2-Domain: Bacteria

According to Bergey's Manual of Systematic Bacteriology, true bacteria are divided into a large number of phyla. In this semester, we will discuss the most important of these phyla.

Phylum: Aquificae

Thought to be deepest (oldest) branch of *Bacteria*. Contains one class, one order, and five genera .Two best studied genera are *Aquifex* and *Hydrogenobacter*.

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Domain	Bacteria
Phylum	Aquificae
Class	Aquificae
Order	Aquificales
Family I	Aquificaeae
Genus 1	Aquifex
Genus 11	Calderobacterium
Genus 111	Hydrogenobacter
Genus IV	Thermocrinis
Family 11	Incertae sedis
Genus	Desulfurobacterium



Genus Aquifex

Are Gram-negative rod-shaped bacteria with a length of 2 to 6 μ m, have a diameter of around 0.5 μ m and are motile. They are non-sporeforming . *Aquifex* means *water-maker* in Latin, and refers to the fact that its method of respiration creates water. *Aquifex* tend to form cell aggregates composed of up to 100 individual cells. Thermophile, growth optimum 85°C - maximum 95°C. Microaerophilic & Chemolithoautotroph (uses hydrogen, thiosulfite, and sulfur as electron donor, uses oxygen as electron acceptor). Genome ~1/3 size of *E. coli*.

Phylum: Thermotogae

Second deepest branch of *Bacteria*. Contains one class, one order, and six genera (best studied genus is *Thermotoga*).

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DomainBacteriaPhylumThermotogaeClassThermotogaeOrderThermotogalesFamilyThermotogaceaeGenusThermotoga



Genus Thermotoga

Gram-negative rods outer sheath-like envelope balloons from ends of cell, called a "toga". Thermophiles (optimum 80°C; maximum 90°C). Grow in active geothermal areas (marine hydrothermal vents/terrestrial solfataric springs). Chemoheterotrophs,have functional glycolytic pathway and can grow anaerobically on carbohydrate/protein digests.

 $\sim 24\%$ of coding sequences similar to archaeal genes(may be due to horizontal gene transfer).

Phylum: Deinococcus-Thermus

Spherical or rod-shaped; in pairs or tetrads. Gram-positive (lack typical Gram-positive cell wall, layered outer membrane similar to Gramnegatives, L-ornithine in peptidoglycan & lacks teichoic acid. Aerobic, catalase positive. Resistant to desiccation and radiation. radiation resistance due to ability to repair genome when it is severely damaged efficient proteins (protected by manganese) and enzymes for DNA repair (Within 12–24 hours can repair chromosomes fragmented by exposure to radiation).

Isolated from ground meat, feces, air, fresh water, and other sources, but natural habitat unknown. Genome consists of two circular chromosomes, a megaplasmid, and a small plasmid.



Contains one class, Two orders &Three genera (genus *Deinococcus* is best studied). 9 of 11 species are mesophilic; 2 are thermophilic.

Domain	Bacteria
Phylum	Deinococcus Thermus
Class	Deinococci
Order 1	Deinococcales
Family	Deinococcaceae
Genus	Deinococcus
Order 11	Thermales
Family	Thermaceae
Genus 1	Thermus
Genus 11	Meiothermus

Class *Mollicutes* (The Mycoplasmas) : Contains five orders and six families. Smallest bacteria capable of self-reproduction. Cannot synthesize peptidoglycan precursors . Lack cell walls and are pleomorphic. Grow as fried egg appearance on agar surface.



Have genomes less than 1000 genes (one of the smallest found in prokaryotes). Most nonmotile, some have gliding motility (2 to 5 microns/second ,self surface proteins surround "neck" of cell ,attach to cytoskeletal proteins; function like microscopic legs, powered by ATP hydrolysis) see the figure below.

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(b) M. pneumoniae

Important Pathogens:

Mycoplasma mycoides	pleuropneumonia in cattle
Mycoplasma gallisepticum	chronic respiratory disease in chickens
Mycoplasma hyopneumoniae	swine pneumonia
Mycoplasma pneumoniae	primary atypical pneumonia in humans
Ureaplasma urealyticum	premature birth, neonatal meningitis and
	pneumonia
spiroplasmas	pathogenic in insects, ticks, and a variety
	of plants

Photosynthetic Bacteria:

There are **5** Phyla of Photosynthetic Bacteria:

1-Phylum *Chlorobi* – green sulfur bacteria

2-Phylum Chloroflexi – green nonsulfur bacteria

3-Phylum Cyanobacteria

4-Purple bacteria divided into 3 groups : purple sulfur (2 γ -proteobacterial families: *Chromatiaceae* and *Ectothiorhodospiraceae*) and purple non-sulfur (distributed between α - and β -proteobacteria). **5-Phylum** *Firmicutes* – heliobacteria

Phylum: Chlorobi

Green sulfur bacteria. Consists of one class, *Chlorobia*; one order, *Chlorobiales*; one family, *Chlorobiaceae*. Representative genera are *Chlorobium*, *Prosthecochloris*, and *Pelodictyon*.

Morphologically diverse. Thrive in sulfide rich areas. Have chlorosomes.

Chlorosomes: Ellipsoidal vesicles attached to plasma membrane, contain accessory photosynthetic pigments.



(a)

Lack flagella; Nonmotile. Some have gas vesicles(adjust depth of cell for light/ H_2S). Obligate anaerobic photolithoautotrophs.

Phylum: Chloroflexi

Green nonsulfur bacteria. Has both : Photosynthetic members :e.g., genus *Chloroflexus* Nonphotosynthetic members: e.g., genus *Herpetosiphon*.

Domain	Bacteria
Phylum	Chloroflexi
Class	Chloroflexi
Order 1	Chloroflexales
Family	Chloroflexaceae
Genus	Chloroflexus
Order 11	Herpetosiphonales
Family	Herpetosiphonaceae
Genus	Herpetosiphon

Genus Chloroflexus

Filamentous (with gliding motility). Thermophilic , often isolated from neutral to alkaline hot springs; grow in orange-reddish mats.



Genus Herpetosiphon

Non-photosynthetic, gliding, rod-shaped, or filamentous. Aerobic chemoorganotrophs with respiratory metabolism. Isolated from fresh water and soil habitats.



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