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First experimental insights of key transmission parameters for Vibrio aestuarianus infection of Pacific oyster, Crassostrea gigas

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Background

- Since 2012 in France, an apparent increase of mortality events was observed in adult Pacific oysters, *Crassostrea gigas*, in association with the detection of the bacteria Vibrio aestuarianus.
- No representative field data.
- Knowledge on transmission parameters of *Vibrio aestuarianus* infection could be useful to limit its spread and establishment in oyster populations.

Objective

To develop and calibrate a simulation model representing the spread of Vibrio aestuarianus within an oyster population

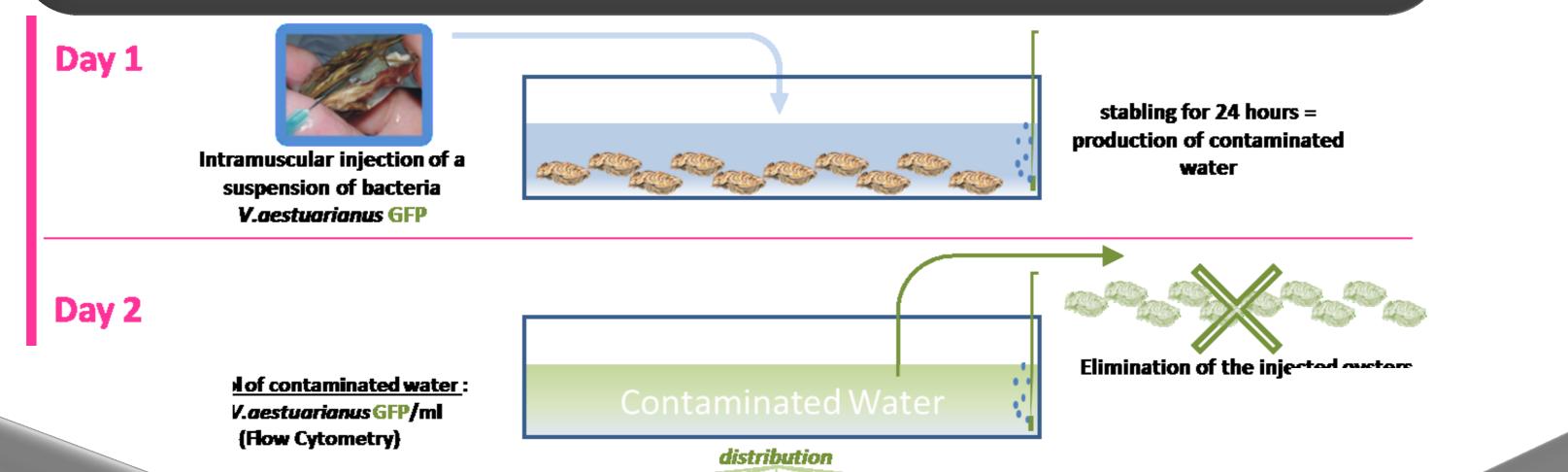
The model

We described the within-population dynamics of Vibrio aestuarianus infection by a deterministic compartmental SWEI model incorporating a water reservoir of Vibrios, which was adapted from an earlier model of Vibrio cholerae (Codeço, 2001, BMC Infectious Diseases, 1:1).

Estimation of model parameters

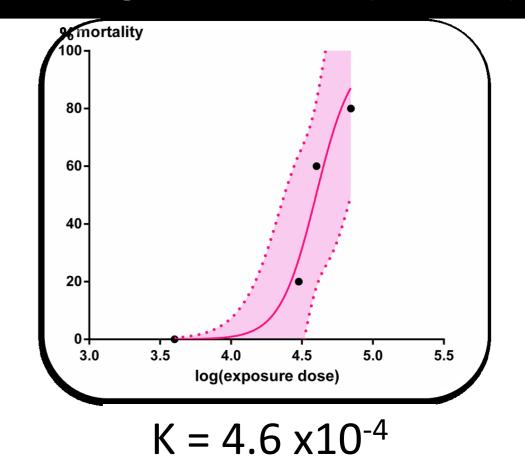
We estimated five parameters based on **experimental assays**:

- exposure rate to the contaminated seawater
- concentration of bacteria in water that yields 50% of chances of catching the infection
- infectious period
- bacteria shedding rate
- pathogen lifetime in water

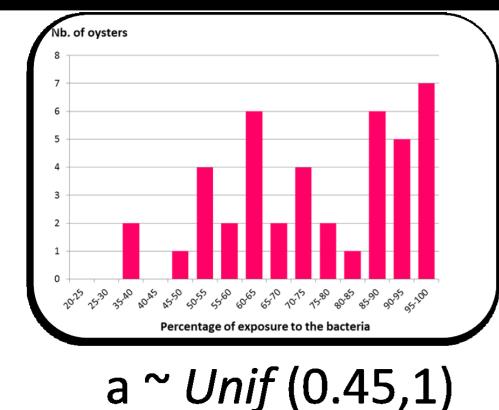


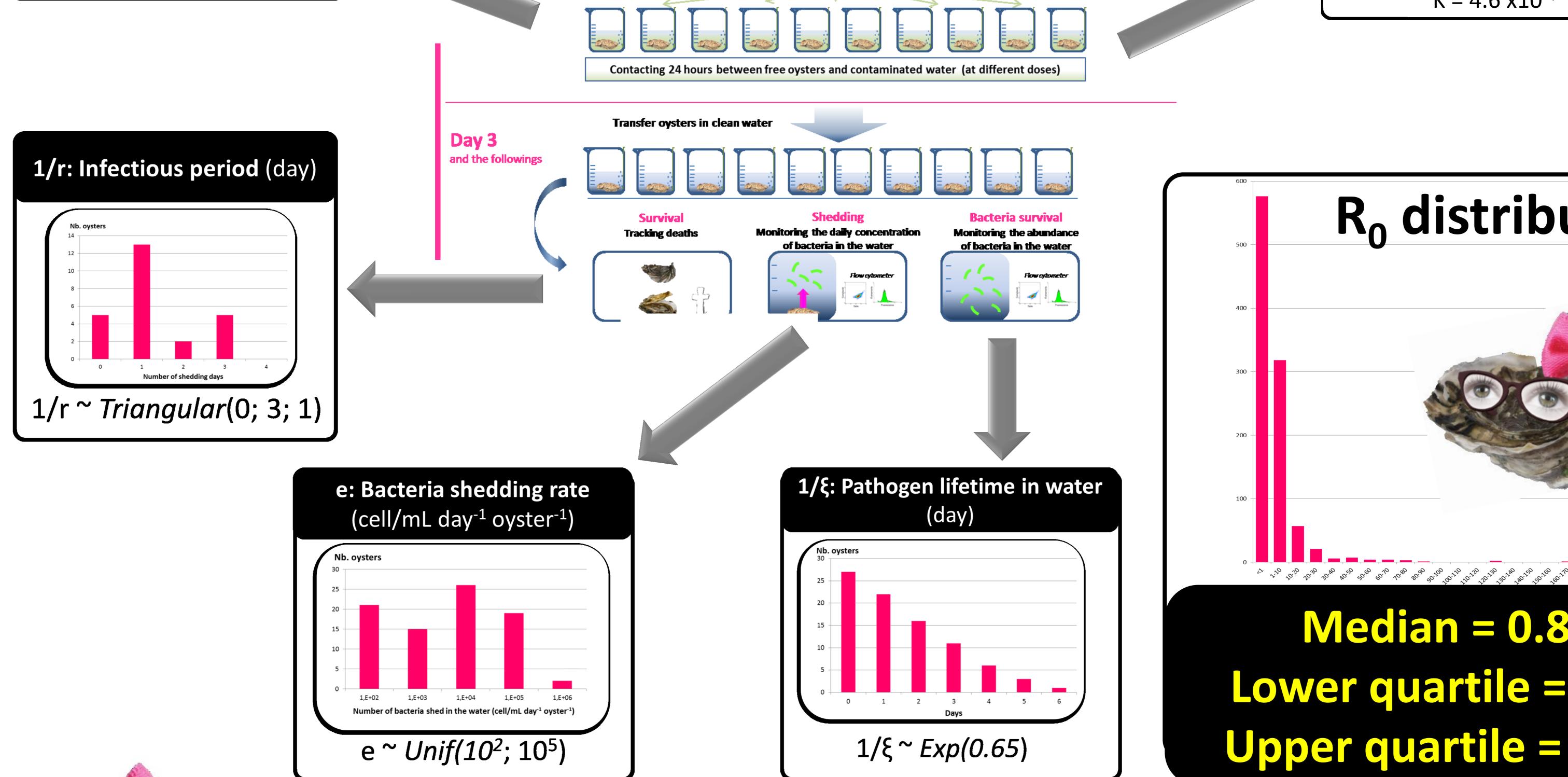


K: Concentration of bacteria in water that yields 50% of chances of catching the infection (cells/mL)

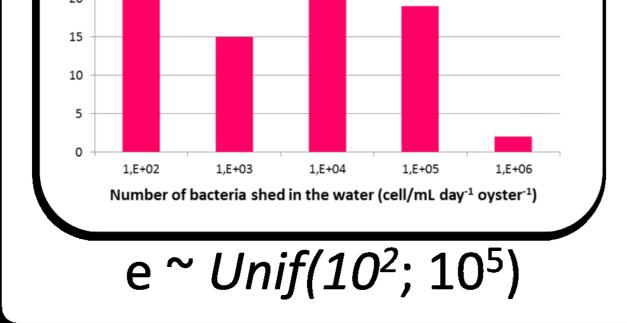


a: Exposure rate to the **contaminated seawater** (day⁻¹)





R₀ distribution



Median = 0.80Lower quartile = 0.12 **Upper quartile = 4.10**

Perspectives

Implement the influence of environmental conditions on the transmission parameters in the model Validate the model against dedicated field data

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