

# Conditionally pathogenic enterobacteria

Among the medically important species of conditionally pathogenic enterobacteria, we include the genera *Escherichia*, *Klebsiella*, *Enterobacter*, *Serratia*, *Citrobacter*, which can be summarized under the name **coliform bacilli** and related genera *Proteus*, *Morganella*, *Providencia*. They are **facultatively anaerobic gram-negative rods**. They are **catalase-positive** and **oxidase-negative**. They reduce nitrates, ferment sugars, **do not form spores**. With the exception of *Klebsiella*, they are able to move with peritrichal flagella. *Klebsiella* are also the only representatives of this category that form shells, other genera do so only very rarely. They commonly occur in the environment and **are part of the physiological intestinal flora**. They have a complex antigenic structure, on which the pathogenicity of each serovar is based. They are classified by more than 150 different thermostable O antigens, more than 100 thermolabile K antigens, and about 50 H antigens. These bacteria are able to cause **intestinal and extraintestinal diseases**, most often in the urinary system.

## *Escherichia* spp.

The genus *Escherichia* includes a total of 8 specified species. The most medically discussed species is *Escherichia coli*. It is a common member of the physiological intestinal flora. It also occurs in small amounts as part of the physiological flora in the upper respiratory tract and genitals. It usually does not cause disease (in case it stays within its normal boundaries), only certain serovars are pathogenic. It typically causes urinary tract infections. *E. coli* are said to be the most common pathogen in this area. Regarding extraintestinal infections, it is also responsible for meningitis, pneumonia, cholecystitis, appendicitis, peritonitis, postoperative wound infections, and sepsis. It is often nosocomial in nature. Intestinal infections are caused by strains EPEC (**enteropathogenic**), ETEC (**enterotoxigenic**), EIEC (**enteroinvasive**), EHEC (**enterohemorrhagic**). Each of these named strains has a different mechanism of pathogenicity. However, everyone must be **able to adhere** to intestinal cells and some produce toxins. **EPEC** adheres tightly to the intestinal cells and causes changes in the intestinal cell, the most significant of which is the **loss of microvilli**. Sometimes it penetrates the cells. It causes severe watery diarrhea in children. It is more of a problem in developing countries. **ETECs** produce LT (thermolabile) or ST (thermostable) enterotoxins. Many serovars produce both types of enterotoxin, logically causing an even more severe course of the disease. It is a common cause of traveler's diarrhea. **EIEC** penetrates the intestinal cells, where it causes an **inflammatory lesion**. The disease mainly affects children in developing countries and tourists. What remains is **EHEC**, enterohemorrhagic also **Shiga-like toxigenic, verotoxigenic *E. coli***. It poses the greatest threat of these phyla. It can cause **hemorrhagic diarrhea**, as well as a hemolytic uraemic syndrome (HUS). Lethality depends on the patient's age and virulence of the given serovar, it reaches up to 10% in *E. coli* type O157:H-, of which the strongest association with HUS has the type O157:H7. It is also dangerous in that it spreads easily. This is evidenced by the case of the EHEC O104:H4 epidemic of 2011 in Northern Germany. At that time, more than 3,800 cases were recorded, 54 people died.



*E. coli* on Endo medium/blood agar

## *Klebsiella* spp.

The genus *Klebsiella* is now defined by ten species. It differs from other enterobacteria in that it regularly forms **enclosures** and is **immobile**. The most important pathogen of this genus is undoubtedly *Klebsiella pneumoniae*. It lives in the intestine, in the oral cavity, in the upper respiratory tract, on the surface of the skin, and also in the surrounding environment. It causes pneumonia and urinary tract infections, especially in debilitated patients in hospitals. Here, it can also become a source of infection for newborns, causing meningitis and sepsis.

## *Enterobacter*, *Serratia*, *Citrobacter* spp.

*Enterobacter*, *Serratia*, and *Citrobacter* are more commonly occurring as infections in hospitals. They attack **bedsores** and the **respiratory and urogenital systems**. They tend to be **resistant** to common antibiotics, thus in hospitals, there is a possibility of transmitting genetic information of resistance to ATB, e.g., to *Klebsiella*. *Enterobacter* and *Serratia* are related to *Klebsiella*, *Citrobacter*, *Salmonella*. Overall, however, they have low pathogenicity. The medically important are *Enterobacter cloacae*, *Enterobacter aerogenes*, *Enterobacter sakazakii*, *Citrobacter freundii*, *Citrobacter koseri*, *Serratia marcescens*.



*Klebsiella pneumoniae* in the electron microscope

## *Proteus*, *Morganella*, *Providencia* spp.

The genera *Proteus*, *Morganella*, *Providencia* have a lot of identical properties. They are characterized by a typical **urea odor** on the cultivation soils. They are a common part of the intestine and can be found in the surroundings. Infections are usually caused endogenously. Their pathogenicity is low. In healthy individuals, they are able to

cause at most infections of the urinary system, in the **weak**, on the other hand, it causes **pneumonia, meningitis** or **sepsis**. Species that cause these diseases include *Proteus mirabilis*, *Proteus vulgaris*, *Morganella morganii*, *Providencia stuartii*, *Providencia rettgeri*.

## Links

### Related articles

- Gram stain
- Bacteria

### References

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