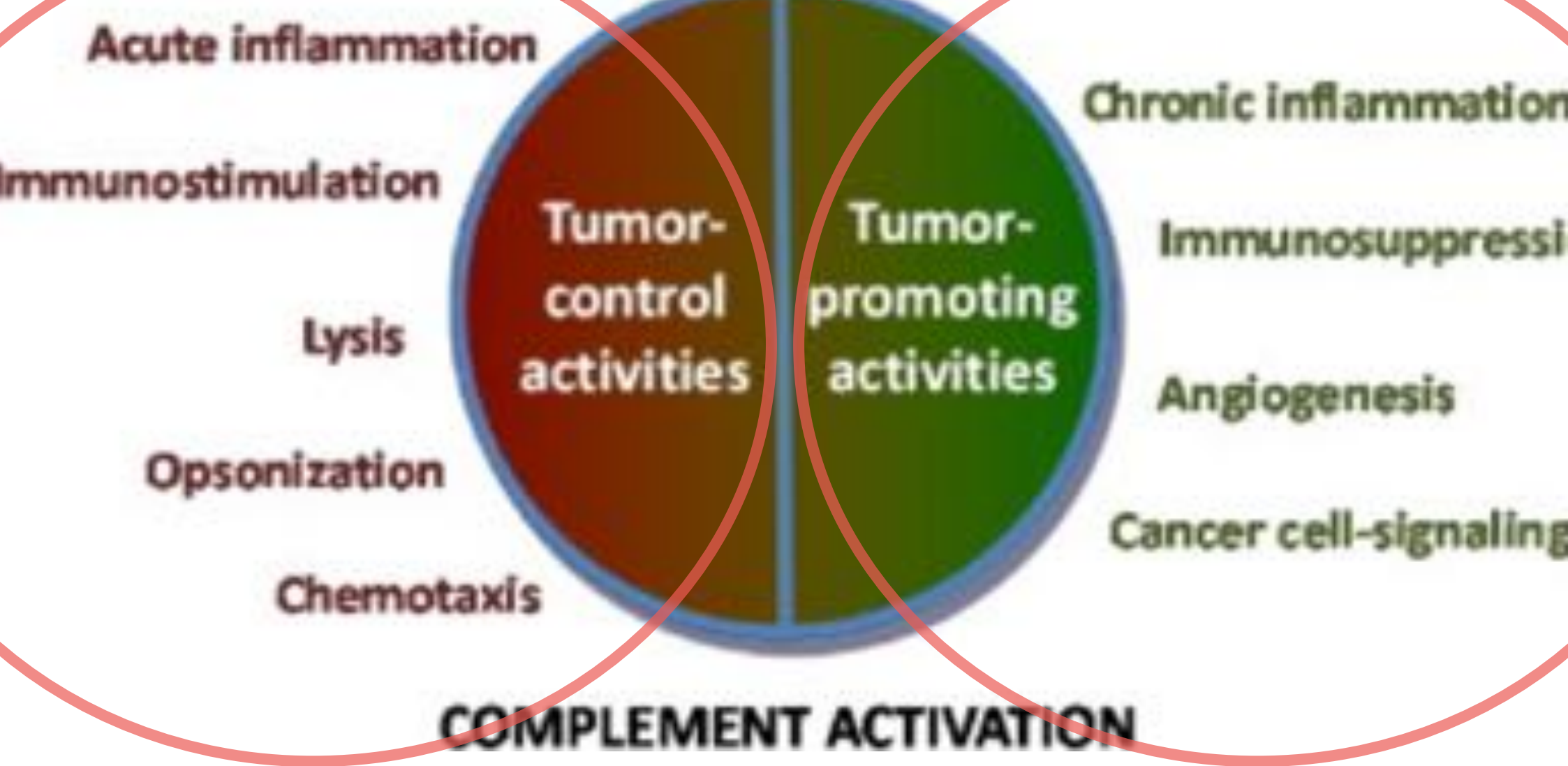
Dependent Inflammation in Cancer.

[a E](#), [Gentile S](#), [Rubino M](#), [Maina V](#), [Papait R](#), [Kunderfranco P](#), [Greco C](#), [Ferraioni M](#), [Laface I](#), [Tartari S](#), [Doni A](#), [Pasqualini F](#), [Barbati E](#), [Basso G](#), [Galdiero M](#), [Roncalli M](#), [Colombo P](#), [Laghi L](#), [Lambris JD](#), [Jaillon S](#), [Garlanda C](#), [Mantovani M](#)

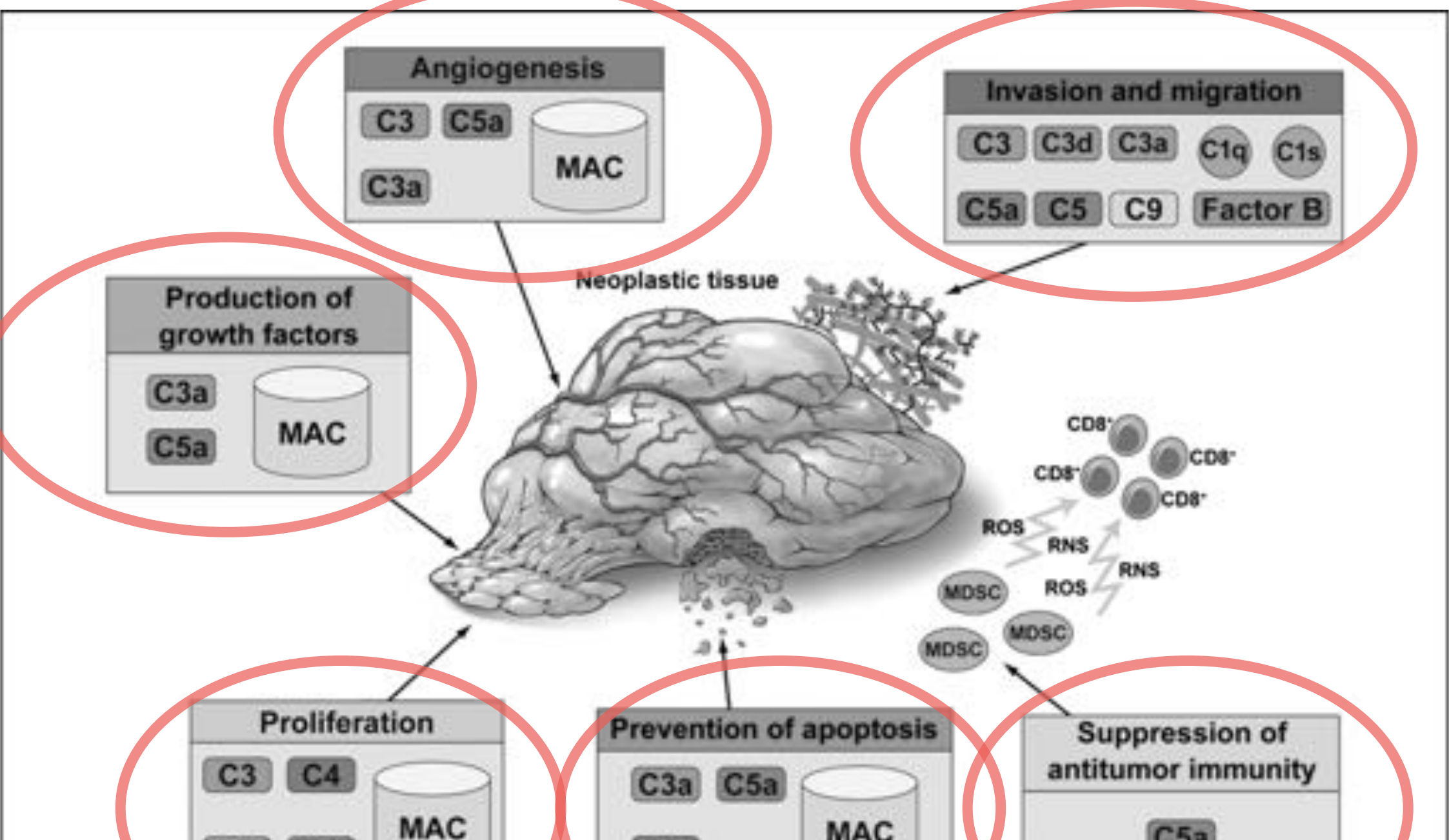
ct

is an essential component of the humoral arm of innate immunity, playing a predominant role in resistance against selected microbes and in the regulation of inflammation. PTX3 activates and regulates the Complement cascade by interacting with Factor H. PTX3 deficiency was associated with increased susceptibility to subcutaneous and epithelial carcinogenesis. Increased susceptibility of Ptx3(-/-) mice was associated with enhanced macrophage infiltration, cytokine production, angiogenesis, and tumor growth. Correlative evidence, gene-targeted mice, and pharmacological blockade experiments indicated that PTX3 deficiency resulted in amplification of Complement activation, CCL2 production, and tumor-promoting macrophage recruitment. PTX3 expression was epigenetically regulated in selected human tumors (e.g., leiomyosarcoma and colorectal cancer) by methylation of the promoter region and of a putative enhancer. PTX3, an effector molecule belonging to the humoral arm of innate immunity, is an intrinsic oncosuppressor gene in mouse and man by regulating Complement

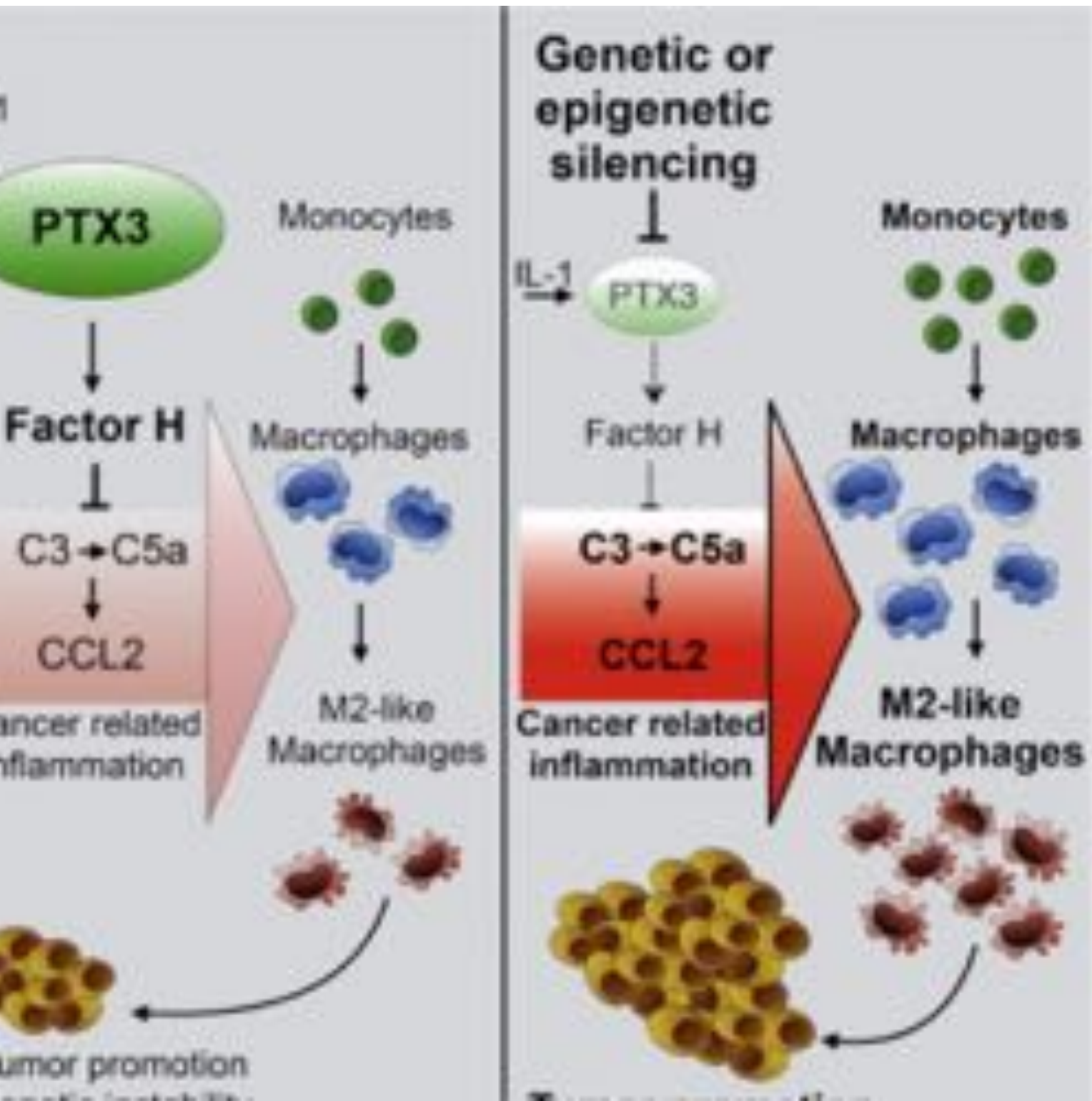
Experimental data support the idea that complement is activated by tumors. However, some studies also suggest that malignant cells evade the harmful complement and make use of some complement effector molecules to promote tumor growth. Unfortunately, the exact mechanisms and consequences of this duality are not very well known!



production, angiogenesis, protection from antigrowth factors and apoptosis, cellular invasion and migration through extracellular matrix, and suppression of antitumor immunity

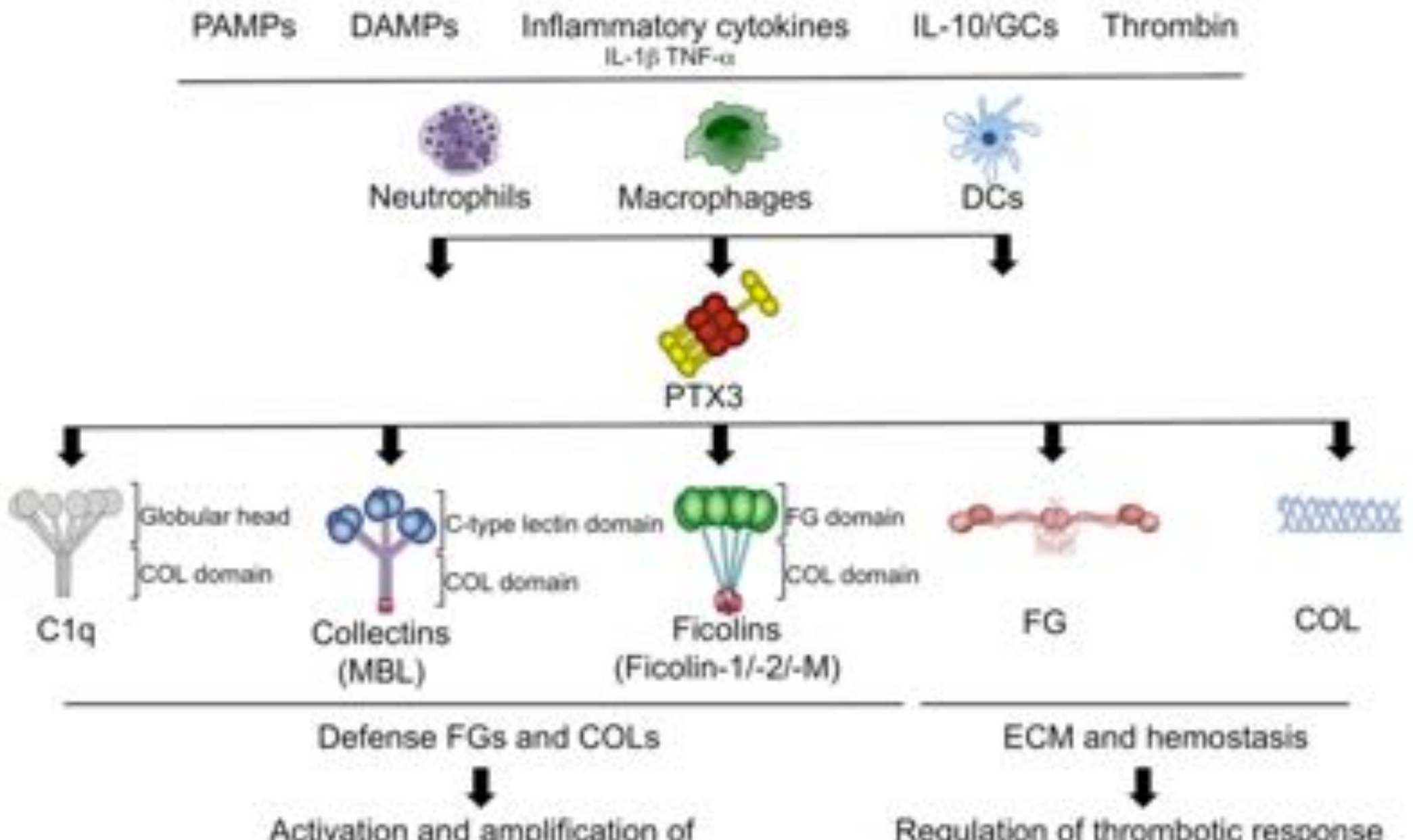


man!

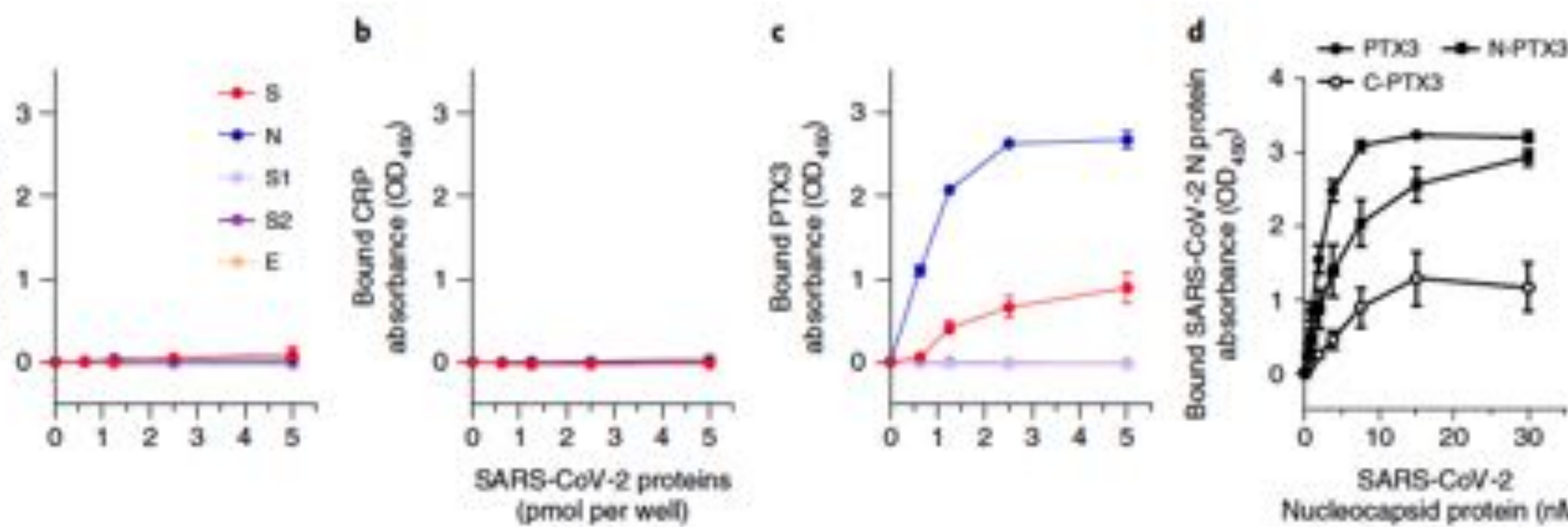


PTX3 gene is silenced by hypermethylation in selected human tumors including colorectal cancer (CRC) and this event occurs early in progression already at the level of adenoma

PTX3 amplifies other effector functions of the innate immune system. At the same time, PTX3 regulates the injury-induced thrombotic response and promotes wound healing by favoring timely fibrinolysis. Therefore, PTX3 interacts with and modulates pathways conserved in innate immunity, hemostasis and extracellular matrix and processes related to both antimicrobial resistance and tissue repair.

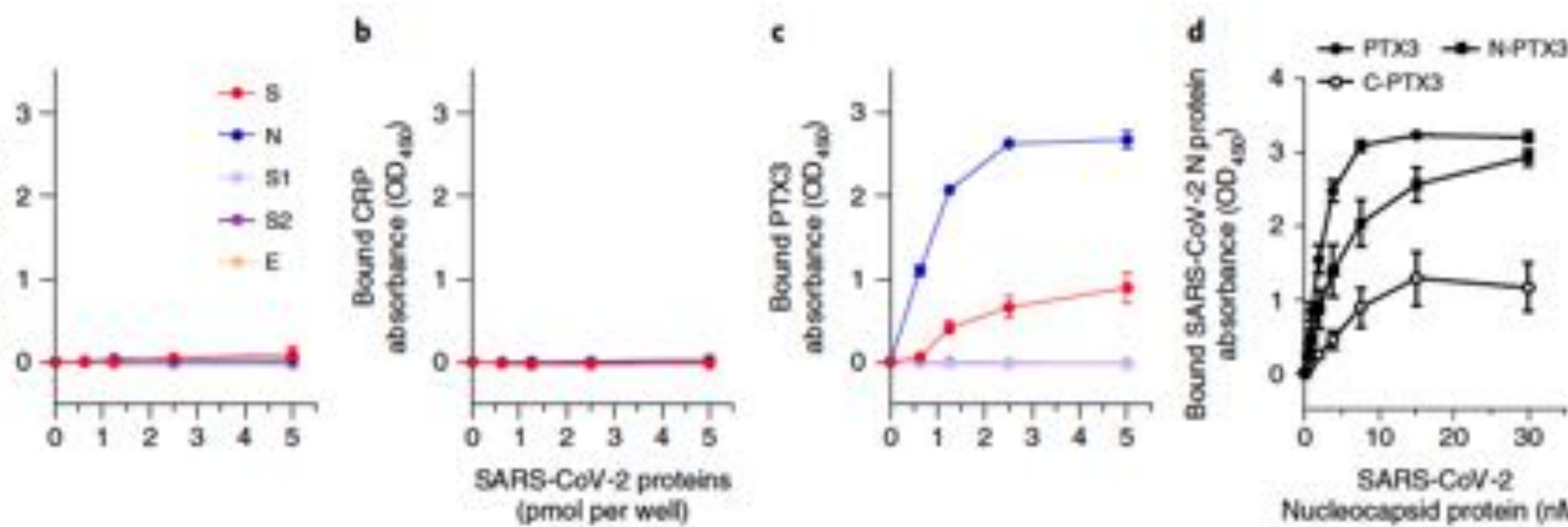


er to the nucleocapsid protein, one of the most abundant proteins of SARS-CoV-2



Interaction between pentraxins and SARS-CoV-2 proteins. a-c, Recombinant His tag SARS-CoV-2 proteins (spike active trimer (S), S1, S2, nucleocapsid (N), envelope (E); the legend refers to **a-c**) were immobilized on 96-well nickel-coated plates at different concentrations. Fixed concentrations of PTX3 (**c**) were incubated over the captured viral proteins. Bound pentraxins were detected by enzyme-linked immunosorbent assay (ELISA) using secondary antibodies. **d**, Full-length PTX3 or the N- or C-terminal domains were captured on 96-well plates. Biotinylated SARS-CoV-2 nucleocapsid protein was incubated at different concentrations. Bound nucleocapsid was detected by ELISA using horseradish peroxidase (HRP)-conjugated streptavidin. Data are presented as mean \pm s.e.m.; $n=3$ independent experiments performed in duplicate; OD₄₅₀, optical density at 450 nm.

er to the nucleocapsid protein, one of the most abundant proteins of SARS-CoV-2

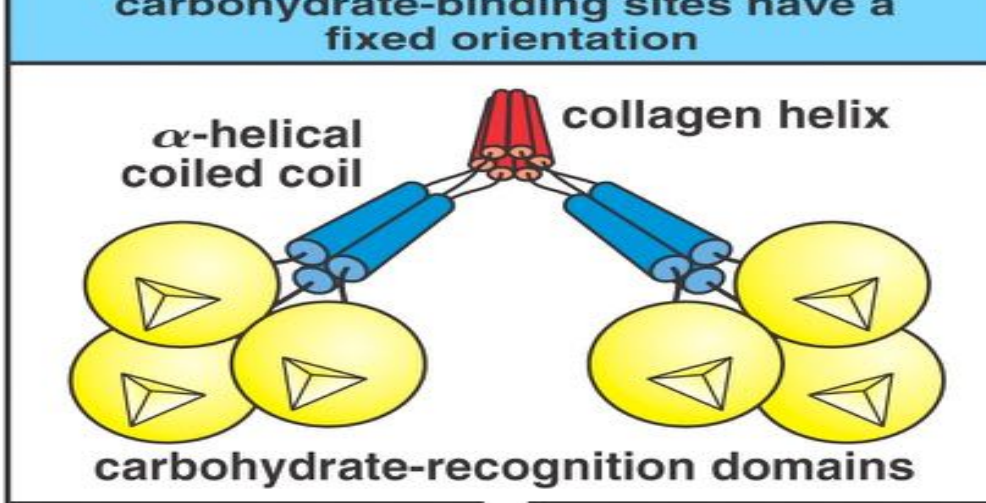


Interaction between pentraxins and SARS-CoV-2 proteins. a-c, Recombinant His tag SARS-CoV-2 proteins (spike active trimer (S), S1, S2, nucleocapsid (N), envelope (E); the legend refers to **a-c**) were immobilized on 96-well nickel-coated plates at different concentrations. Fixed concentrations of PTX3 (**c**) were incubated over the captured viral proteins. Bound pentraxins were detected by enzyme-linked immunosorbent assay (ELISA) using secondary antibodies. **d**, Full-length PTX3 or the N- or C-terminal domains were captured on 96-well plates. Biotinylated SARS-CoV-2 nucleocapsid protein was incubated at different concentrations. Bound nucleocapsid was detected by ELISA using horseradish peroxidase (HRP)-conjugated streptavidin. Data are presented as mean \pm s.e.m.; $n=3$ independent experiments performed in duplicate; OD₄₅₀, optical density at 450 nm.

- **MBL**
- **SPA**
- **SPD**
- **FICOLINs**

is structurally related to the C1q and is part of the family of protein collectins, together with proteins A and D of pulmonary surfactant.

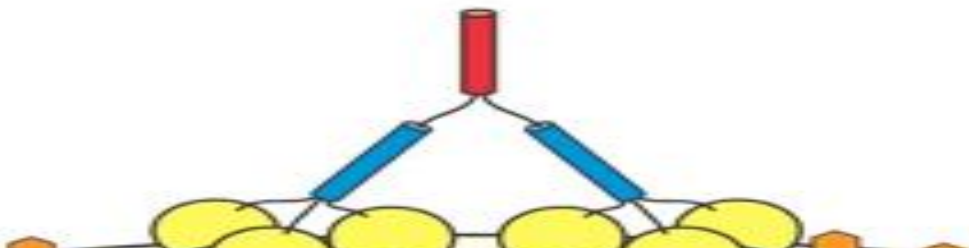
It was recently discovered that a second group of proteins called ficolins, which includes the L-ficolin, the M-ficolin, and the D-ficolin, possesses lectin activity.



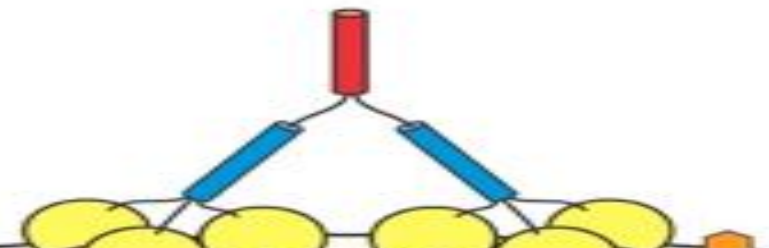
Mannose-binding lectin (MBL), or mannose-binding protein, recognizes different carbohydrates on the surface of many microorganisms, including bacteria, viruses, protozoa and fungi as an oligomeric structure (400-700 kDa), formed by subunits in their turn consist of three polypeptide chains, each of 32 kDa and containing from two to six clusters "of carbohydrate recognition sites" that can bind mannose, maltose, N-acetylglucosamine, N-acetylgalactosamine and glucose.

It has been shown that the mere presence of these sugar residues is not sufficient for the binding of MBL, but their orientation is critical, as they are related only the residues that have a correct arrangement. The bond has a low affinity (K_d 10⁻³) and, in order to be effective, it is essential that more "carbohydrate recognition sites" bind simultaneously.

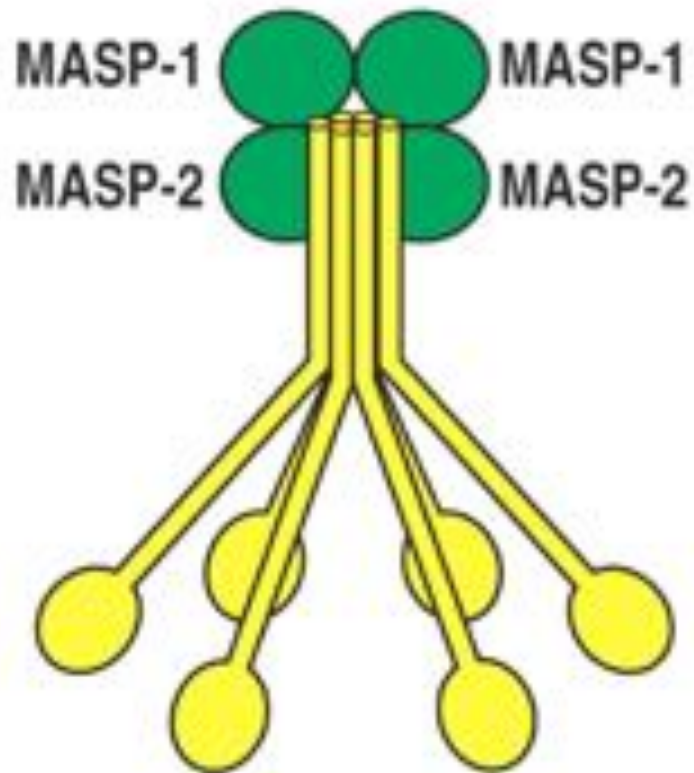
MBL binds with high affinity to mannose and fucose residues with correct spacing



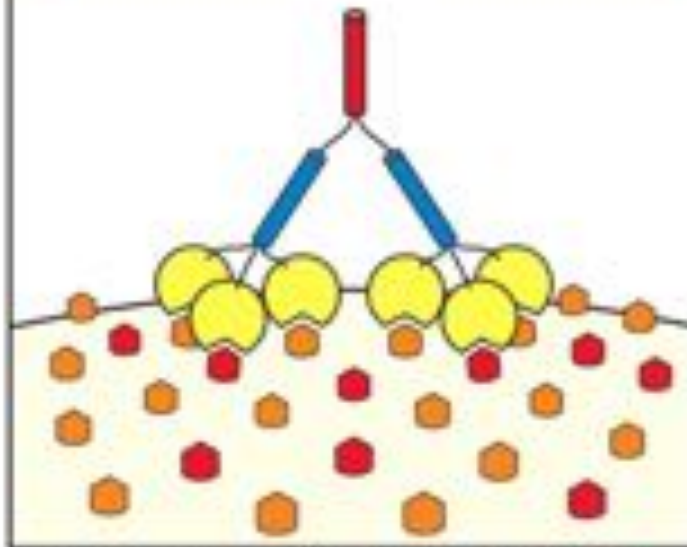
Mannose and fucose residues that have different spacing are not bound by MBL



The lectin-binding pathway of complement activation



MBL binds with high affinity to mannose and fucose residues with correct spacing



Mannose and fucose residues that have different spacing are not bound by MBL

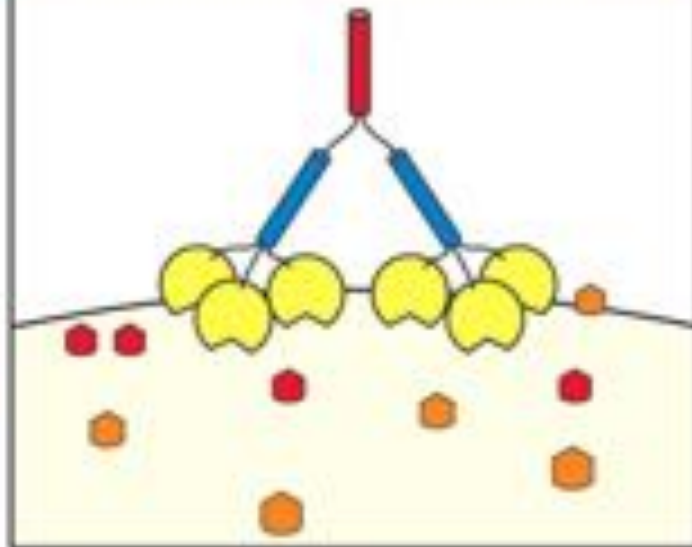
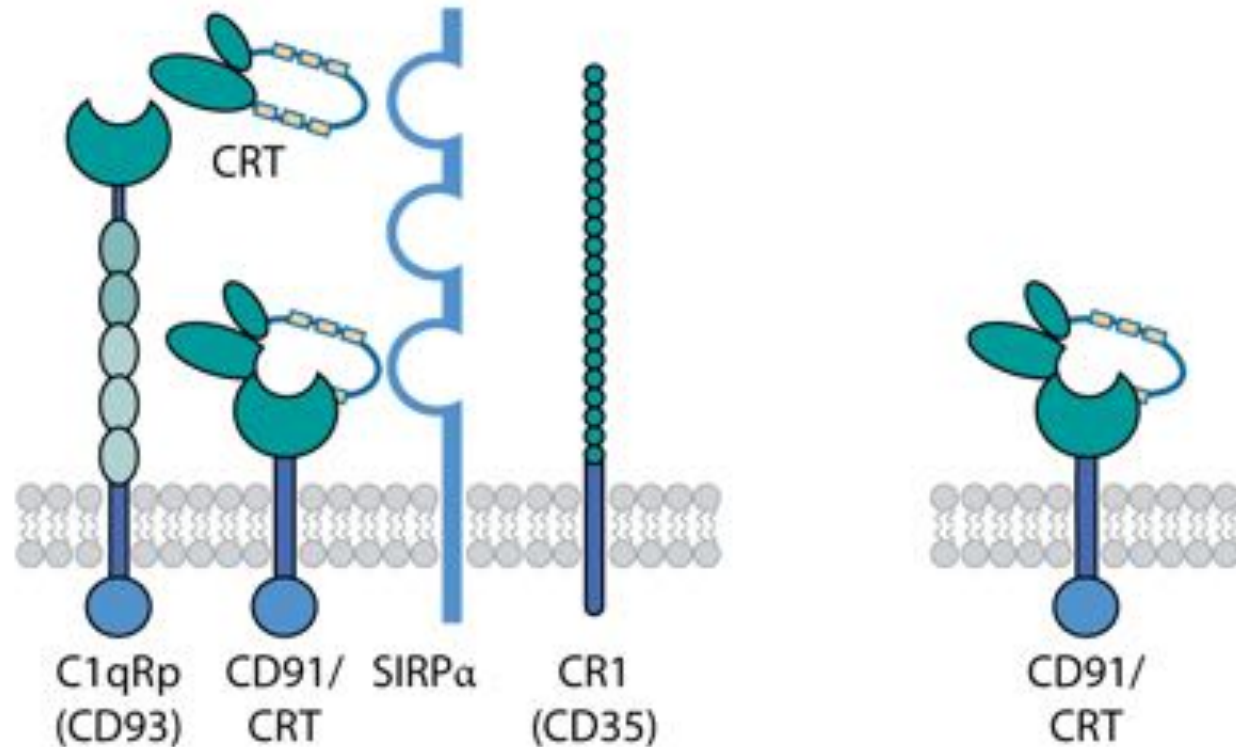
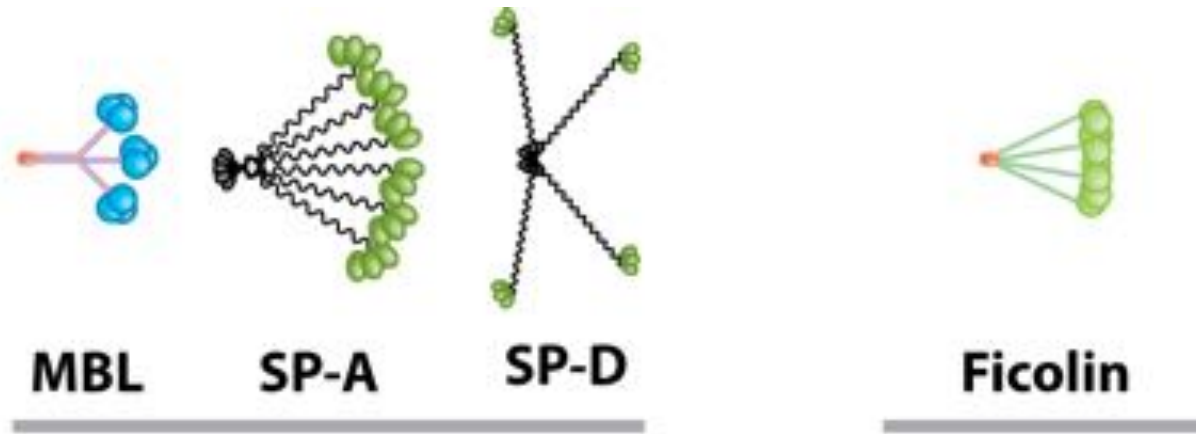


Figure 2-11 part 2 of 2 Immunobiology 4/e. © Garland Science 2005

MBL
SPA
SPD
FICOLINs

- It interacts with **MASP1** and **MASP2** (Mannan Associated Serine Protease)



The humoral collectins activate

The clinical MBL!

Plasma concentration of MBL immediately after birth from 1000 to a maximum of 2500 ng/mL within a few weeks, after which it falls usually up to 1700 ng/mL in adults where it can increase up to 20 times during infections and inflammatory processes.

Changes in the plasma concentration reveal the physiological importance of the MBL that is present at every age during the early stages of contact with the pathogens prior to the increase in the concentration of IgM but it is crucial in the period following childbirth, when the concentration of antibodies decreases maternally and starts the production of those of the newborn.

Low concentrations of MBL are genetically controlled polymorphisms and/or mutations in the promoter and coding region of the MBL2 gene and **MBL deficiency, the incidence of which is estimated to be approximately 20-25% of the world population, are associated with increased susceptibility to infections such as ear infections, pneumonia, gastroenteritis, meningitis, and conjunctivitis.**

In children, a heterozygous mutation of the MBL gene doubles the risk of hospitalization due to infectious diseases than children with normal MBL levels; in case of a homozygous mutation, the risk of infection is also increased and the disease is worse.

Patients with cystic fibrosis with a mutation in the MBL gene are more susceptible to infections with *Pseudomonas aeruginosa* and have a lower life expectancy than patients with cystic fibrosis without this mutation.

For patients undergoing chemotherapy and consequently to that, develop neutropenia, they are more susceptible to long periods of fever if they have a low concentration of MBL.

In children under one year suffering from Kawasaki disease, whether MBL deficiency is present, it is a risk factor for coronary aneurysms.

In HIV-positive patients the progression of AIDS is faster in case of deficiency of MBL.

MBL deficiency is associated with the occurrence of recurrent spontaneous abortions and, in addition, to intrauterine infections.

MBL deficiency is also associated with autoimmune diseases such as systemic lupus erythematosus (SLE) and rheumatoid arthritis (RA)

Case Binding Lectin Deficiency: More than Meets the Eye **Susanne Halbrich, Moshe Ben-Zur, and Christine McCusker**

The report describes a 5-year-old boy who presented to the emergency department with symptoms and chest X-ray findings of pneumonia. Further history revealed recurrent infections, and workup for immunodeficiency revealed a deficiency of mannose-binding lectin (MBL), a pattern recognition receptor involved in activation of the innate immune system. Innate immunodeficiency may be more common than currently appreciated, with MBL deficiency affecting up to 50% of individuals in certain populations. While pneumonia is a common presentation in the Pediatric Emergency Department, clinical presentations of children with defects in innate immunity can be unpredictable. These cases pose particular challenges to physicians, and the level of suspicion for immunodeficiency must remain high. It is crucial to identify patients with such impairments to better manage them and prevent future complications.



X-ray from the emergency department demonstrating a

Roma,

Sig..... PAZIENTE
(Cognome e Nome)

Prelievo del.....

Provenienza ...DAI Pediatria.....

DOSAGGIO LECTINA LEGANTE IL MANNOSIO (MBL) PER DEFIC

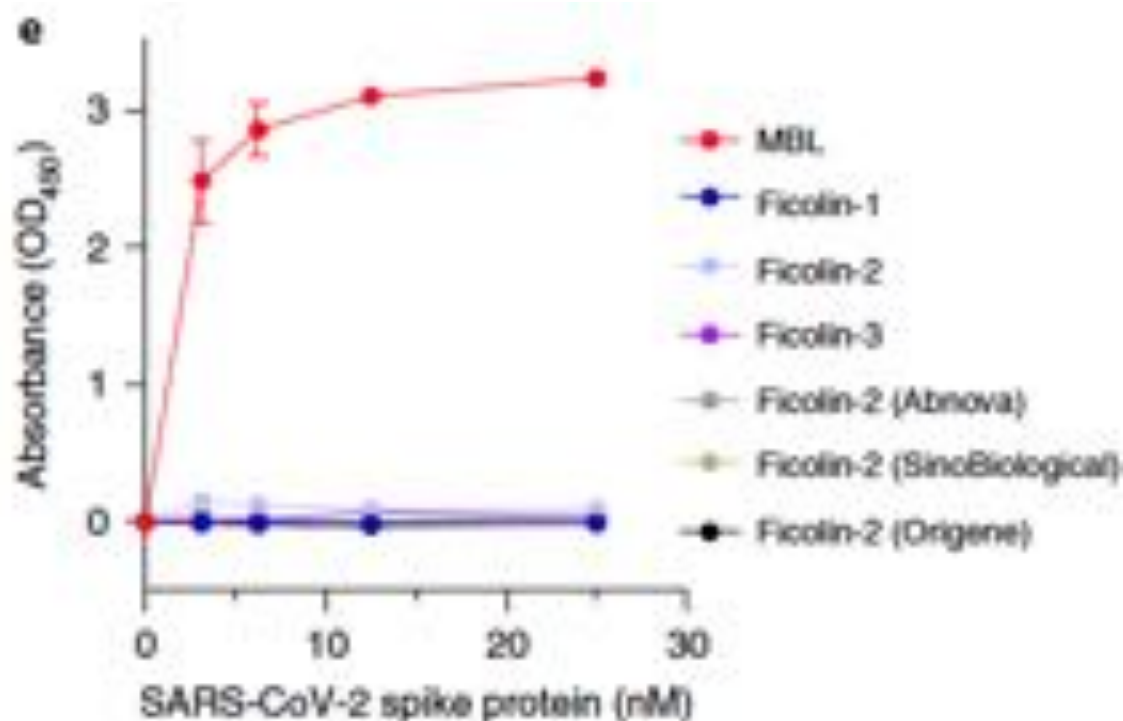
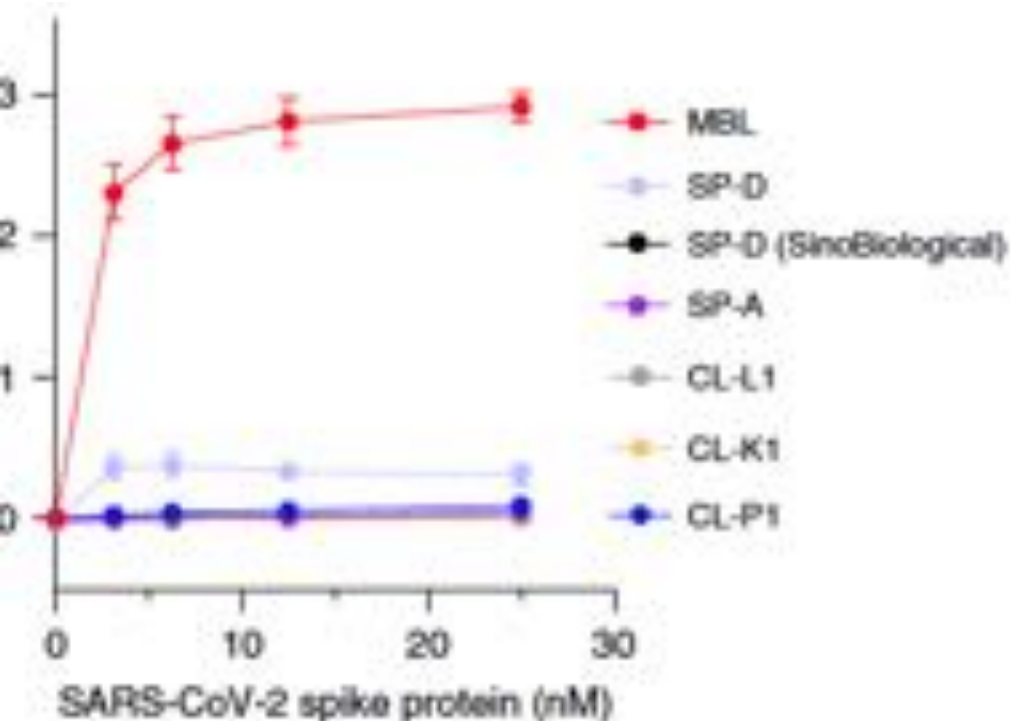
MBL 1470 (>100 ng/ml V.N.)

Il test è stato eseguito mediante MBL Oligomer ELISA kit (BioPorto Diagnostics).

Il Responsabile

Fabrizio Mainero

specifically to the spike protein of SARS-CoV-2!



Interaction of C1q, MBL, ficolins and surfactant proteins with SARS-CoV-2 proteins. **a-c**, Recombinant His tag SARS-CoV-2 proteins (S, S2, nucleocapsid (N) and envelope (E)) were immobilized on 96-well nickel-coated plates at different concentrations. Fixed concentrations of MBL (**b**) were incubated over the captured viral proteins. Recombinant SARS-CoV-2 spike proteins tested were expressed in different cell lines. MBL (2 $\mu\text{g ml}^{-1}$; 6.7 nM) was incubated over the captured viral proteins. In **a-c**, bound proteins were detected by ELISA with specific antibodies. Data in **a** and **b** are presented as mean \pm s.e.m.; $n=3$ independent experiments performed in duplicate. Data in **c** are presented as mean \pm s.e.m.; $n=2$ independent experiments, one performed in duplicate and one in triplicate. **d,e**, MBL-, CL-L1, CL-K1, CL-P1-, SP-D- and SP-A-coated plates (**d**) and ficolin-1-, ficolin-2- and ficolin-3-coated plates (**e**) were incubated with various concentrations of biotinylated SARS-CoV-2 spike protein. Bound spike protein was detected by ELISA with HRP-conjugated streptavidin (mean \pm s.e.m.; $n=3$ independent experiments in duplicate). **f**, Surface plasmon resonance (SPR) analysis of the interaction of recombinant full spike protein trimer to immobilized MBL (dissociation constant (K_D) = 34 nM, left). No binding was detected in the control experiment (right). RU, resonance units.

The SPA and SPD!

Proteins A and D of the pulmonary surfactant or SPA and SPD are also defined as pulmonary collectins found mainly in bronchial fluids where they are synthesized by alveolar macrophages and Clara cells.

Their serum concentrations are around 100 ng/ml but may change in relation to the lung pathology in progress.

Their concentration may increase in pulmonary disorders such as pulmonary alveolar proteinosis (PAP) and decrease in pulmonary fibrosis and acute respiratory distress syndrome (ARDS).

Humans have been identified three ficolins: the H-ficolin (Ficolin-2), also known as the MBL-associated antigen, the L-ficolin/P35 (Ficolin-3) both serum proteins, and a third ficolin called MBL/P35-related (Ficolin-1), not present in the serum.

Most recently these ficolins has been associated with a new membrane protein called CD11b, which binds groups of acetylated sugar residues expressed by pathogens and damaged cells.

Similar to the MBL, the three human ficolins bind carbohydrates on the surface of pathogens and activates the lectin pathway of complement through the MASPs.

The H-Ficolin is synthesized in both the liver in the lung and is present in serum at a concentration average of 15 µg/ml.

H-ficolin may be absent in patients with SLE, probably due to the presence of autoantibodies anti-H-Ficolins, and in liver diseases where the serum levels decrease with increasing severity of cirrhosis.

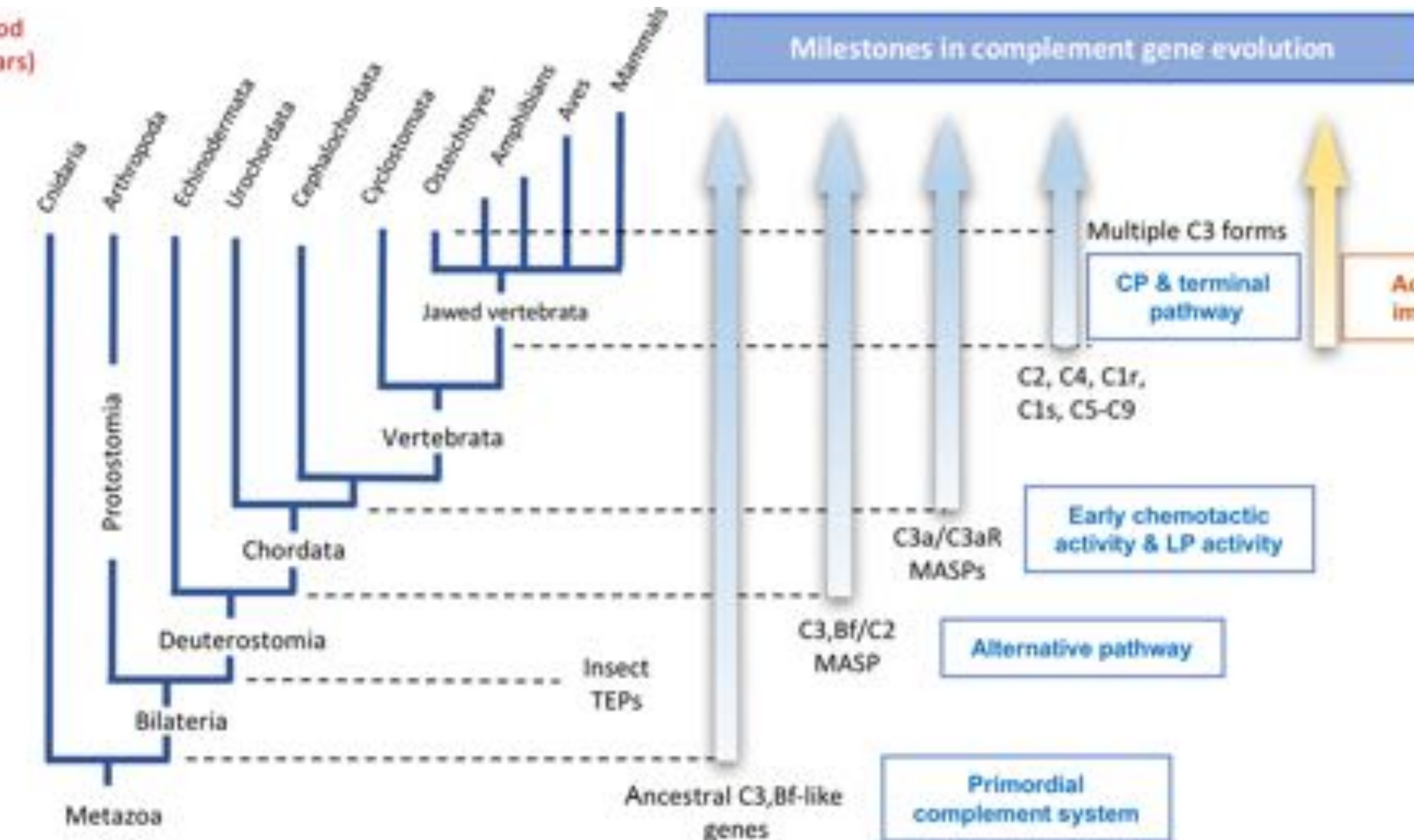
Recently has been described a mutation of the gene FCN3, coding for the H - ficolin, which cause defects of complement activation, and gene polymorphisms FCN1 which

Proteins of the complement system!

Functional protein classes in the complement system	
Binding to antigen:antibody complexes and pathogen surfaces	C1q
Binding to mannose on bacteria	MBL
Activating enzymes	C1r C1s C2 Bb D MASP-1 MASP-2
Membrane-binding proteins and opsonins	C4b C3b
Peptide mediators of inflammation	C5a C3a

Functional protein classes in the complement system	
Membrane-attack proteins	C5b C6 C7 C8 C9
Complement receptors	CR1 CR2 CR3 CR4 C1qR
Complement-regulatory proteins	C1INH C4bp CR1 MCP DAF H I P CD59

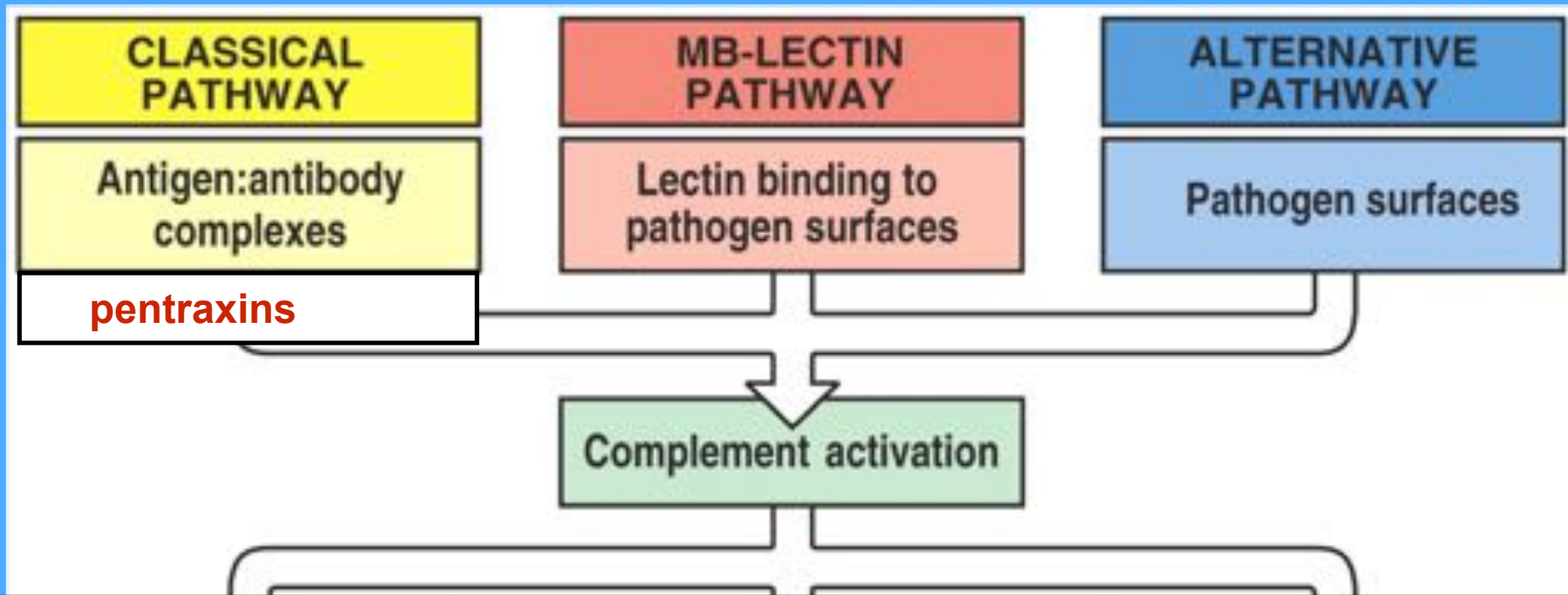
THE COMPLEMENT SYSTEM IS THE OLDEST DEFENSE SYSTEM



Name	MW	mg/dl	fragments
------	----	-------	-----------

C1q	410	0,7-3	
C1r	83	0,34-1	
C1s	85	0,3-0,8	
C4	204	15-53	C4a, C4b, C4c, C4d
C2	102	0,15-0,3	C2a, C2b
C3	190	55-120	C3a, C3b, C3c, C3d, C3e, C3f, C3g, C3dg, iC3b
C5	196	0,70-0,85	C5a, C5b
C6	125	0,6-0,7	
C7	120	0,55-0,7	
C8	150	0,55-0,8	
C9	66	0,5-1,6	
Fattore B	100	1,4-2,4	Ba, Bb
P	224	0,2-0,3	
Fattore D	24	0,01-0,02	
MBL	540	0,01	
MASP-1	94	0,005	
MASP-2	76	0,005	
C1IH	105	1,8-2,75	
C4BP	550	2,5	
Fattore H	150	3-5, 6	
Fattore I	100	0,34-0,55	
CD59	20	0,005	

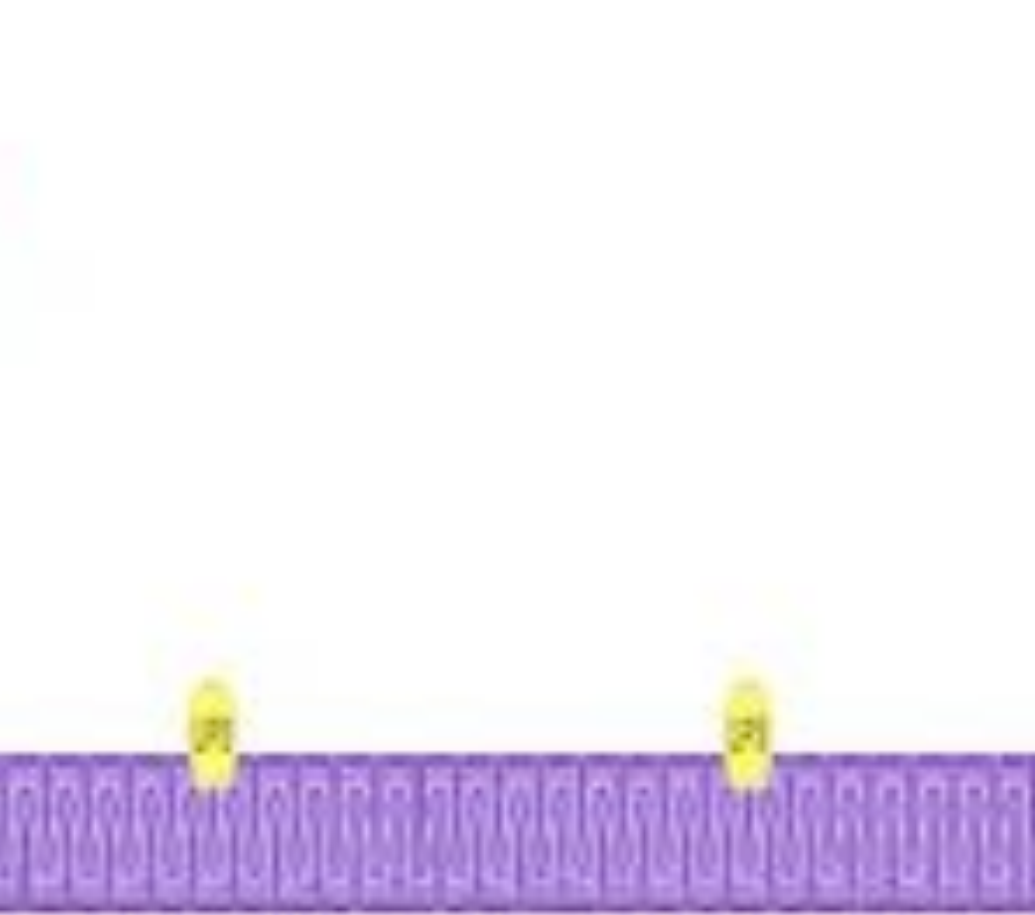
THE THREE MECHANISMS OF COMPLEMENT ACTIVATION



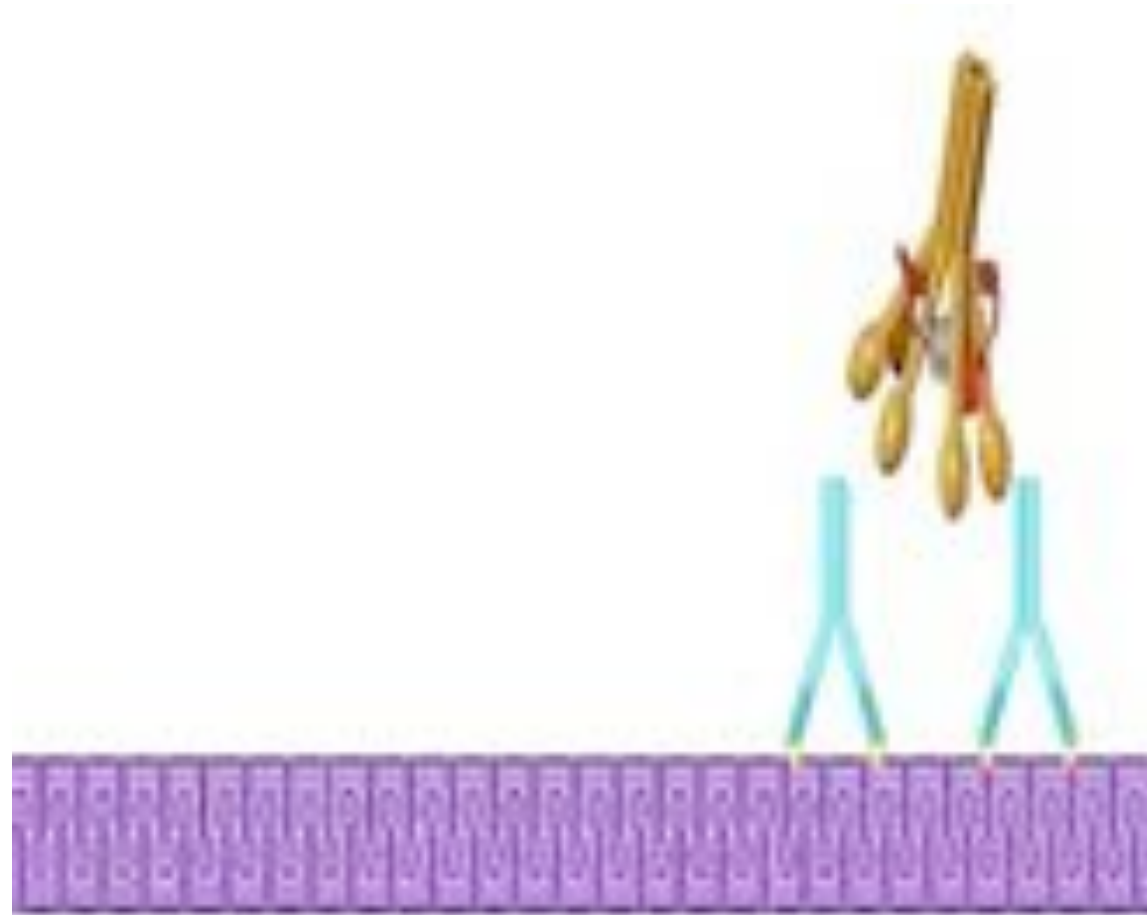
**ALL THREE MECHANISM HAVE C3
AS CENTRAL PROTEIN
AND CONVERGE
IN THE ACTIVATION OF C5!!!**

OF COMPLEMENT ACTIVATION!

ALTERNATIVE PATHWAY



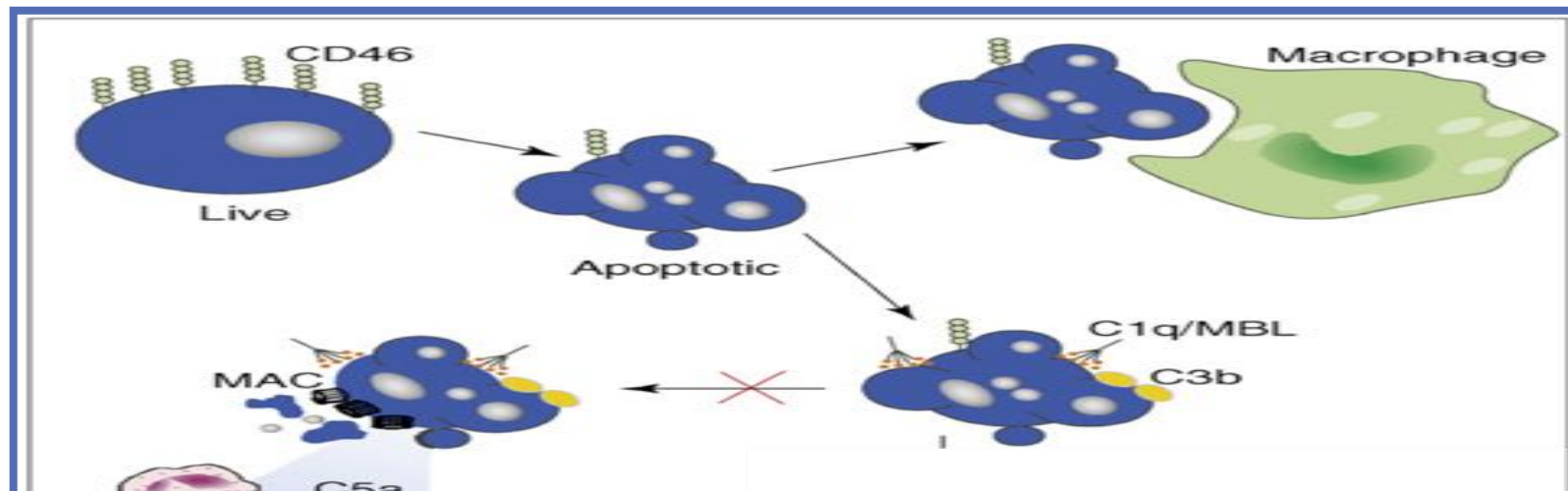
CLASSIC PATHWAY



activated by pentraxins !

CLASSICAL COMPLEMENT ACTIVATION IN THE NATURAL IMMUNITY AND INFLAMMATION!

In addition to recognizing the Fc portion of antibodies, C1q binds to pentraxins (CRP, SAA, AAI, AAI3) through its gC1q domain. gC1q also binds directly to many gram-negative bacteria through Omp, LPS, or lipid A and to viruses (e.g., gp41 of HIV-1 or gp21 of HTLV-1). C1q also interacts with misfolded proteins, such as amyloid A β peptide and prion proteins found in neurodegenerative diseases and with several ECM proteins (such as fibronectin, vitronectin, heparin, heparan sulfate, chondroitin sulfate, and laminin). Finally, C1q binds via the globular head domain to surface blebs on apoptotic cells and to necrotic cells directly or through **pentraxins!**



MEMBRANE ATTACK COMPLEX IN CELL MEMBRANE or MAC, WHICH DESTROYS PATHOGENS

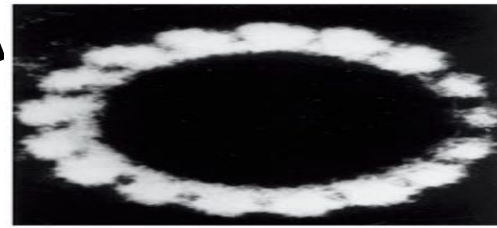
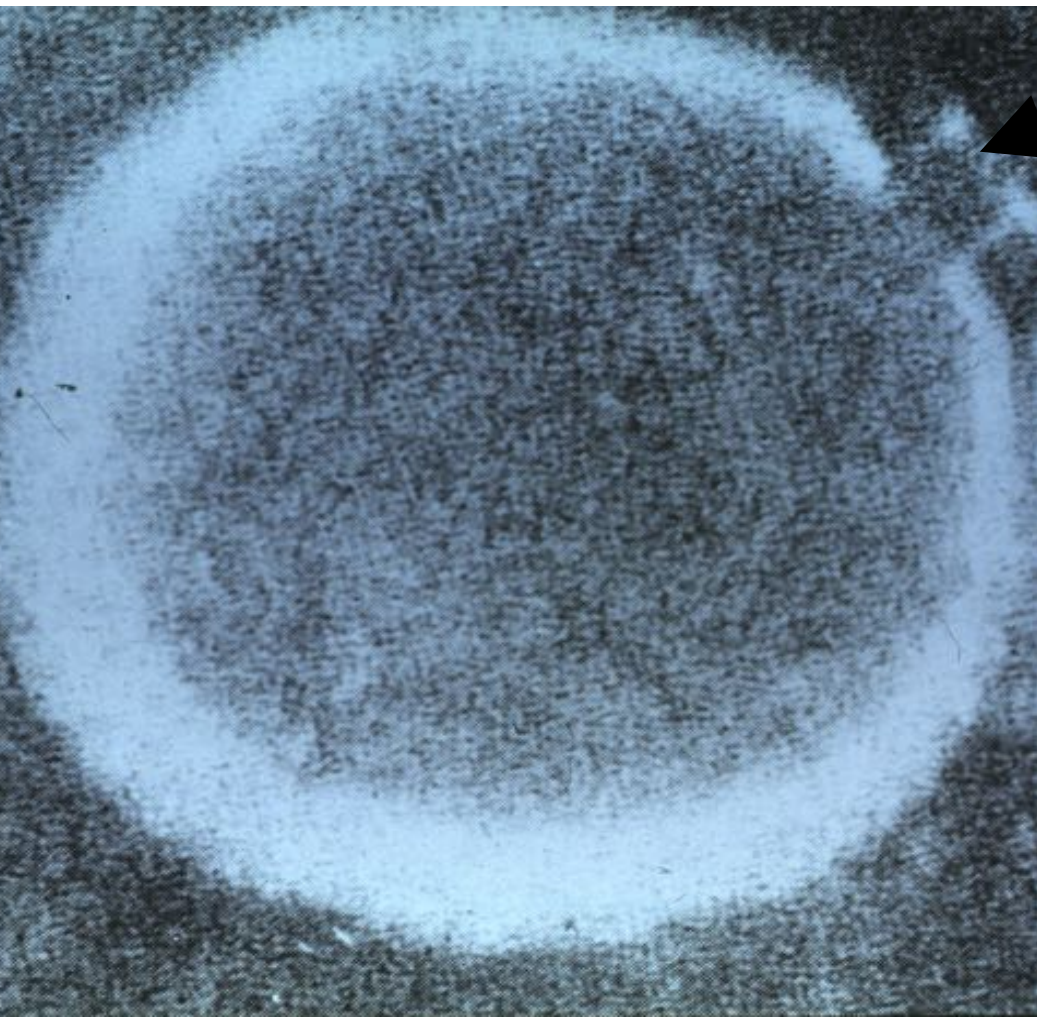
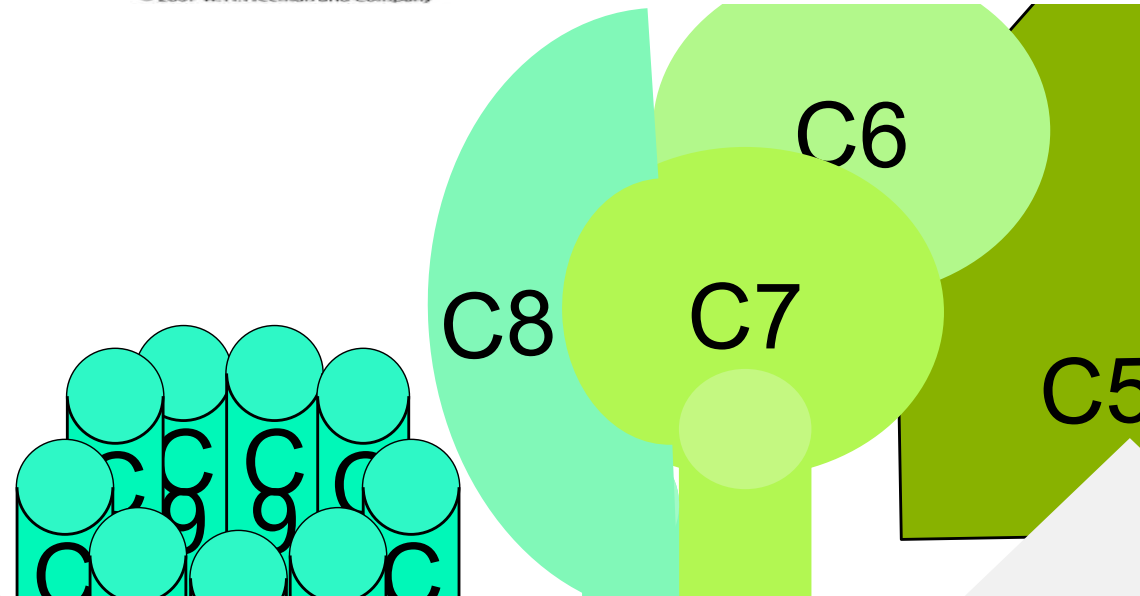
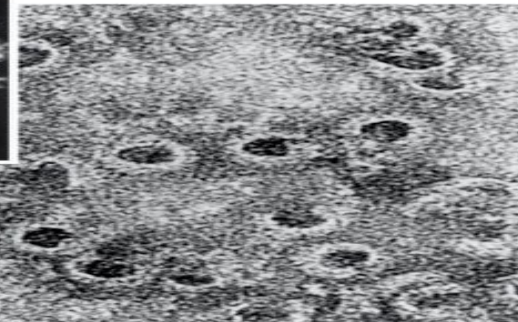
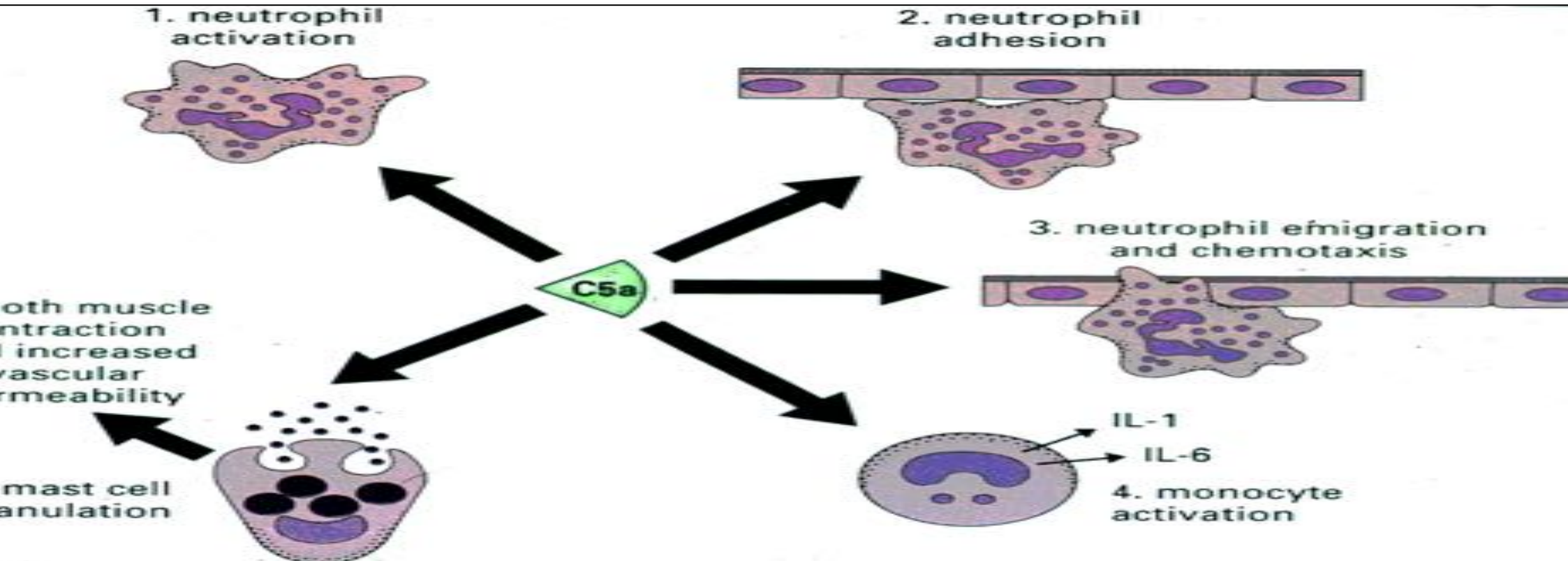


Figure 7-8
Ruby IMMUNOLOGY, Sixth Edition
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AND OF THE PHAGOCYTOSIS OSPONINS!

APHYLATOXINS (C5a-C3a)

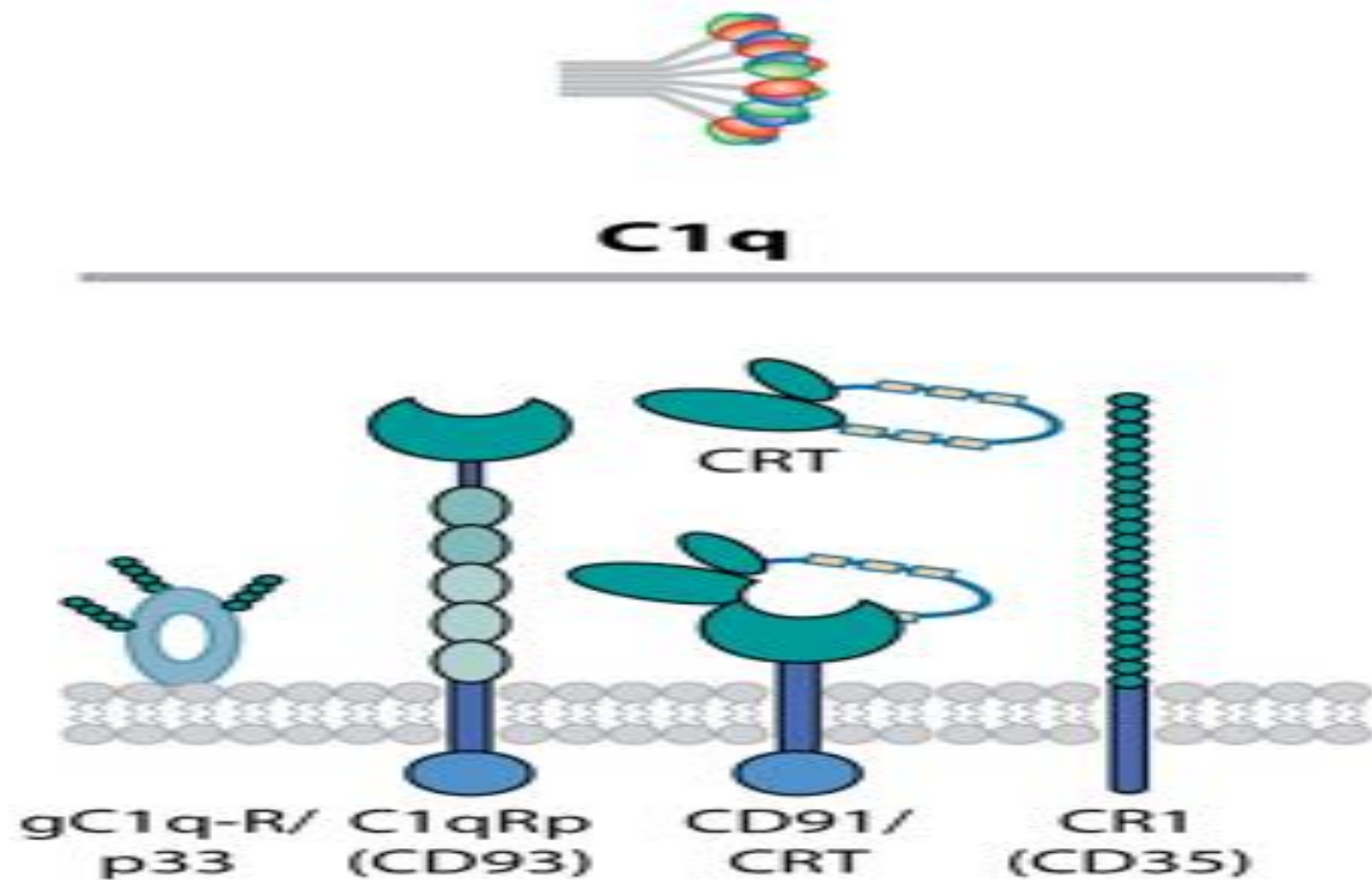


OPSONINS (C4b-C3b)-PHAGOCYTOSIS



binds to a wide range of cell types (PMN, monocytes, lymphocytes, DCs, ECs, etc), resulting in the induction of cell-specific biological responses, which include phagocytosis, chemotaxis, the generation of procoagulant activity, activation of and enhancement of FcγR- and CR1-mediated phagocytosis and superoxide production.

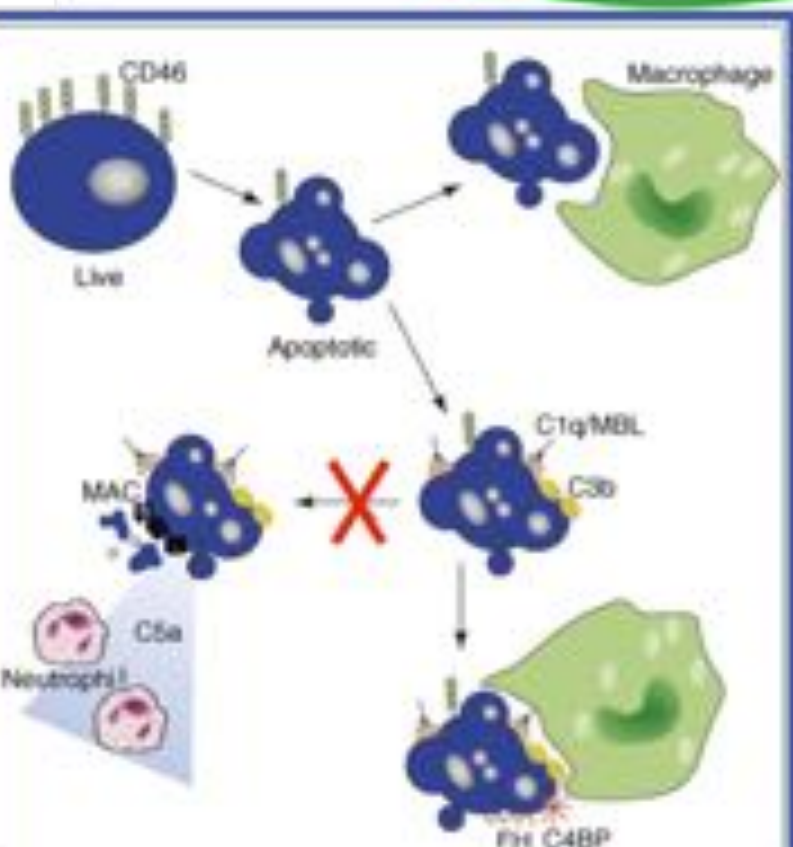
**Receptors for
C1q humoral
complement
factor !**



Recently, investigators have described four types of C1q-binding proteins/receptors expressed on the cell surface. These include cC1q-R/calreticulin (CRT), a 60-kDa protein ; gC1q-R, a 30-kDa homotrimeric protein; C1q-Rp (CD93), a 120-kDa O-sialoglycoprotein; and CR1 (CD35), the receptor for C3b. In addition to C1q, CRT reportedly serves as a receptor for

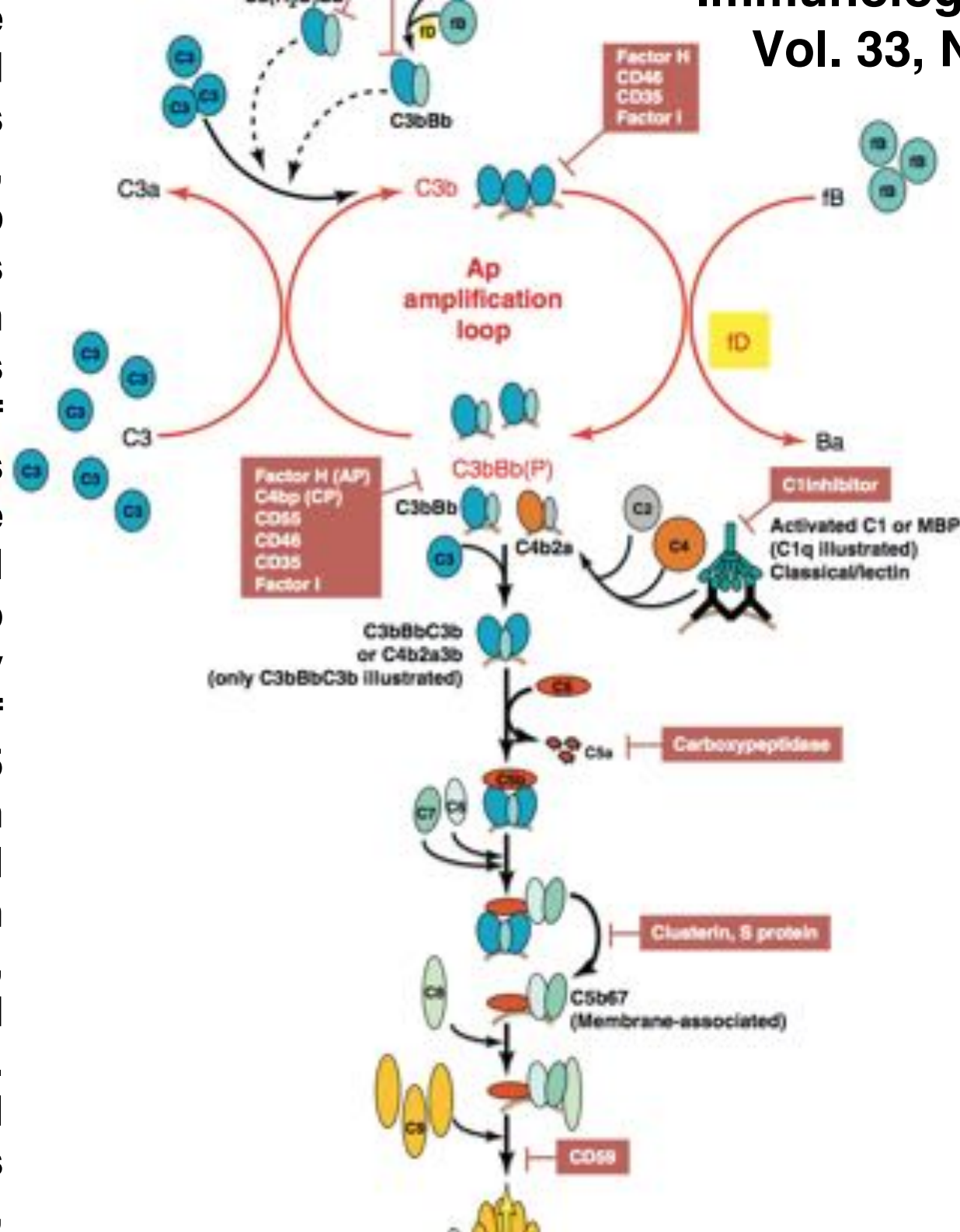
Table 1 | Complement interactions with pathogens and self

	Activation profiles	Outcomes	Examples
Pathogen	Robust and unrestricted	Inflammation and immunity	Bacteria and viruses
Altered self	Limited and targeted	Mild inflammation and no immunity	Apoptotic and injured cells and tissues; lipid and proteinaceous debris
Normal self	Baseline (through tickover)	No inflammation and no immunity	Healthy cells and tissues

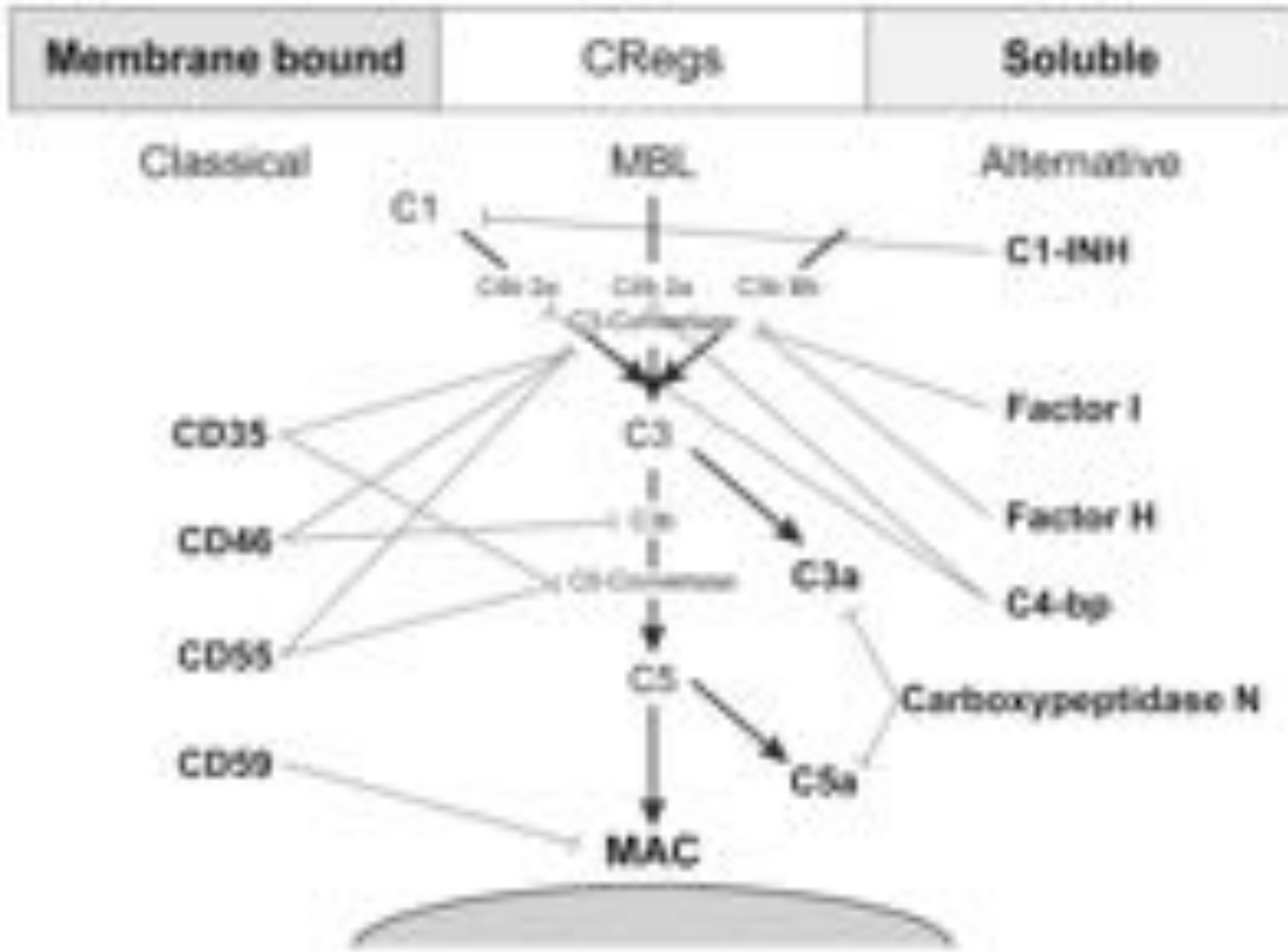


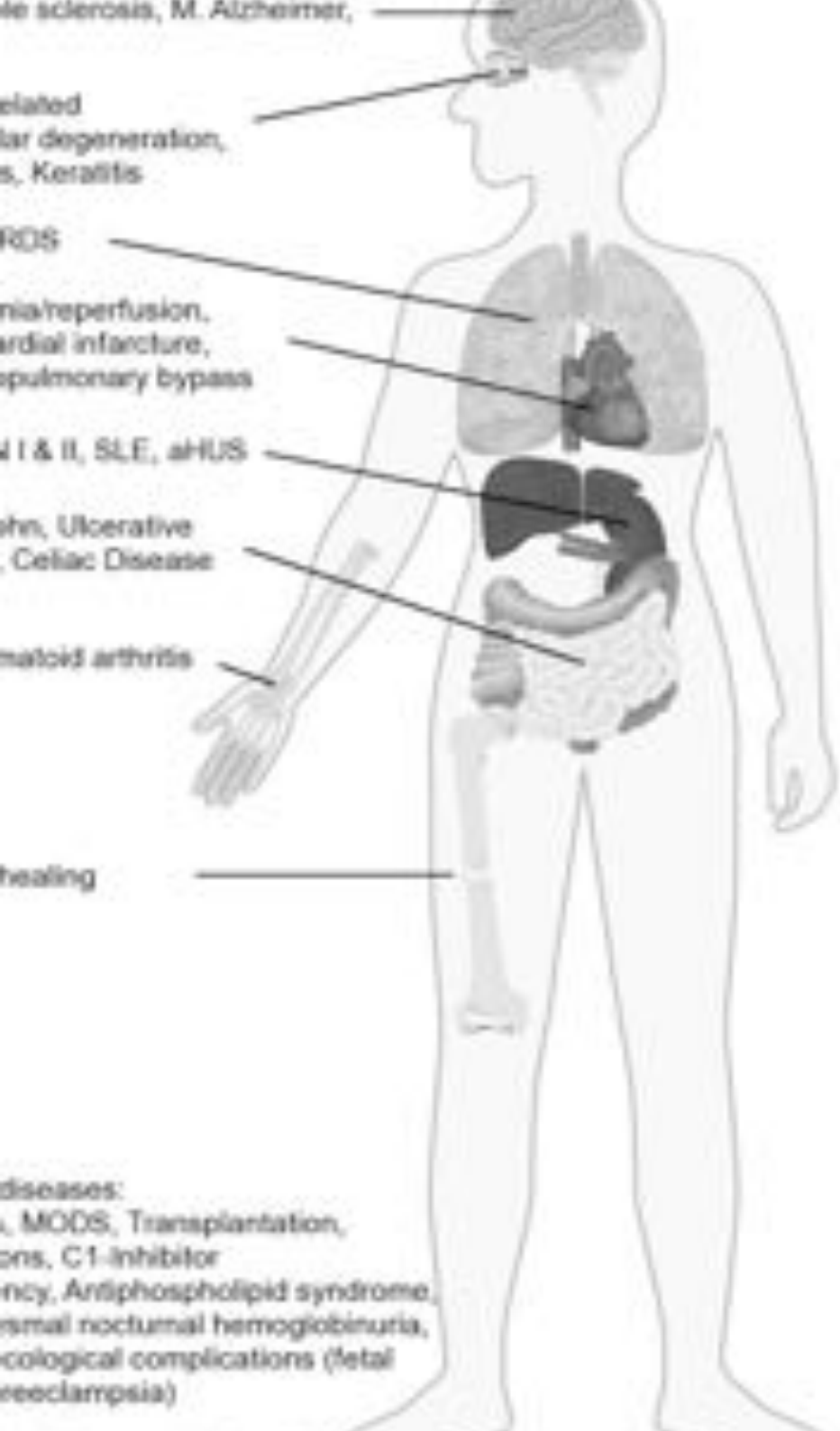
Complement activation on apoptotic cells depends on recognition by C1q and C3b/iC3b; the binding of Factor H and C4BP allows phagocytosis, without substantial activation of the terminal complement pathway and inflammation

lysis of the C3 thioester, or cleavage to C3b by plasma proteases. Fluid production of either molecule results in formation of the AP C3 convertase, C3b, and production of further C3b. C3b either binds a surface or remains in solution. Each newly produced C3b can form a convertase, which cleaves C3 resulting in exponential production of C3b. This self-propagation, referred to as the 'amplification loop' and indicated here, is responsible for amplifying a small amount of C3 to yield large responses. C3b can enter through any activation pathway into the amplification loop. Binding of C3b to C3 convertase creates C5 convertase; cleavage of C5 and generation of C5b marks the start of the terminal pathway. C6 and C7 bind C5b to form C567, which is released from convertase, incorporates C8 and C9 molecules to form the MAC. Complement damage is protected from accidental damage by regulatory proteins present in plasma and on membranes,

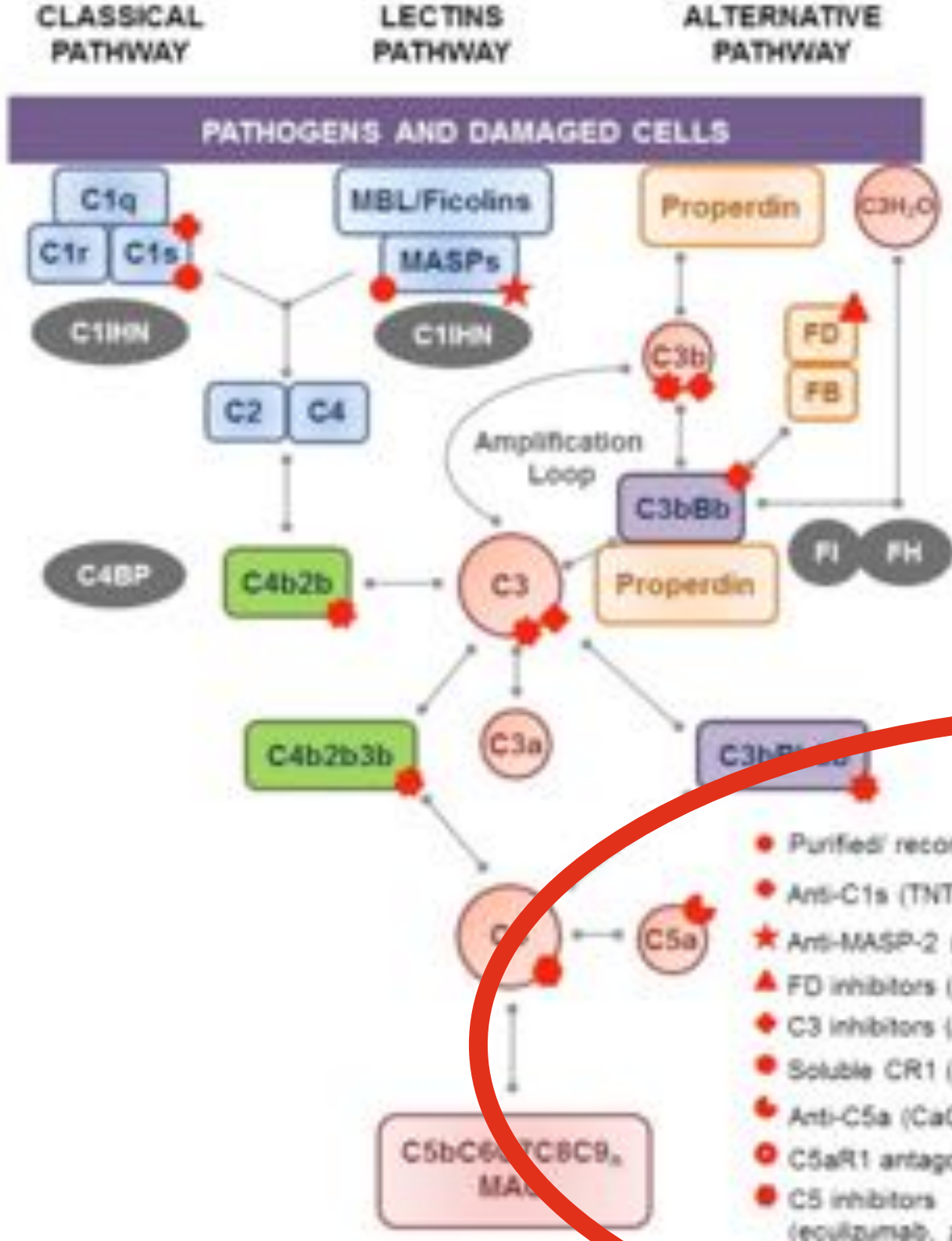


Membrane regulators acting on different stages of the complement cascade!





ALS, amyotrophic lateral sclerosis; ALI, acute lung injury; ARDS, adult respiratory distress syndrome; MPGN, membranoproliferative glomerulonephritis; SLE, systemic lupus erythematosus; aHUS, atypical hemolytic uremic syndrome; MODS, multiple organ dysfunction syndrome.

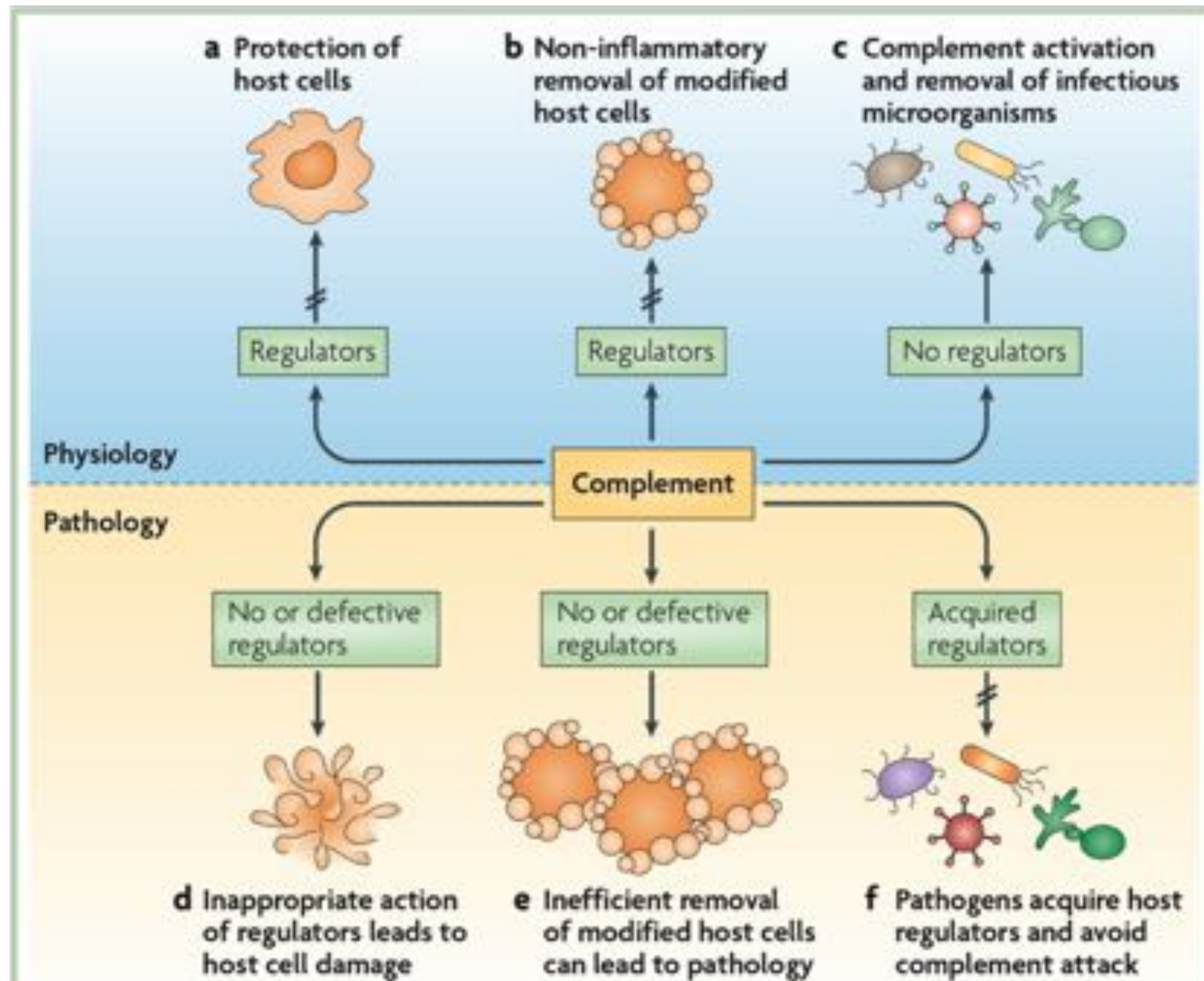


Clin Immunol. 2015 Sep 1;161(2):225-240.

Applying complement therapeutics to rare diseases.

Reis ES, Mastellos DC, Yancopoulos D, Risitano AM, Ricklin D, Lambris

Complement activation has multiple effects, which can either benefit OR be detrimental to the host and possibly lead to pathology



Hemolytic assay or CH50 (or AH50)!

CH50: defining the amount of complement required to induce 50% lysis of sensitized erythrocytes.

Is expressed as the reciprocal of the dilution serum that provides 50% lysis.

Serum sample

+

Sheep erythrocytes pre-sensitized with specific antibodies.

Electrophotometric measurement of the hemoglobin released.

Correlation between hemoglobin released.

CH50 reduction correlated with the reduction of the levels

observed reduction of complement by:

consumption of C for the formation of immune-comple

creased synthesis of C;

creased catabolism of C.

Phase of infection	CH50	C4	C3	Factor B	Conditions with Activation Pathway
Acute	Decreased	Decreased	Decreased	No change	SLE, SS, RA, and cryoglobulinemia
Subacute	Decreased	No change	Decreased	Decreased	Endotoxemia; II MPGN
Chronic and relapsing	Decreased	Decreased	Decreased	Decreased	SLE, shock, and immune complex diseases
Phase II— relapsing	Decreased	Decreased	No change	No change	Hereditary angioedema; malarial infection (<i>P. vivax</i>)
Phase III	Significantly increased	Significantly increased	Significantly increased	Significantly increase	Acute and chronic inflammation; pregnancy

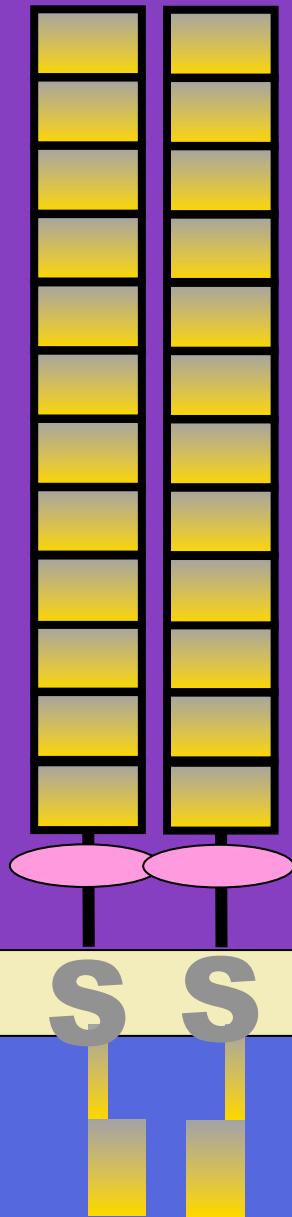
SLE, systemic lupus erythematosus; SS, Sjogren syndrome; RA, rheumatoid arthritis; MPGN, membranoproliferative glomerulonephritis.

The LPS binding protein or LBP was identified in 1990 and is present **in serum at a concentration of less than 0.5 $\mu\text{g/ml}$ but which reaches 50 $\mu\text{g/ml}$ at 24 hours during an APF**

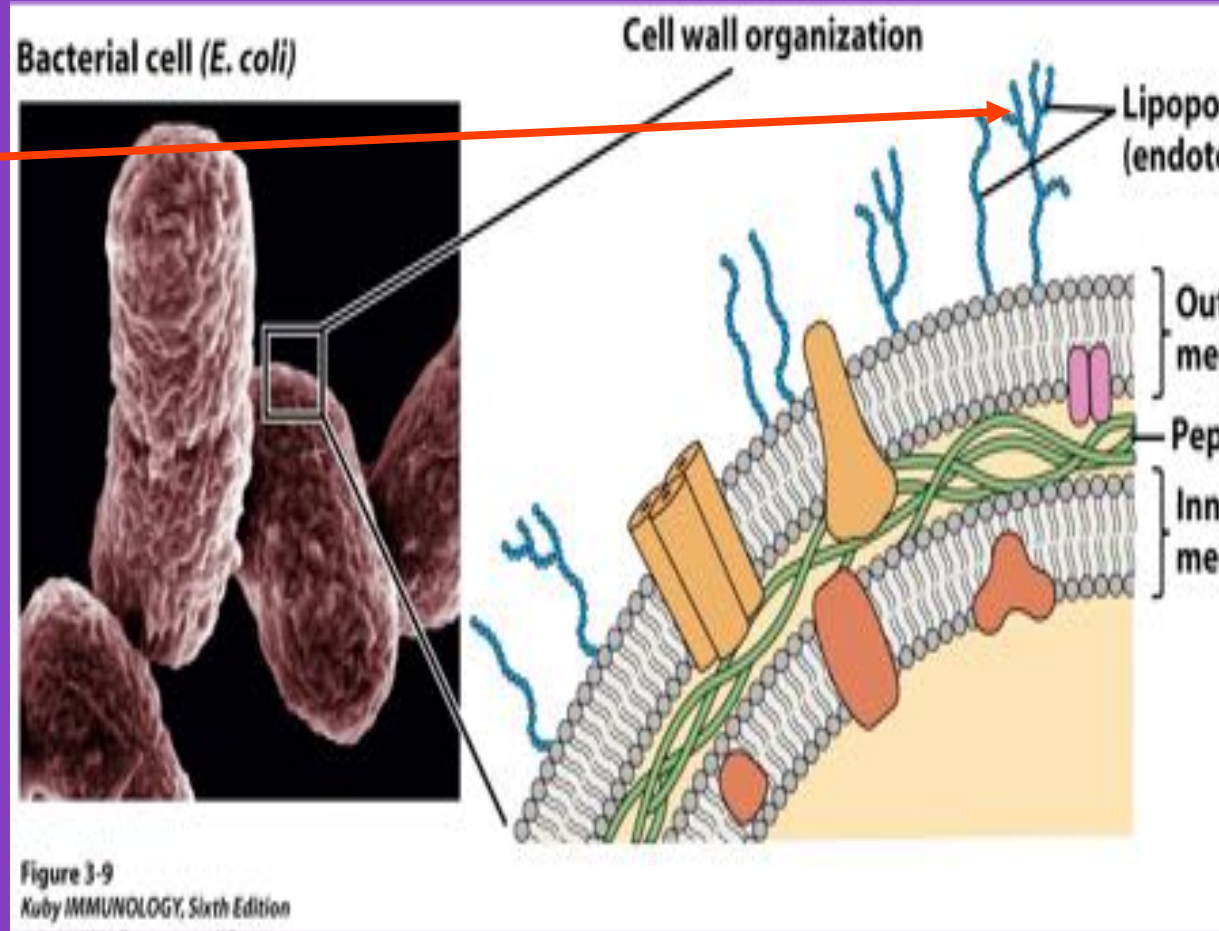
It is a 60 kDa protein synthesized by hepatocytes, has two binding sites for the lipid A of LPS, which binds with high affinity to CD14 and transports of phagocytes to the site of subsequent binding with TLR4 and activation of the production of inflammatory cytokines.

Although deficits have not been found in humans, the importance of LBP is underscored by the fact that **knockout mice (KO) to LBP are much more susceptible to Salmonella**

TLR4

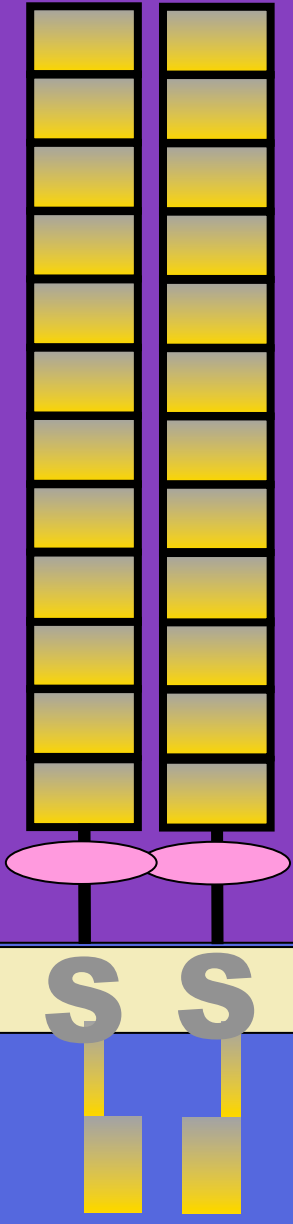
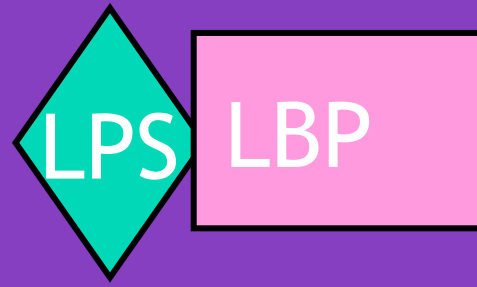


LPS

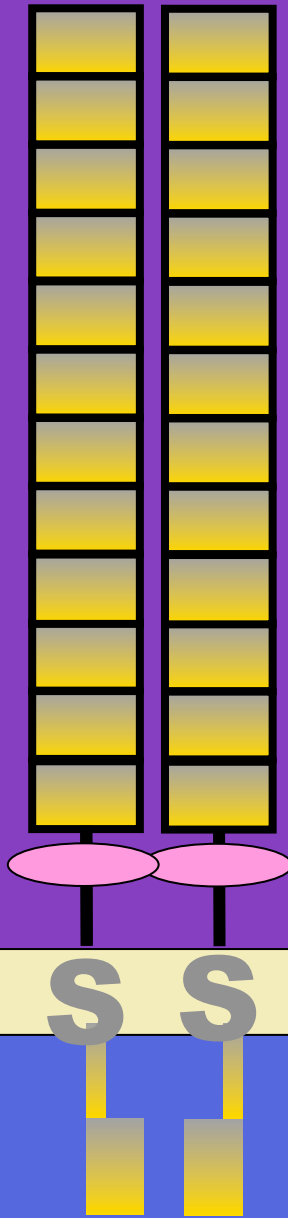


LPS binding prote

TLR4



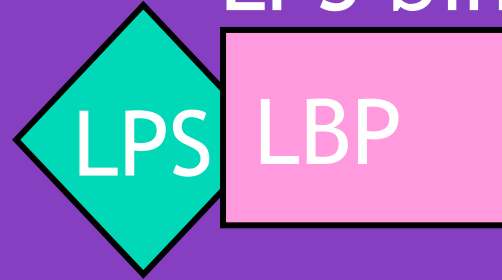
TLR4



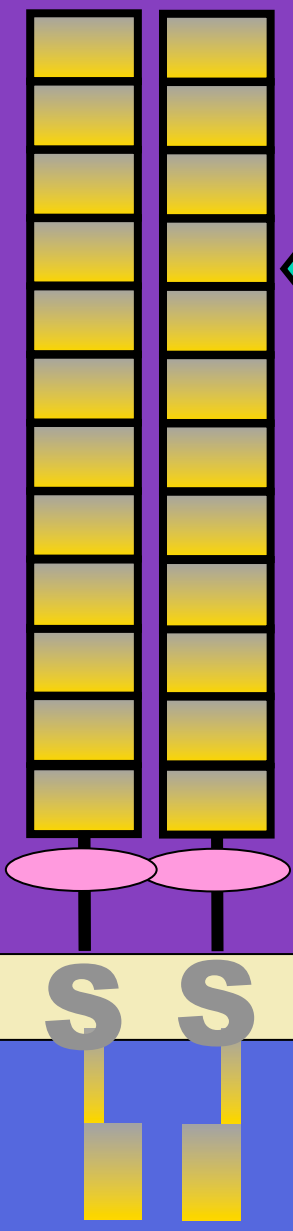
CD14



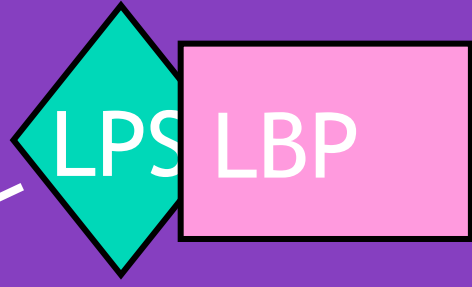
LPS binding protein



TLR4

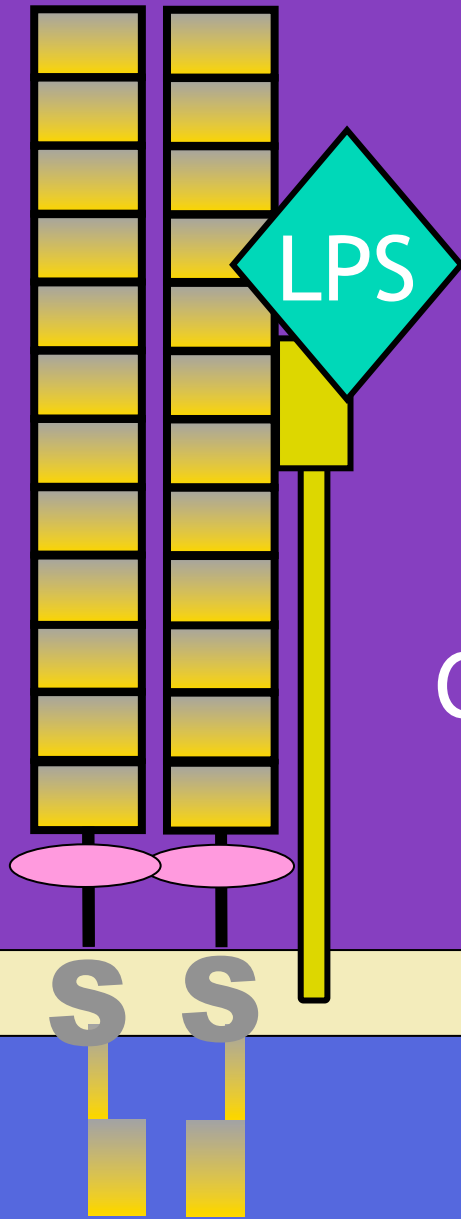


CD14



LPS binding protein

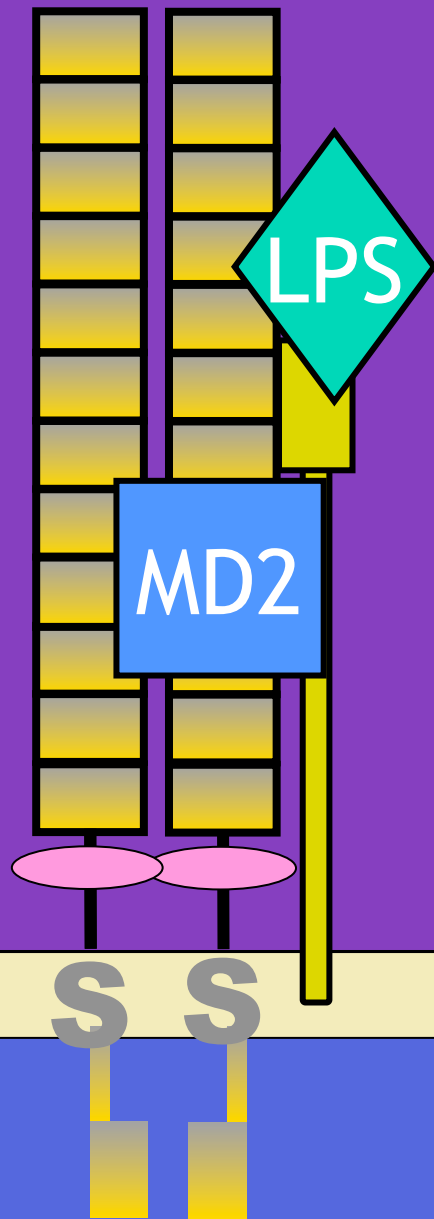
TLR4



LPS

CD14

TLR4



LPS

MD2

CD14

S

S

TLR4



LPS

MD2

CD14

S S



SIGNAL TRANSDUCTION PATHWAYS

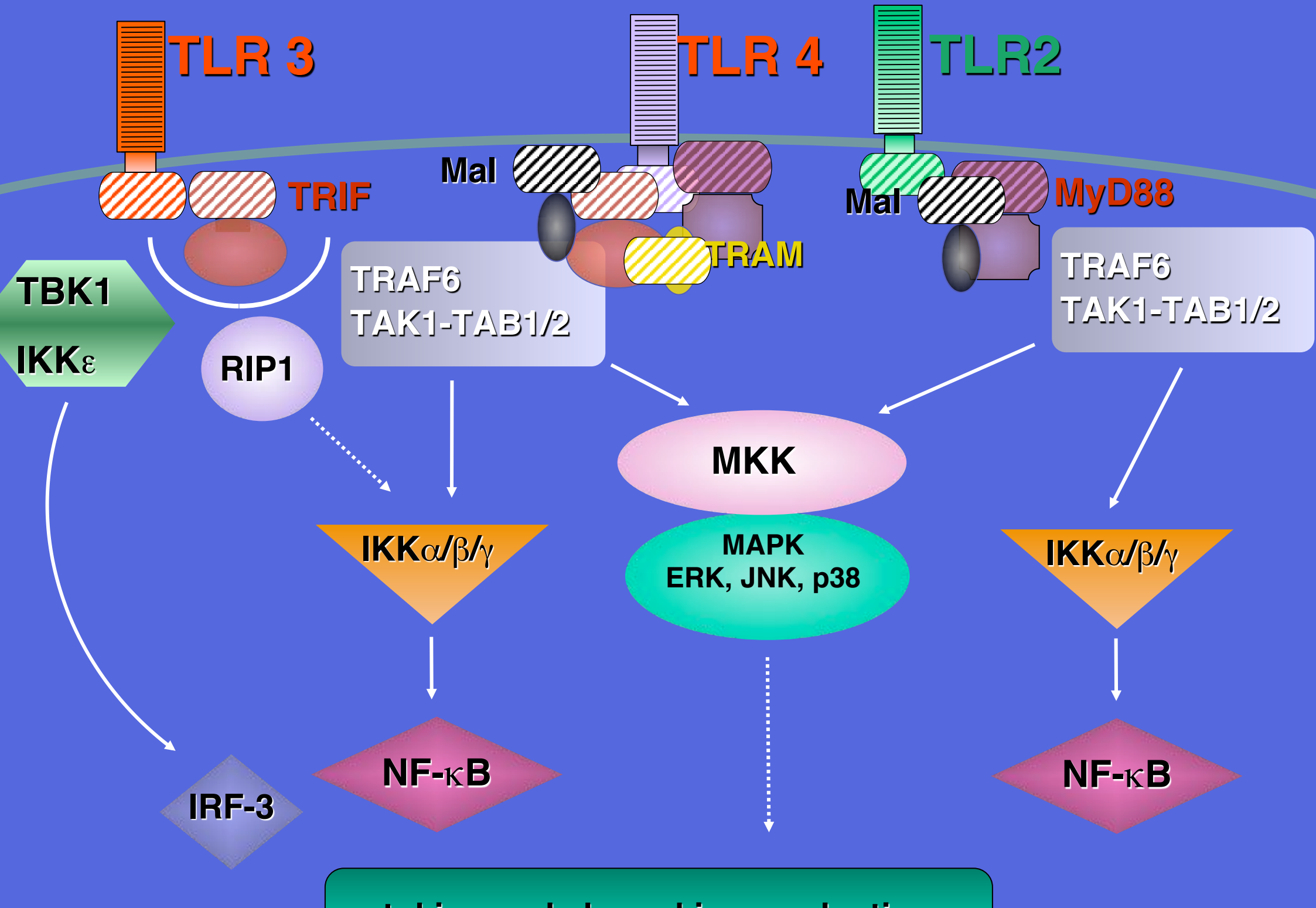


Table 1 Ligands recognized by pattern-recognition molecules

Receptor	Producers ¹	Ligands
Mincle	Liver (hepatocytes)	• Complement components (C1q, Factor H, L-ficolin, M-ficolin)
		• Microorganisms (bacteria, viruses, fungi, parasites)
		• Phosphorylcholine, carbohydrates
		• Modified LDLs
		• ECM protein (fibronectin, collagen IV, laminin, proteoglycans)
		• Amyloid fibrils
		• DNA
MD-2/CD14	Monocytes, M Φ , PMN, EC, DC, fibroblasts, epithelial cells	• Complement components (C1q, Factor H, L-ficolin) • Microorganisms (bacteria, viruses, fungi) and microbial moieties (OmpA) • ECM protein (Iol, TSG-6) • Apoptotic cells • FGF2
	M Φ , DC, EC	• Fc portion of immunoglobulin
		• Pentraxins (CRP, SAP, PTX3)
		• Microorganisms and microbial moieties (LPS, lipid A, Omps)
		• A β peptide of prions
		• ECM protein (fibronectin, laminin, fibromodulin, osteoadherin)
		• Apoptotic cells
TL, SP-1	Liver (hepatocytes), lung (type II alveolar cells), M Φ	• Carbohydrates • Microorganisms and microbial moieties (LPS, LOS, LTA, PDG)
	Liver (hepatocytes), lung (type II alveolar cells), PMN, monocytes	• Carbohydrates • Microorganisms and microbial moieties (LTA, PDG, 1,3- β -D-glucan)
	Monocytes, M Φ , PMN, mast cells	• Complement components (C3b)
		• Microorganisms, zymosan
	Liver (hepatocytes, monocytes, M Φ)	• Microorganisms and microbial moieties (OmpA)

proteins of the coagulation system and fibrinolysis:

**fibrinogen, plasminogen, tissue plasminogen
activator, Protein S.**

gen is the most abundant plasma contains from 100 to 400 mg/dl. With an
lar weight of 340 kDa, fibrinogen is a dimer composed of three pairs of peptide
(and gamma-B) linked by disulfide bridges, multiple proximate to the N-terminal.
ains extend outside in two other identical domains (D) at the C-terminal in w
hains are intertwined. Thrombin detaches the fibrinopeptides of A and B from
al ends, forming a fibrin monomer, which polymerizes into fibrils, and
dinally, which in turn form the clot macroscopic.

Fibrinogen levels become elevated in acute phase!

ogen levels increase during pregnancy and the use of contraceptives.

levels generally indicate an extensive activation of coagulation with consump
ogen.

are several variants of hereditary fibrinogen pathologies, some with relative alt
gulation and bleeding diathesis others with an increased tendency to thrombosi

Erythrocyte Sedimentation Rate

ESR measures the rate at which erythrocytes fall or settle in the plasma of a randomly selected blood specimen over a specified period of time (usually 60 minutes) in millimeters per hour; however, newer methods involving centrifugation can generate results in approximately 5 minutes. This phenomenon was first observed by Edmund Faustyn Biernacki in 1888, who noted that the rate at which blood settled varied among individuals and that red blood cells (RBCs) settle more quickly in the presence of increased levels of fibrinogen.

In 1897, Dr Robert Fahraeus noted that ESR differed in pregnant versus nonpregnant women and proposed the test as a possible indicator of pregnancy. In 1921, Dr Alf Vilhelm Albertsson Westergren introduced the test as a laboratory indicator of the prognosis of patients with pulmonary tuberculosis. Westergren defined the measurement standards for the ESR test that still are used widely today, including the utilization of sodium citrate as an anticoagulant.

ESR can be confounded by many factors, leaving this widely used test vulnerable to misinterpretation in clinical practice. Aggregation of erythrocytes promotes falling and increased sedimentation; however, RBCs are negatively charged and tend to repel one another. Thus, the presence of positively charged, large, asymmetric acute phase proteins such as fibrinogen and immunoglobulins increases the ESR. The rate of erythrocyte settlement can be influenced by a wide variety of immunologic and non-immune factors, including alterations of the quality and quantity of the RBCs, as well as variations in the normal patterns and amounts of various plasma proteins.

rinogen levels become
ted in acute phase up to
s of occasional **over 1.0**
In this case also becomes
kedly elevated **the**
rocyte sedimentation
(ESR): it is believed that
% of the increase of ESR
ue to the fibrinogen
ralizing effect on the sialic
residues of red blood cells
are known to inhibit the
rocyte aggregation!

Erythrocyte sedimentation rate

Expense: Low

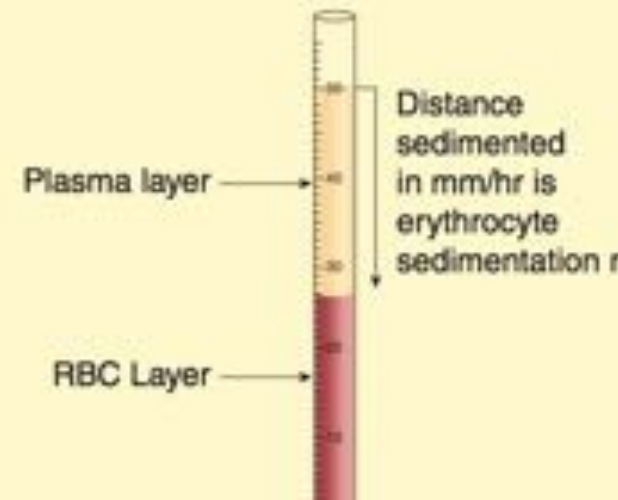
Manual or semi-auto

Goal of the test is to measure the height of sedimented RBC after an incubation, often 1 hour

Whole blood placed in a cylindrical vessel with markings to assess column height



RBC allowed to sediment undisturbed within cylindrical vessel



Antiproteases:

-antitrypsin (AAT) and the α 1-anti-chymotryps

alpha-1-antitrypsin is also referred to as alpha-1 proteinase inhibitor (AAT) because it inhibits a wide variety of proteases.

reference range in blood of 1.5 - 3.5 gram/liter.

Although AAT is a powerful APP, its increase in inflammatory processes has low clinical specificity.

Normally there are no appreciable amounts of trypsin in the circulation. However, it and other similar proteases, such as collagenases, are produced predominantly by leukocytes in response to inflammatory stimuli from necrotic or damaged cells. The AAT is able to neutralize these proteases, thereby preventing tissue damage, and from this derives its physiological function of homeostatic control of endogenous proteolysis in the body.

function is very important and it was revealed by the discovery that the serum of some young adults with pulmonary emphysema and cirrhosis of the children were deficient AAT.

The majority of individuals are homozygous for M, the functional allele of the AAT gene, and has the MM phenotype. About 10% of the Caucasian population is heterozygous for M and other alleles of AAT, as the PiZ. More than 2% are carriers of the allele PiZ and has the MZ phenotype. Although these individuals are asymptomatic, their descendants ZZ are susceptible to lung disease or liver disease.

Serum protein electrophoresis can be used for screening for AAT deficiency, but it is necessary to perform confirmatory testing complex, such as **trypsin inhibitory capacity (TIC)**, so the phenotype seeking to cross electrophoresis or isoelectrofocusing in order to exclude the presence of some other allele as PiS or PiF that migrates differently. The ZZ phenotype ICT has a very low which corresponds to very low concentrations of AAT. **It is essential that such persons should avoid cigarette smoke as this activates alveolar macrophages to release proteases**

α 1-anti-chymotrypsin!

is not only highly specific for chymotrypsin, a protease that cleaves the peptide bonds at the carboxyl site of tyrosine and phenylalanine, but it is the only APP.

, which has a molecular weight of 68 kDa with approximately 5% of the carbohydrate content and a normal serum concentration from 40 to 60 mg/dL, can rapidly increase up to 100 mg/dL during and for the duration of inflammation.

Transport proteins:

ceruloplasmin, haptoglobin and hemopexin.

ceruloplasmin consists of a single polypeptide chain, can bind six atoms of copper. The presence of copper in ceruloplasmin give a blue color to the protein and "in vitro" activity manifests oxidase activity. The level of ceruloplasmin at birth is lower, its serum level ranges from 20 to 40 mg/dL in young adults, increasing to twice in the treatment of contraception and pregnancy. Ceruloplasmin is an acute-phase reactive.

Ceruloplasmin is a glycoprotein essential for the body to transport the copper. The removal of iron from the tissues through the activity of the enzyme ceruloplasmin ferroxidase.

Aceruloplasminemia is a genetic disease with an autosomal recessive inheritance pattern caused by a mutation of a gene located on chromosome 3. Unlike Wilson's disease, which is inherited in an autosomal recessive and caused by mutations in the ATP7B gene, which codes for ATPase that controls the transport of copper into the bile and its excretion. In aceruloplasminemia, there are no apparent defects in the gene for ceruloplasmin ferroxidase, there are no apparent defects in

Hemopexin!

The hemopexin binds heme released after haemoglobin degradation. In this way the small molecule porphyrin, with iron atom, is protected in respect of excretion, preserving the organic deposit of iron.

The normal serum concentration is from 50 to 120 mg/dl.

Haptoglobin has two heavy chains and two light chains that, of different molecular weight and joined by disulfide bridges, determine three hapto-phenotypes: (1-1), (2-1) and (2-2).

mean serum HTG concentration is (2.5 +/- 1.2 g/L).

Binds hemoglobin released by lysis of erythrocytes in order to preserve the protein reserves. The hemoglobin - haptoglobin complexes are removed by macrophages in the liver and spleen of the reticuloendothelial system to ensure recovery of the heme-iron. Therefore, the physiological function of haptoglobin is to allow the recovery of iron when red blood cells, at the end of their circulation are destroyed (hemolysis saline).

Under normal conditions the concentration of haptoglobin in the circulation is the result of the balance between its synthesis in the liver and its elimination.

concentration of haptoglobin is therefore inversely proportional to the hemolysis. The serum haptoglobin also increases in response to stress, infection,

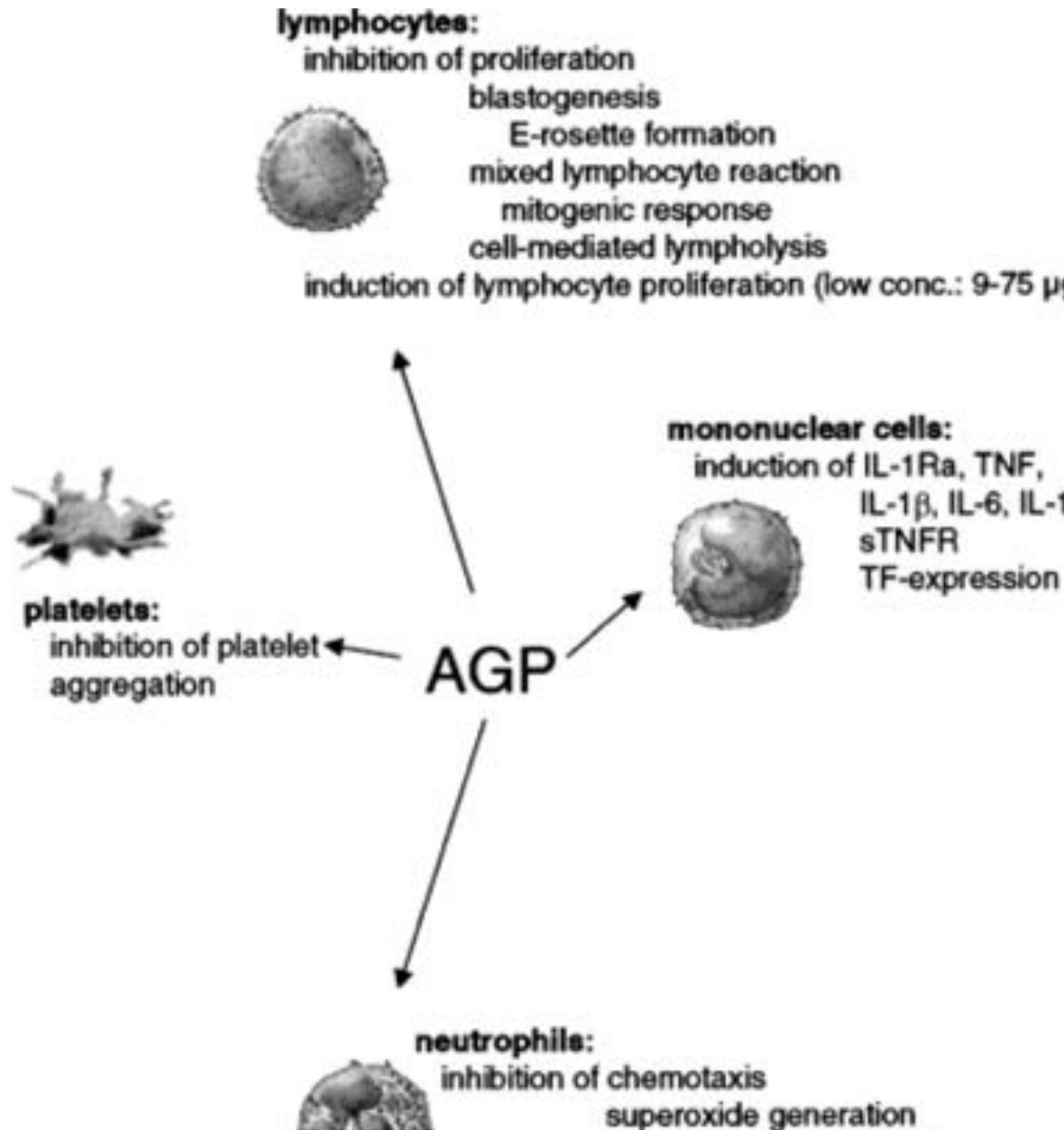
**new APP that can be used as a marker of
systemic or localized inflammation:
 α 1-acid glycoprotein, soluble CD14 or CD14S,
monocyte-specific S100 calcium-binding proteins and
procalcitonin!**

the α -1 acid glycoprotein (AGP) or **Orosomucoid (ORM)** protein with a molecular weight of 41-43 kDa and glycosylated (45%).

P serum concentrations are between 0.6-1.2 mg/dL and increase considerably in the case of acute phase inflammatory response.

is known as the primary carrier of basic drugs (where albumin carries acidic drugs), steroids, and protease inhibitors.

Overview of effects of AGP on lymphocytes, platelets, mononuclear cells and neutrophils.



The soluble CD14 (sCD14) is the soluble form of CD14, a protein of the membrane of monocytes-macrophages, anchored by glycosyl bond-phosphatidilico-inositol and that functions as coreceptor for the LPS.

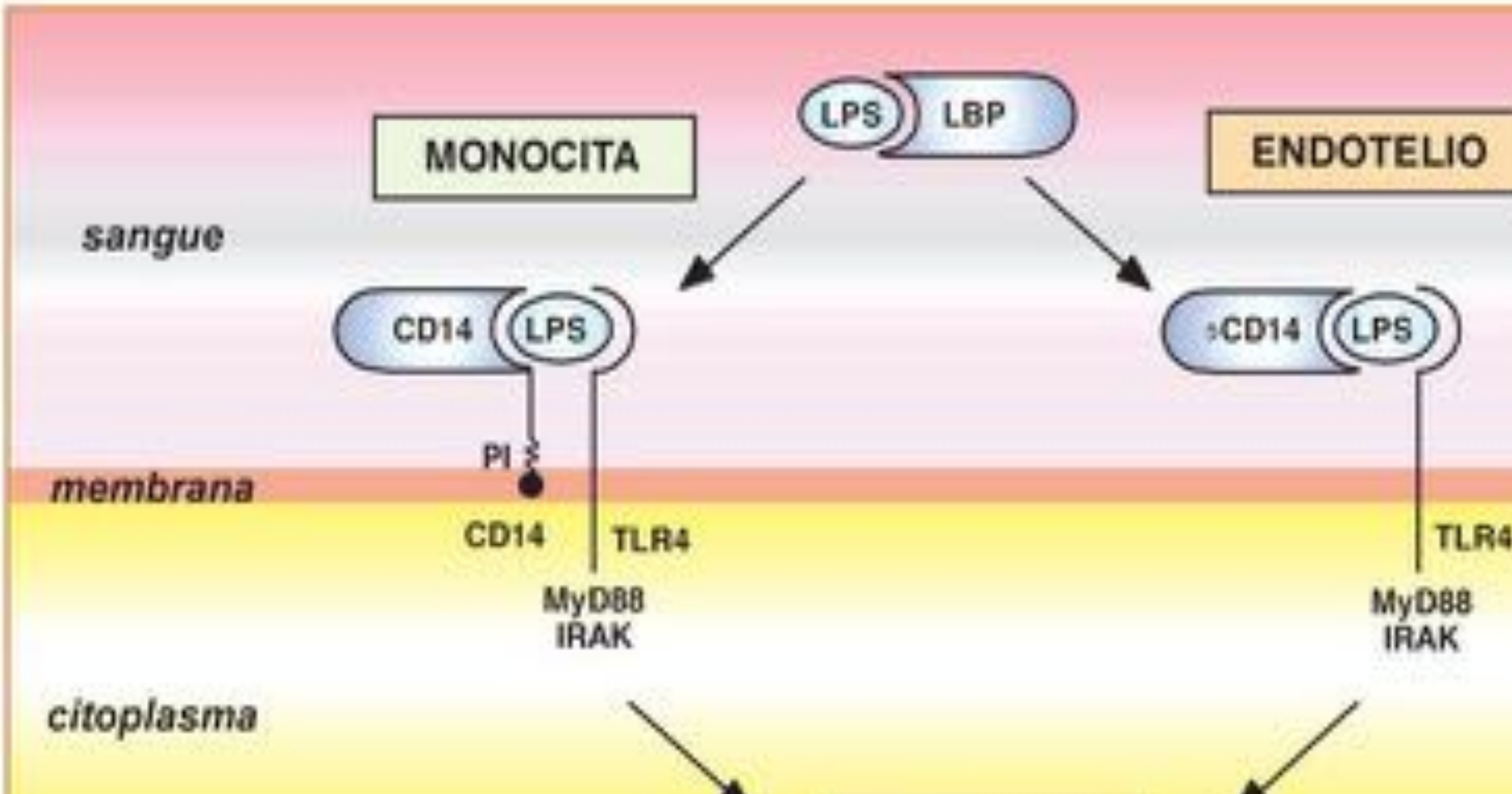
Its **serum levels are low (around 100 ng/ml)** but increase as consequence of the activation LPS-mediated monocytes-macrophages that release large amounts of sCD14.

If it was initially postulated that the sCD14 was released to desensitize monocytes-macrophages and to limit the production of inflammatory cytokines. Recently, it has been detected its presence in breast milk, where it seems that enhances the differentiation of B lymphocytes. Plasma levels

CHEMOKINES

MONOCYTES

ENDOTHELIUM



Presepsin as a potential marker for bacterial infection relapse in critical care patients

A preliminary study.

Mini V, Ceccarelli G, D'Alessandro M, Colleparado D, Morelli A, D'Egidio A, Mariotti S, Nicoletti A, Evangelista B, D'Ettorre G, Angeloni A, Venditti M, Bachetoni A.

Bacterial infection carries a high risk of mortality in critical care patients. Improved diagnostic procedures are required for effective management of sepsis. Recently, **the soluble CD14-derived presepsin, has been suggested as a reliable marker of sepsis**, and we set out to compare its diagnostic performance with that of procalcitonin (PCT). We focused on a cohort of septic patients who, during hospitalization, relapsed after a period of clinical relief from symptoms.

In total 21 adult patients were studied during their hospitalization in the Critical Care Unit of Umberto I hospital; 74 plasma samples were collected at multiple time points, and **presepsin and PCT levels were measured using a PATHFAST® analyzer.**

Presepsin and PCT were significantly lower in healthy controls than in sepsis or severe sepsis. Both enabled a significant difference to be detected between systemic inflammatory response syndrome (SIRS) and severe sepsis ($p < 0.05$). The area under the curve (AUC) calculated from the receiver operating characteristic (ROC) curve analysis was 0.888 for presepsin and 0.910 for PCT. **In those patients in whom clinical recurrence of sepsis was observed, while PCT levels normalized during the relapse phase, presepsin levels (>1000 pg/mL) remained high.**

Conclusions: This study confirms the importance of monitoring a combination of several biomarkers for a reliable diagnosis. **Maximal presepsin levels could alert clinicians not to suspend antibiotic treatment and to carefully monitor septic patients' state of health even after clinical symptoms**

soluble presepsin (soluble CD14 subtype) as a marker of host response in patients

with severe sepsis or septic shock: data from the multicenter, randomized ALBIOS trial

de Santis G, Dell'Acqua R, Ferreri L, et al. *Crit Care Med*. 2018;46(12):e1000-1007. doi:10.1097/CCM.0000000000002931

Presepsin is a soluble fragment of the cluster-of-differentiation marker protein 14 (CD14) involved in pathogen recognition by the innate immune system. We evaluated the relation between its circulating concentration, host response, appropriateness of antibiotic therapy, and mortality in patients with severe sepsis.

Presepsin was measured 1, 2, and 7 days after enrollment of 997 patients with severe sepsis or septic shock in the Albumin Italian Outcome Sepsis (ALBIOS) trial. They were randomized to albumin or crystalloids. We tested in univariate and adjusted models the association of single measurements of presepsin or changes over time with clinical outcomes, appropriateness of antibiotic therapy, and ICU or 90-day mortality.

Baseline presepsin concentration (946 [492-1,887] ng/L) increased with the SOFA score, the number of prevalent organ failures, and the incidence of new failures of the respiratory, coagulation, liver, and kidney systems. Presepsin concentration decreased in ICU over 7 days in patients with negative blood cultures, and in those with positive blood cultures receiving appropriate antibiotic therapy; it increased with inappropriate antibiotic therapy ($p = 0.0009$). Baseline presepsin was independently associated with, and correctly reclassified, the risk of ICU and 90-day mortality. Increasing concentration from day 1 to day 2 predicted higher ICU and 90-day mortality (adjusted $p < 0.0001$ and 0.01 , respectively). There was no effect on presepsin concentration.

CONCLUSIONS:

Presepsin is an early predictor of host response and mortality in septic patients. Changes in presepsin concentration over time predict mortality and ICU length of stay.

Presepsin as a novel sepsis biomarker

Ji Zou, Wei Wen, Xin-chao Zhang

Emergency Medicine Department, Beijing Hospital, Beijing 100730, China

Corresponding Author: Xin-chao Zhang, Email: xinchaoz@163.com

BACKGROUND: In 2004, a new biomarker sCD14-subtypes (presepsin) was found and its value was shown in the diagnosis and evaluation of sepsis. This article is a brief overview of the new biomarker.

DATA SOURCES: A literature search using multiple databases was performed for articles, especially meta-analyses, systematic reviews, and randomized controlled trials.

RESULTS: Compared with other markers, presepsin seems to have a better sensitivity and specificity in the diagnosis of sepsis. Presepsin as a biomarker is not only suitable for the early diagnosis of sepsis, but also for the assessment of its severity and prognosis.

CONCLUSIONS: Presepsin has a higher sensitivity and specificity in the diagnosis of sepsis as a new biomarker, and is a predictor for the prognosis of sepsis. More importantly, presepsin seems to play a crucial role as a supplemental method in the early diagnosis of sepsis. Since there is no multicenter study on the relationship between presepsin and sepsis, further studies on the clinical values of presepsin are needed.

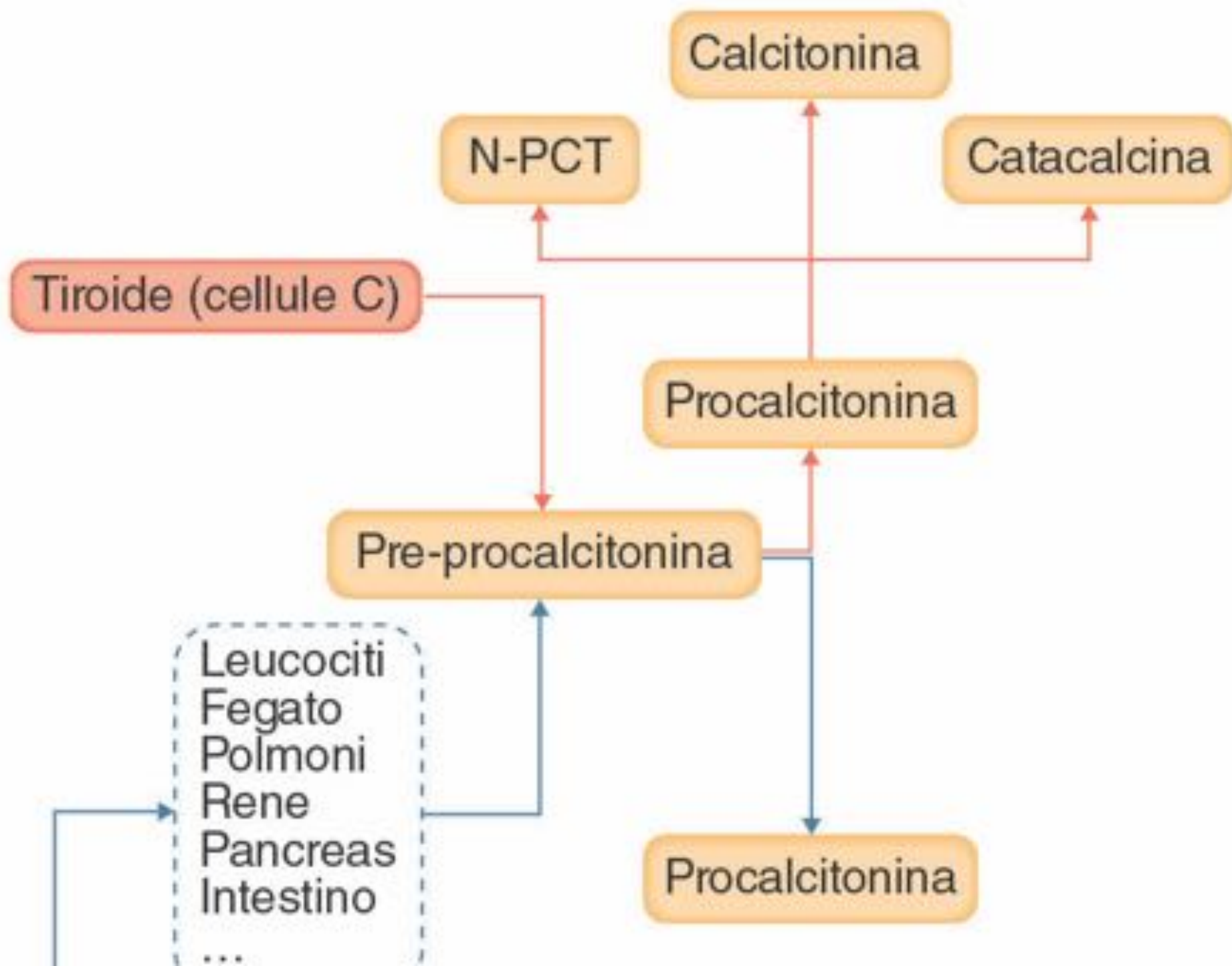
KEY WORDS: Presepsin; Sepsis; Diagnosis

Procalcitonin, or PCT, a precursor of calcitonin identified in 1975, is produced by the C cells of the thyroid gland and released into the circulation in normal plasma levels of about 5-50pg/ml.

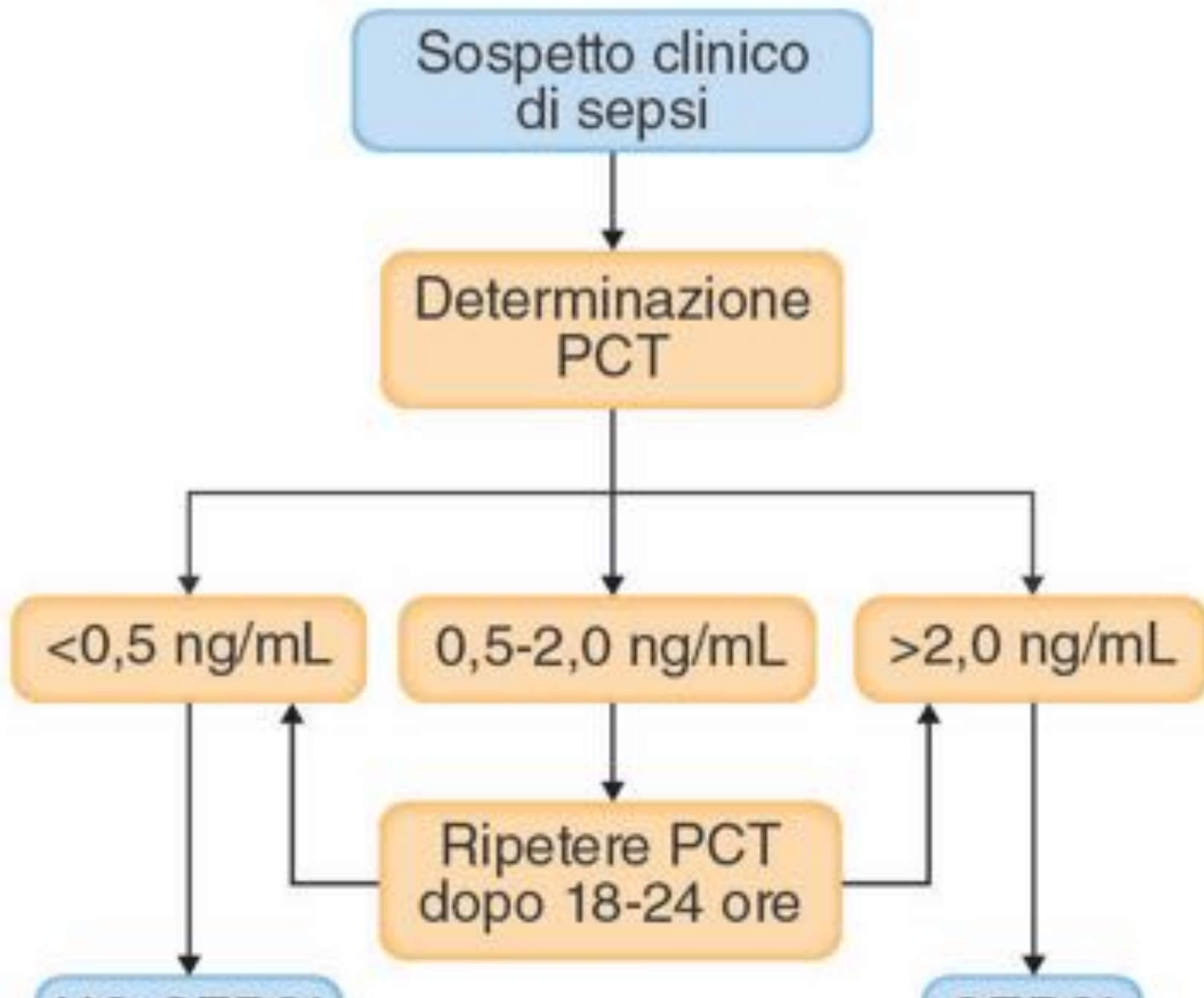
The course of inflammation is produced by the liver and PMNs and is a useful diagnostic marker of severe bacterial infections, but not viral infections and measurement may be useful to discriminate infectious SIRS from non-infectious because PCT levels increase in meningitis patients with bacterial infections. PCT levels are around 2.4 ng/ml in patients with pneumonia without sepsis and 31 ng/ml in patients with pneumonia and sepsis.

Procalcitonin appears to be a good indicator of early onset and helps to identify patients at risk of sepsis.

PCT can also be used to distinguish between septic ARDS and ARDS not secondary to sepsis. Because of these characteristics, the PCT is currently used in intensive care in



TO CHECK IN TIME THE SEPSIS!!



SEPSIS HISTORY AND DEFINITIONS!



William Osler

American College of Chest Physicians;
Society of Critical Care
Medicine Consensus
Conference

SCCM/ESICM/ACCP/
ATS/SIS
International Sepsis
Definitions Conference

The Third
International
Consensus Defi
for Sepsis and
Shock Sepsis

500 a.c.

1913

1992

2001

2016

Sepsis: infiammazione, ipotensione o ipertensione, coscienza o incoscienza, come una condizione ad alto rischio di morte, correlata a un'infezione

"con alcune eccezioni, sembra che il paziente muoia a causa della reazione del corpo all'infezione, piuttosto che a causa dell'infezione stessa"

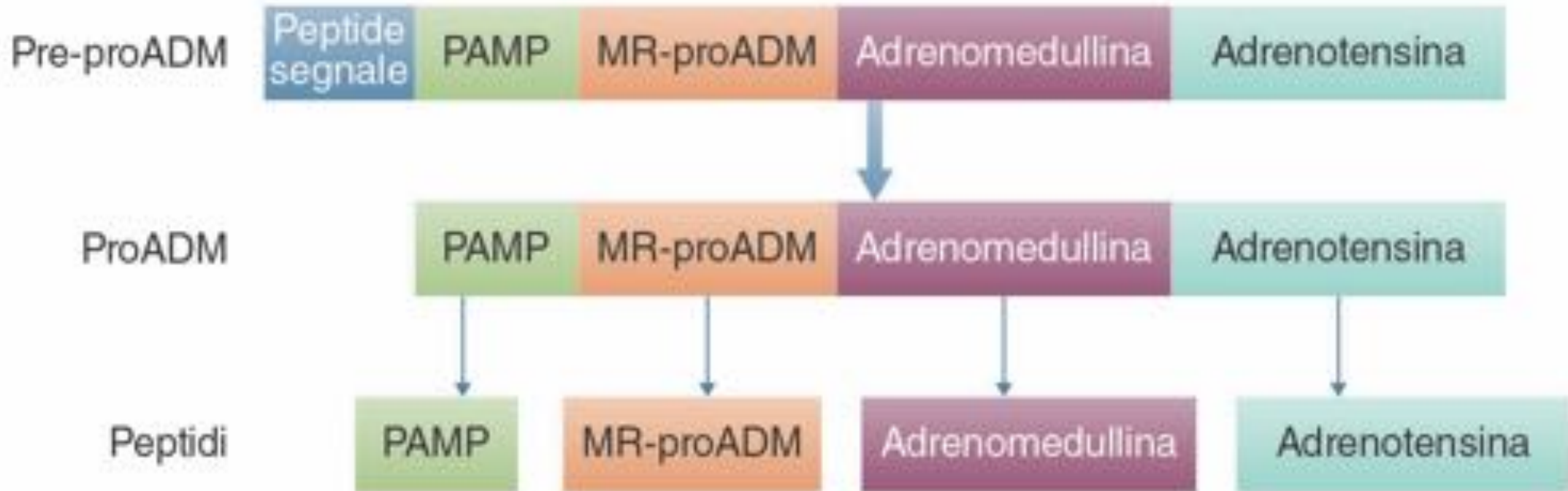
Introduzione del termine SIRS.
La sepsi è definita come la presenza contemporanea di SIRS e infezione

This document reflects a process whereby a group of experts and opinion leaders revisited the 1992 sepsis guidelines and found that apart from expanding the list of signs and symptoms of sepsis to reflect clinical bedside experience, no evidence exists to support a change to the definitions.

Disfunzione d'organo pericolosa per la vita, causata da una disordinata risposta dell'ospite a un'infezione

(ADM).

BIOSINTESIS OF ADRENOMODULIN



H LEVELS OF ADM ARE PRESENT IN SEPSIS, BUT TH
ARE UNDERESTIMATED BECAUSE BIND VARIOUS
RECEPTORS OR ARE DEGRADED BY PROTEASE.
ERNATIVE, A FRAGMENT OF 48 aa, MR-proADM, H
EEN IDENTIFIED AND IS RELEASED BY FOUIMOLAR

Test di laboratorio	Scopo
Procalcitonina	Diagnosi di sepsi e guida alla terapia antibiotica
Presepsina	Diagnosi di sepsi
MR-proADM	Indicatore prognostico
Proteina C reattiva	Diagnosi di sepsi
Emocoltura	Diagnosi eziologica di sepsi
Esame emocromocitometrico	Diagnosi di anemia e piastrinopenia
Bilirubina	Compromissione epatica
Aminotransferasi	Compromissione epatica
Creatinina	Compromissione renale
Glicemia	Scompenso metabolico
Test di coagulazione	Diagnosi precoce di CID (coagulazione intravascolare disseminata)
Troponine cardiache	Compromissione cardiaca

Clin Chim Acta. 2004 Jun;344(1-2):37-51.

Leukocyte-specific calcium-binding S100 proteins as clinical laboratory markers of inflammation

Foell D1, Frosch M, Sorg C, Roth J.

The first member of the S100 family of proteins, which are part of the larger group of calcium-binding proteins, was isolated in 1965.

S100 proteins comprise the group of calgranulins, are pro-inflammatory molecules synthesized and secreted by phagocytes. The three members of this group, S100A8, S100A9, and S100A12 are over-expressed at the site of inflammation.

The heterodimer of S100A8 and S100A9, known as "leukocyte protein L1", is now still referred to as **calprotectin** by some research groups.

While the complex S100A8/S100A9 that S100A12 are useful diagnostic factors for inflammation especially in the case of arthritis, chronic inflammation of the lung and other internal diseases.

are index of activation of phagocytes more than any other parameter of inflammation.

E, Van Assche G.

Gastroenterol Belg. 2013 Sep;76(3):322-8.

[Diagnosis and prognostics of inflammatory bowel disease with fecal neutrophil-derived biomarkers calprotectin and lactoferrin.](#)

Gastroenterol Belg. 2013 Sep;76(3):322-8.

[Fecal calprotectin in gastrointestinal disorders.](#)

Di Gennaro M, Gallo A, Santoro L, D'Onofrio F, Landolfi R, Gasbarrini A.

World J Gastroenterol. 2013 Jun;17(12):1569-82.

[Diagnostic Accuracy of Fecal Calprotectin During the Investigation of Suspected Pediatric Inflammatory Bowel Disease: A Systematic Review and Meta-Analysis.](#)

Chen P, Anderson NH, Wilson DC.

Gastroenterol. 2013 May 14.

[Inflammatory Bowel Disease: small bowel motility impairment correlates with inflammatory-related markers C-reactive protein and calprotectin.](#)

Chen P, Pazahr S, Chuck N, Blume I, Froehlich JM, Cattin R, Raible S, Bouquet H, Bill U, Rogler G, Patak MA.

Gastroenterol Motil. 2013 Jun;25(6):467-73.

[Diagnostic utility of calprotectin and lactoferrin in patients with inflammatory bowel disease: is there some value in the literature?](#)

Di Gennaro M, D'Inca R, Pathak S, Sturniolo GC.

World J Clin Immunol. 2012 Aug;8(6):579-85.

Fecal calprotectin (FC) has been proposed as a useful and non-invasive marker of acute intestinal inflammation.

Summarize recent evidences on FC, providing practical perspectives on its diagnostic role in different gastrointestinal conditions.

S:

Relevant data derived from studies on inflammatory bowel disease (IBD). FC concentration showed a good diagnostic precision for separating organic and functional intestinal diseases. FC levels correlated with IBD activity. FCCs were higher in subjects with NSAID enteropathy. The correlation between FC and endoscopy is under investigation.

CONCLUSIONS:

FC has been widely proposed as a filter to avoid unnecessary endoscopies. Nevertheless, it should not be considered as a marker of organic intestinal disease at all; rather it represents a marker of "eosinophilic intestinal inflammation". In IBD, more and larger studies are needed to confirm its ability to correlate with IBD extent, to predict response to therapy and relapse, and the presence of clinical intestinal inflammation in asymptomatic first degree relatives of patients with IBD.

di infiammazione stinale

Il test per individuare pazienti con possibile
Malattia Infiammatoria Cronica (MIC) e il test per
la sindrome dell'intestino irritabile (SII).



DIAGNOSI E TEST

Test

Test

Test

Test

DIAGNOSI
STINALE (SII)

ROSA

SONO

INTESTINO

Calprest

CHE COS'È CALPREST

Calprest è il test immunoenzimatico di Eurospital che consente di verificare, in modo accurato e non invasivo, la presenza di uno stato infiammatorio a carico del tratto intestinale.

Calprest permette di effettuare una diagnosi differenziale tra patologie di tipo organico (Malattia Infiammatoria Cronica Intestinale - MIC, note anche come Inflammatory Bowel Disease - IBD) e di tipo funzionale (Sindrome dell'Intestino Irritabile - SII, Irritable Bowel Syndrome - IBS). Se Calprest fornisce un risultato negativo, si può, con quasi assoluta certezza, escludere un'infiammazione a carico della mucosa intestinale.



UN TEST SEMPLICE E ACCURATO

Fino ad oggi, per valutare lo stato infiammatorio della mucosa intestinale era necessario ricorrere ad esami invasivi (colonoscopia e conseguente esame istologico). Di recente, però, ha trovato sempre più credito l'uso di marcatori non invasivi. Tra questi, uno dei più attendibili e sicuri è rappresentato dalla determinazione della concentrazione fecale della calprotectina, una proteina antimicrobica presente nei neutrofilo che, in presenza di processi infiammatori a carico dell'intestino, viene rilasciata nel lume intestinale e pertanto può essere rilevata nelle feci.

Il principio diagnostico di Calprest si basa sulla determinazione quantitativa nelle feci della calprotectina: nei pazienti affetti da Malattia Infiammatoria Cronica Intestinale il livello di calprotectina è infatti generalmente molto elevato. Nei soggetti con Sindrome dell'Intestino Irritabile (SII) il livello di calprotectina è invece decisamente inferiore a quello riscontrato nei pazienti con malattia attiva, talvolta superiore al limite di riferimento ma in ogni caso sempre superiore rispetto a quello rilevabile nei soggetti sani.

Calprest permette di utilizzare questo marcatore per selezionare i pazienti con infiammazione da avviare a ulteriori esami e risulta in tal senso maggiormente accurato rispetto ai normali test biochimici (VES, PCR).

SENSIBILITÀ E SPECIFICITÀ

La determinazione della calprotectina fecale viene impiegata per la diagnosi differenziale tra IBD ed IBS grazie al suo elevato valore predittivo negativo che permette di escludere un'eventuale patologia organica.

SENSIBILITÀ DIAGNOSTICA	SPECIFICITÀ DIAGNOSTICA	VALORE PREDITTIVO NEGATIVO
99%	97%	99%

INTERPRETAZIONE DEI RISULTATI

I campioni con una concentrazione di calprotectina superiore a 50 mg calprotectina/kg devono essere considerati positivi al test. Nei soggetti adulti sani il valore medio della calprotectina è di 25 mg calprotectina/kg.

Un risultato positivo di Calprest è indice di infiammazione intestinale e permette di selezionare con sicurezza i pazienti da avviare a ulteriori indagini diagnostiche.

DIPARTIMENTO ASSISTENZIALE INTEGRATO
MEDICINA DIAGNOSTICA

U.O. IMMUNOLOGIA- IMMUNOPATOLOGIA DELCS
Responsabile F.F. Prof. Fabrizio Mainiero
Tel. 06-49970966

Roma,

Sig..... PAZIENTE
(Cognome e Nome)

Prelievo del

Provenienza ...DAI Pediatria.....

DOSAGGIO CAPROTECTINA FECALE

< 50 mg/kg di feci Ne

50 - 100 mg/kg di feci
Zona Grigia, si consi

> 100 mg/kg di feci P

Il kit Calprest (Eurospital, Trieste, Italia) è utilizzato x l'analisi della calprotectina fecale. Calprest è un test immunoenzimatico che sfrutta l'uso di anticorpi polivalenti (riconoscimento del massimo numero di epitopi) diretti contro la calprotectina e per un dosaggio quantitativo di essa.

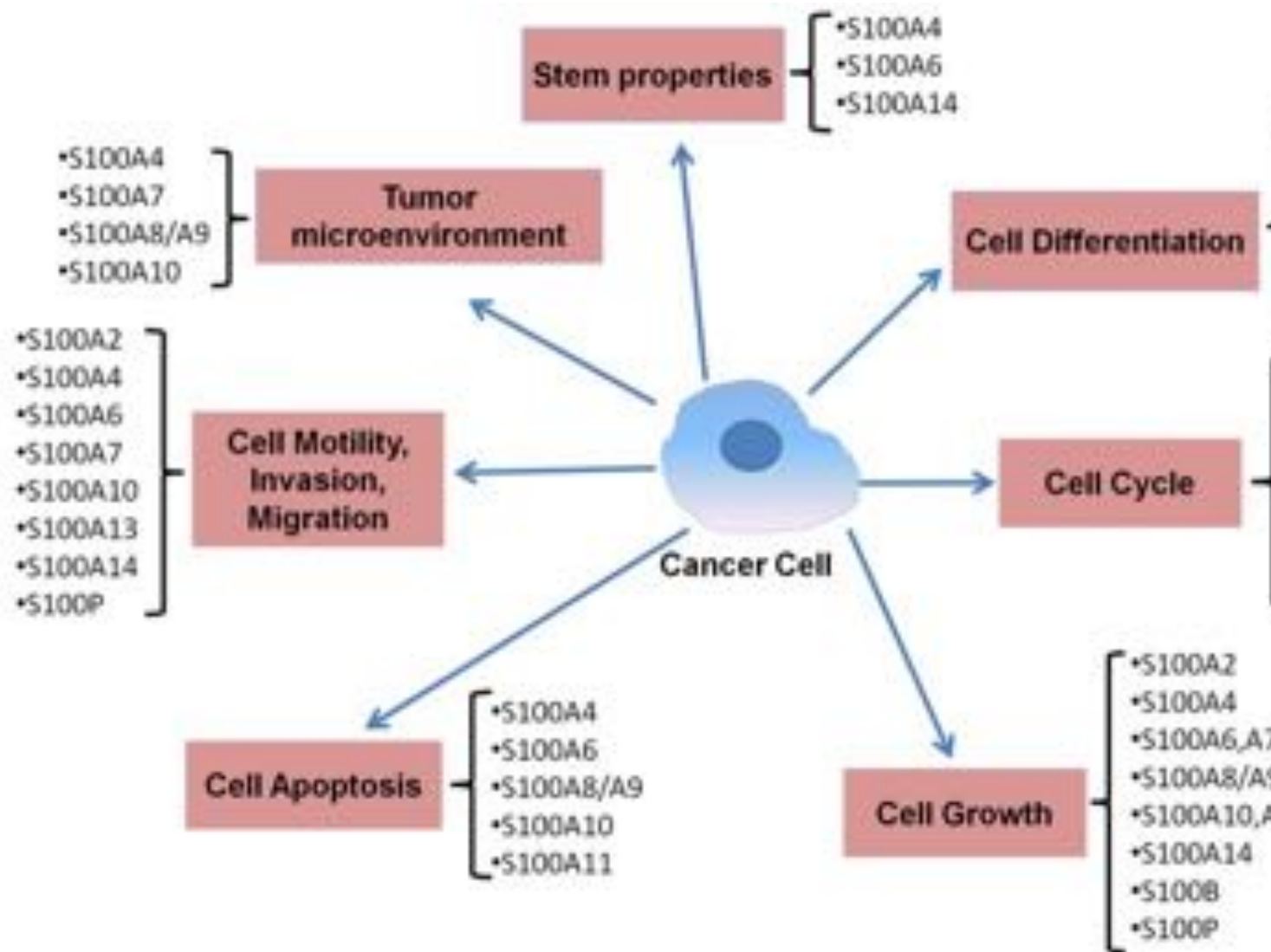
Il Responsabile

Fabrizio Mainiero

and therapeutic target for cancer.

S100P expression is described in many different cancers, and its expression is associated with drug resistance, metastasis, and poor clinical outcome. S100P is member of the S100 family of small calcium-binding proteins that have been reported to have either intracellular or extracellular functions, or both. Extracellular S100P can bind with the receptor for advanced glycation end products (RAGE) and activate cellular signaling. Through RAGE, S100P has been shown to mediate tumor growth, drug resistance, and metastasis. S100P is specifically expressed in cancer cells in the adult. Therefore, **S100P is a useful marker for differentiating cancer cells from normal cells, and can aid in the diagnosis of cancer by cytological examination.** The expression of S100P in cancer cells has been related to hypomethylation of the gene. Multiple studies have confirmed the beneficial effects of blocking S100P/RAGE in cancer cells, and different blockers are being developed including small molecules and antagonist peptides.

cated in multiple stages of
 rigenesis and progression.
 ng the S100 genes, 22 are
 ered at chromosome locus
 , a region frequently
 anged in cancers. S100
 in possesses a wide range
 racellular and extracellular
 ions such as regulation of
 um homeostasis, cell
 feration, apoptosis, cell
 sion and motility,
 keleton interactions,
 ein phosphorylation,
 ation of transcriptional
 ors, autoimmunity,
 otaxis, inflammation and
 otency. Many lines of
 nce suggest that altered
 ession of S100 proteins
 associated with tumor
 ression and prognosis.
 efore, S100 proteins might



Electrophoresis is the diagnostic tool for the detection of APP

ey can also be detected by ELISA, nephelometry, immunoturbidimetry radioimmunoassay and molecular biology.

Technical problems with the testing of APP include changes in concentrations of APP observed with the use of different anticoagulants and in the presence of hemolysis and lipemia.

Although APP are considered to be stable at -20 °C, the long term storage at -70 °C is recommended.

Laboratory evaluation of APP!

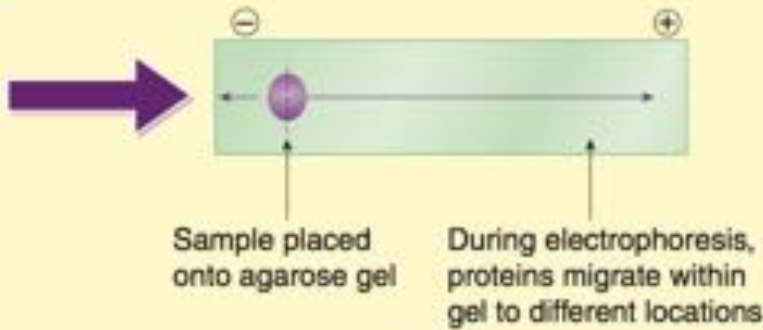
Protein electrophoresis (PEP)

Expense: Moderate

Semi-automated

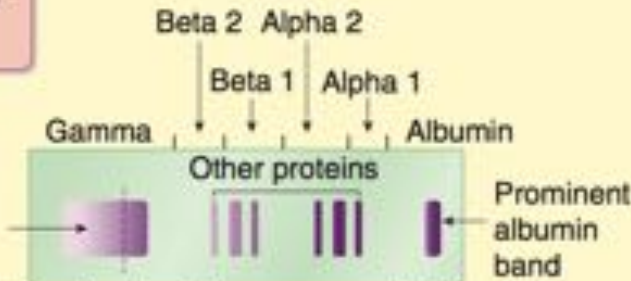
Sample can be:
 Serum for SPEP analysis
 Cerebrospinal fluid (CSF) for UPEP analysis

Serum and CSF are usually concentrated prior to electrophoresis to increase the concentration of proteins in the sample



Sample with an additional monoclonal antibody, which can appear in multiple forms, for example, shows a dense band of protein not present in a sample from a healthy individual

Area of gel:



Broad band of IgG immunoglobulins

Prominent albumin band

Bands of proteins are generated by electrophoresis and made visible by staining the gel



Normal serum

Serum

Nephelometry for quantitation of selected proteins and other compounds

Expense: Moderate

Semi-automated

Sample of any body fluid is incubated with an antibody to the compound being measured

When the compound present, antigen-antibody complexes form

Antibody to the compound is the reagent added to the sample



Antigen is compound being measured

The amount of scattered light is proportional to the

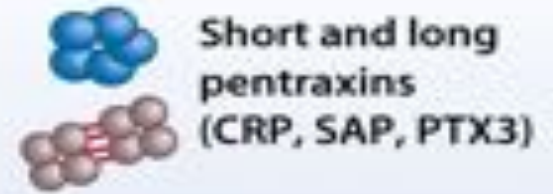
Antigen-antibody complex scatters light from a beam

THE MAIN FUNCTION OF THE APP POSITIVE IS TO ACTIVATE THE COMPLEMENT AND THE PHAGOCYTOSIS,

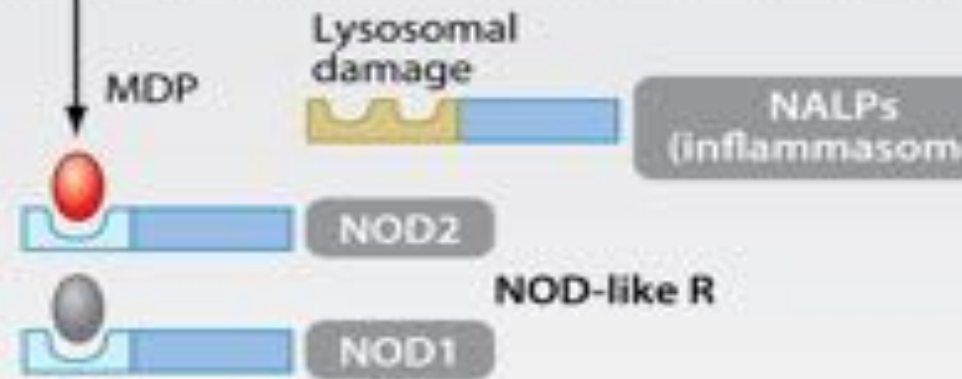
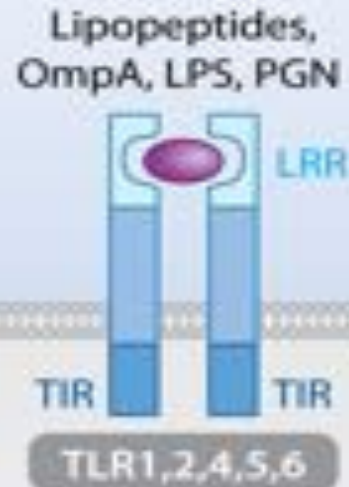
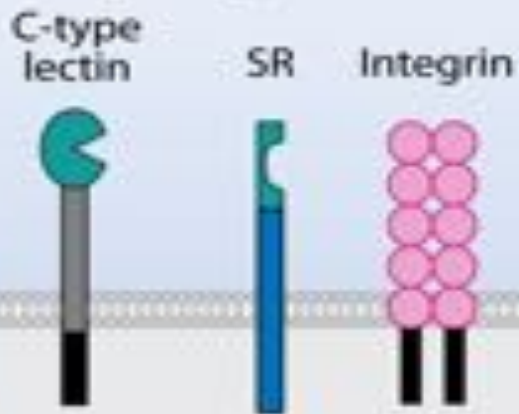
THEY CAN BE NAMED:

THE SOLUBLE DAMAGE RECEPTORS OF NATURAL IMMUNITY AND INFLAMMATION!

LOGICAL
IDS



Capturing R



Endosome/
phagosome



CYTOSOL

SIGNALLING!



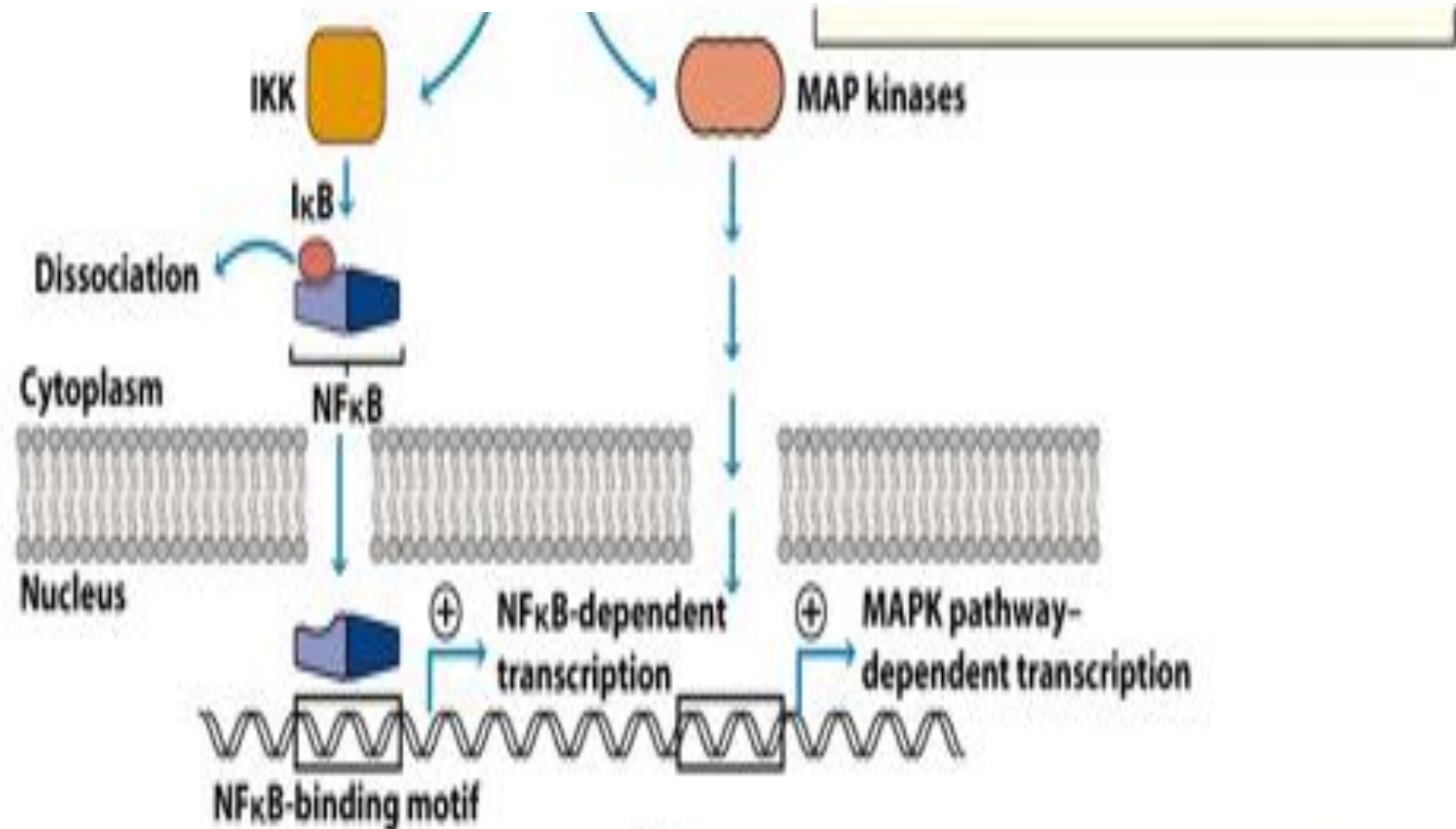
**THE APP POSITIVE AS THE SOLUBLE
DAMAGE RECEPTORS**

**CAN COOPERATE WITH THE
CYTOPLASMIC AND MEMBRANE
RECEPTORS**

TO ACTIVATE

**THE TRANSCRIPTIONAL
PROGRAM OF NATURAL IMMUNITY
AND INFLAMMATION!**

INVOLVE NFκB AND MAPK ACTIVATION!



6a

The freed NFκB translocates from the cytoplasm into the nucleus, where it

6b

The MAPK cascade results in translocation of a transcriptional activator from the



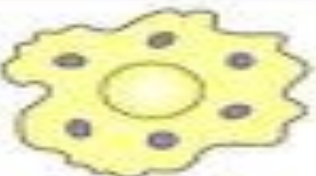
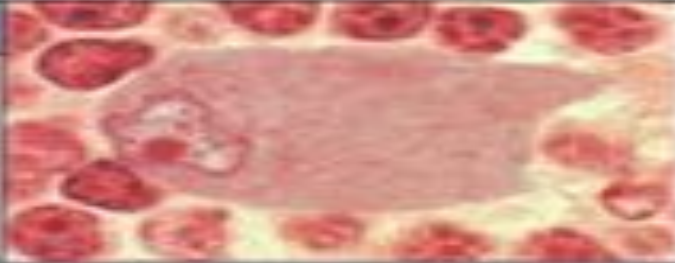



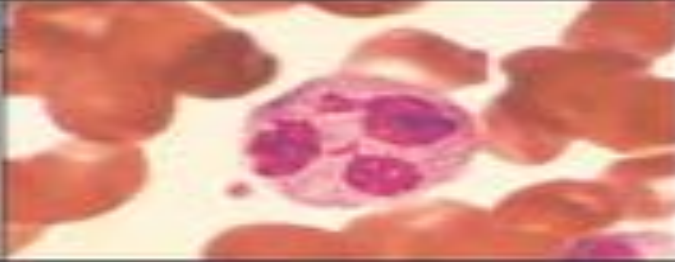

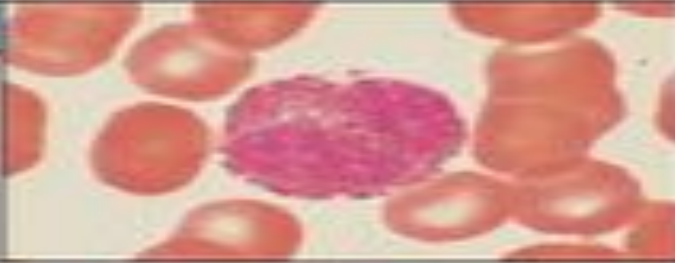

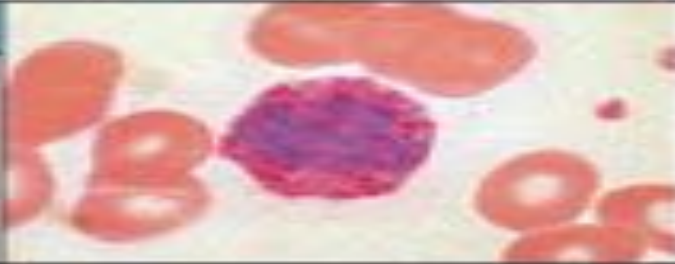


ШАРККА

**THE TRANSCRIPTIONAL
PROGRAM OF NATURAL IMMUNITY AND
INFLAMMATION
CONTROLS**

**EXPRESSION AND PRODUCTION OF
MOLECULES INVOLVED IN MULTIPLE
FUNCTIONS SUCH AS:**

Monocytes/

DEGRANULATION

Cell		Activated function
Macrophage		 Phagocytosis and activation of bactericidal mechanisms Antigen presentation
Dendritic cell		 Antigen uptake in peripheral sites Antigen presentation in lymph nodes
Neutrophil		 Phagocytosis and activation of bactericidal mechanisms
Eosinophil		 Killing of antibody-coated parasites antiviral
Basophil		 Release of histamine
Mast cell		 Release of granules

DEGRANULATION

PHAGOCYTOSIS:

SENSING

To defend the body against

bacteria, human neutrophils

SWALLOWING

(white blood cells) ingest

invading pathogens like

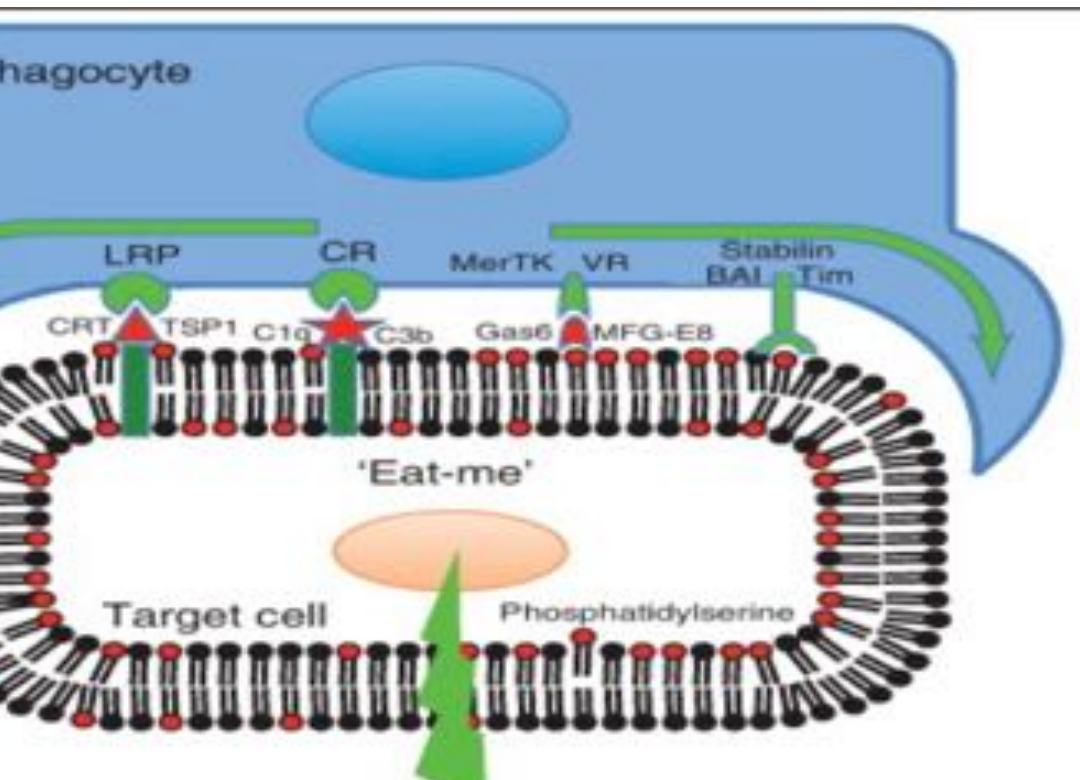
this *E. coli*

DIGESTING

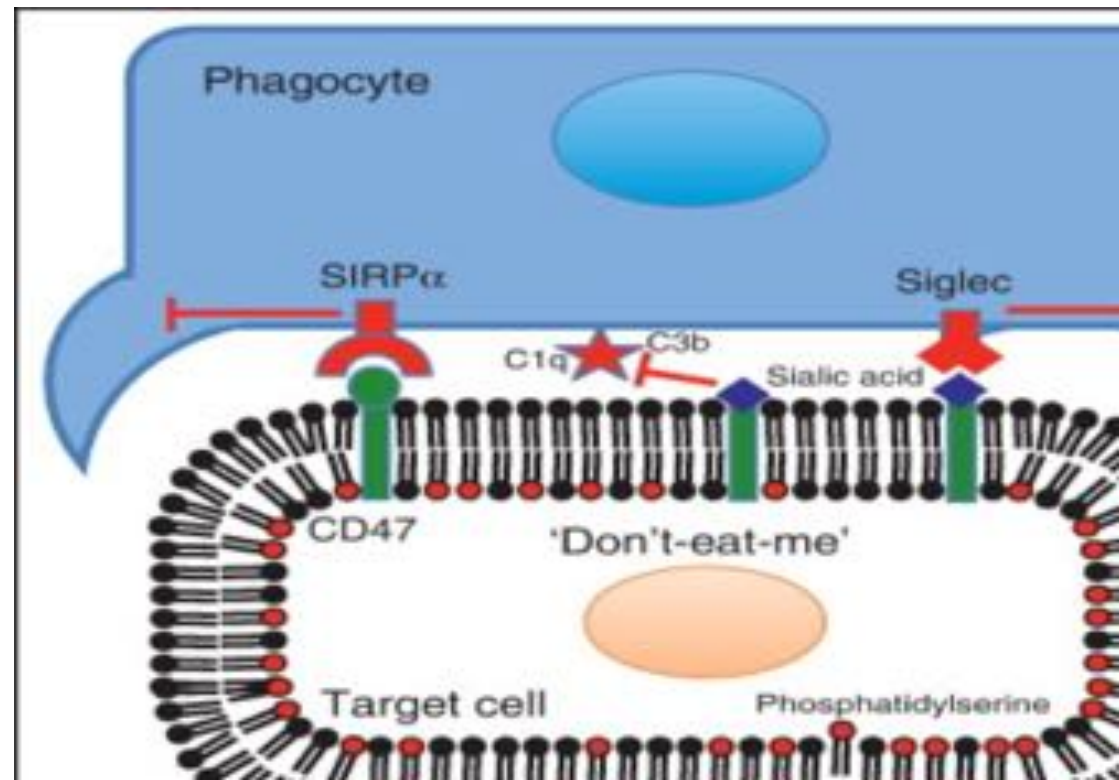
RECENTLY IT HAS BEEN DEMONSTRATED THAT A TYPE OF PHAGOCYTOSIS, CALLED **PHAGOPTOSIS**, ELIMINATES ALSO CELLS ALIVE!

The term is created by combining phago-, which is derived from the ancient Greek 'phagein' to devour, and -ptosis, which is from the ancient Greek 'ptosis' meaning to fall; using the connotation of dying; therefore, phagoptosis would connote 'devouring-induced death caused by being devoured'.

'Eat-me' signalling!



Don't eat-me' signalling

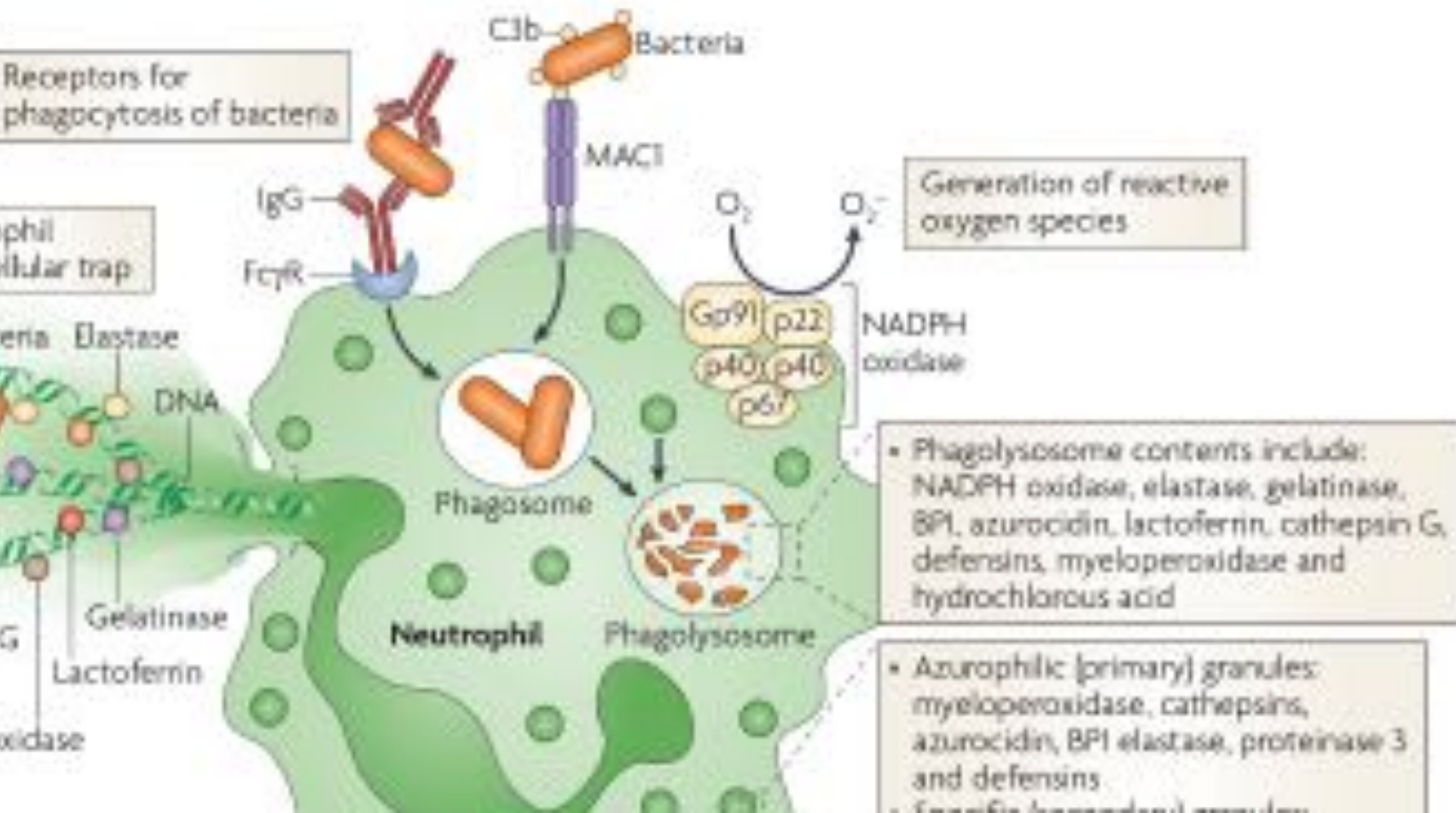


atrophins and other cells, and thus is quantitatively one of the main forms of cell death in the body!

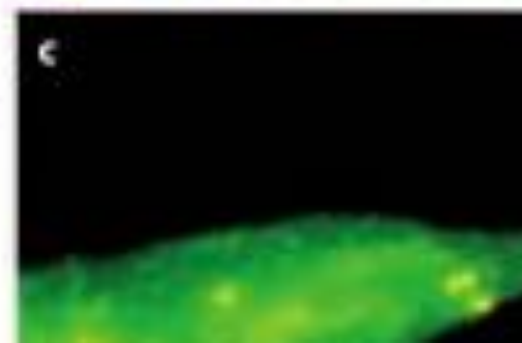
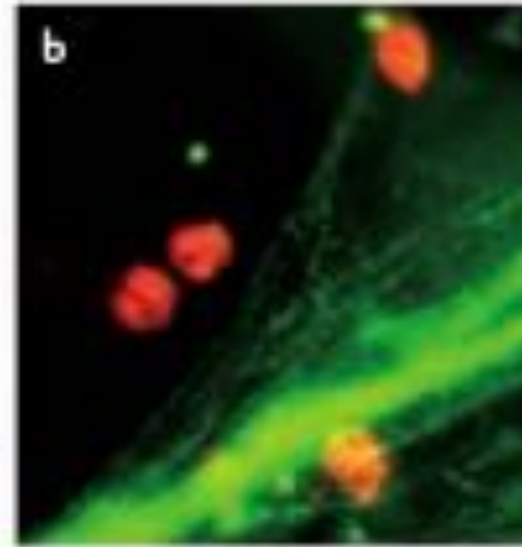
Table I. Rough estimates of the physiological rates of cell turnover by different forms of cell death in humans

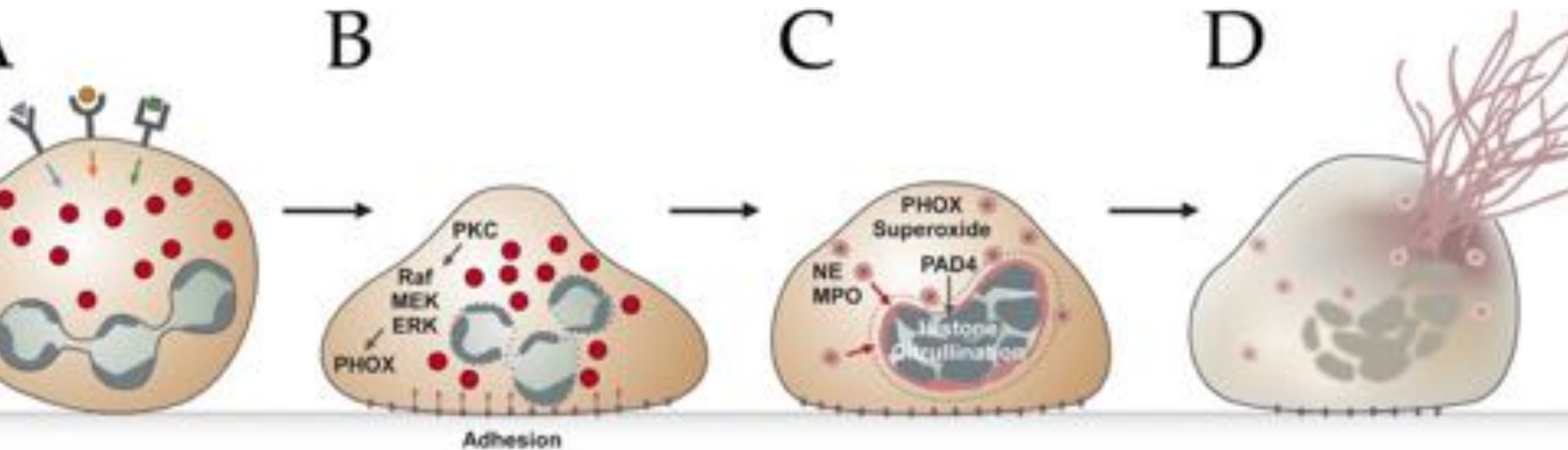
Type of cell death	Cells	Rate (thousands of cells/second)
Phagoptosis	Erythrocytes	2000
	Neutrophils	500–1000
Shedding	Enterocytes	80
Cornification	Keratinocytes	40
Necrosis	Enterocytes	10
Apoptosis	T cells and B cells	1
Autophagy		None known

Neutrophil Extracellular Traps: an additional antibacterial weapon!



DNA





stimulation of receptors (A), neutrophils adhere to the substrate (B) and mobilize granule components, namely NE and MPO (C). Granules are depicted as red circles. Histones in the nucleus get processed, and the intracellular membranes disintegrate. Finally, the cell membrane ruptures, and the mixture of cytoplasm and nucleoplasm is extruded to form NETs (D).

It has been reported that peptidylarginine deiminase 4 (PAD4), an enzyme that converts L-arginine to monomethyl-Arg to citrulline in histones, is essential for NET formation. The dense chromatin decondensation along the NETs were rich in histone citrullination.

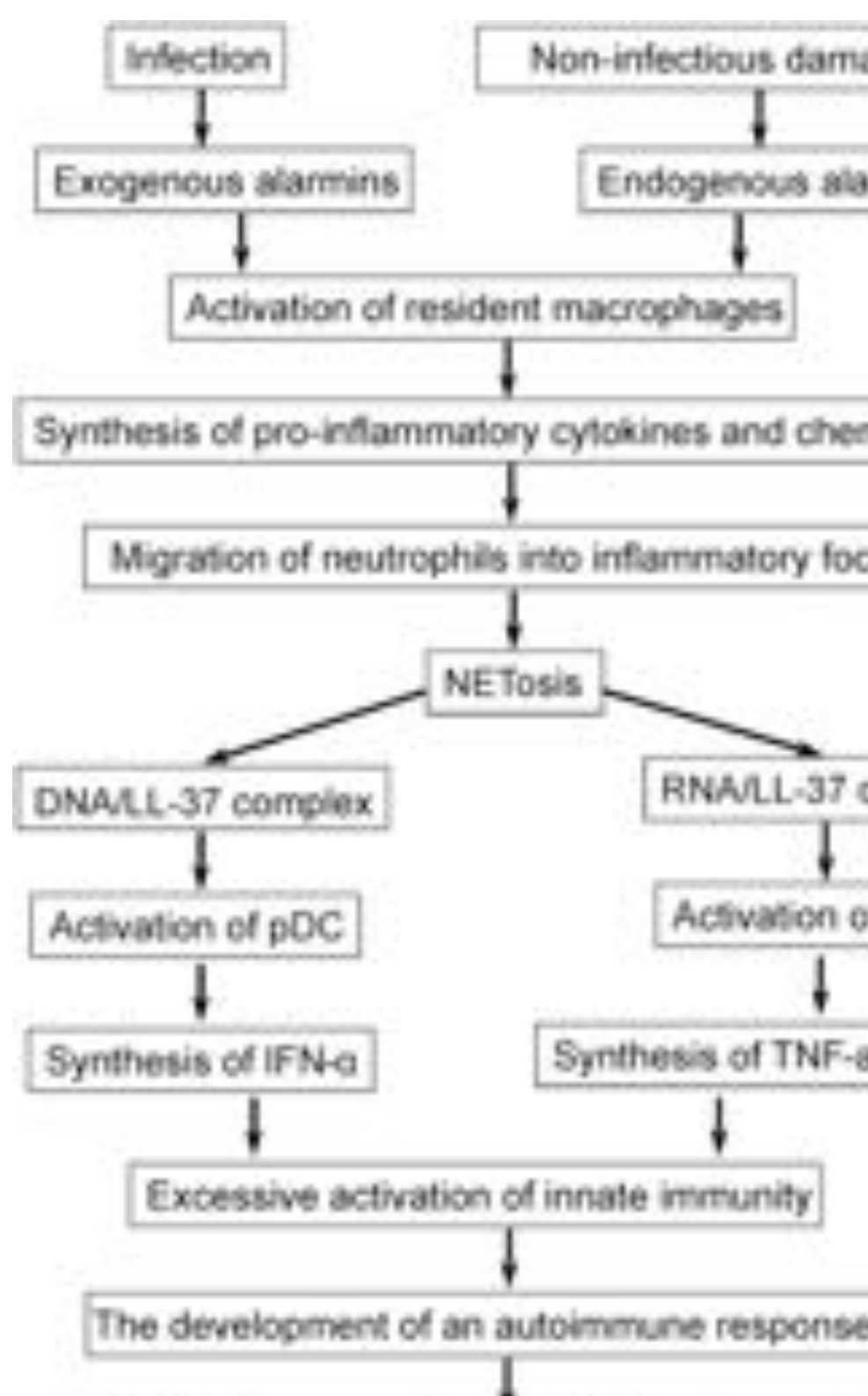
[Front Immunol.](#) 2012;3:307.

PAD4 mediated histone hypercitrullination induces heterochromatin decondensation and chromatin unfolding to form neutrophil extracellular trap-like structures

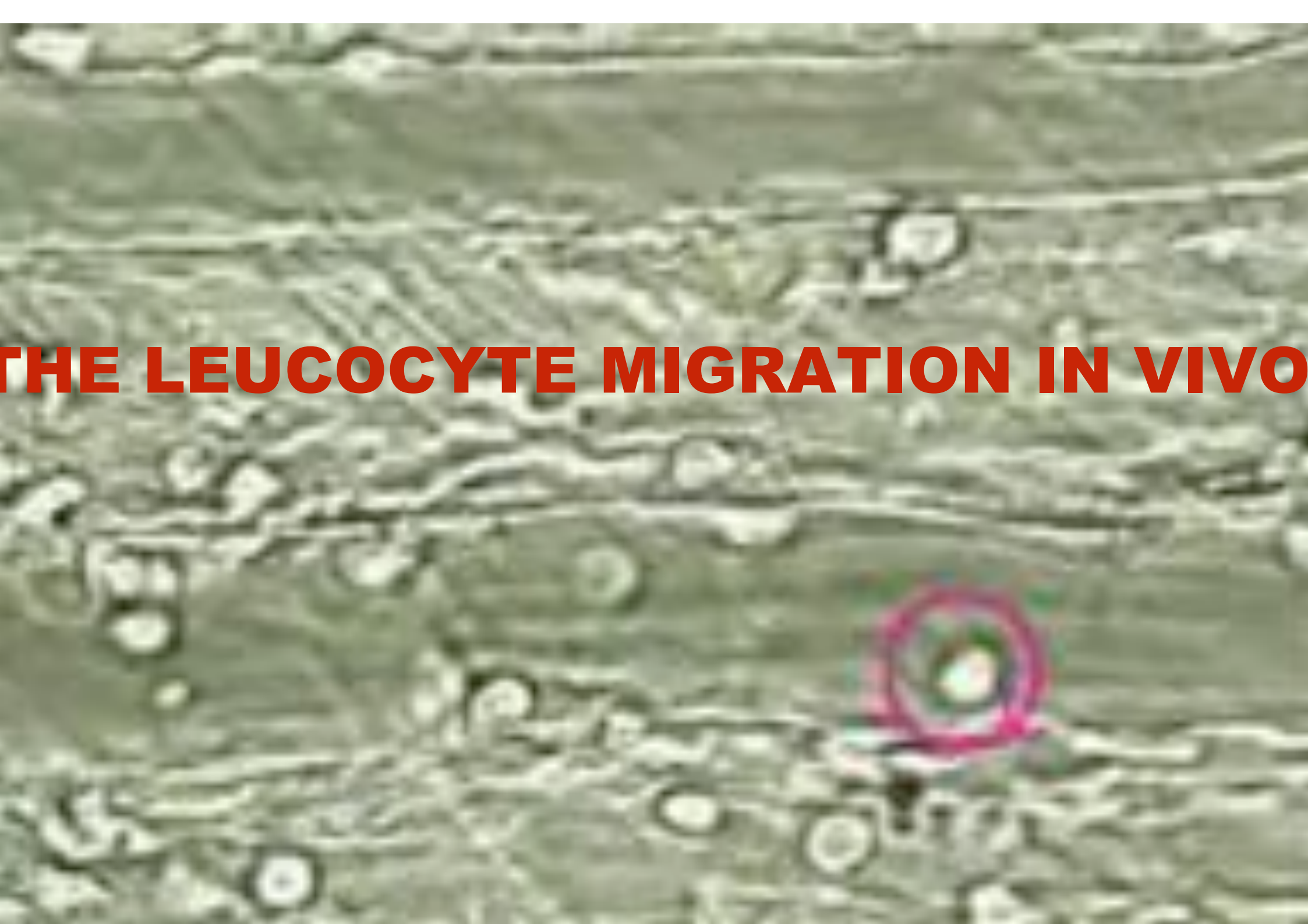
Pathogenesis of many autoimmune diseases is initially based on a redundant or exaggerated activation of the innate immune system. It was suggested that an excessive activation of the innate immunity is often the hallmark of a chronic inflammatory process in the system. This inflammation can be induced by exogenous and endogenous alarm factors, or pathogens. **We believe that the recently discovered neutrophil extracellular traps, or NETs, completely meet the criteria of alarmins.**

This review summarizes current knowledge concerning the general characteristics of NETs, their antimicrobial properties, and their role in the development of chronic inflammatory diseases that underlie the pathogenesis of diabetes and atherosclerosis.

Insights on the NETosis can provide the foundation for developing new diagnostic

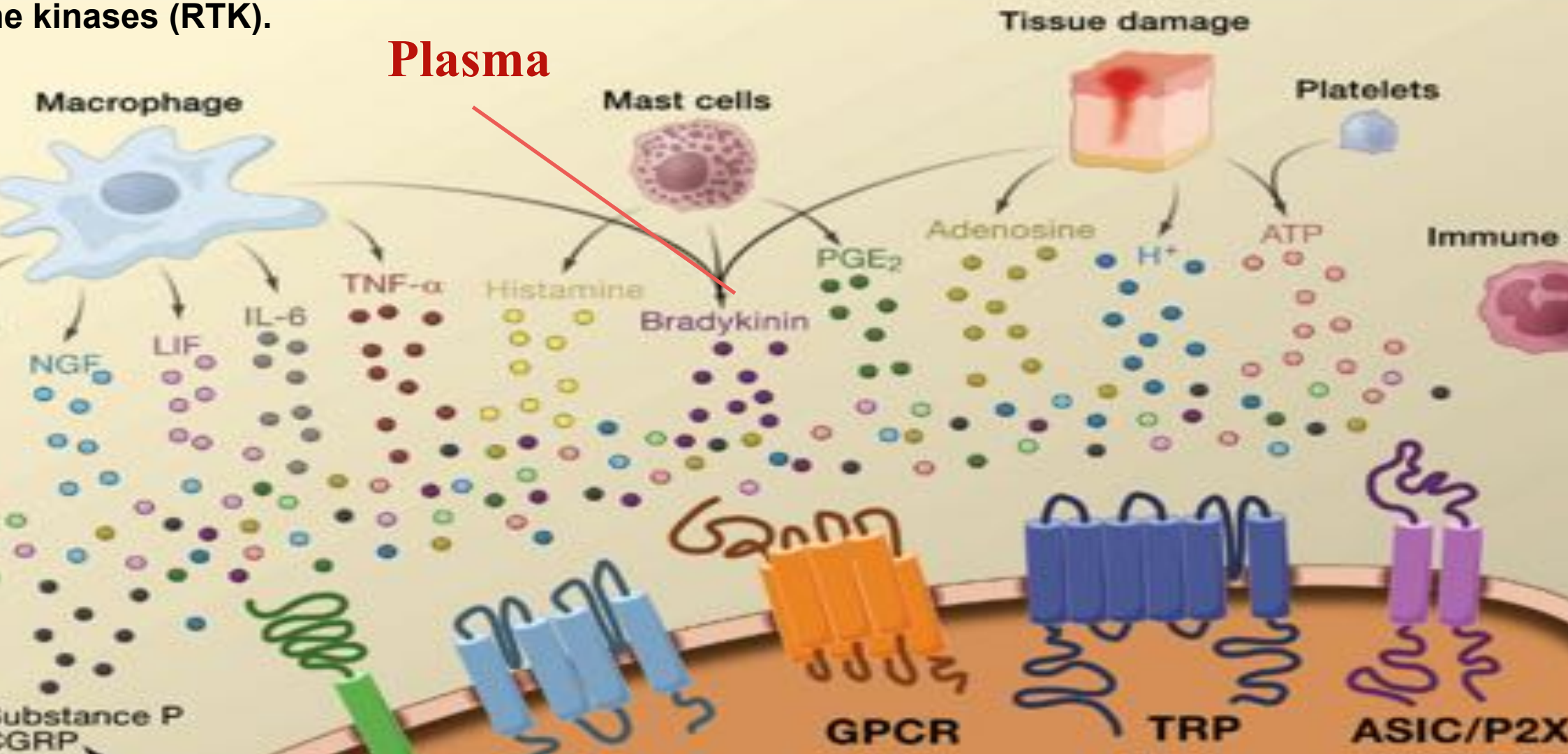


THE LEUCOCYTE MIGRATION IN VIVO



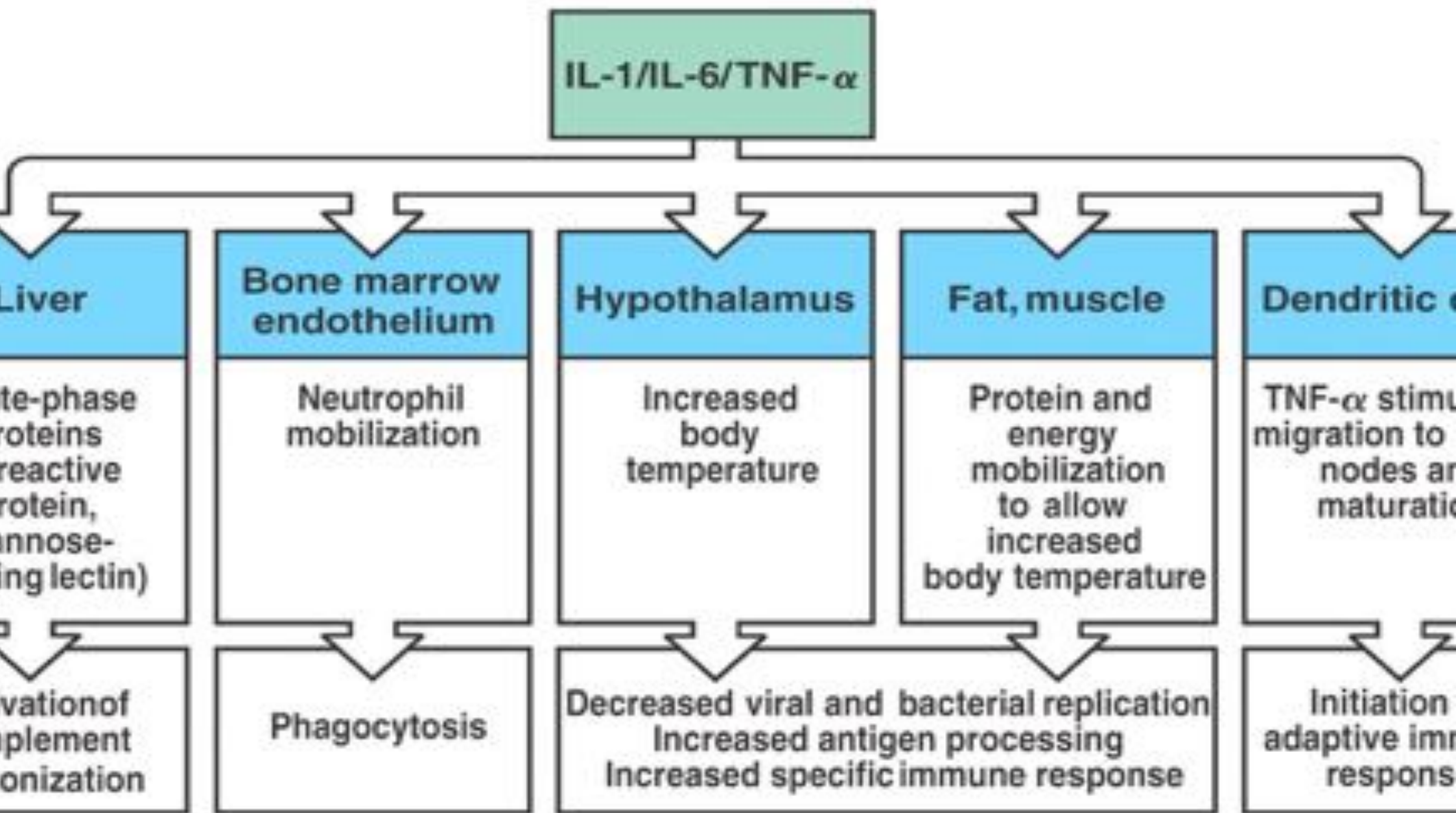
MEDIATORS....

damage leads to the release of inflammatory mediators by activated nociceptors or nonneuronal cells such as mast cells, basophils, platelets, macrophages, neutrophils, endothelial cells, keratinocytes, and fibroblasts. This "inflammatory soup" of signaling molecules includes histamine, ATP, adenosine, substance P, calcitonin-gene related peptide (CGRP), bradykinin, extracellular prostaglandins, thromboxane, and leukotrienes, nerve growth factor (NGF), tumor necrosis factor α (TNF- α), interleukin-1 β (IL-1 β) etc. These mediators act directly by binding to one or more cell surface receptors, including G protein-coupled receptors (GPCR), TRP channels, acid-sensitive ion channels (ASIC), two-pore potassium channels (K2P), and receptor tyrosine kinases (RTK).



Cytokine	Main producer	Acts upon	Effect
IL-1	Macrophages Keratinocytes	Lymphocytes	Enhances responses
		Liver	Induces acute-phase protein secretion
IL-6	Macrophages Dendritic cells	Lymphocytes	Enhances responses
		Liver	Induces acute-phase protein secretion
CXCL8 (IL-8)	Macrophages Dendritic cells	Phagocytes	Chemoattractant for neutrophils
IL-12	Macrophages Dendritic cells	Naive T cells	Diverts immune response to type 1, proinflammatory, cytokine secretion
TNF- α	Macrophages Dendritic cells	Vascular endothelium	Induces changes in vascular endothelium (expression of cell-adhesion molecules (E- and P-selectin), changes in cell-cell junctions with increased fluid loss

...WHICH CAN HAVE VARIOUS EFFECTS...



TNF

Low quantities
(plasma conc. $<10^{-9}$ M)

Moderate quantities

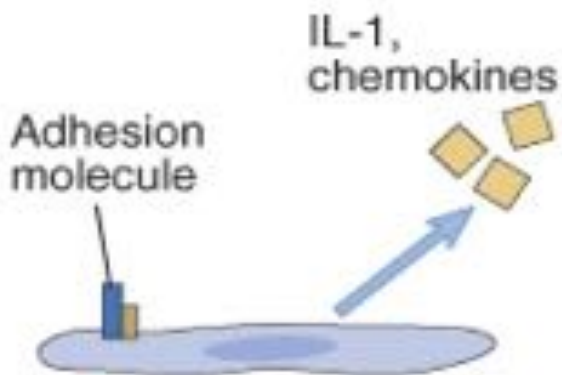
High quantities
(plasma conc. $\geq 10^{-7}$ M)

Local inflammation

Systemic effects

Septic shock

Endothelial cell



Leukocyte

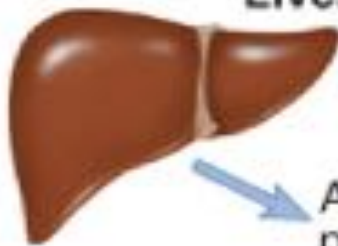


Brain



Fever

Liver



Acute-phase proteins

Bone marrow

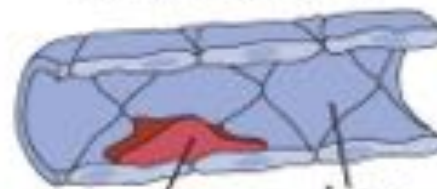


Heart



Low output

Blood vessel



Thrombus

Low resistance

Liver

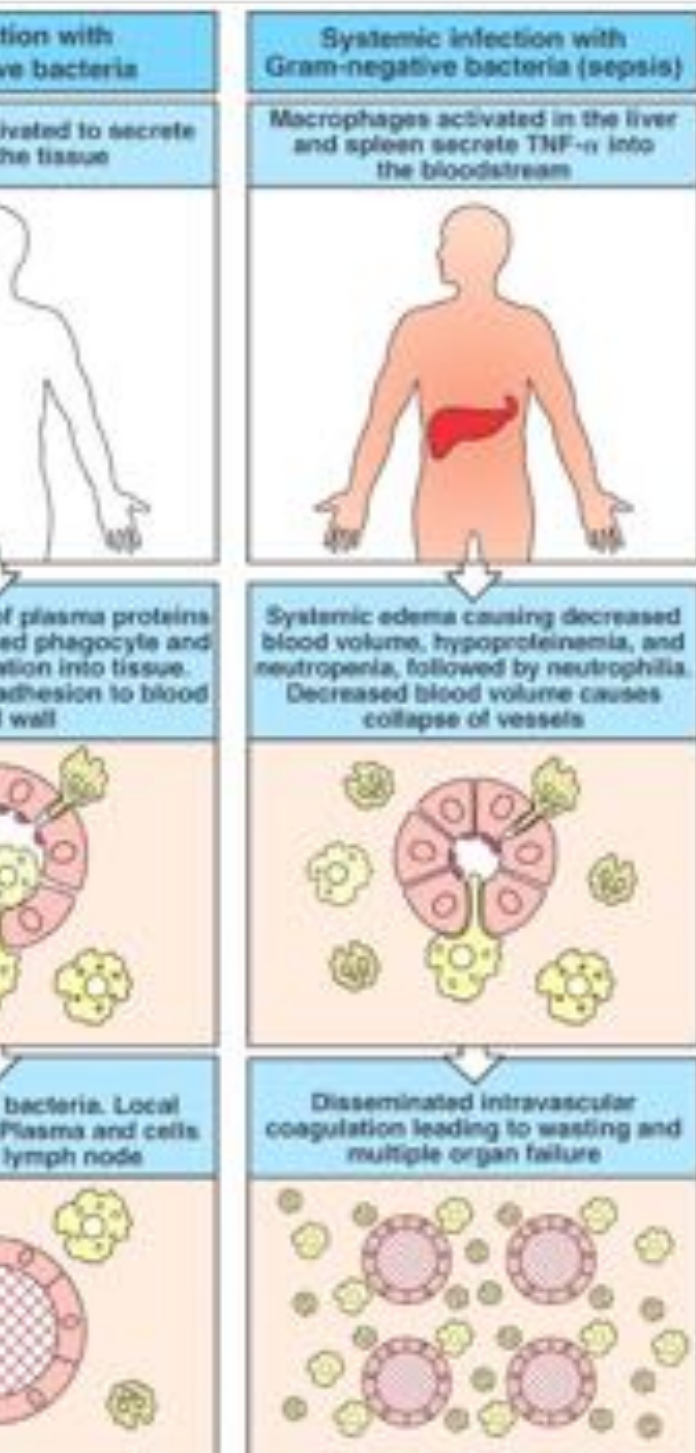


Pathological consequences of
Systemic Inflammatory response
(SIRS)
are:

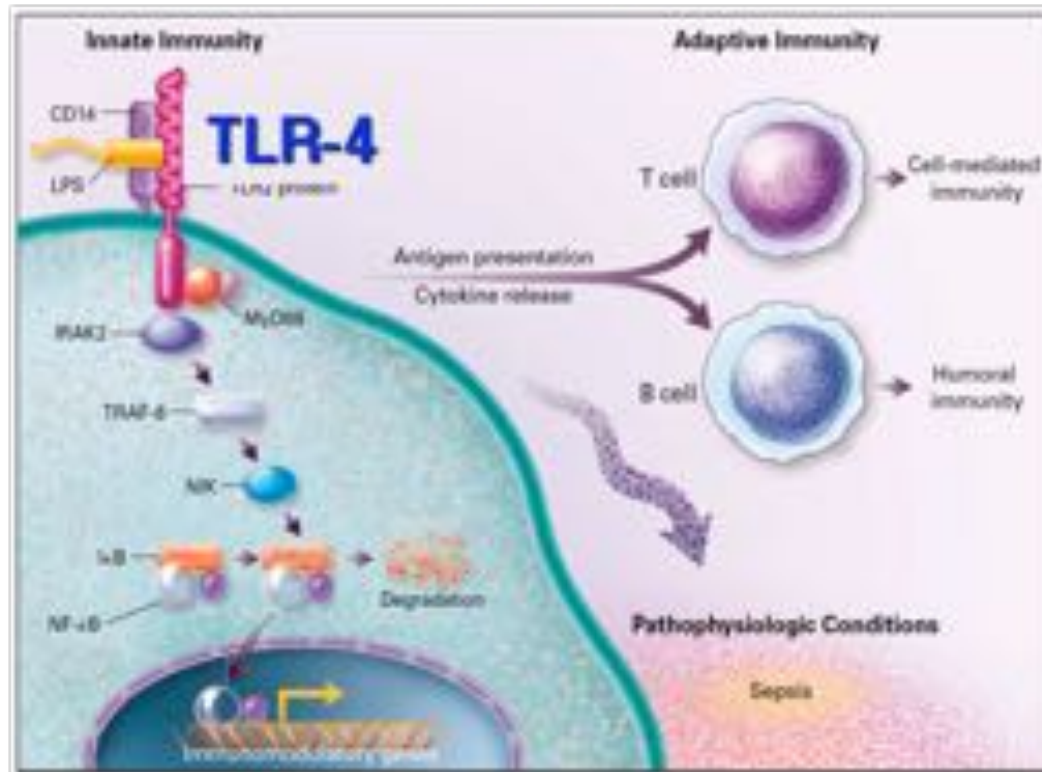
Septic shock;

ARDS;

Multiple organ dysfunction (MOD)



Pathological consequences of inflammatory response to systemic LPS: the septic shock



Septic shock is classically triggered by Gram- bacteria (TLR-4/LPS); Gram+ bacteria too can induce a systemic

• endotox

• esotos

vasodilation

low cardiac output

**formation of
thrombi**

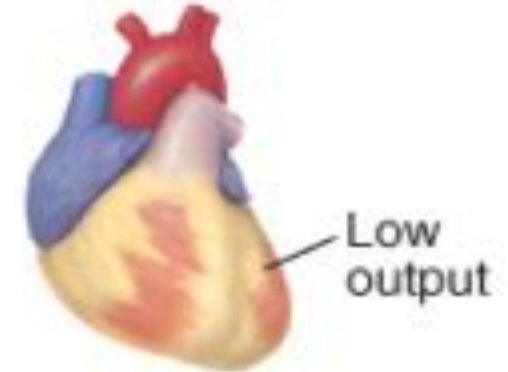
**intravascular
coagulation**

TNF

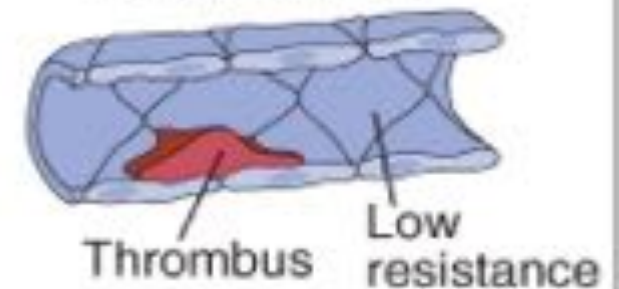
High quantities
(plasma conc. $\geq 10^{-7}$ M)

Septic shock

Heart



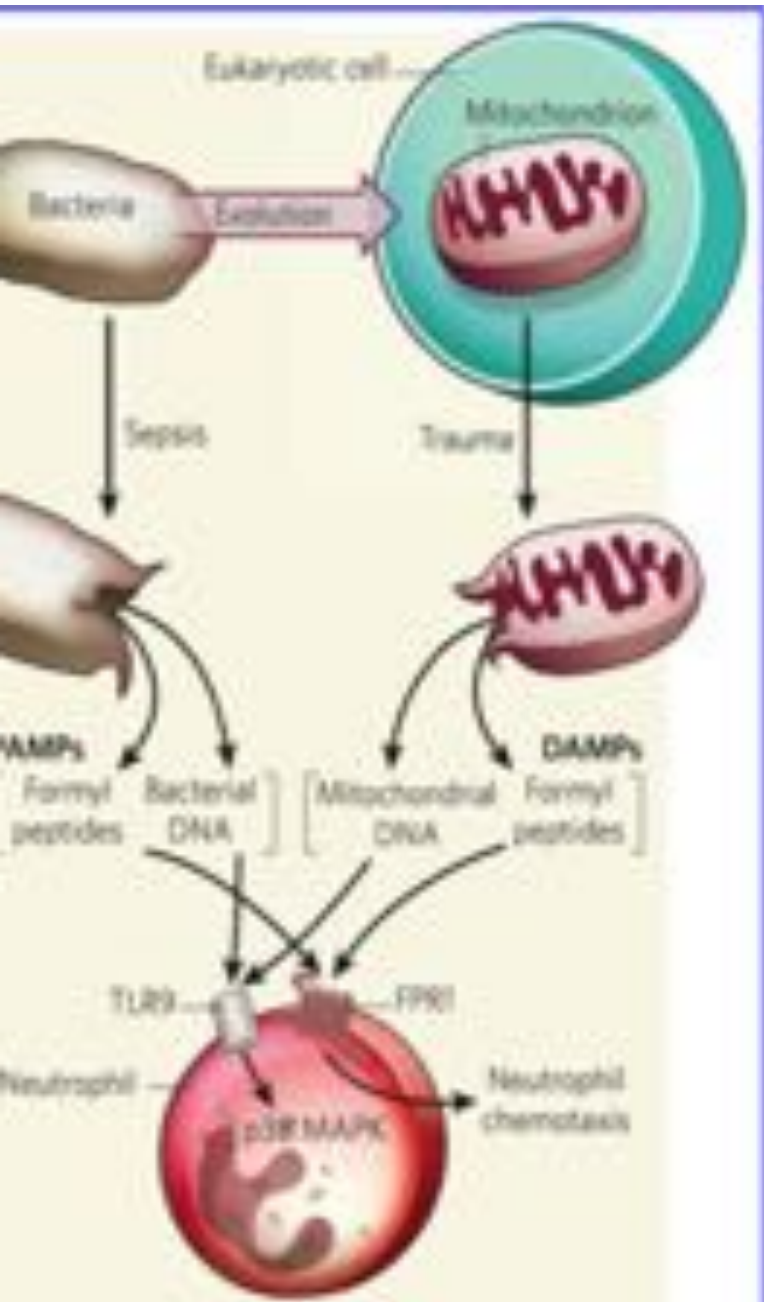
Blood vessel



Liver

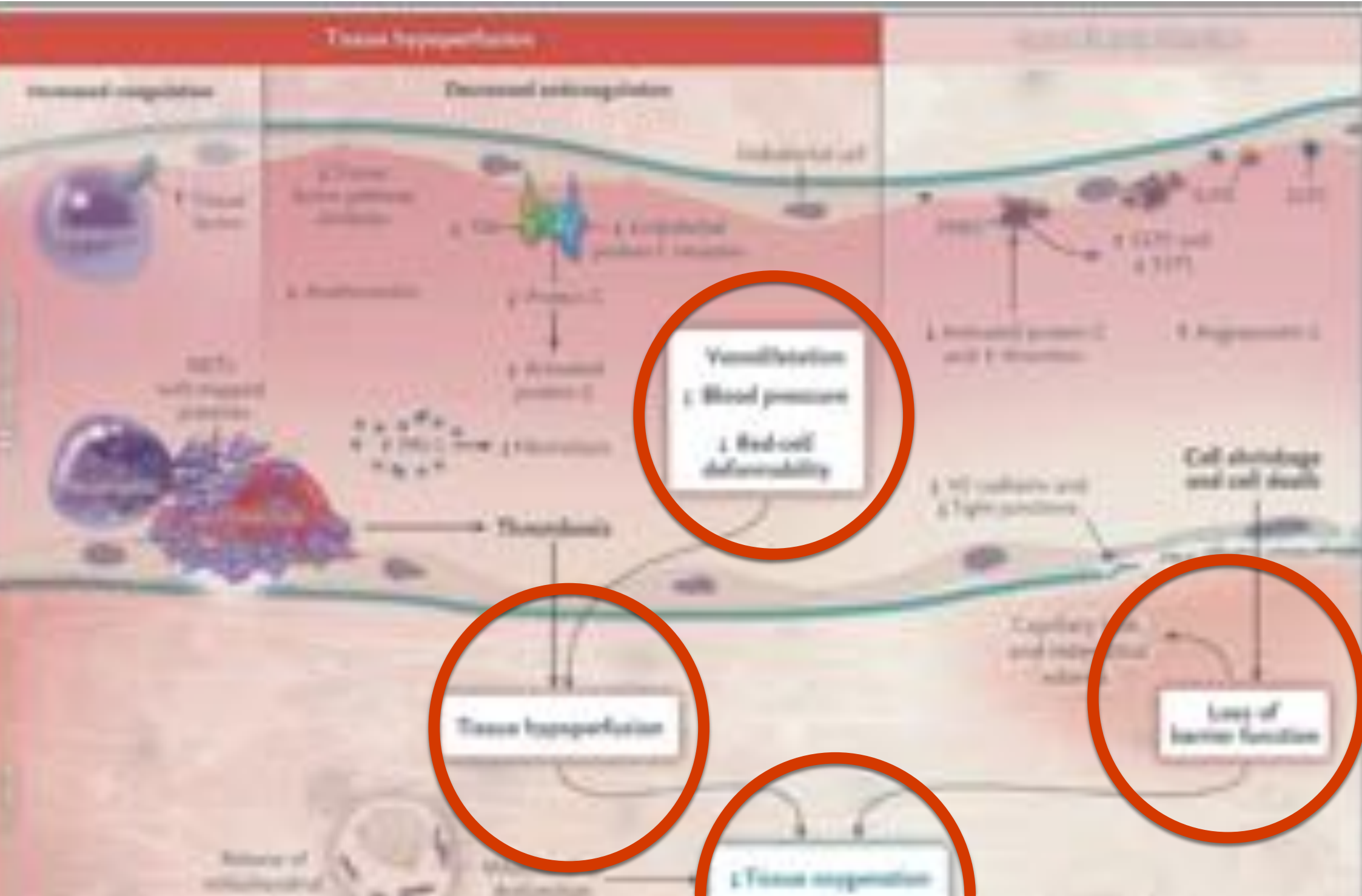


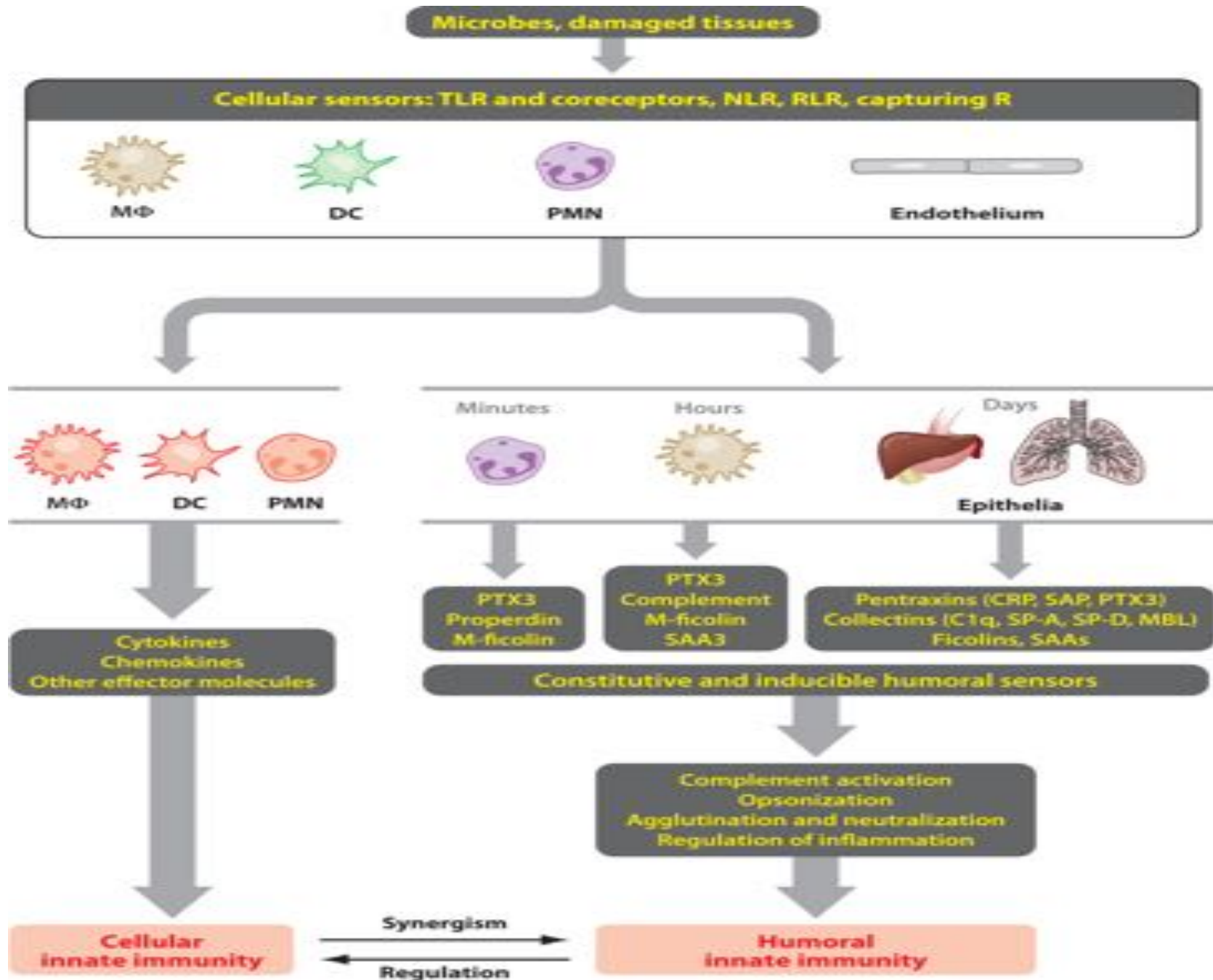
RESPIRATORY DISTRESS SYNDROME (ARDS)



	ARDS		
	Mild	Mod/severe	Severe
Timing	Acute onset within 1 week of a known clinical insult, new/worsening respiratory symptoms		
Hypoxemia	PaO_2/FiO_2 201-300 with PEEP/CPAP \geq 5	PaO_2/FiO_2 \leq 200 with PEEP \geq 5	PaO_2/FiO_2 \leq 100 with PEEP \geq 5
Origin of Edema	Respiratory failure associated to known risk factors, not fully explained by cardiac failure or fluid overload. No objective assessment of cardiac failure or fluid overload. No risk factor are present		
Radiological Abnormalities	Bilateral opacities*	Bilateral opacities*	Opacities in at least 3 quadrants
Additional Physiological Derangement	N/A	N/A	$V_{E,Curr} > 10$ L or $C_{60} < 40$ m

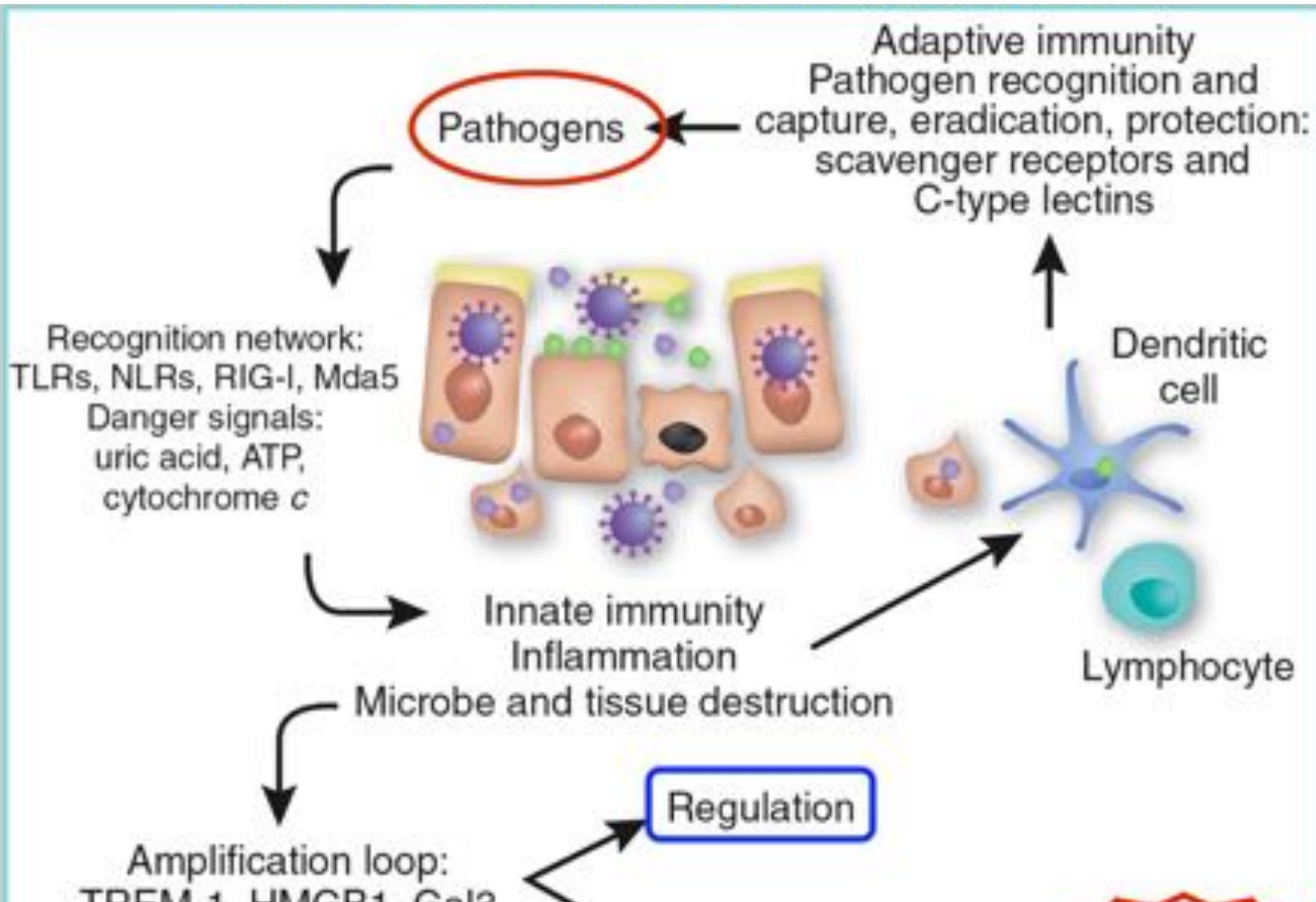
Multiple organ failure (MOF)!





Humoral and Cellular sensors share fundamental mechanisms of effector function: complement activation and regulation

UNDER NORMAL CONDITIONS AND SEPSIS!



The humoral and cellular arms of innate immunity form an integrated system with synergism in deciphering pathological patterns and regulating the innate and inflammatory response!