

Immune Response

Primary

- Lag phase longer (differentiates into plasma & memory B. cells)
- Logarithmic phase increase in serum antibodies

Secondary

- Lag phase shorter
- Memory B cells more
- Higher affinity of memory B cells

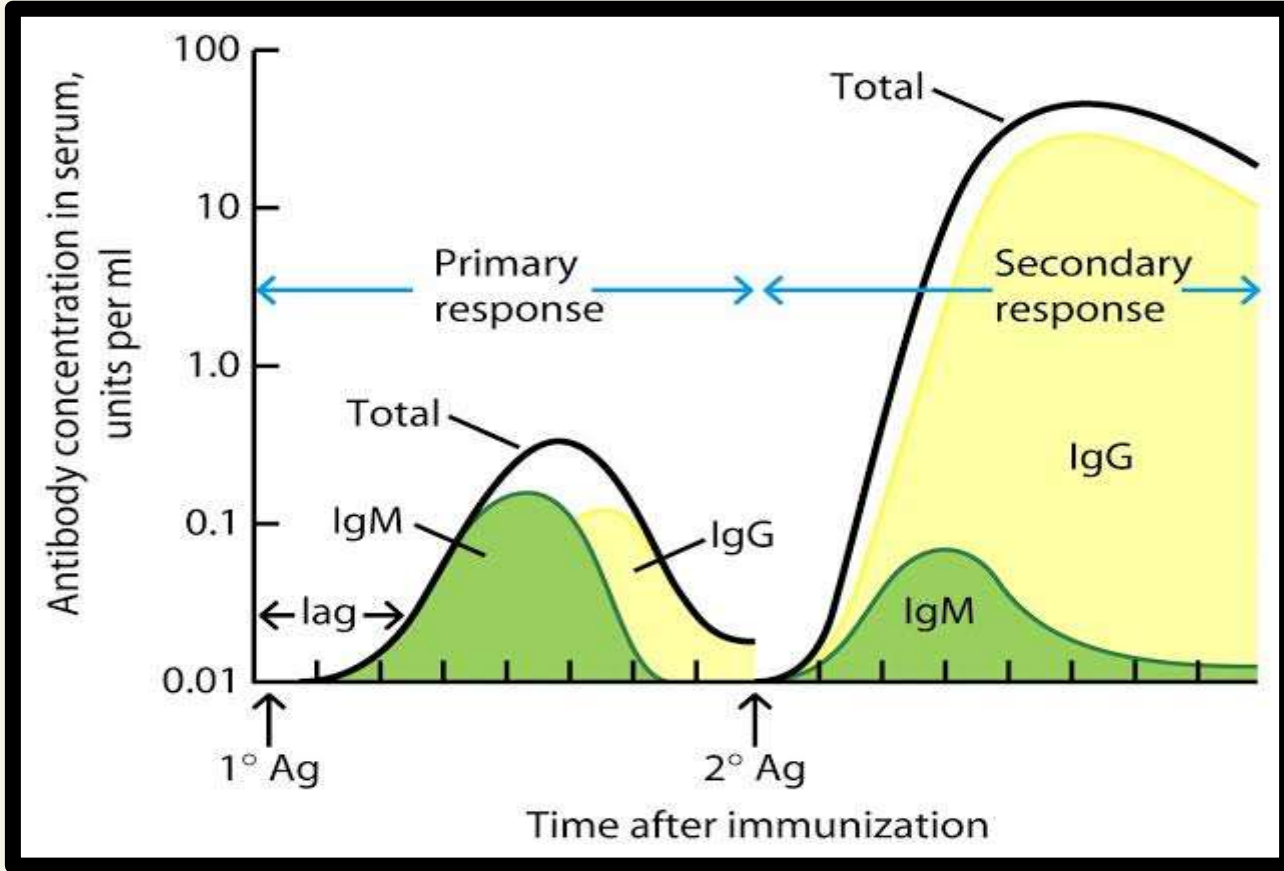


TABLE 11-4**Comparison of primary and secondary antibody responses**

Property	Primary response	Secondary response
Responding B cell	Naive (virgin) B cell	Memory B cell
Lag period following antigen administration	Generally 4–7 days	Generally 1–3 days
Time of peak response	7–10 days	3–5 days
Magnitude of peak antibody response	Varies depending on antigen	Generally 100–1000 times higher than primary response
Isotype produced	IgM predominates early in the response	IgG predominates
Antigens	Thymus-dependent and thymus-independent	Thymus-dependent
Antibody affinity	Lower	Higher



02
The
Complement
System

- **Complement is the name given to a complex series of some 20 proteins which, along with blood clotting, fibrinolysis and kinin formation, forms one of the triggered enzyme systems found in plasma**
- **Present in inactive form in serum**
- **Produce a rapid, highly amplified response to a trigger stimulus mediated by a cascade phenomenon**
- **Complement components are designated by the letter 'C' followed by a number.**
- **Most pivotal component is C3 .**

- **Peptide fragments formed by activation of a component are denoted by small letters like C3a, C3b etc.**
- **Complement fragments interact with one another to form functional complexes (C452a) which have enzymatic activity**
- **Functions:**
 - **-Lysis of cells, bacteria, and viruses**
 - **- Opsonization- Inflammation**
 - **- Immune clearance**

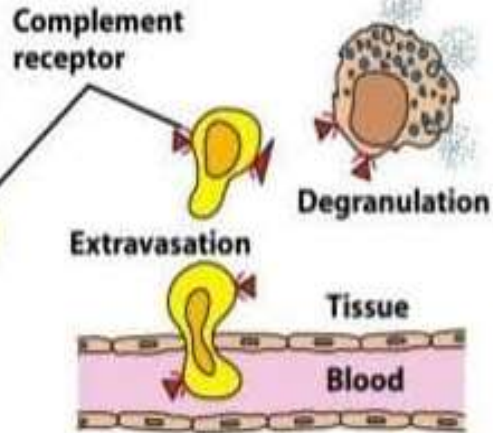
LYSIS



OPSONIZATION



ACTIVATION OF INFLAMMATORY RESPONSE



CLEARANCE OF IMMUNE COMPLEXES

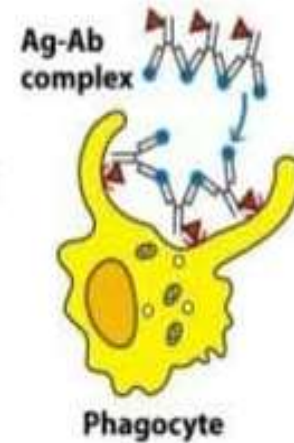


Figure 7-1
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Complement activation

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graph TD; A[Complement activation] --> B[Classical (Ab dependent)]; A --> C[Alternative (Ab independent)]; A --> D[Lectin]
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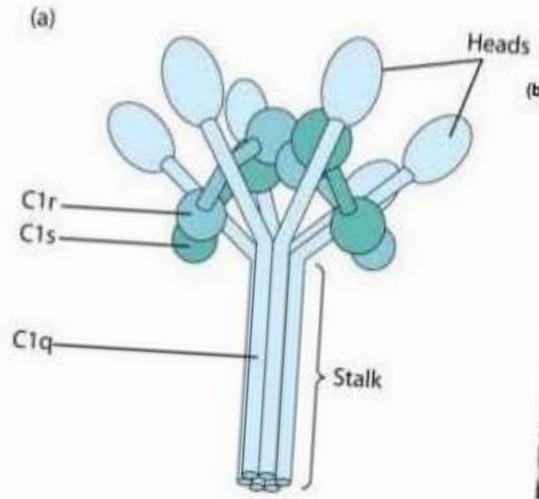
Classical (Ab dependent)

Alternative (Ab independent)

Lectin

Classical Pathway

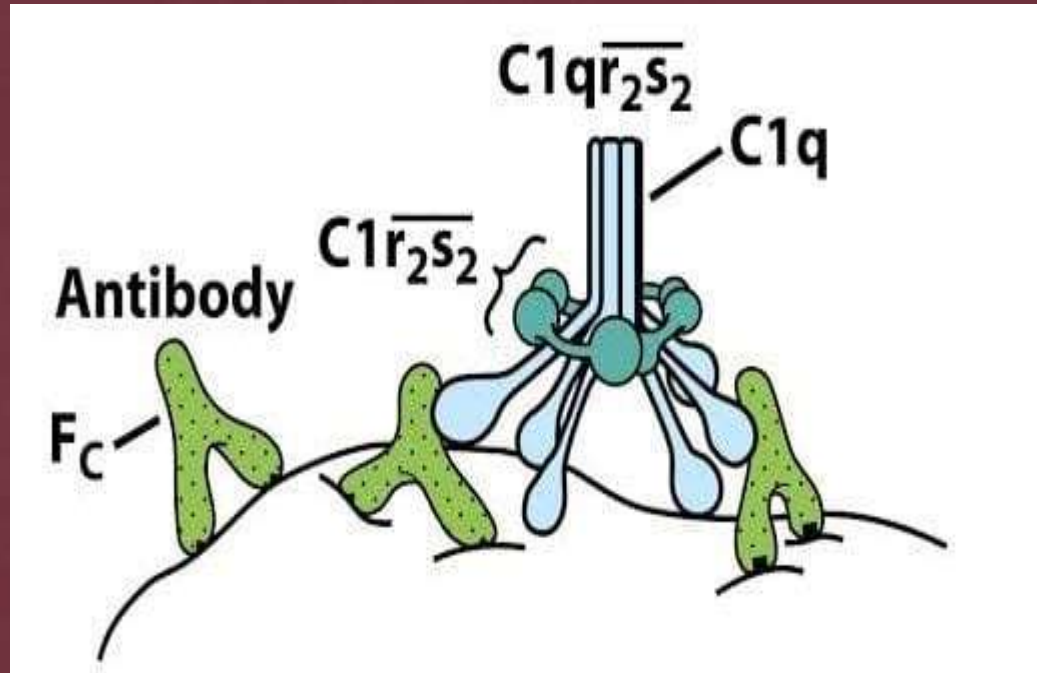
- Begins with the formation of soluble antigen-antibody complexes (immune complexes) or with the binding of antibody to antigen on a suitable target, such as a bacterial cell
- IgM and certain subclasses of IgG (human IgG1, IgG2, and IgG3) can activate the classical complement pathway
- Formation of an antigen-antibody complex induces conformational changes in the Fe portion of the IgM molecule that expose a binding site for the C1 component
- This further activates C1s component of C1 complement



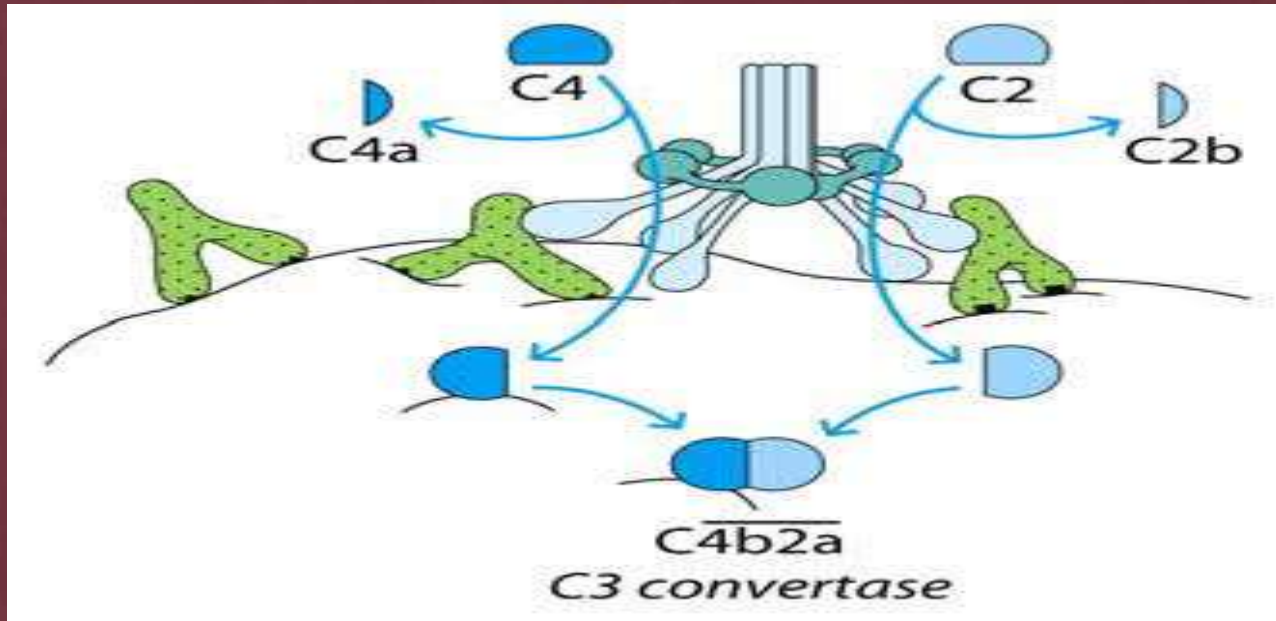
C1q molecule

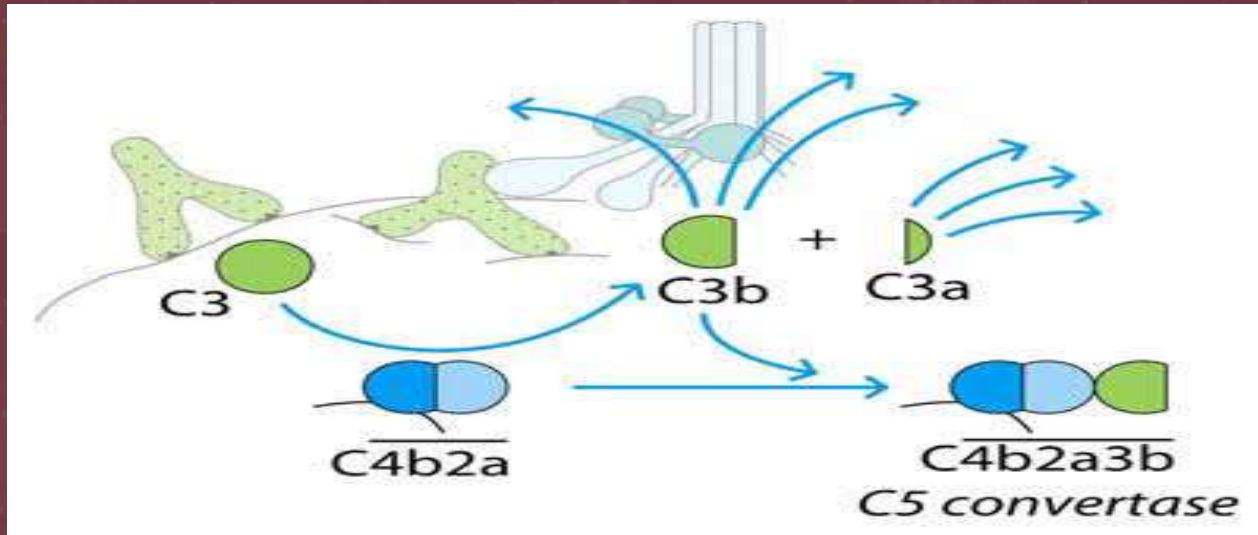


Kelly J et al., Immunology 2003



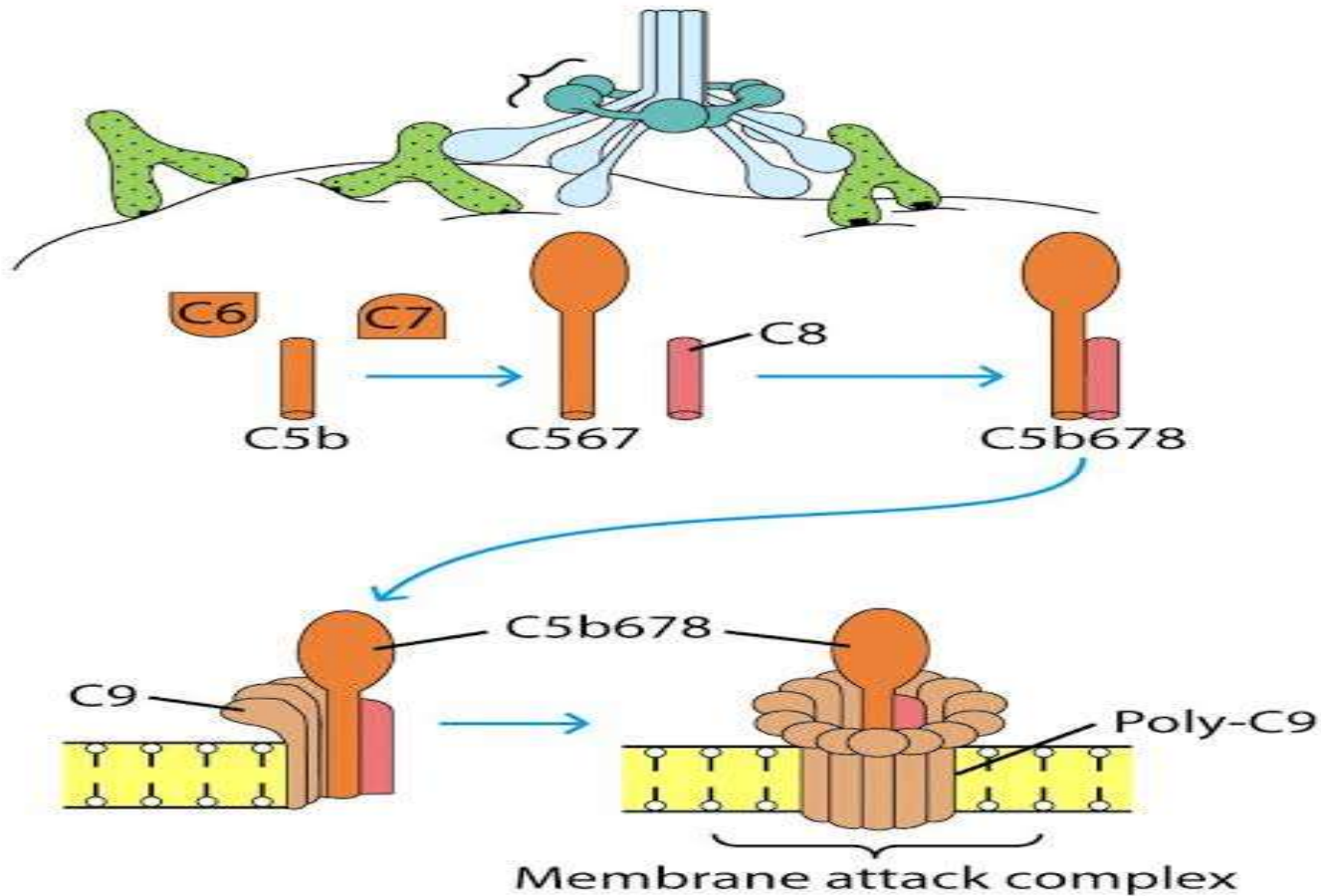
- C4 is activated when CTS hydrolyzes a small fragment (C4a) from the amino terminus of the chain, exposing a binding site on the larger fragment (C4b)
- C4b fragment attaches to the target surface in the vicinity of C1, and the C2 proenzyme then attaches to the exposed binding site on C4b.
- C2 is then cleaved by the neighboring CTS, forming C4b2a complex is called C3 convertase

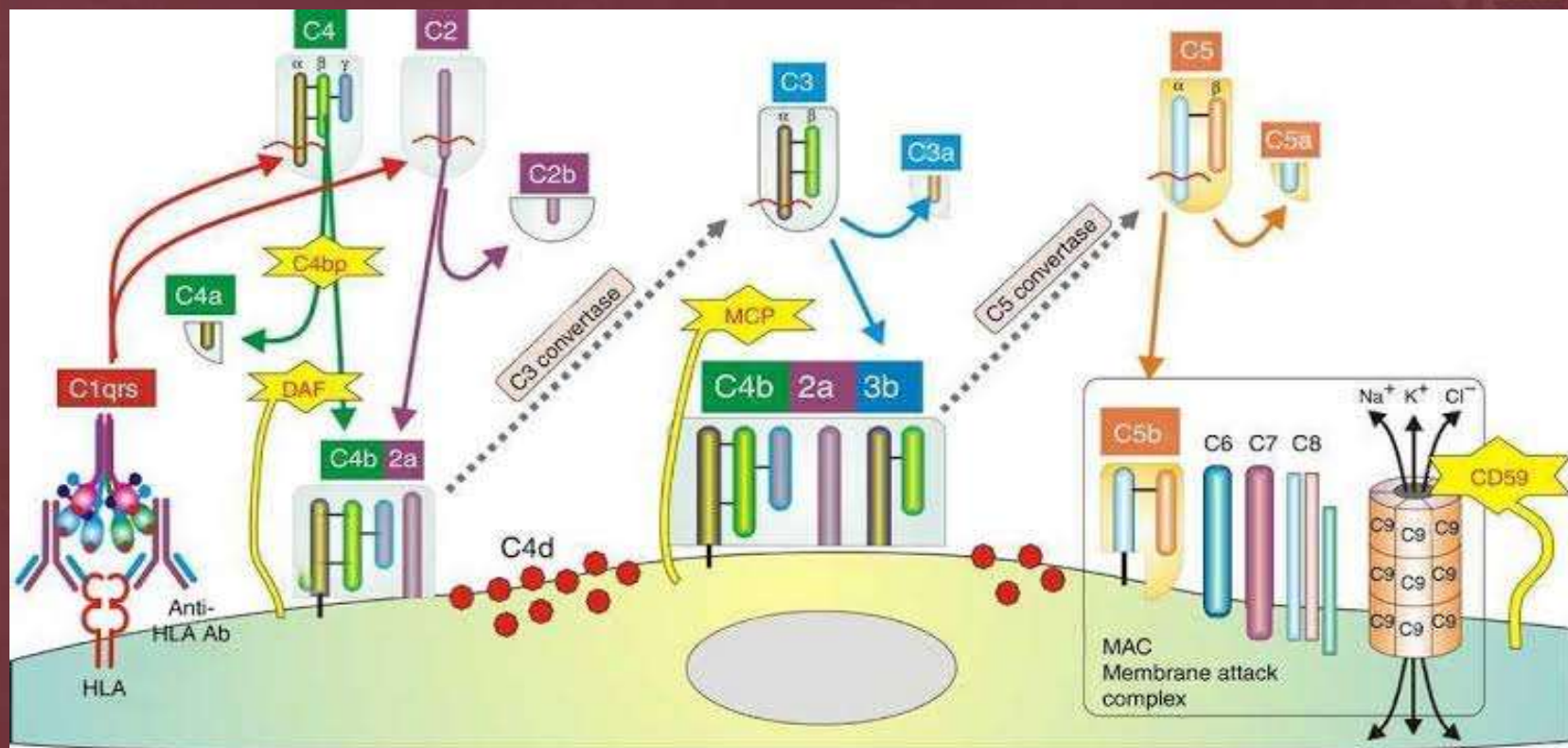




- *C3 convertase molecule can generate over*
- *200 molecules of C3b, which binds to*
- *C4b2a to form a tri molecular complex*
- *C4b2a3b, called C5 convertase*

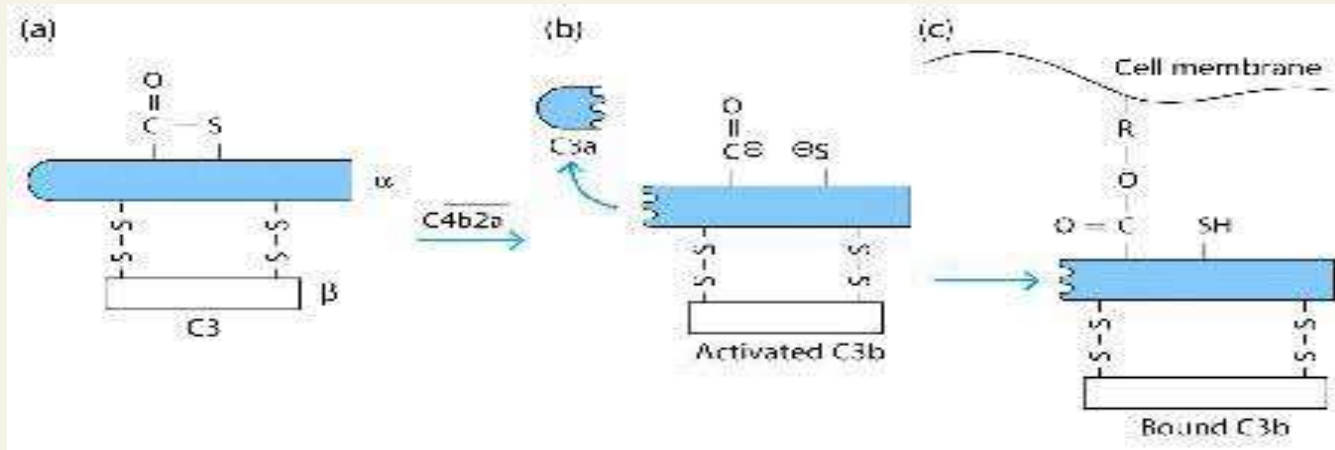
- **C5 convertase breaks C5 complement, generating C5b, which attaches to the surface of the target cell and C6 and initiates formation of the membrane- attack complex**
- **C5b6 binds to C7, the resulting complex undergoes a hydrophilic-amphiphilic structural transition which serve as binding sites for membrane phospholipids**
- **Binding of C8 to membrane-bound C5b67, forming C5b678 complex which creates a small pore, 10 Å in diameter**
- **Final step involves polymerization of C9, a perforin-like molecule, to the C5b678 complex completing MAC formation (functional pore size of 70-100 Å).**



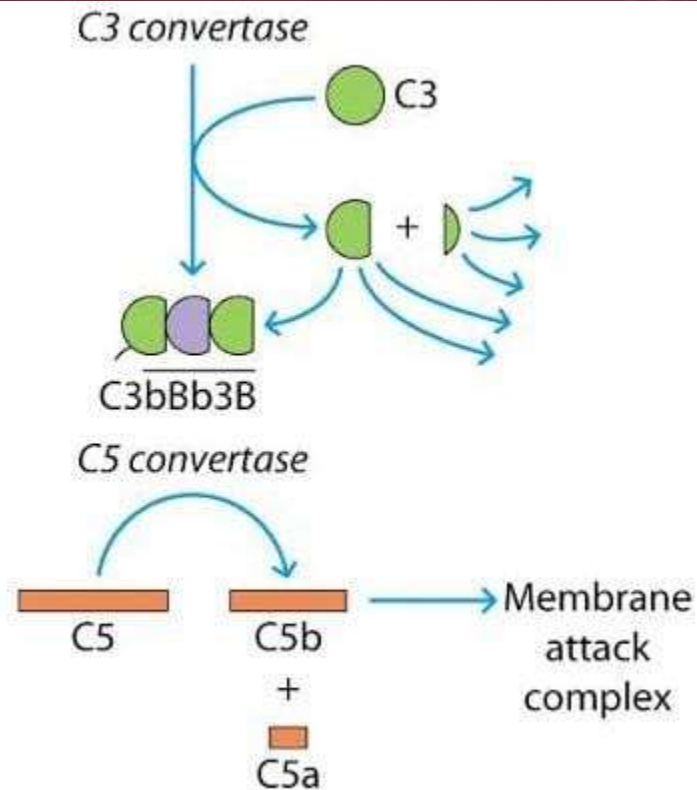
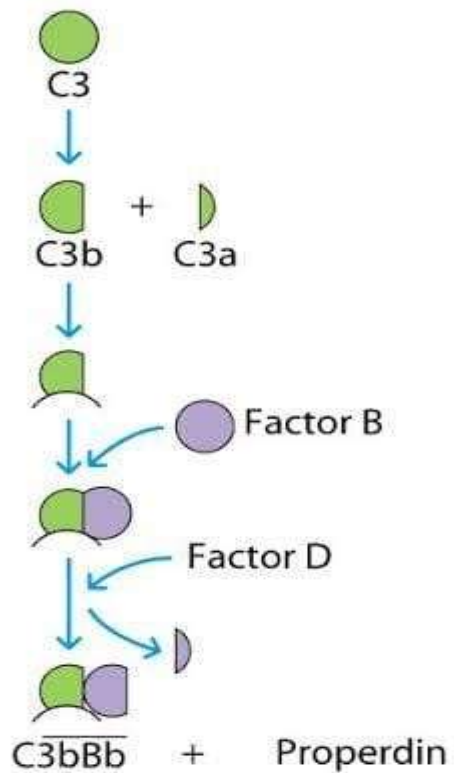


Alternate Pathway

- Alternative pathway does not need for antigen-antibody complexes for initiation
- Serum C3, is subjected to slow spontaneous hydrolysis to yield C3a and C3b

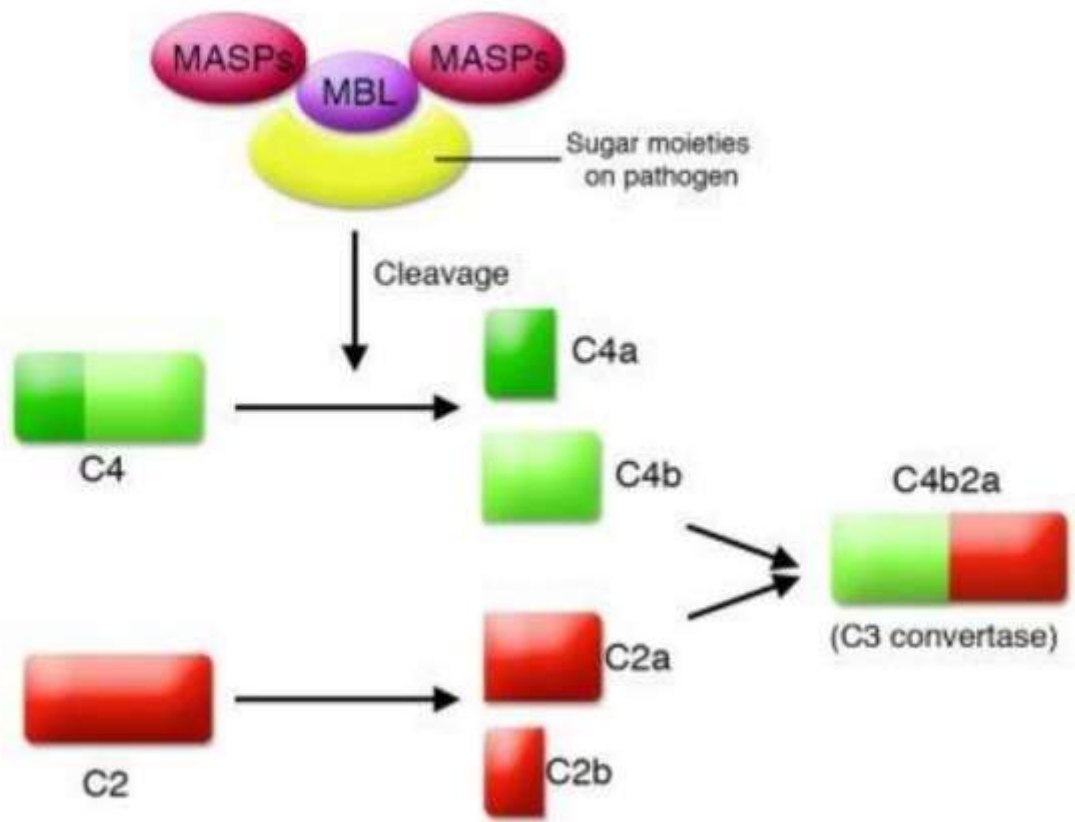


- **C3b component can bind to foreign surface antigens, which can bind another serum protein called factor B**
- **Factor D cleaves the C3b-bound factor B, generating C3bBb -C3 convertase..**
- **Further serum protein properdin binds to it and stabilizes it.**
- **C3 convertase cleaves C3 complement generating the C3bBb3b complex, which exhibits C5 convertase activity**



Lectin Pathway

- Lectin pathway is activated by the binding of mannose-binding lectin (MBL) to mannose residues on glycoproteins or carbohydrates on the surface of microorganisms
- Next MBL-associated serine proteases, MASP-1 and MASP 2, bind to MBL
- This association causes cleavage and activation of C4 and C2
- After this follows the classical pathway to generate C5 convertase



Biological Effects of Complements

Effect	Complement product mediating*
Cell lysis	C5b-9, the membrane-attack complex (MAC)
Inflammatory response	
Degranulation of mast cells and basophils [†]	C3a, C4a, and C5a (anaphylatoxins)
Degranulation of eosinophils	C3a, C5a
Extravasation and chemotaxis of leukocytes at inflammatory site	C3a, C5a, C5b67
Aggregation of platelets	C3a, C5a
Inhibition of monocyte/macrophage migration and induction of their spreading	Bb
Release of neutrophils from bone marrow	C3c
Release of hydrolytic enzymes from neutrophils	C5a
Increased expression of complement receptors type 1 and 3 (CR1 and CR3) on neutrophils	C5a
Opsonization of particulate antigens, increasing their phagocytosis	C3b, C4b, iC3b
Viral neutralization	C3b, C5b-9 (MAC)
Solubilization and clearance of immune complexes	C3b

Regulation Of Complement System

TABLE 13-2 Proteins that regulate the complement system

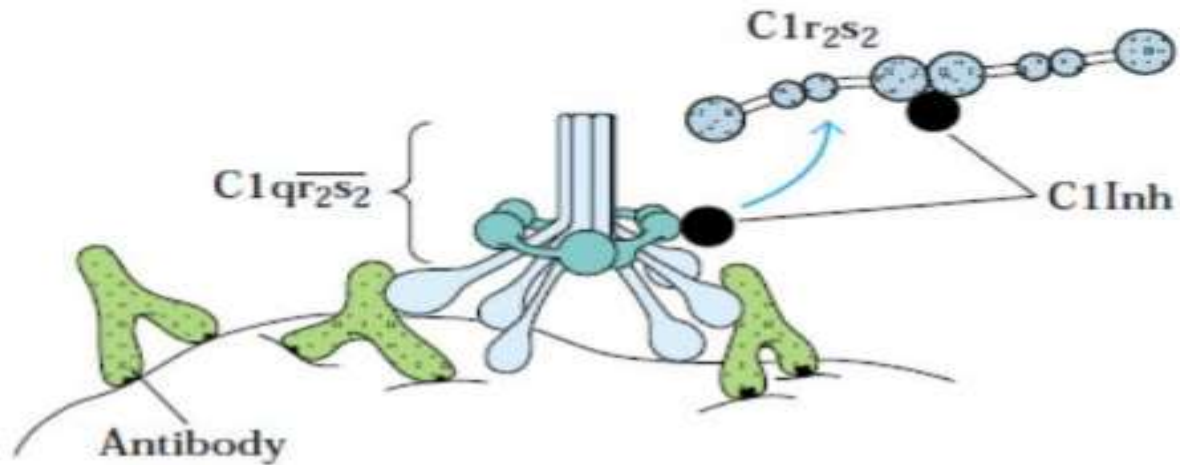
Protein	Type of protein	Pathway affected	Immunologic function
C1 inhibitor (C1Inh)	Soluble	Classical	Serine protease inhibitor; causes C1r ₂ s ₂ to dissociate from C1q
C4b-binding protein (C4bBP)*	Soluble	Classical and lectin	Blocks formation of C3 convertase by binding C4b; cofactor for cleavage of C4b by factor I
Factor H*	Soluble	Alternative	Blocks formation of C3 convertase by binding C3b; cofactor for cleavage of C3b by factor I
Complement-receptor type 1 (CR1)*	Membrane bound	Classical, alternative, and lectin	Block formation of C3 convertase by binding C4b or C3b; cofactor for factor I-catalyzed cleavage of C4b or C3b C3bBb
Membrane-cofactor protein (MCP)*			
Decay-accelerating factor (DAF or CD55)*	Membrane bound	Classical, alternative, and lectin	Accelerates dissociation of C4b2a and C3bBb (classical and alternative C3 convertases)

*An RCA (regulator of complement activation) protein. In humans, all RCA proteins are encoded on chromosome 1 and contain short consensus repeats.

(a) Before assembly of convertase activity

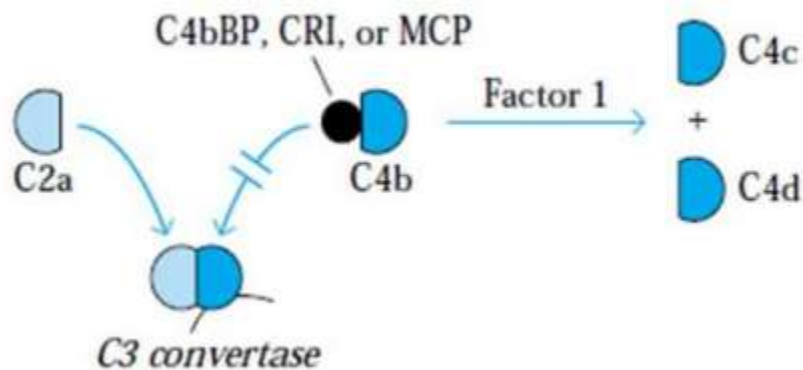
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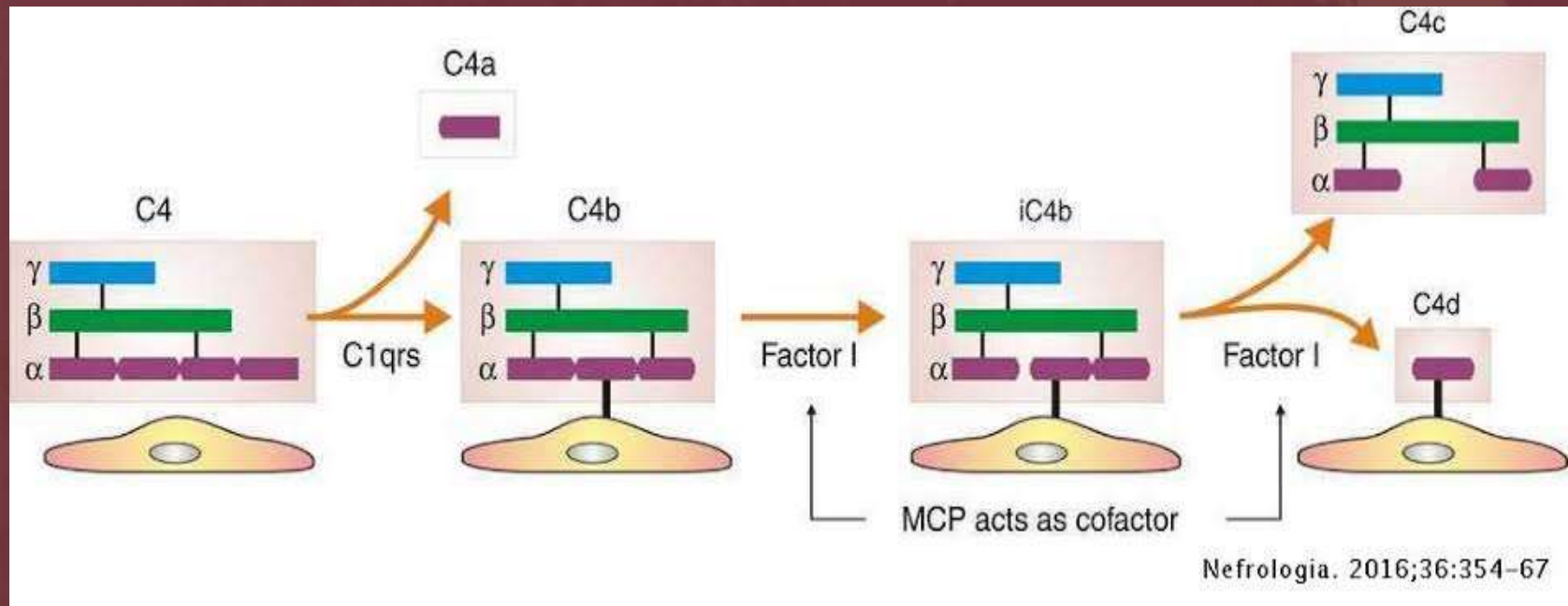
C1 inhibitor (C1Inh) binds $C1r_2s_2$, causing dissociation from $C1q$



2 Association of C4b and C2a is blocked by binding C4b-binding protein (C4bBP), complement receptor type I, or membrane cofactor protein (MCP)

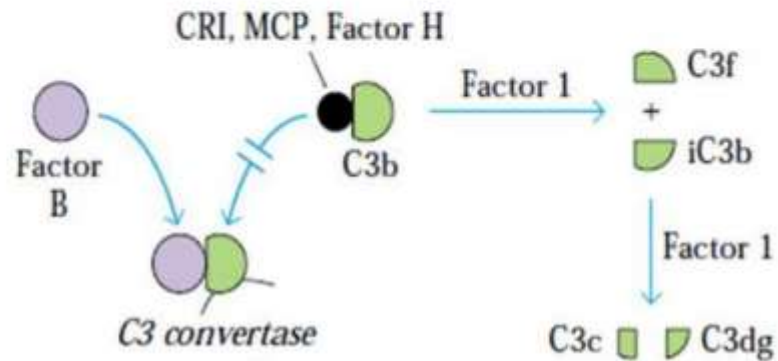
3 Inhibitor-bound C4b is cleaved by Factor 1





4 In alternative pathway, CR1, MCP, or Factor H prevent binding of C3b and Factor B

5 Inhibitor-bound C3b is cleaved by Factor 1



(b) After assembly of convertase

C3 convertases are dissociated by C4bBP, CR1, Factor H, and decay-accelerating Factor (DAF)



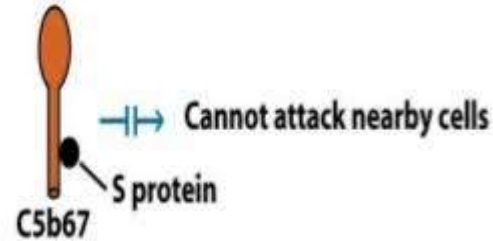
C4bBP, CR2
Factor H, DAF



Dissociation of convertase;
remaining C4b or C3b
cleaved by Factor 1

Regulation at assembly of membrane-attack complex (MAC)

1 S protein prevents insertion of C5b67 MAC component into the membrane.



2 Homologous restriction factor (HRF) or membrane inhibitor of reactive lysis (MIRL or CD59) bind C5b678, preventing assembly of poly-C9 and blocking formation of MAC.

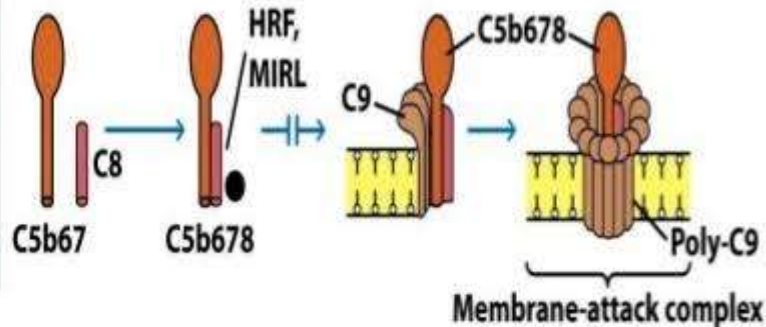


Figure 7-10c

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