Deltaproteobacteria

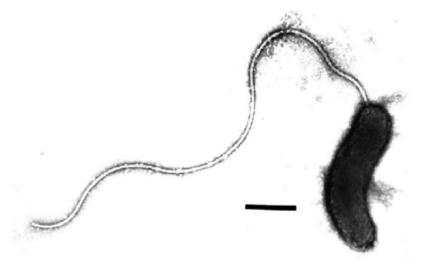
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δ -proteobacteria

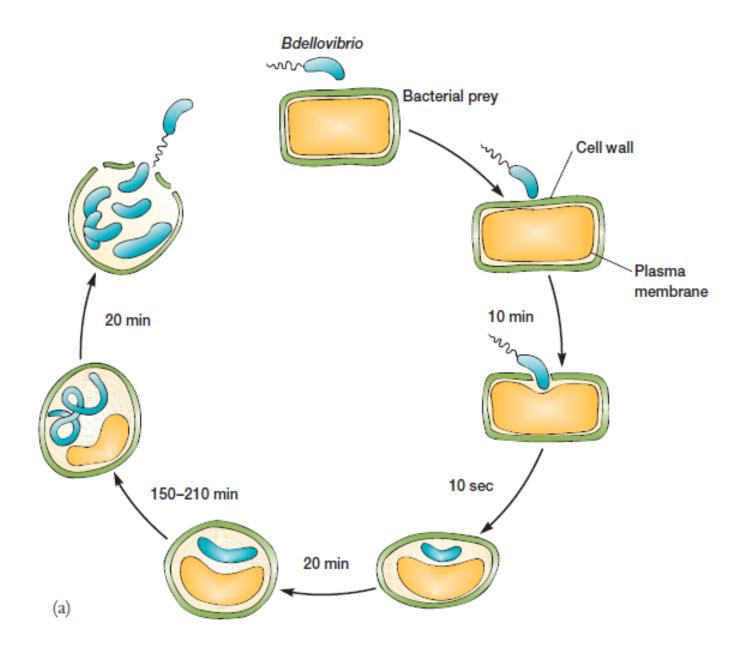
- Although the δ -proteobacteria are not a large assemblage of genera, they show considerable morphological and physiological diversity.
- These bacteria can be divided into two general groups, all of them chemoorganotrophs.
- Some genera are predators such as the bdellovibrios and myxobacteria.
- Others are anaerobes that generate sulfide from sulfate and sulfur while oxidizing organic nutrients.
- The class has seven orders and 17 families.

Order Bdellovibrionales

- The order has only the family Bdellovibrionaceae and three genera.
- The genus *Bdellovibrio* [Greek bdella, leech] contains aerobic gram-negative, curved rods with polar flagella (**figure 22.32**).
- The flagellum is unusually thick due to the presence of a sheath that is continuous with the cell wall.
- *Bdellovibrio has a distinctive* life-style: it preys on other gram-negative bacteria and alternates.



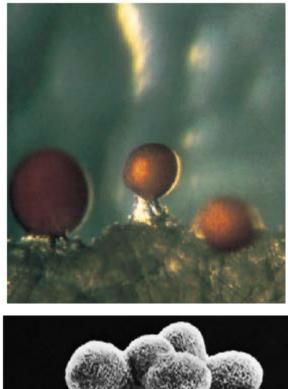
- The life cycle of Bdellovibrio is complex although it requires only 1 to 3 hours for completion.
- The free bacterium swims along very rapidly (about 100 cell lengths per second) until it collides violently with its prey.
- It attaches to the bacterial surface, begins to rotate at rates as high as 100 revolutions per second, and bores a hole through the host cell wall in 5 to 20 minutes with the aid of several hydrolytic enzymes that it releases.
- Its flagellum is lost during penetration of the cell.
- After entry, *Bdellovibrio takes control of the host cell and* grows in the space between the cell wall and plasma membrane while the host cell loses its shape and rounds up.
- The predator inhibits host DNA, RNA, and protein synthesis within minutes and disrupts the host's plasma membrane so that cytoplasmic constituents can leak out of the cell.
- The growing bacterium uses host amino acids as its carbon, nitrogen, and energy source.
- It employs fatty acids and nucleotides directly in biosynthesis, thus saving carbon and energy. The bacterium rapidly grows into a long filament under the cell wall and then divides into many smaller, flagellated progeny, which escape upon host cell lysis.

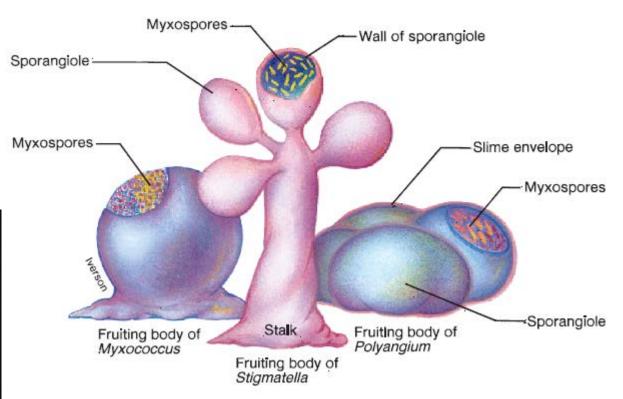


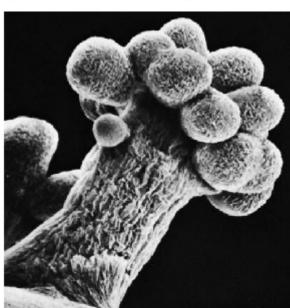
Order Myxococcales

- The myxobacteria are gram-negative, aerobic soil bacteria characterized by gliding motility, a complex life cycle with the production of fruiting bodies, and the formation of dormant myxospores.
- In addition, their G C content is around 67 to 71%, significantly higher than that of most gliding bacteria.
- Myxobacterial cells are rods, about 0.6 to 0.9 by 3 to 8 μm long, and may be either slender with tapered ends or stout with rounded, blunt ends.
- The order *Myxobacteriales is divided into* four families based on the shape of vegetative cells, myxospores, and sporangia.

- Most myxobacteria are micropredators or scavengers.
- They secrete an array of digestive enzymes that lyse bacteria and yeasts.
- Many myxobacteria also secrete antibiotics, which may kill their prey.
- The digestion products, primarily small peptides, are absorbed.
- Most myxobacteria use amino acids as their major source of carbon, nitrogen, and energy.
- All are chemoheterotrophs with respiratory metabolism.
- In the presence of a food supply, myxobacteria migrate along a solid surface, feeding and leaving slime trails.
- When their nutrient supply is exhausted, the myxobacteria aggregate and differentiate into a **fruiting body.**
- Some cells develop into dormant **myxospores that frequently are** enclosed in walled structures called sporangioles or sporangia.
- Each species forms a characteristic fruiting body.







- Myxospores are not only dormant but desiccation-resistant, and they may survive up to 10 years under adverse conditions.
- They enable myxobacteria to survive long periods of dryness and nutrient deprivation.
- The use of fruiting bodies provides further protection for the myxospores and assists in their dispersal.
- Myxobacteria are found in soils worldwide.
- They are most commonly isolated from neutral soils or decaying plant material such as leaves and tree bark, and from animal dung.
- Although they grow in habitats as diverse as tropical rain forests and the arctic tundra, they are most abundant in warm areas.