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TROPICAL SKIN INFECTIONS

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Abstract Free Paper Presentations

Molecular Mechanism of Antibiotic Resistance in Gram Positive and Gram Negative Bacteria Infection

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Abstract

Antibiotic resistance is increasing worldwide at an accelerating pace, reducing the efficacy of therapy for many infection and causing health cost, morbidity and mortality related to infectious diseases. There are five main target sites for antibacterial action that are cell wall synthesis, protein synthesis, nucleic acid synthesis, metabolic pathway and cel membrane function. Bacteria are classified according to their cell wall as Gram positive or Gram negative. The main structural component of the cell wall is a peptidoglycan that is targeting for antibacterial action. Resistance of bacteria to antibiotic may be intrinsic or acquired. Intrinsic resistance is part of bacteria genetic make up, encoded on the chromosome. Development of resistance is the major limiting factor of antibacterial. It arises through random mutation of bacterial chromosomal gene or through acquisition of resistance genes on integrons, transposons and plasmid. Conjugation is the most common method of resistance transfer in clinically important bacteria. Conjungative plasmid, which are capable of self-transmission to other bacterial host, are common in Gram-negative enteric bacilli whereas non-conjungative plasmid are common in Gram-positive cocci.

Mutated or acquired gene confer resistance by altering the target site of the antibacterial, altering the uptake of the drug or producing drug-destroying enzymes.