Basic Immunology

Lecture 16

Complement system

Why is complement system important?

- Major <u>effector</u> system of the humoral IR
- Component of the <u>innate</u> (non-specific) immune IR
- Results immediate response
- Connection to the specific IR

Discovery:

1890: Jules Bordet's experiment:

- Immune serum against *Vibrio cholerae* caused lysis of the bacteria
- Heating the antiserum destroyed this activity
- Addition of a fresh serum to the antiserum restored its killing ability

Paul Ehrlich:

- 2 components of the ANTISERUM:
- → heat stable: specific antibody
- → heat sensitive: responsible for the lytic activity →

COMPLEMENT

Components:

- <u>Inactive factors</u> in the serum and body fluids which can activate each other in an <u>enzyme cascade</u>
- Cell surface receptors (CR) for binding the activated complement components
- Regulatory proteins: soluble and cell surface bound to prevent uncontrolled complement activation

Molecular mediators of inflammation

Plasma enzyme mediators:

- kinin kallikrein system
- Fibrinolytic system
- Complement cascade
- Clotting cascade

<u>Lipid mediators</u>:

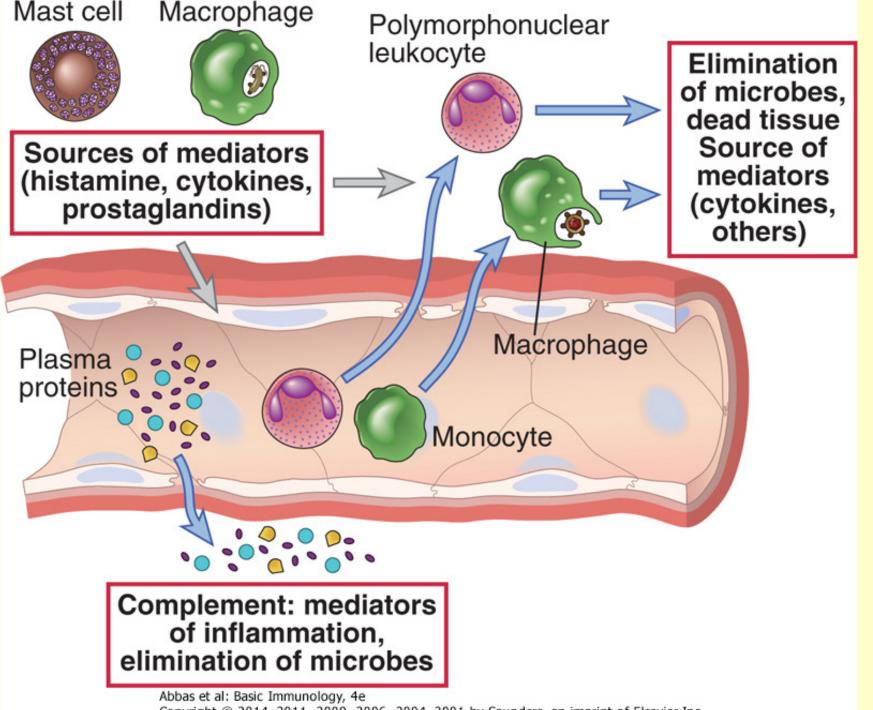
leukotrienes, prostaglandins (PGE)

Chemoattractants:

- -Chemokines: IL-8
- -Complement components
- PAF (platelet activating factor)

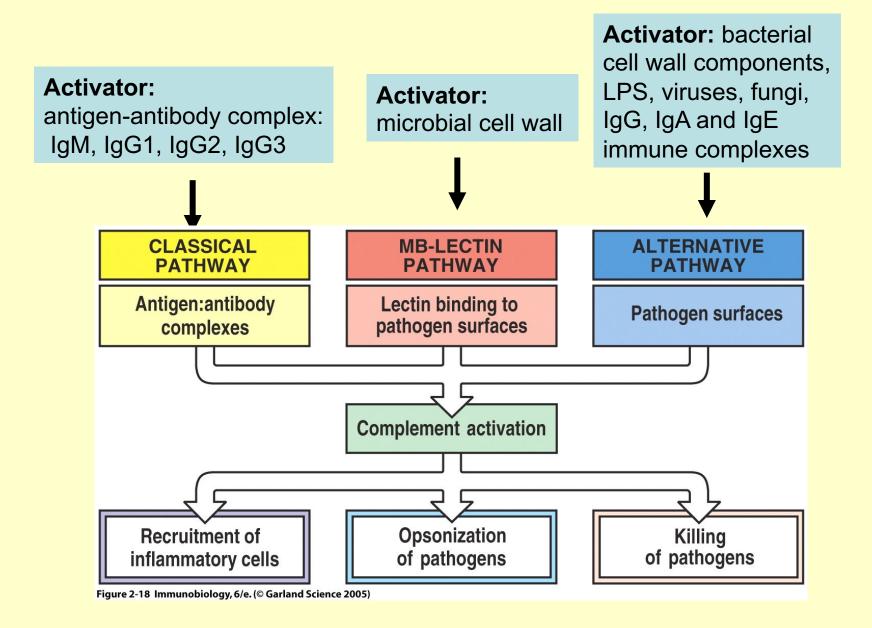
Inflammatory cytokines:

IL-1, IL-6, TNFalpha



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Activation of the complement enzyme cascade



Early steps of classical pathway activation

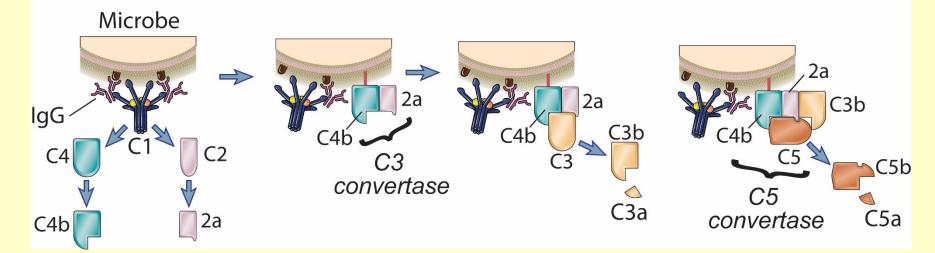
Classical Pathway

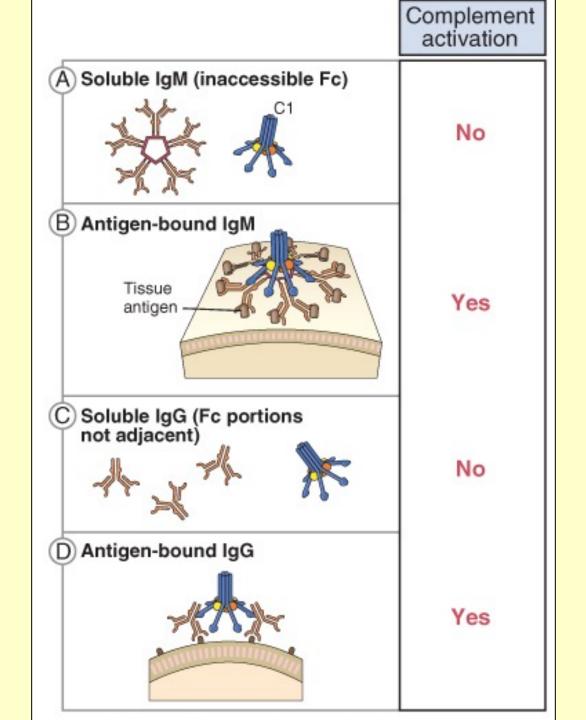
Binding of complement proteins to microbial cell surface or antibody

Formation of C3 convertase

Cleavage of C3

Formation of C5 convertase





First components of lectin pathway

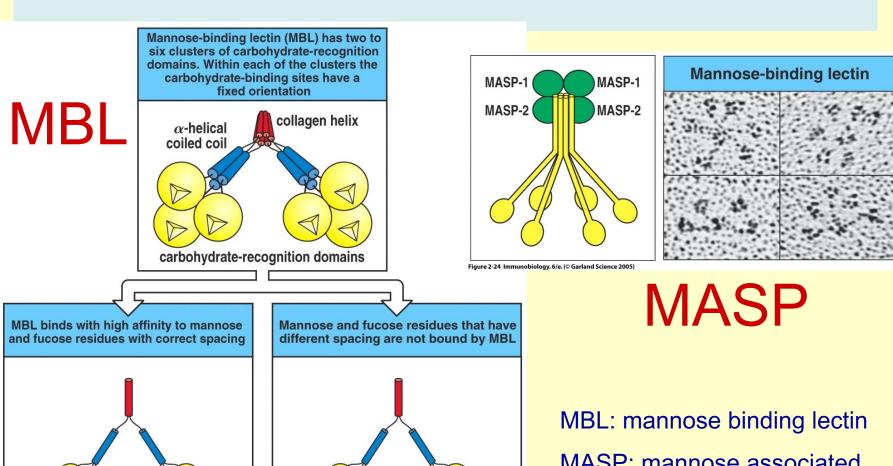
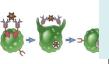


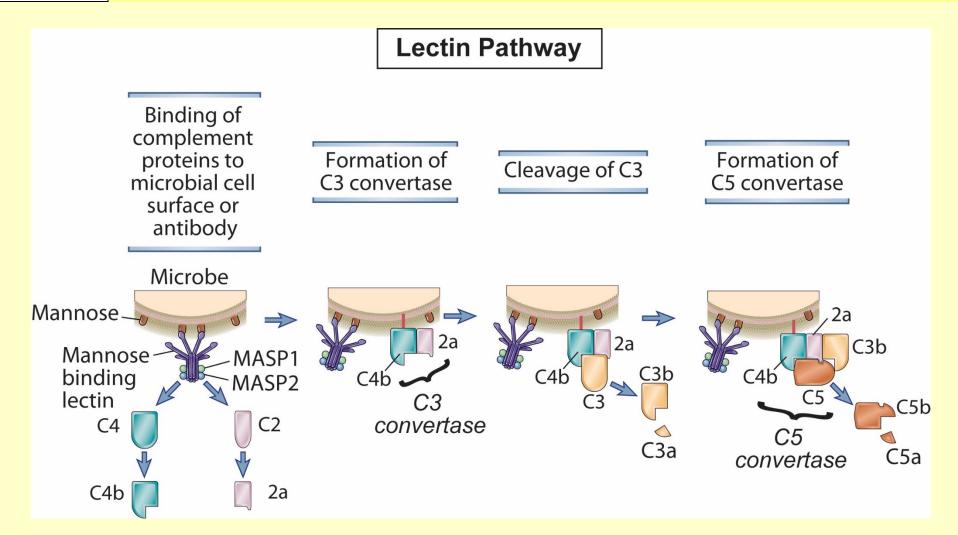
Figure 2-11 Immunobiology, 6/e. (© Garland Science 2005)

MASP: mannose associated

serine protease



Early Steps of Lectin pathway activation



Early steps of alternative pathway activation

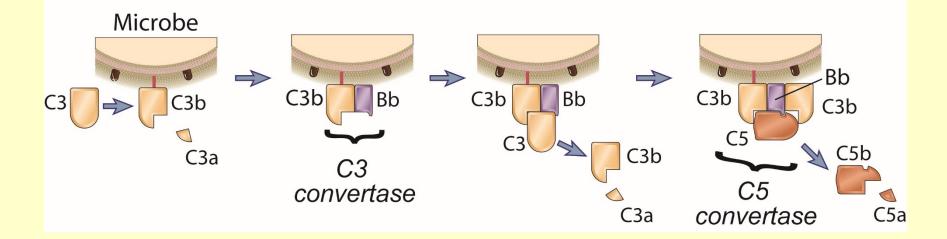
Alternative Pathway

Binding of complement proteins to microbial cell surface or antibody

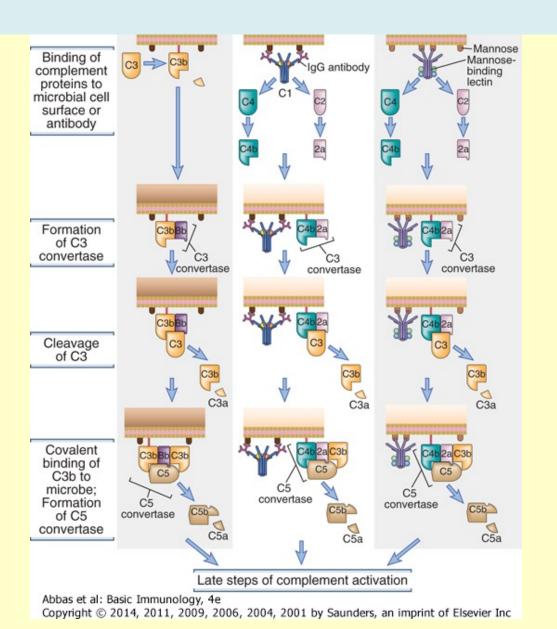
Formation of C3 convertase

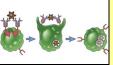
Cleavage of C3

Formation of C5 convertase



Comparison of pathways activation





Late Steps of Complement Activation: MAC

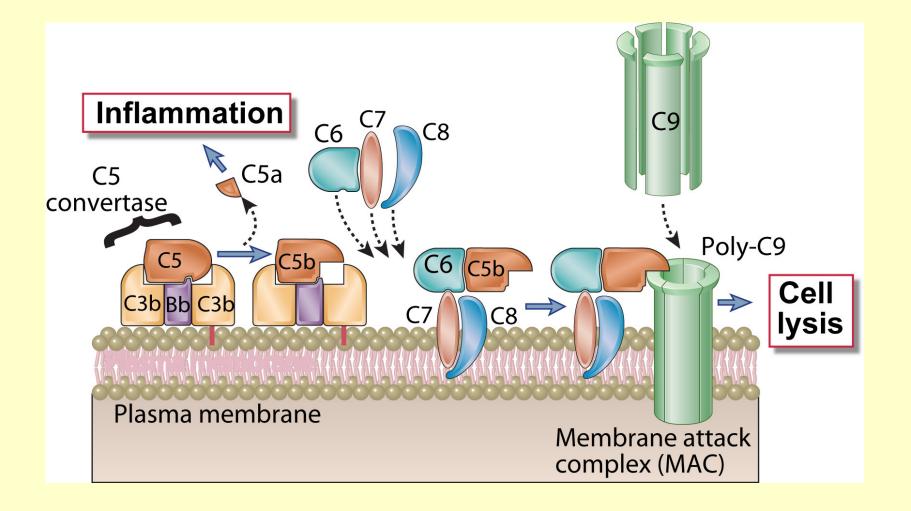
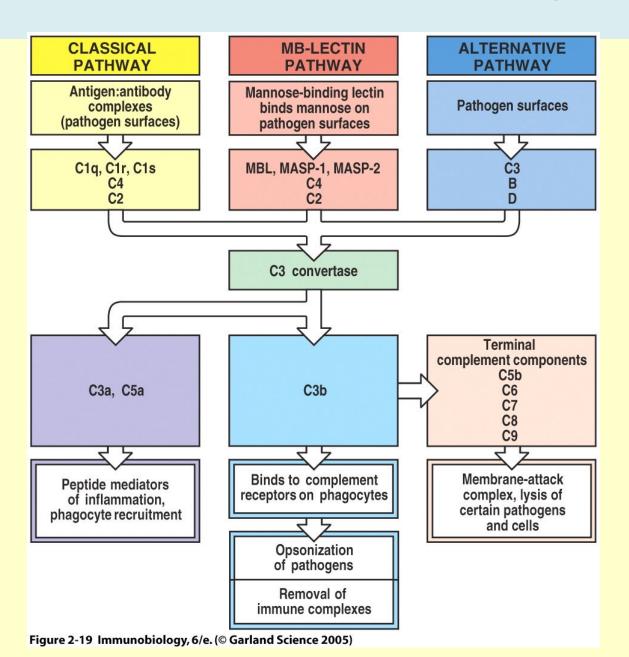
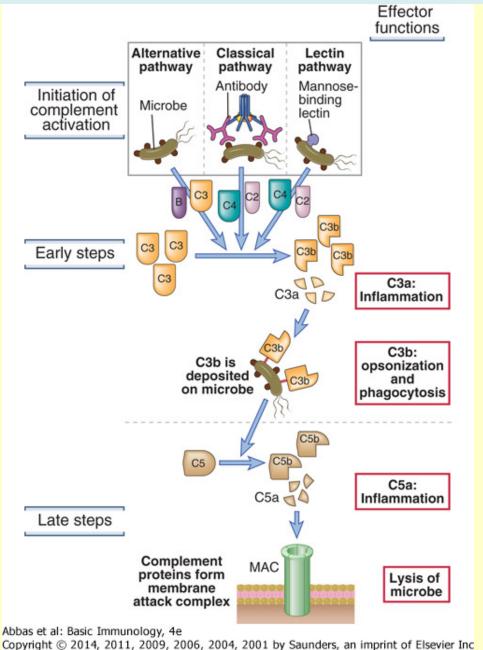


Fig. 12-12

Components and effector actions of complement



Effector functions of complement



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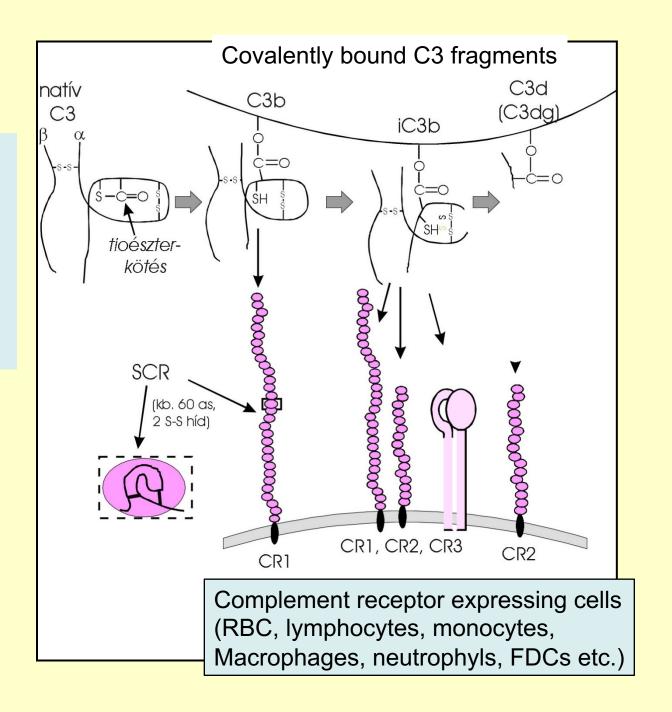
Complement receptors

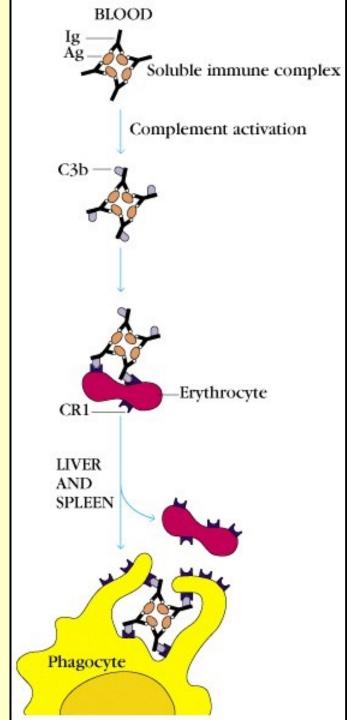
| Receptor | Specificity | Functions | Cell types | |
|---------------------------------------|--|---|---|--|
| CR1 (CD35) | C3b, C4b iC3b | Promotes C3b and C4b decay Stimulates phagocytosis Erythrocyte transport of immune complexes | Erythrocytes, macrophages, monocytes, polymorphonuclear leukocytes, B cells, FDC | |
| CR2 (CD21) | C3d, iC3b, C3dg Epstein– Barr virus | Part of B-cell co-receptor Epstein-Barrvirus receptor | B cells, FDC | |
| CR3 (Mac-1) (CD11b/ CD18) | iC3b | Stimulates phagocytosis | Macrophages, monocytes, polymorphonuclear leukocytes, FDC | |
| CR4 (gp150,95) (CD11c/ CD18) | iC3b | Stimulates phagocytosis | Macrophages, monocytes, polymorphonuclear leukocytes, dendritic cells | |
| C5a receptor | C5a | Binding of C5a activates G protein | Endothelial cells, mast cells, phagocytes | |
| C3a receptor | СЗа | Binding of C3a activates G protein | Endothelial cells, mast cells, phagocytes | |

Figure 2-31 Immunobiology, 6/e. (© Garland Science 2005)

C3b-binding receptors

Complement receptors



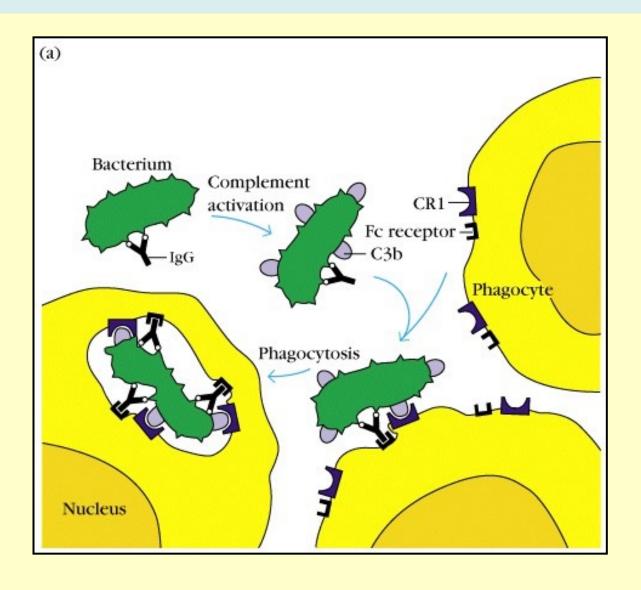


Clearance of immuncomplexes from blood

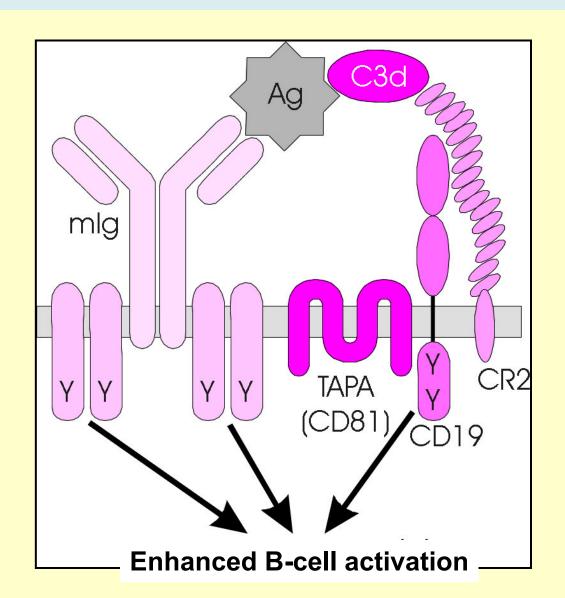
- 1. Immuncomplex formation
- 2. Complement activation C3b binding
- 3. Binding of IC to CR1 of the RBCs
- 4. Transport to the spleen and liver
- 5. Macrophages bind immuncomplexes and take them up by phagocytosis

Inefficient clearance: immuncomplex deposition

Opsonization with C3b → CR mediated phagocytosis

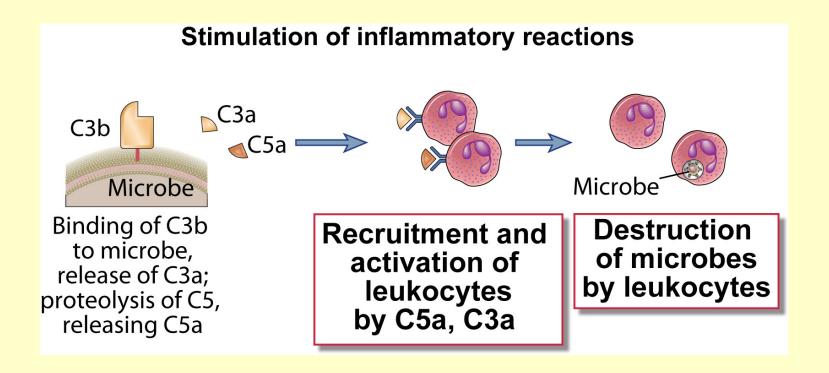


B-cell co-activation through CR2



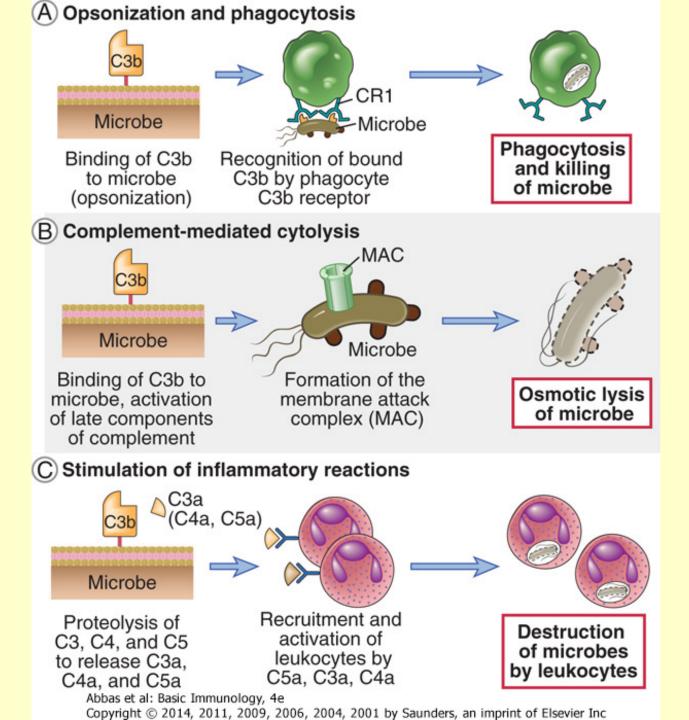


Functions of C3a and C5a



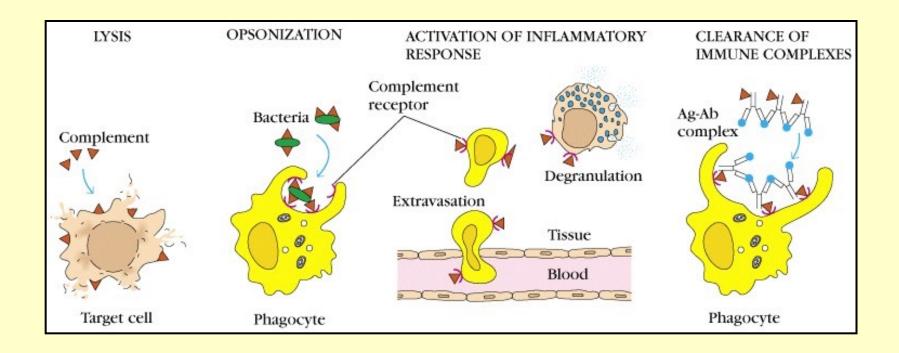
Chemotaxis of granulocytes
Enhancing blood vessel permeability
Mast cell and basophil granulocyte degranulation
Smooth muscle contraction

Fig. 12-17B

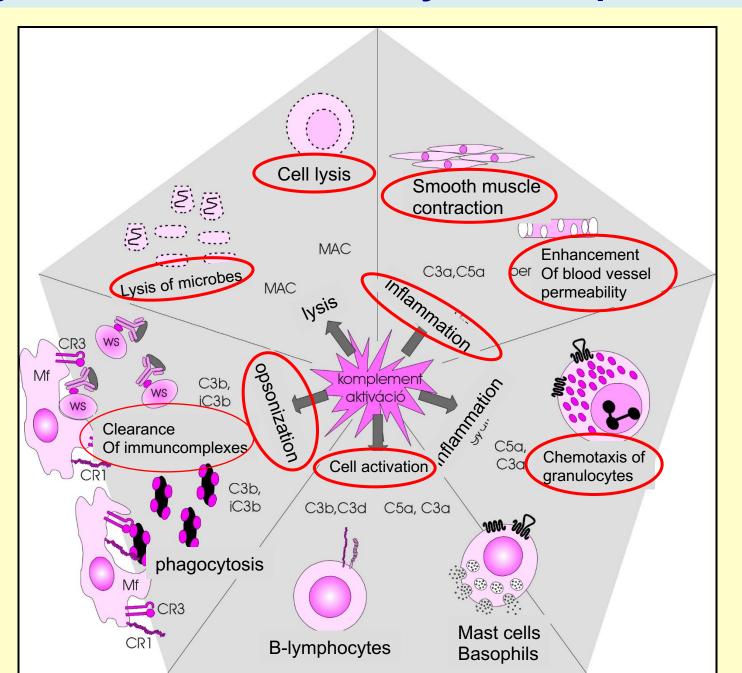


Functions of the complement:

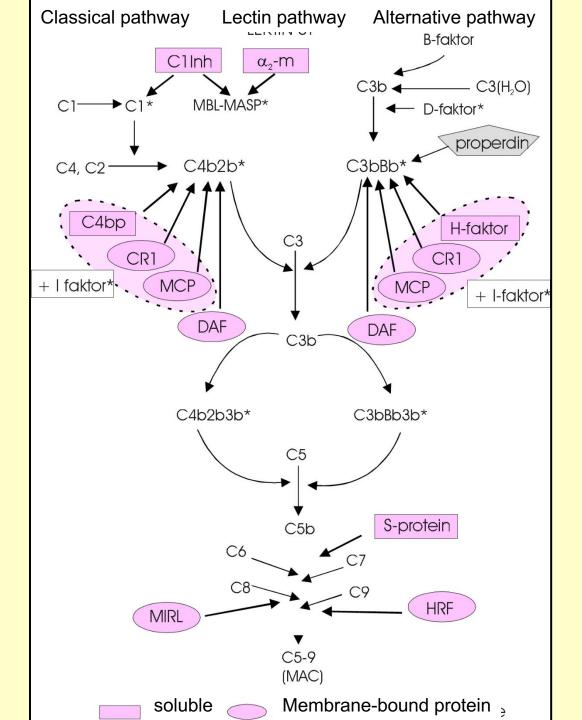
- 1. Lysis of cells, bacteria, viruses
- 2. Opsonization, which promotes phagocytosis of particulate antigens
- 3. Binding to complement receptors results activation of the inflammatory response and specific IR
- 4. Immune clearance of immune complexes from circulation



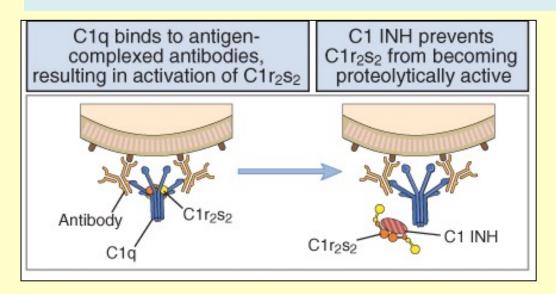
Biological effects, mediated by the complement



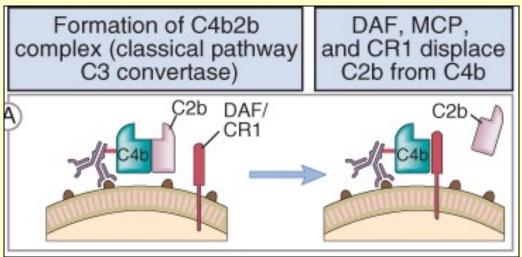
Regulatory proteins



Regulatory proteins of classical pathway



C1 INHIBITOR



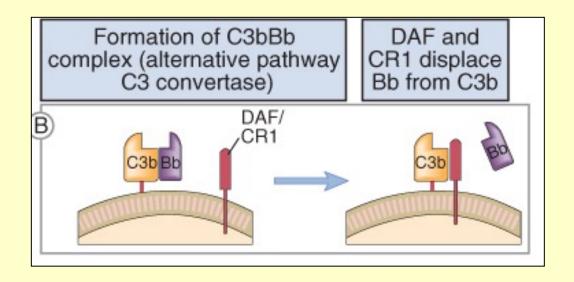
DAF: Decay accelerating factor

MCP: Membrane cofactor

Protein

CR1: complement receptor-1

Regulation of alternative pathway



Regulatory proteins of the classical and alternative pathways

| Name (symbol) | Role in the regulation of complement activation | | |
|---------------------------------|---|--|--|
| C1 inhibitor (C1INH) | Binds to activated C1r, C1s, removing them from C1q | | |
| C4-binding protein (C4BP) | Binds C4b, displacing C2b; cofactor for C4b cleavage by I | | |
| Complement receptor 1 (CR1) | Binds C4b, displacing C2b, or C3b displacing Bb; cofactor for I | | |
| Factor H (H) | Binds C3b, displacing Bb; cofactor for I | | |
| Factor I (I) | Serine protease that cleaves C3b and C4b; aided by H, MCP, C4BP, or CR1 | | |
| Decay-accelerating factor (DAF) | Membrane protein that displaces Bb from C3b and C2b from C4b | | |
| Membrane cofactor protein (MCP) | Membrane protein that promotes C3b and C4b inactivation by I | | |
| CD59 (protectin) | Prevents formation of membrane-attack complex on autologous or | | |

There is a close relationship between the factors of the three complement activations pathways

| Step in pathway | Protein serving function in pathway | | | Polotionship |
|----------------------------------|-------------------------------------|-------------|------------|------------------------------|
| Otop in patriway | Alternative (innate) | MB-lectin | Classical | Relationship |
| Initiating serine protease | D | MASP | C1s | Homologous (C1s and MASP) |
| Covalent binding to cell surface | C3b C4b | | Homologous | |
| C3/C5 convertase | Bb | C2b | | Homologous |
| Control of activation | CR1 H | CR1 C4BP | | Identical Homologous |
| Opsonization | C3b | | | Identical |
| Initiation of effector pathway | tor pathway C5b | | | Identical |
| ocal inflammation C5a, C3a | | | Identical | |
| Stabilization | P None | | Unique | |

Figure 2-29 Immunobiology, 6/e. (© Garland Science 2005)

Complement and diseases

- MAC deficiency C8 mutation: → autoimmune diseases
- → Neisseria infections (Gram- bacteria)
- Faktor H és MCP mutation: atypic HUS (haemolytic uremic syndrome)
- C3, B factor, I factor H factor polymorphisms → unregulated complement activation on cell surfaces: macula degeneration
- C1 inhibitor mutation: <u>hereditary angioedema</u> (HAE)
- Diagnostics: CH50 measurement = total complement activity test

Hereditary angioedema (HAE)

- is an autosomal dominant disease caused by either a lack of C1-inhibitor protein or dysfunctional C1-inhibitor protein.
- HAE manifests with symptoms related to angioedema of the upper airway, skin, and/ or gastrointestinal tract.
- Autosomal dominant inheritance or a new mutation
- Deficiency of complement C1-inhibitor (C1-INH), a plasma protein that is an important inhibitor of several serine proteases, specially of the complement system and the contact activation/kallikrein-kinin pathway, but also the fibrinolytic system
- Treatment: C1-inhibitor plasma concentrate or recombinant C1-INH replacement therapy (rhC1INH; conestat alfa) in Europe