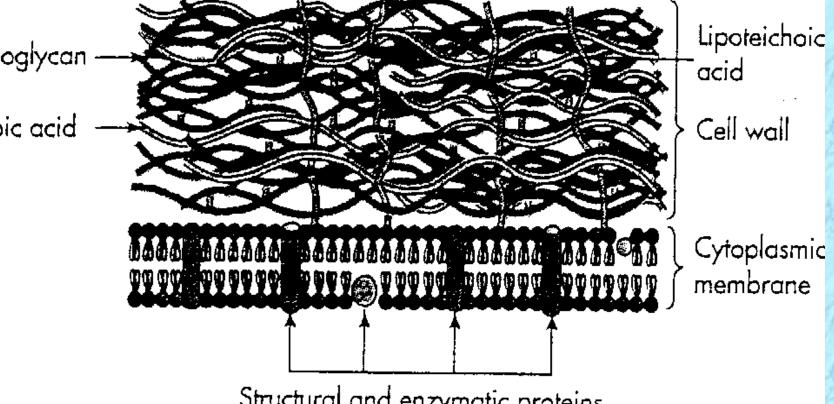
Lipopolysaccharide structure

Gram-Positive Cell Wall

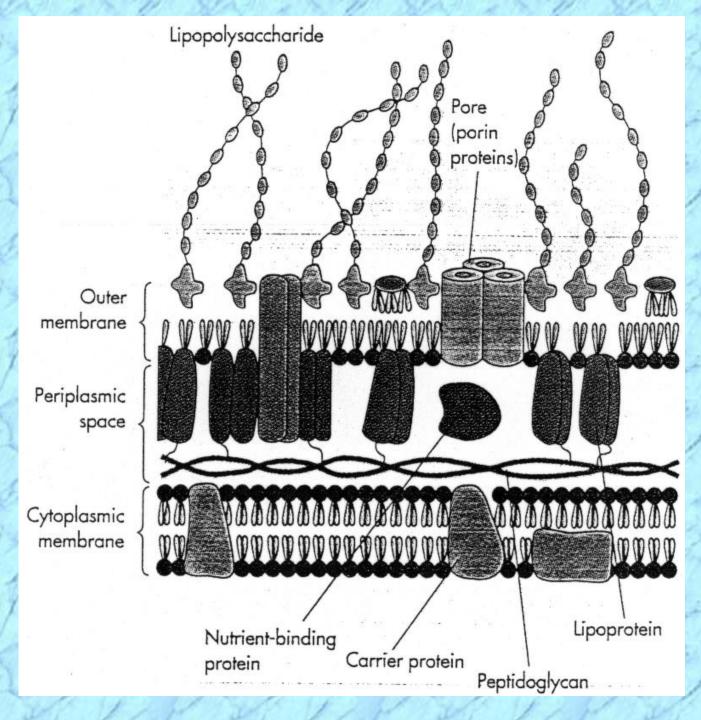
Peptidoglycan

Teichoic acid

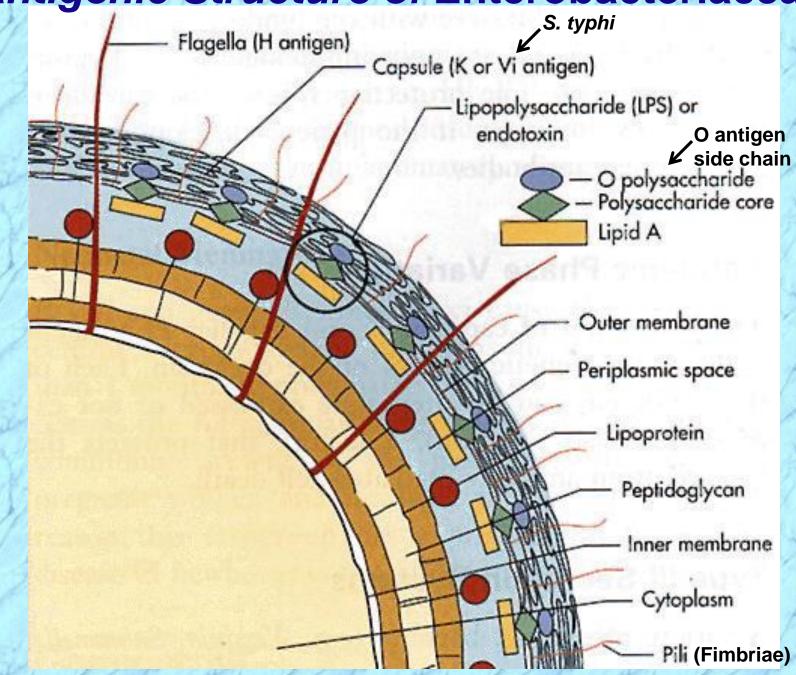


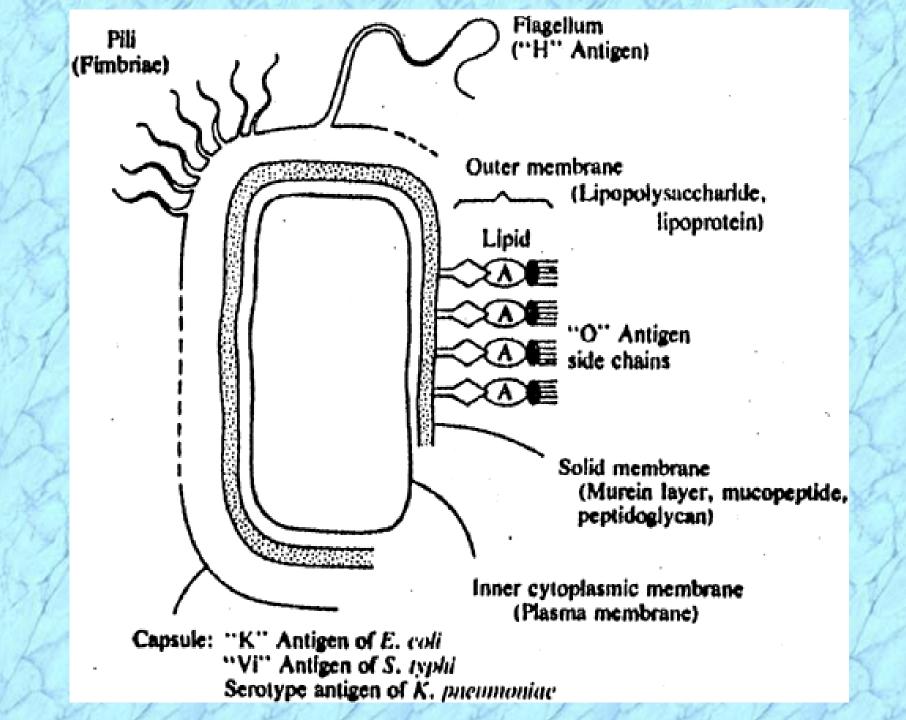
Structural and enzymatic proteins

Gram-Negative Cell Wall

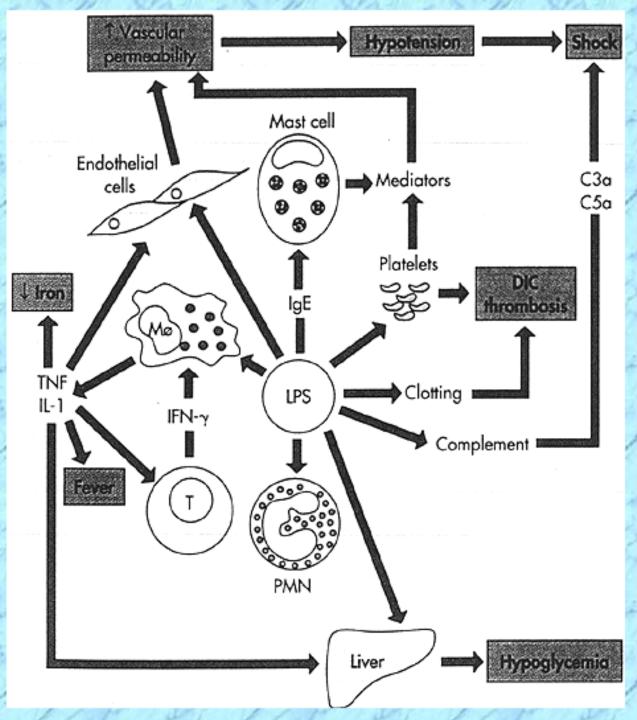


Antigenic Structure of Enterobacteriaceae





Diversity of Activities Associated with LPS



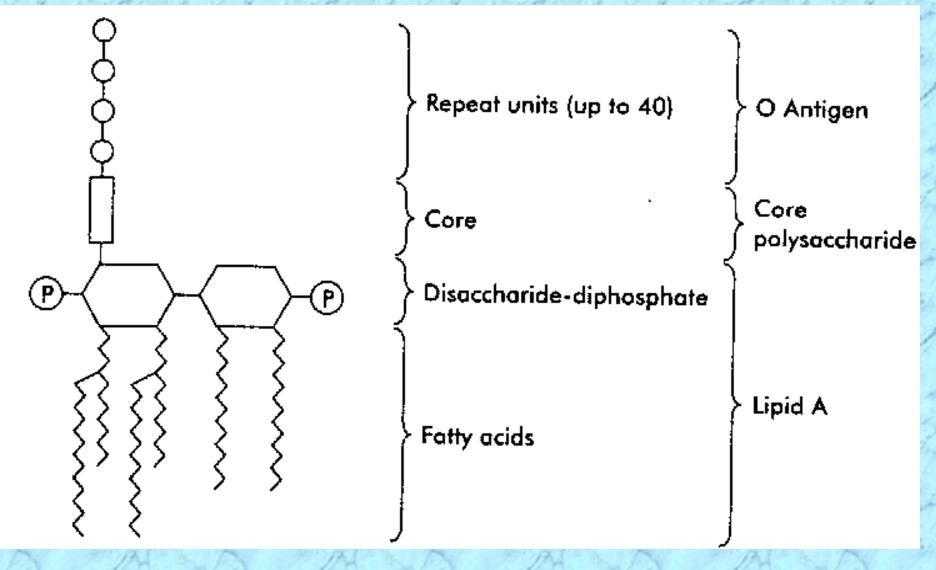




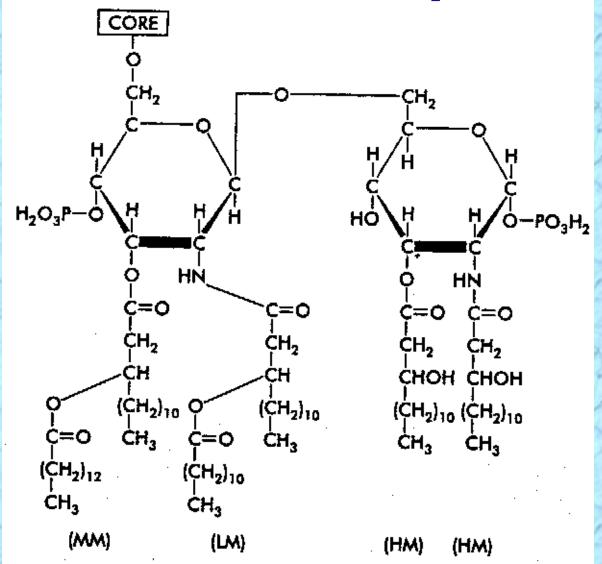
- 1. Released from the cell before 1. Integral part of cell wall or after lysis
- 2. Protein
- 3. Heat labile
- 4. Antigenic and immunogenic
- 5. Toxoids can be produced
- 6. Specific in effect on host
- 7. Produced by gram-positive and gram-negative organisms

- 2. Endotoxin is LPS; Lipid A is toxic component
- 3. Heat stable
- 4. Antigenic; ??immunogenicity
- 5. Toxoids cannot be produced
- 6. Many effects on host
- 7. Produced by gram-negative organisms only

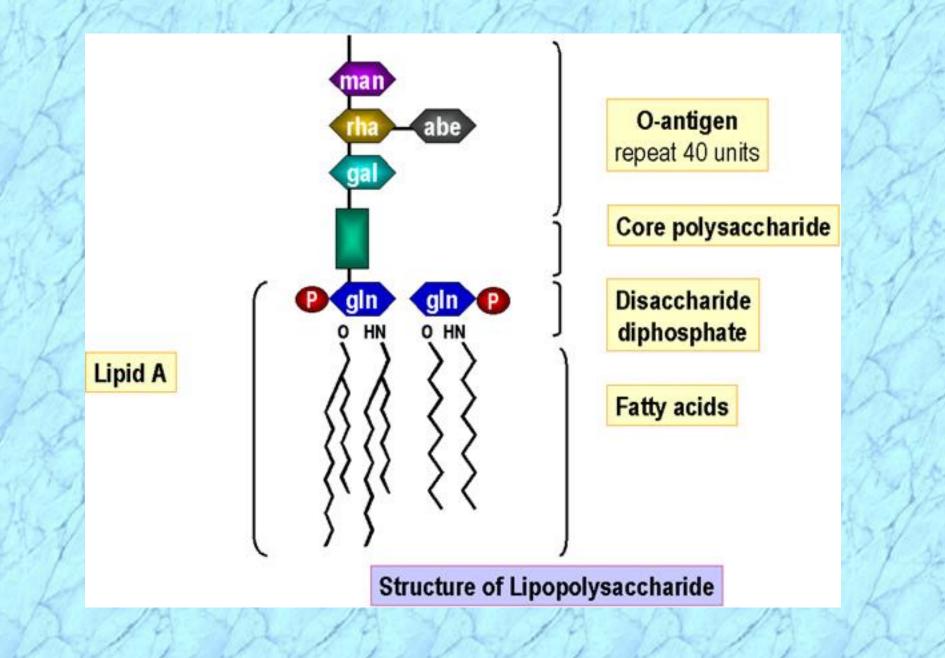
Structure of Lipopolysaccharide



Structure of Lipid A



Hydrophobic Lipid A is endotoxic component



Structure of Core Polysaccharide

l Ģlu-GlcNAc	KDO = Keto-deoxy-octulonate
Ģal	Hep = L-Glycero-D-mannoheptose
Giu-Gal	HM = β -Hydroxymyristic acid (C ₁₄) LM = Lauroxymyristic acid
Hep	MM = Myristoxymyristic acid
Hep-P-P-Eth-N	Eth•N = Ethanolamine Glu = Glucose
KDO KDO DEL	GlcNAc = N-Acetyglucosamine
KDO-KDO-P-Eth•N	Gal = Galactose

KDO is distinctive sugar moiety in core polysaccharide

Repeat Units of O Antigen Side Chain

Example: (Repeated up to 40 times) Mannose — Abequose Rhamnose Galactose

Heat stable O antigen is often used to serotype

Lipooligosaccharide (LOS) 10-100kda

e.g. Naiseria spp. Causative agent for gornorroea

Virulence factors/Mechanism of Disease

Consider an analogy of an INTRUDER (pathogen/microbe) breaking into a HOUSE (body) and either causing damage and or not

Pathogen- Microbe **Pathogenicity-** Ability to cause disease **Determinants of virulence** Adhesion **Evading host defense** (phagocytosis) Damage- Cause damage to the cells

Flagella

bacterial mobility & chemotaxis to colonize under mucosa

Urease

neutralize gastric acid gastric mucosal injury (by ammonia)

Lipopolysaccharides

adhere to host cells inflammation

Outer proteins

adhere to host cells

host cell

Exotoxin(s) * *

 vacuolating toxin (vacA) gastric mucosal injury

Type IV secretion system

pilli-like structure for injection of effectors

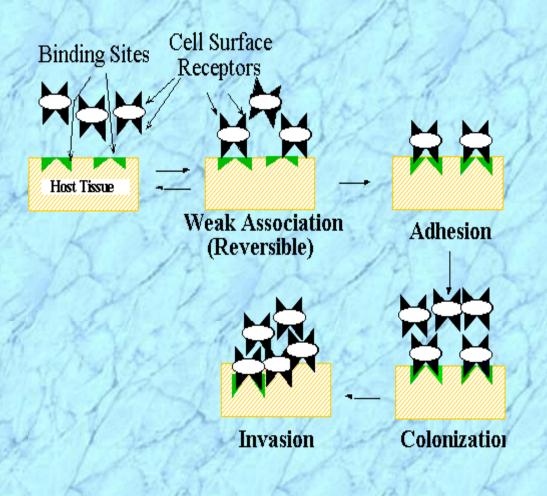
Secretory enzymes

- mucinase, protease, lipase gastric mucosal injury

Effectors (cagA e.t.c)

actin remodelling, IL-8 induction, host cell growth and apoptosis inhibition

TheMechanisms of Bacterial Pathogenecity



EXAMPLES OF SPECIFIC ATTACHMENTS OF BACTERIA TO HOST CELL OR TISSUE SURFACES

Bacterium	Adhesin	Receptor	Attachment site	Disease
Streptococcus pyogenes	Protein F	Amino terminus of fibronectin	Pharyngeal epithelium	Sore throat
Enterotoxigen ic E. coli	Type-1 fimbriae	Species- specific carbohydra te	Intestinal epithelium	Diarrhea
Bordetella pertussis	Fimbriae ("filament ous hemaggluti nin	Galactose on sulfated glycolipids	Respirator y epithelium	Whooping cough

Bacterial VF can be divided into several groups on the basis of the mechanism of virulence and function: Membrane Proteins

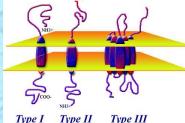
Adhesion, colonization and invasion Promote adherence to the host cell surface Responsible for resistance to antibiotics Promote intercellular communication

Polysaccharide Capsules

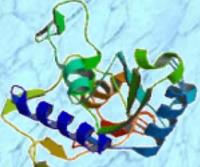
surround the bacterial cell and have anti-phagocytic properties

Secretory Proteins

can be toxins







can modify the host cell environment and are responsible for some host cell-bacteria interactions

To understand HOW pathogenic bacteria interact with their host to produce clinical disease is fundamental

Discovering Virulence Factors is the first step in understanding bacterial pathogenesis and their interactions with the host, which may also serve as a novel targets in drugs and vaccine development

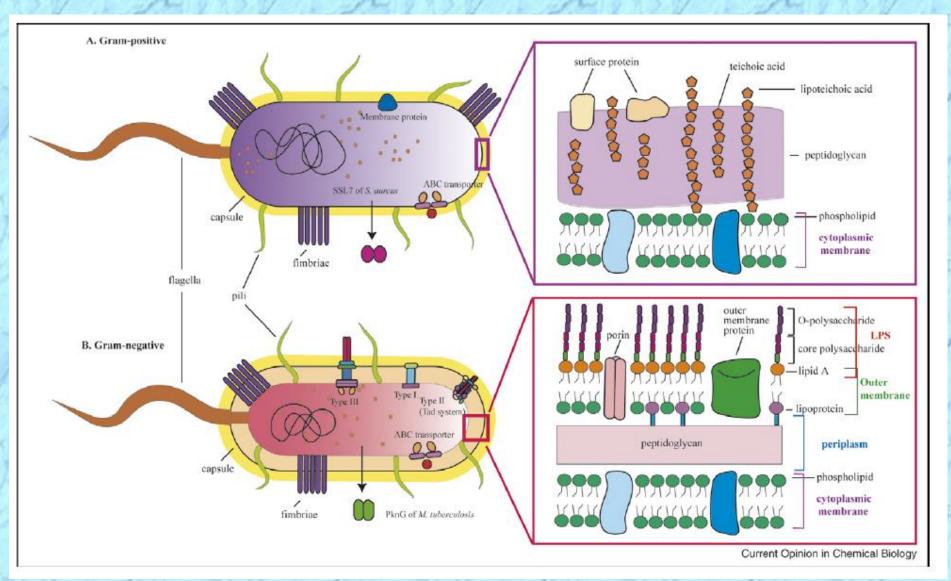
Comparative Genomics

& Transcriptomics Proteomics

Important Tools in discovering VF in bacterial pathogens



Major Virulence Factors of Pathogenic Bacteria



Evading host immune system

Avoid phagocytosis

Capsule- Most pathogens have capsule Enzyme synthesis e.g. Coagulase in S. aureus (boils, abscess, cellulitis).

Waxy cell wall

e.g. Mycobacterium- prevents phagocytosis and destruction

Calming signal

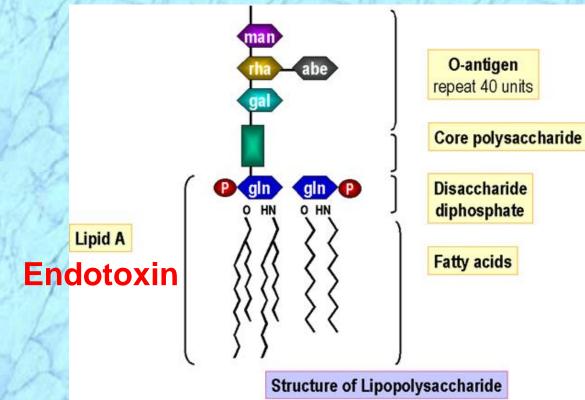
Signals the immune system not to produce Abs against the invading pathogen

Antigenic variation

Most common factors used by many pathogens

Cause Damage

Endotoxin- Found within the bacteria Exotoxin- Secreted outside of the cell



Endotoxin

- 1. Only found in G-ve bacteria
- 2. Known to stimulate cytokines- (super-antigen)
 - Cytokines storm
 - Fever
 - Body aches
 - Leaky blood vessels
 - Reduced blood pressure- Septic shock
 - Organ failure
 - DIC- Disseminated Intravascular Clotting

Superantigens

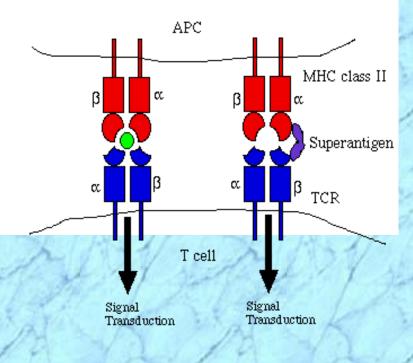
Special type of Exotoxin

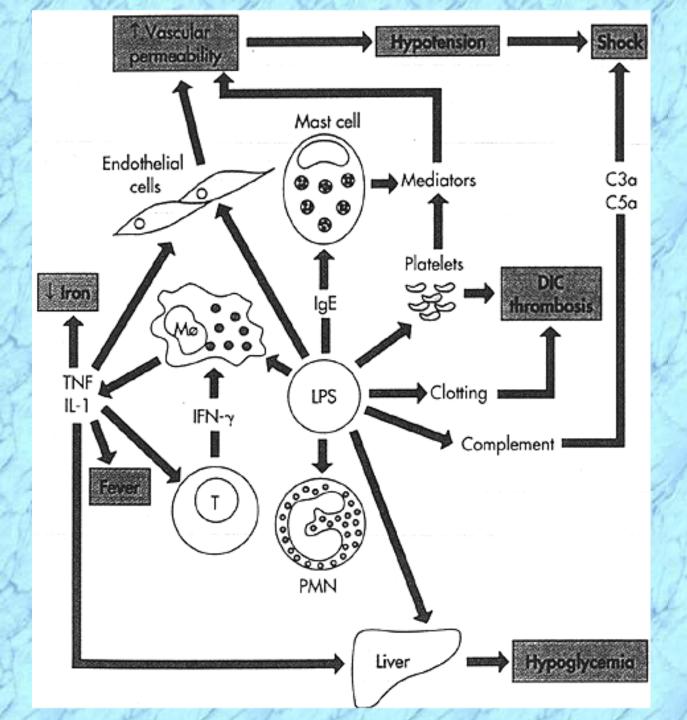
Nonspecifically stimulate T-cells.

Cause intense immune response due to release of cytokines from host cells.

Fever, nausea, vomiting,

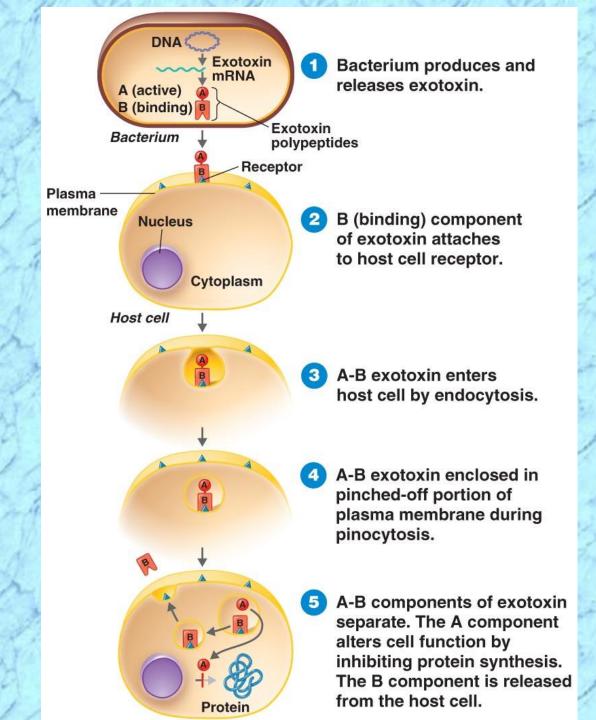
diarrhea, shock, and death.





Type of Exotoxins:

A-B Exotoxins



Exotoxin

Commonly found in G+ve bacteria Cytotoxin

- Destroy RBC and sometimes WBC
- e.g. *S. pyogenes* is beta-hemolytic **Neurotoxin**
 - Blocks the transmission of chemical across the synaptic cleft E.g. C. botulinum (Flaccid paralysis)
 - Stimulates the transmission leading to increased contraction. E.g. C. tetani (Spastic paralysis)

Enterotoxin

Promote leakage of potassium into the GIT and influx of water leading to Diarrhea and Dehydration

Streptococcus Mechanism of disease

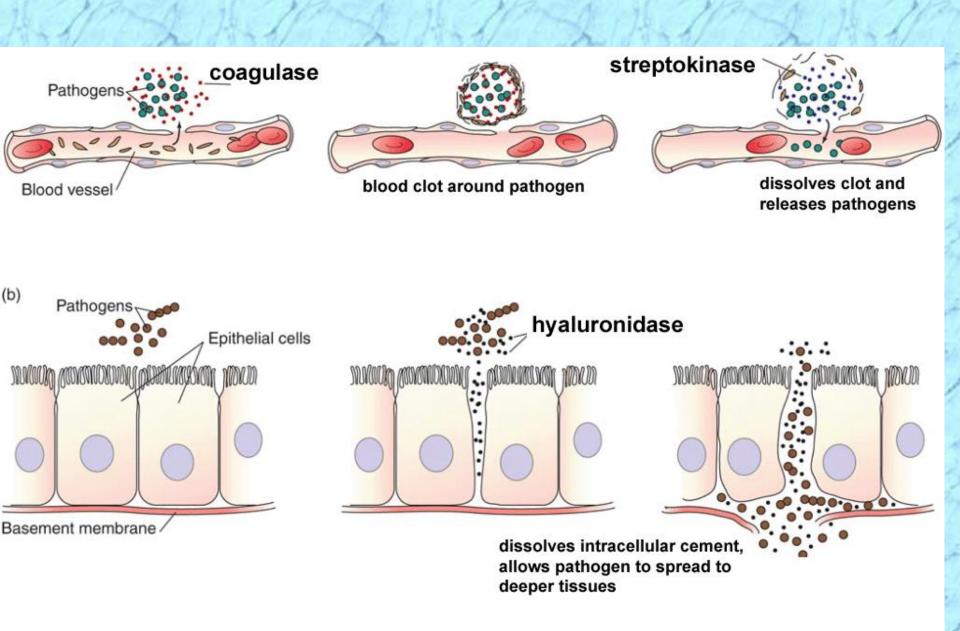
Alpha-hemolytic

S. pneumoniae- otitis media, sinusitis, pneumonia S. mutans- Caries/cavities (Plaque), Atherosclerotic plaque

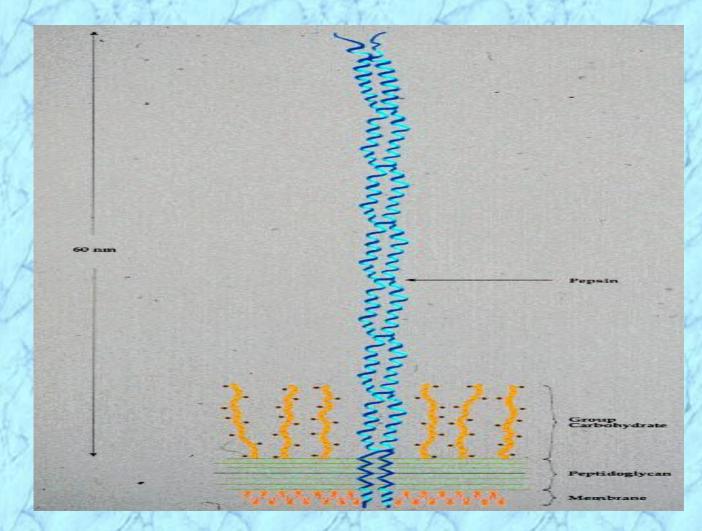
Beta- hemolytic (Group A&B) Group A- S. pyogenes Strep throat Otitis media Scarlet fever Beta- hemolytic (Group B) S. agalactiae Meningitis, Pneumoniae

S. pyogenes Mechanism of disease

Exotoxins Invasins Help bacteria move through tissues **Streptolysins** Hemolytic to RBC **Streptokinase** Digest blood clot and move through tissues **Pyogenic exotoxins** Inflammatory response **M** proteins Attacks Abs (Ig)



M PROTEIN STRUCTURE



* Apoptosis as Bacterial virulense

factor programmed cell death: activation of apoptosis to destroy cells,

• utilization of apoptosis to initiate inflammation

 The induction of apoptosis by Bordetella pertussis adenylate cyclase hemolysin

P. aeruginosa can induce apoptosis by producing exotoxin

Quorum Sensing as a virulence factor

An important factor in the pathogenesis of some infections is a bacterial cell-to-cell signaling mechanism, referred to as "quorum sensing", which enables bacteria to coordinately turn on and off specific virulence genes through the production of autoinducer molecules.

 Interference or blocking of quorum-sensing systems lead to inhibition of a variety of virulence factors.

Genomics and Bacterial Pathogenesis

- Complete knowledge of an organism's genetic makeup allows exhaustive identification for virulence genes, vaccine and antimicrobial targets, and diagnostics.
- Understanding the genomes of pathogenic bacteria is the solution.