

Master's degree in Biology - Chemistry-Biology Department

Master 2 internship project Year 2025-2026

Laboratory/Institute: IBS and CHU Grenoble **Director:** W. Weissenhorn **Team:** PB&RC (bacterial pathogenesis & cellular responses) **Head of the team:** I. Attrée

Name and status of the scientist in charge of the project: Yvan CASPAR and Sylvie ELSEN

HDR: yes ⊠ (S. Elsen) no ⊠ (Y. Caspar)

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Program of the Master's degree in Biology:
 ☑ Microbiology, Infectious Diseases and Immunology ☐ Physiology, Epigenetics, Differentiation, Cancer ☐ Neurosciences and Neurobiology
<u>Title of the project</u> : Contribution of cell surface polysaccharides to the resistance of Klebsiella pneumoniae to human innate immunity
Objectives:

Evaluate the contribution of the capsule, LPS and Enterobacteral Common Antigen polysaccharides to the resistance of *Klebsiella pneumoniae* to human innate immunity (impact on bacterial lysis by the complement system and on opsonophagocytosis).

Abstract:

Klebsiella pneumoniae (Kp) is becoming highly multi-drug resistant to several antibiotic classes and is thus considered a WHO priority pathogen. At early infection stages, primary human defense is mediated by innate immunity, mainly through the complement system in blood and opsonophagocytosis in tissues. We carried out Tn-Seq experiments on eight different clinical strains of Kp to decipher their mechanisms of plasma resistance and confirmed cell surface polysaccharides as main factors contributing to the susceptibility of this species to lysis by the complement system. This project will assess the individual contribution on complement resistance and opsonophagocytosis of genes of the cps locus (capsule), wec operon (ECA synthesis and first step of LPS synthesis) or LPS biosynthesis pathway in several strains of Kp. This work will provide key information about potential targets for monoclonal antibody therapies or vaccine development.

Methods:

Basic bacterial genetics (mating, transformation, cloning), phenotypic profiling, serum bactericidal assay, molecular biology, biochemistry (western blots, SDS-PAGE, Silver Stain PAGE, polysaccharides analysis), Flow cytometry, automated microscopy and image analysis (High Content Screening), Bioinformatics

Up to 3 relevant publications of the team:

- Pont S, et al. Bacterial behavior in human blood reveals complement evaders with some persister-like features. PLoS Pathog. 2020 Dec 16;16(12):e1008893. doi: 10.1371/journal.ppat.1008893. PMID: 33326490
- Janet-Maitre M, et al. Genome-wide screen in human plasma identifies multifaceted complement evasion of Pseudomonas aeruginosa. PLoS Pathog. 2023 Jan 25;19(1):e1011023. doi: 10.1371/journal.ppat.1011023. PMID: 36696456

Requested domains of expertise:

Bacteriology, molecular biology, cloning, microscopy and bioinformatics