## Lecture Series & Workshops 2019-2021

From Single Organisms to Systems Ecology and Evolution

## Brucella abortus, a polarized bacterial pathogen with a peculiar envelope

## **Prof. Xavier DE BOLLE**

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- 7<sup>th</sup> of October 2020, 12 PM
- Link to Webex Event

For individual meetings with Prof. De Bolle, please **register** by mail to <u>secretary@microbiology.lu</u> so that we can initiate the contact.

Brucella abortus is a pathogen causing bovine brucellosis, a worldwide zoonosis. These bacteria belong to the alpha-Proteobacteria class, like the model bacterium Caulobacter crescentus. Although they have very different lifestyles and morphologies, Brucella and Caulobacter share a well conserved cell cycle control pathway, modulating the CtrA two-component regulator. In B. abortus, CtrA controls cell division, which is consistent with the role of the orthologues in other alpha-Proteobacteria. Interestingly, CtrA also binds many promoters of genes producing proteins involved in envelope biogenesis. Within the alpha-Proteobacteria class, the Rhizobiales order display several examples of unipolar growth. In B. abortus, we showed that outer membrane proteins and lipopolysaccharide are inserted at a unique pole and at the division site, the positions at which immature peptidoglycan is also localized. The outer membrane of B. abortus was found to be static and heterogeneous. Recently, we found that many integral outer membrane proteins are covalently bound to peptidoglycan. Altogether, these results reveal that B. abortus cells are highly organized at the molecular level.

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