

**Master 2 internship project  
Year 2021-2022**

**Laboratory/Institute:** TIMC (CNRS UMR 5525-Université Grenoble Alpes).

**Director:** Alexandre Moreau-Gaudry

**Team:** TRanslational microbial Evolution & Engineering (TREE).

**Head of the team:** Prof. Dominique Schneider and Prof. Bertrand Toussaint

**Name and status of the scientist in charge of the project:** Béatrice Schaack and Corinne Mercier. **HDR:** yes

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**Program of the Master's degree in Biology:**

- |   |   |
|---|---|
| <input checked="" type="checkbox"/> Immunology, Microbiology, Infectious Diseases | <input checked="" type="checkbox"/> Structural Biology of Pathogens |
| <input type="checkbox"/> Physiology, Epigenetics, Differentiation, Cancer         | <input type="checkbox"/> Neurosciences and Neurobiology             |

**Title of the project: Evolution of the function and structure of extracellular vesicles during a long-term experiment with bacteria**

**Abstract (up to 10 lines):** Many living organisms including bacteria produce extracellular vesicles involved in cell-cell communication and molecular transport. However, many of these vesicles' traits are still poorly characterized including their structure, function, regulation, and dynamics. We will use the long-term evolution experiment (LTEE) with *Escherichia coli* to study the dynamics and function of outer membrane vesicles (OMVs). The LTEE consists in 12 populations that are propagated since > 70,000 generations from a common *E. coli* ancestor in a glucose-limited environment. Our preliminary results have shown that 1/ OMVs are produced by both the ancestor and evolved clones over time; 2/ the biochemical traits of OMVs markedly differ between the ancestor and evolved clones sampled at 50,000 generations; 3/ the composition of the OMV membrane differs from that of the producing bacterial cells, suggesting a regulated secretion process. The goal is to investigate both the function of the different types of OMVs produced by the ancestor and evolved clones and their impact on bacterial evolution.

**Objectives (up to 3 lines):**

- Investigate the fusion capability of OMVs produced by the LTEE-derived strains with bacterial cells
- Investigate the type of information that is potentially exchanged between the LTEE strains via OMVs
- Investigate the properties of recipient cells upon integration of OMVs.

**Methods (up to 3 lines):**

Bacterial cultures, fluorescence assays, dynamic light scattering, high-performance thin layer chromatography, fluorescence activating cell sorting, microscopy (light, confocal, transmission electron microscopy), genome analyses, molecular cloning.

**Up to 3 relevant publications of the team:**

1/ Cell-free expression of the outer membrane protein OprF of *Pseudomonas aeruginosa* for vaccine purposes. Mayeux G, Gayet L, Liguori L, Odier M, Martin DK, Cortès S, **Schaack B**, Lenormand JL. Life Sci Alliance. 2021 May 10;4(6):e202000958.

2/ Physicochemical evidence that *Francisella* FupA and FupB proteins are porins. Siebert C, **Mercier C**, Martin DK, Renesto P, **Schaack B**. Int J Mol Sci. 2020 Jul 31;21(15):E5496.

3/ Mutator genomes decay, despite sustained fitness gains, in a long-term experiment with bacteria. Couce A, Caudwell LV, Feinauer C, Hindré T, Feugeas JP, Weigt M, Lenski RE, Schneider D, Tenaillon O. 2017. Proc Natl Acad Sci U S A. 114: E9026-E9035.

**Requested domains of expertise (up to 5 keywords):**

biochemistry, microbiology, cell biology, molecular biology, statistics