## Chapter 10 Identification and Classification of Prokaryotes

## **Summary Outline**

## 10.1. Taxonomy

- A. **Taxonomy** consists of three interrelated areas:
  - 1. Identification
  - 2. Classification
  - 3. Nomenclature
- B. **Identification** of prokaryotes
  - 1 microscopic examination
  - 2. cultural characteristics
  - 3. biochemical tests
  - 4. nucleic acid
  - 5. patient's disease symptoms
- C. Taxonomic categories in a hierarchical order include species, genus, order, class, division or phylum, kingdom and domain.
- 10.2. Using phenotypic characteristics to identify prokaryotes
  - A. Microscopic morphology: Size, shape, and staining characteristics of a microorganism
    - 1. The **Gram stain** is a **differential stain** that distinguishes the Gram-positive and Gramnegative bacteria.
    - 2. Certain microorganisms have unique identifying characteristics that can be detected by using **special staining procedures**.
  - B. Metabolic differences
    - 1. The use of **selective** and **differential**.
    - 2. Most **biochemical tests** rely on a pH indicator or chemical reaction that results in a color changer when a compound is degraded.
    - 3. Identification using biochemical tests relies on the use of a dichotomous key.
  - C. Serology: The proteins and polysaccharides that make up a prokaryote are sometimes unique enough to be considered identifying markers.
  - D. Fatty acid analysis: Cellular fatty acid composition can be used as an identifying marker.
- 10.3 Using genotypic characteristics to identify prokaryotes
  - A. Nucleic acid probes to detect specific DNA sequences
  - B. Amplifying specific DNA using the polymerase chain reaction
  - C. Sequencing ribosomal RNA genes
- 10.4 Characterizing strain differences
  - A. **Biochemical typing:** A strain that has a characteristic **biochemical variation** is called a **biovar** or **biotype**.
  - B. Serological typing: A strain that differs serologically from other strains is called a serovar or serotype.
  - C. Genomic typing: Genomic differences detected by probes
    - 1. Different isolates of the same species that have different restriction fragment length polymorphisms (RFLPs) are considered different strains. RFLPs can be detected by pulsed-field gel electrophoresis and by ribotyping.
  - D. **Phage typing** based on the patterns of susceptibility to various types of **bacteriophage** can be used to demonstrate strain differences.
  - E. Antibiogram: Antibiotic susceptibility patterns can be used to distinguish strains.

## 2 Chapter 10

- 10.5 Difficulties in classifying prokaryotes: **Prokaryotes** have **few differences in size and shape** and **do not undergo sexual reproduction** so it is **difficult to determine their genetic relatedness**.
- 10.6 Numerical taxonomy relies on a battery of phenotypic characteristics and classifies bacteria based on their similarity coefficient.
- 10.7 Using genotypic characteristics to classify prokaryotes
  - A. Based on the comparison of the nucleotide sequence of the DNA of different organisms.
  - B. **DNA base composition** usually expressed as the G + C content. If the G + C content of two organisms differs by more than a few percent, then they are not closely related.
  - C. **DNA hybridization**: The extent of nucleotide sequence similarity can be determined by measuring how completely single strands of their DNA will anneal to one another.
  - D. Comparing the sequences of 16S ribosomal nucleic acid