



Emulation of an individual-based model simulation of microbial communities

Oluwole Oyebamiji

School of Mathematics & Statistics,
Newcastle University, UK

Supervisor: **Darren Wilkinson**

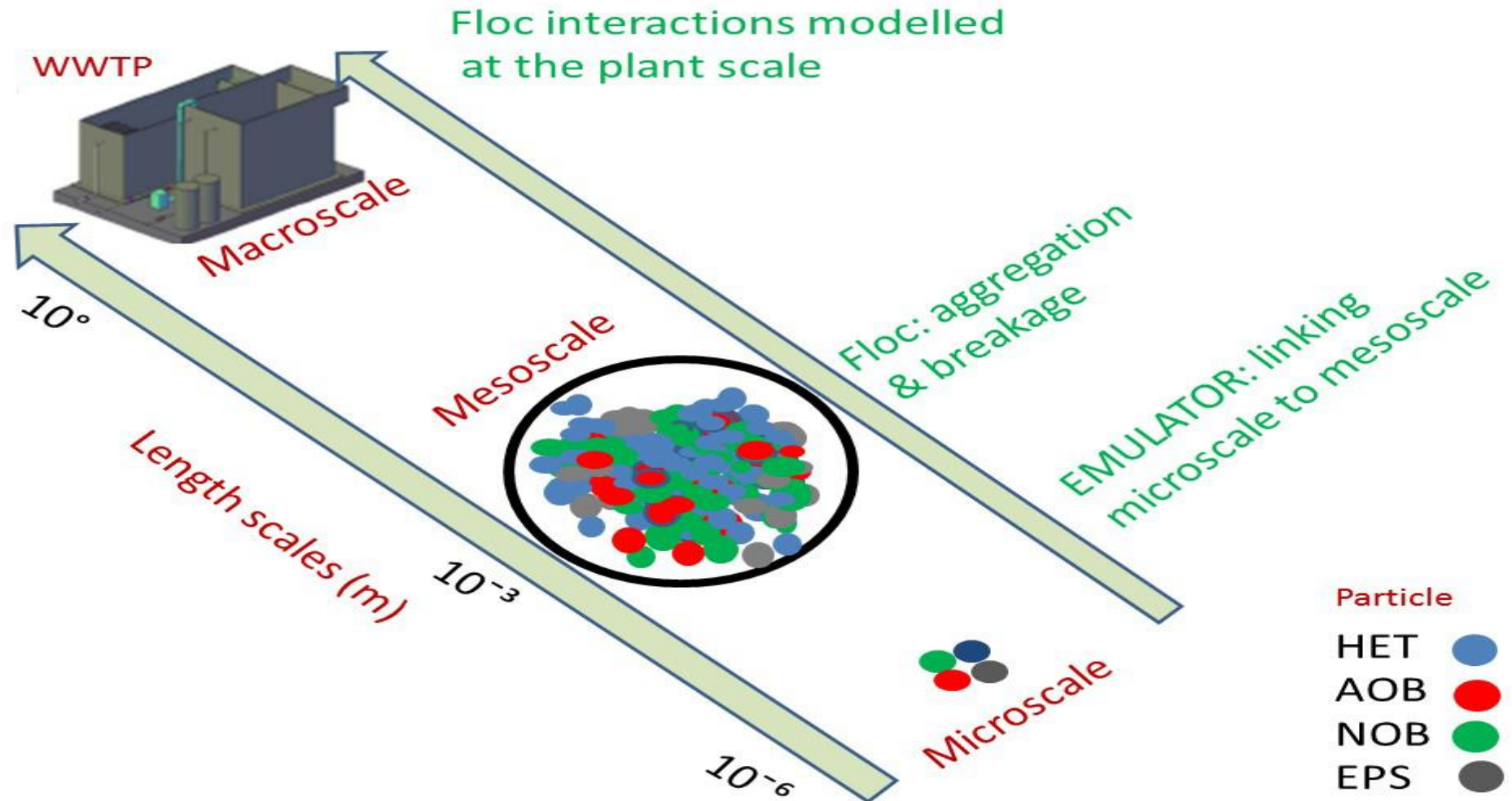
**This project was funded by the Newcastle University
Frontiers in Engineering Biology (NUFEB)**

<http://research.ncl.ac.uk/nufeb>

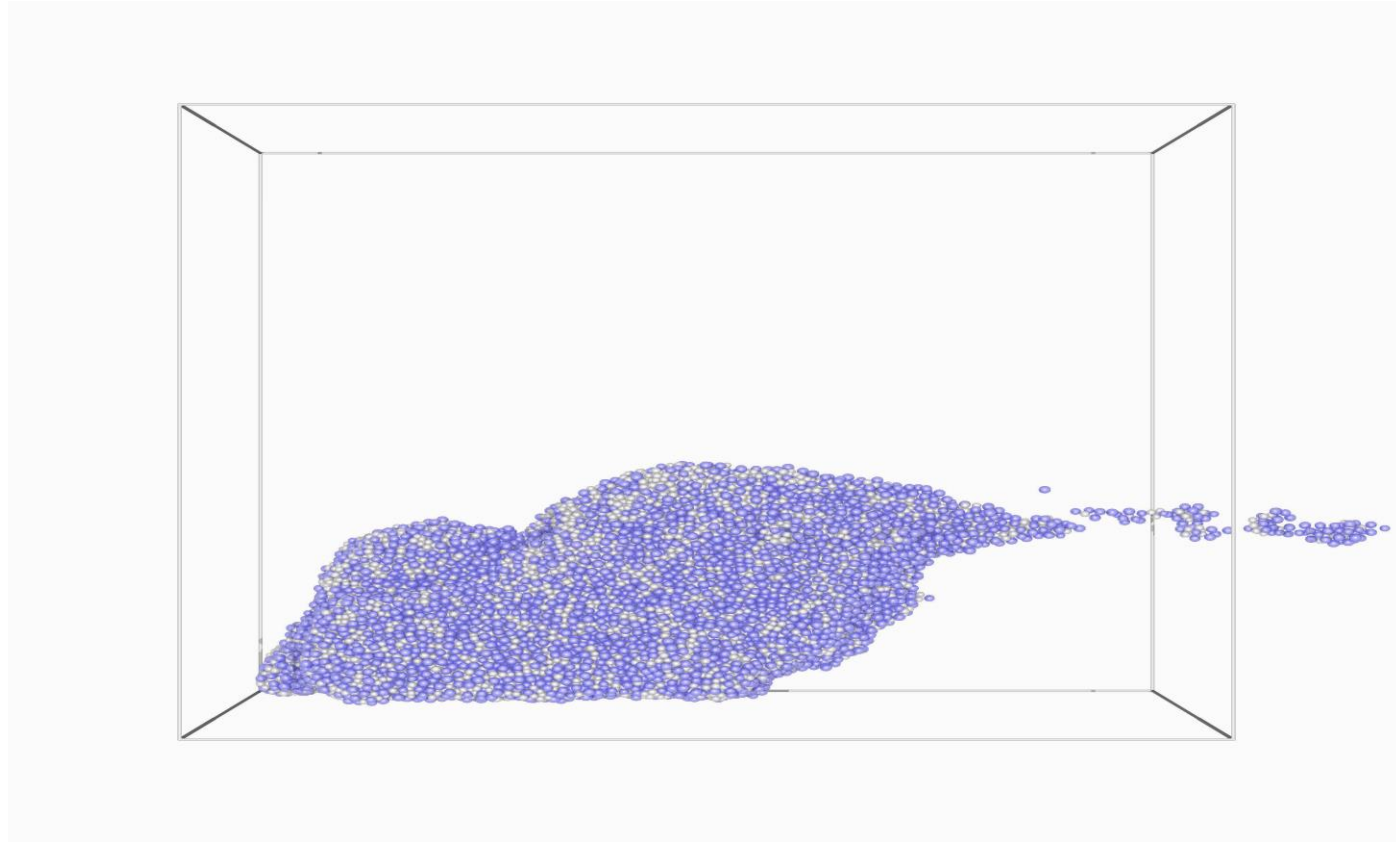
Introduction

- ❖ Wastewater treatment plants are open systems that depend on many species of bacteria
 - ❖ Methods that require many simulator runs become impractical if a single simulator run at one input value takes a long time
 - ❖ The influence of hydrodynamic shear force on biofilm deformation
 - ❖ Outputs: number of shear events, volume of detached cluster
- Inputs: biofilm height, surface roughness, mass and EPS composition
- ❖ Bayesian Poisson regression and dynamic linear models (DLMs)

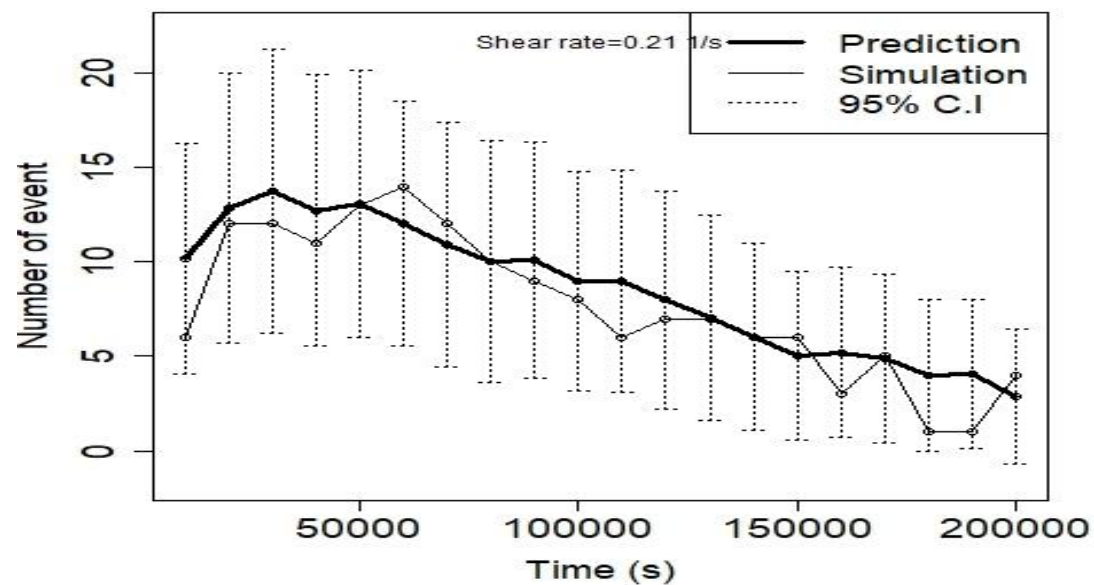
The problem



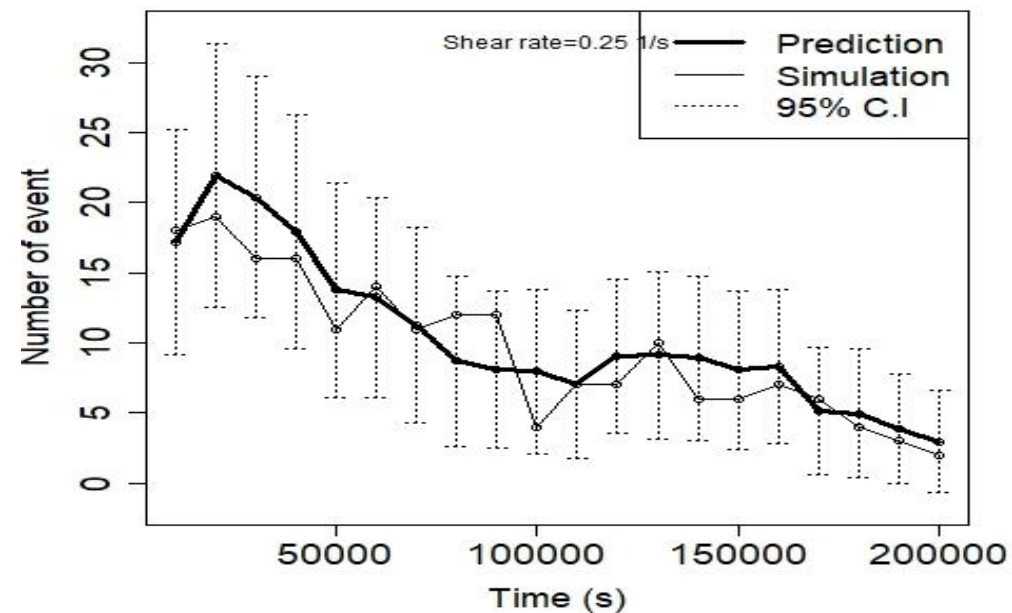
Biofilm simulation under flow fields



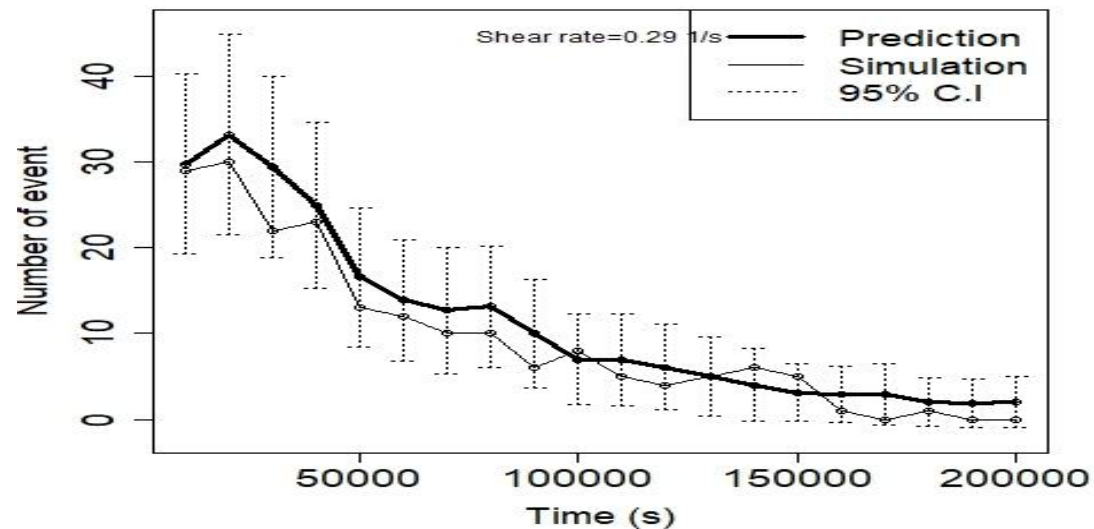
Number of shearing event per 10000s



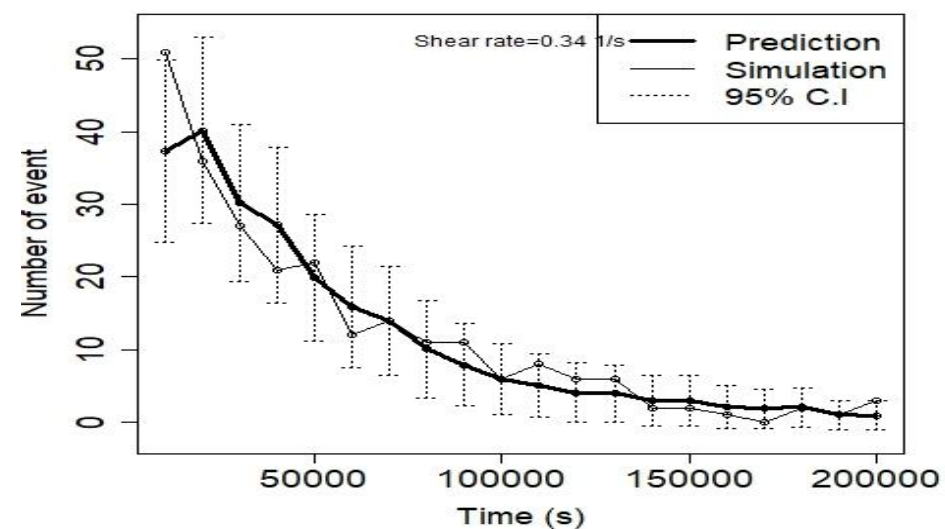
Number of shearing event per 10000s



Number of shearing event per 10000s



Number of shearing event per 10000s



Summary

- Evaluate the strategy for emulating high-level summary from the individual-based simulation of microbial communities using the NUFEB 1.1
- Measure aggregated characteristics on the microscale simulation
- Apply a dynamic Bayesian surrogate model (GP emulator) for modelling the interesting morphological characteristics that are essential for the design and performance of wastewater reactor.
- Perform Bayesian sensitivity analysis to identify the most influential input parameters.
- Quantify and model detachment patterns of a biofilm in response to hydrodynamic shear stress.

Future plans

We have made considerable progress on the parameter calibration using the recently released **NUFEB 2.0** version of the model

- ❖ Continue with the NUFEB 2.0 parameter calibration using IdynoMICS and experimental data.
- ❖ Plan to properly investigate the use of emulators as an upscaling strategy to build mesoscale models.