

---

# Identification and characterization of Mabs\_4780, a new determinant required for intracellular survival and pathogenicity of *Mycobacterium abscessus*

Iman Halloum<sup>\*†1</sup>, Séverine Carrère-Kremer<sup>2</sup>, Vipul Singh<sup>2</sup>, Audrey Bernut<sup>2</sup>, Georges Lutfalla<sup>3</sup>, and Laurent Kremer<sup>4</sup>

<sup>1</sup>Laboratoire de Dynamique des Interactions Membranaires Normales et Pathologiques (DIMNP) – CNRS : UMR5235 – CNRS-Université de Montpellier II 34 095 Montpellier cedex 5, France

<sup>2</sup>Laboratoire de Dynamique des Interactions Membranaires Normales et Pathologiques (DIMNP) – CNRS : UMR5235 – France

<sup>3</sup>Laboratoire de Dynamique des Interactions Membranaires Normales et Pathologiques – CNRS : UMR5235 – France

<sup>4</sup>Laboratoire de Dynamique des Interactions Membranaires Normales et Pathologiques (DIMNP) – CNRS : UMR5235 – Université Montpellier 2 Bât 24 2<sup>e</sup>étage Place Eugène Bataillon Montpellier 34095 cedex 5 France, France

## Résumé

*Mycobacterium abscessus* (Mabs) is an emerging rapid-growing mycobacteria causing severe lung infections, particularly in cystic fibrosis patients. The smooth morphotype displays surface expression of glycopeptidolipids (GPLs) whereas the rough morphotype is characterized by the loss of surface GPL. Rough variants are involved in more severe clinical forms although the underlying physiopathological mechanisms remain obscure. We have recently developed a zebrafish embryo model to decipher the pathogenesis of Mabs and the chronology of the infection process.

Herein, we evaluated the contribution of MABS\_4780 in rough Mabs virulence. A mutant was constructed in which MABS\_4780 was disrupted by a hygromycin cassette. This strain exhibited a higher susceptibility to thiacetazone, a second-line antitubercular drug, compared to the parental strain and higher sensitivity to detergents, presumably due to alterations of cell wall composition/structure. Consistent with hypothesis, solving the three-dimensional structure of the *M. smegmatis* orthologue revealed a MaoC-like structure of known dehydratases, potentially involved in cell wall lipid biosynthesis. Since *Amoeba* may represent the environmental reservoir of Mabs, we also assessed the intracellular fate of the mutant in *Acanthamoeba Castellanii*. The mutant failed to replicate intracellularly but this growth defect was not due to a general metabolic abnormality since it grew similarly to parental strain *in vitro*. In addition, unlike the R variant, the mutant strain was extremely attenuated in infected zebrafish embryos and was unable to produce abscesses within the central nervous system and to kill the embryos.

Our findings demonstrate the unanticipated role of MABS\_4780 in physiopathology of Mabs infection, emphasizing its potential as an attractive drug target.

---

\*Intervenant

†Auteur correspondant: iman.halloum@gmail.com

**Mots-Clés:** Mycobacterium abscessus, Zebrafish embryos, Mabs\_4780, pathogenesis