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18/08/20

Topic: Complement System, Activation, Functions & Deficiencies (Part II)

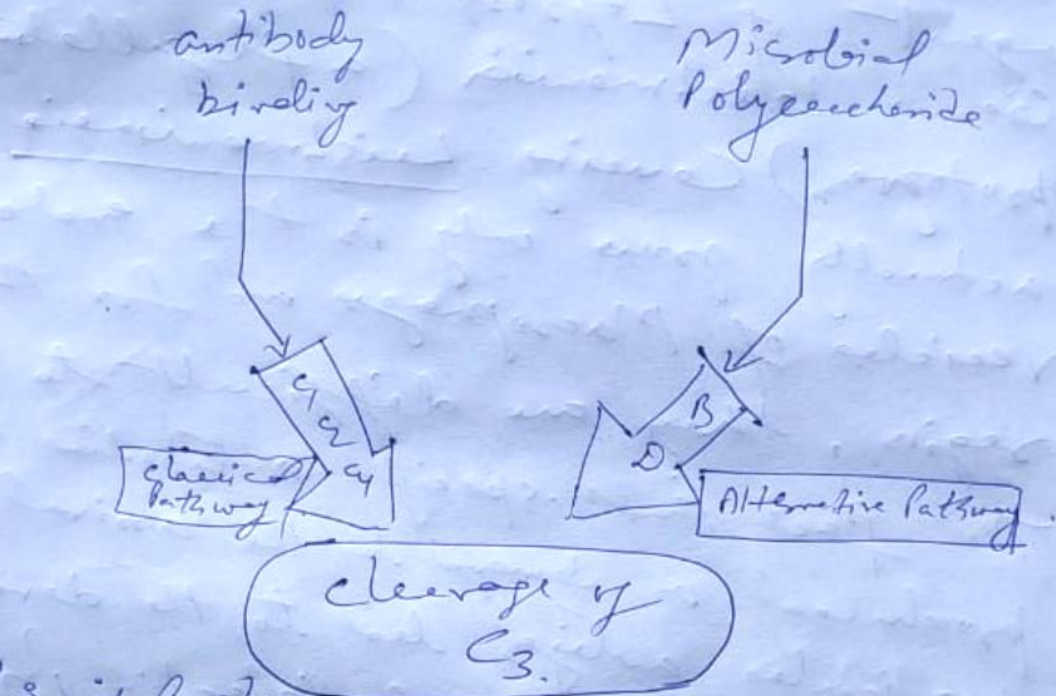
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Contd. from Part I -

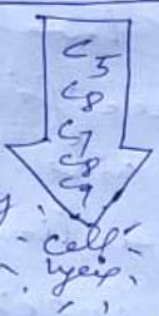
The activation of  $C_3$  by cleavage is the central reaction in the Complement Activation Sequence, and it is here that the classical and alternative pathways converge. In both pathways,  $C_3$  is cleaved by an enzyme complex called  $C_3$  Convertase. A different  $C_3$  Convertase is produced by each pathway formed by spontaneous assembly of two of the Complement Components activated earlier in the cascade. Both types of  $C_3$  Convertase cleave  $C_3$  into two fragments. The larger of ( $C_3b$ ) binds covalently to the target cell membrane and binds  $C_5$ . Once bound the  $C_5$  protein is cleaved by the  $C_3$  Convertase to initiate the spontaneous assembly of the rest of the membrane attack complex (C5 through C9). That leads to the membrane attack complex. Since each activation

enzyme cleaves many molecules of the next protoenzyme is the chain, the activation of the early components consists of an amplifying proteolytic cascade; each molecule activated at the beginning of the sequence lead to the production of many membrane attack complexes. The classical pathway usually activated by IgG or IgM antibodies bound to the antigens on the surface of a micro-organism or the target cell. The binding of antigens by these antibodies enable their constant regions to bind in turn to the first component is the classical pathway, C<sub>1</sub> which is the large complex composed of three sub-components C<sub>1q</sub>, C<sub>1r</sub>, and C<sub>1s</sub>. The molecule of C<sub>1q</sub> protein is large sized (4,50,000 daltons) and made up of six identical subunits; each composed of three different polypeptide chains. The carboxyl terminal (-COO) halves of each of the polypeptides in a subunit are folded in a globular structure, the terminal halves; here a NH<sub>2</sub> amino acid sequence wound together to form a

collagen like triple stranded helix. The six subunits are linked together by disulphide bonds between their triple helical stems; forming a structure that resembles a bunch of Tulips.



The principal stages  
is Complement  
Activation by classical  
& alternative pathway



Such as Activator  
occurs at the surface  
of an invading bacterium  
(After Albert et al. 1989)

The binding of a globular head of C<sub>1</sub> to an IgG or IgM antibody bound to antigen activate C<sub>1</sub> to start the early proteolytic cascade of the classical pathway. Activation of the C<sub>1</sub> activates C<sub>4</sub> to become proteolytic and C<sub>4</sub> is then cleaved and activates C<sub>3</sub>. Activated C<sub>3</sub> then cleaves C<sub>5</sub> into two fragments. C<sub>5a</sub> and C<sub>5b</sub> are smaller designated as small and large respectively. C<sub>5b</sub> immediately binds cover

(Pg 4)

-tently to the membrane and binds  $C_2$ .  
Once bound  $C_2$  is also cleared by  $C_5$ .  
There are some microbes, which are ingested but not killed by macrophages. e.g. Mycobacterium leprae, the causative agent of Leprosy and parasite of Genus Leishmania (a flagellated protozoan) causing Leishmaniasis, actually grows only in the endocytic vesicles of Macrophages and for this reason, they are difficult to kill.

The fact that specific antibodies are produced on demand by the arrival of particular antigens lead to the development of the Clonal Selection Principle.

This theory specifies the idea that an enormously wide range of  $\beta$  lymphocytes, each potentially capable of producing a specific kind of antibody, is present in the body before birth. When an antigen gets into the body after birth, it selects a lymphocyte of appropriate type, adheres to its surface and causes it to proliferate into a clone of cells, all of which proceed to produce appropriate antibodies. This is further aided by the Complement System, which speeds up the rate of proliferation.