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Klebsiella pneumoniae: Shipping out virulence factors.

VENDREDI, 21 AOÛT 2026 à 12h30

Pavillon Charles-Eugène-Marchand, salle Hydro-Québec (1210)

Abstract:

Understanding how bacteria interact with their surrounding environment is key to understanding the infectious cycle of one health pathogens. *Klebsiella pneumoniae* is considered a quintessential one health pathogen as it impacts the environment, animals and humans. In particular, the detection of novel antimicrobial resistance genes in *K. pneumoniae* often precedes worldwide spread. The emergence and spread of multi-drug resistant and hypervirulent *K. pneumoniae* pathotypes have led to renewed interest in identifying and characterizing virulence factors and identifying potential vaccine targets. In this seminar, I will discuss the molecular assembly and function of a nanomachine used by *K. pneumoniae* to secrete proteins.

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