

Quantification of Innate Immune Function in Whole-blood Infection Assays

Ines Leonhardt^{1,2}, Teresa Lehnert^{2,3}, Kerstin Hünninger^{1,4}, Sandra Timme^{3,5}, Daniel Thomas-Rüddel^{2,6}, Marc Thilo Figge^{2,3,5} and Oliver Kurzai^{1,2,4}

¹Research Group Fungal Septemics, Leibniz Institute for Natural Product Research and Infection Biology - Hans Knöll Institute, Jena. ²Center for Sepsis Control and Care (CSCC), Jena University Hospital, Jena. ³Research Group Applied Systems Biology, Leibniz Institute for Natural Product Research and Infection Biology - Hans Knöll Institute, Jena. ⁴Institute for Hygiene and Microbiology, University of Würzburg, Würzburg. ⁵Institute for Biology and Pharmacy, Friedrich Schiller University Jena, Jena. ⁶Department of Anesthesiology and Intensive Care Medicine, Jena University Hospital, Jena.

Introduction

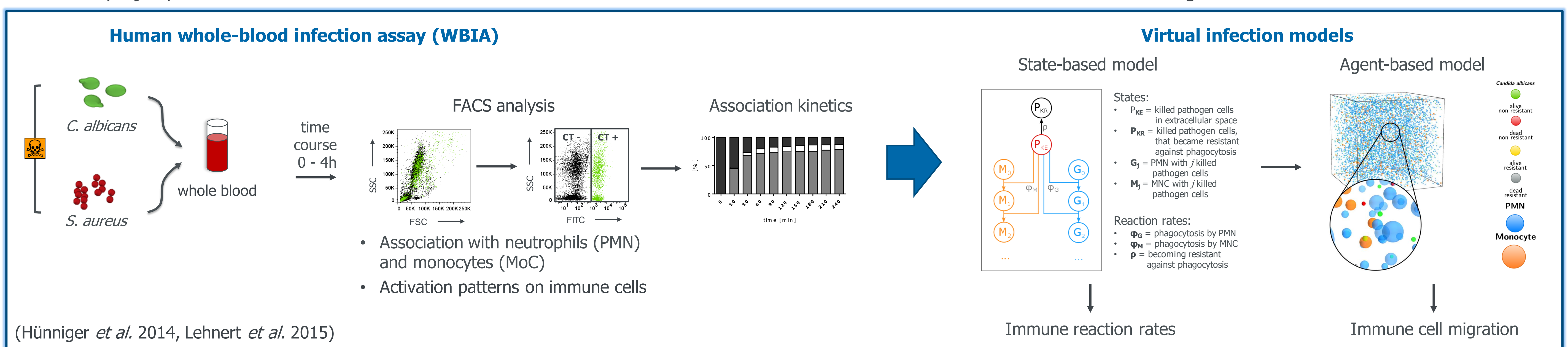
- Marked heterogeneity of sepsis as a clinical syndrome
- Caused by highly diverse pathological conditions and shows variable kinetics in individual patients
 - Classification of sepsis patients by their immune status is necessary for immunomodulatory therapy approaches

Objectives

- Individualised quantification of altering immune effector functions of septic patients
 - Are there pathogen-specific patterns of immune activation during whole blood infection?
 - Are there immune effector functions that allow stratification of sepsis patients?

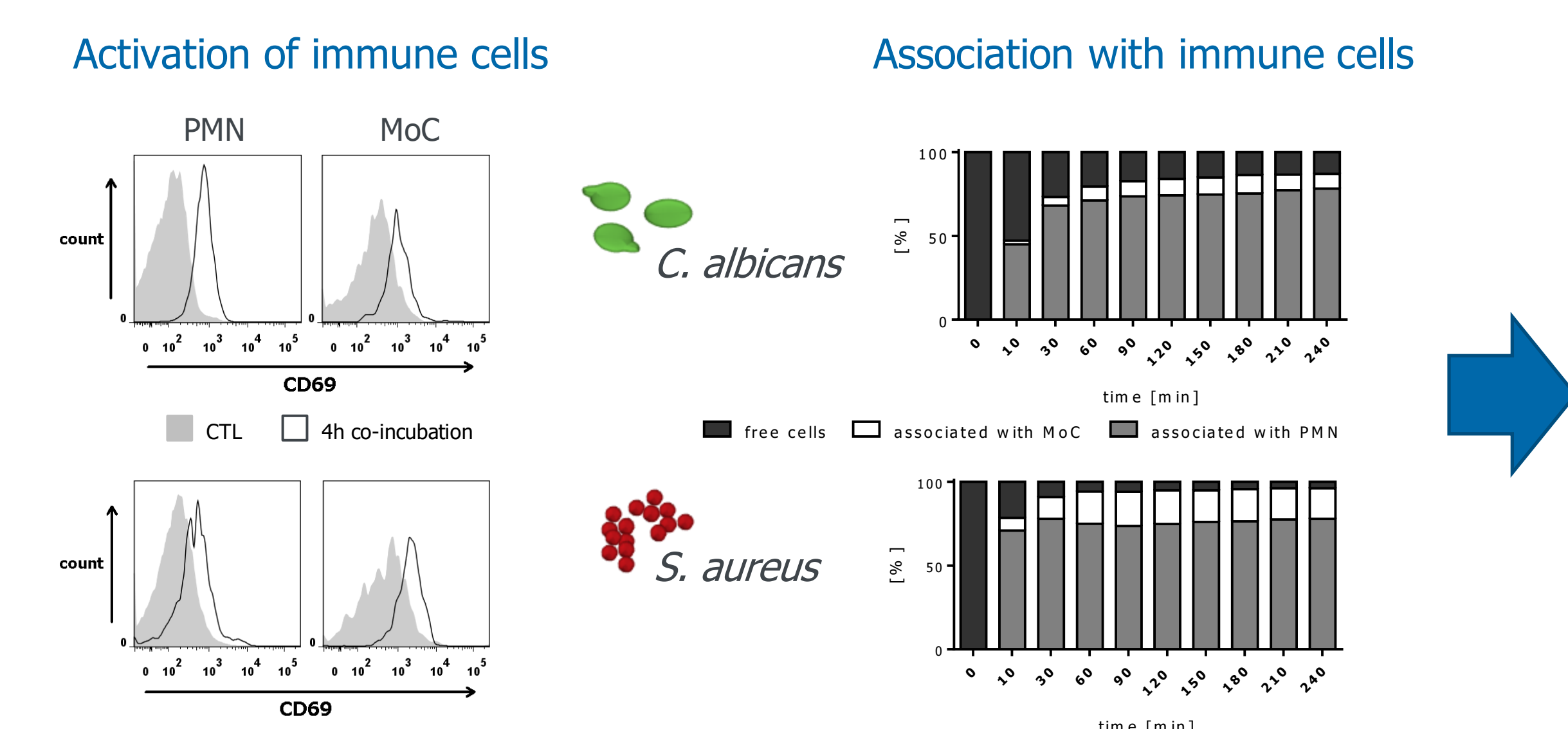
Approach

Within this project, we will use data from a human whole blood model of infection combined with advanced mathematical modeling.

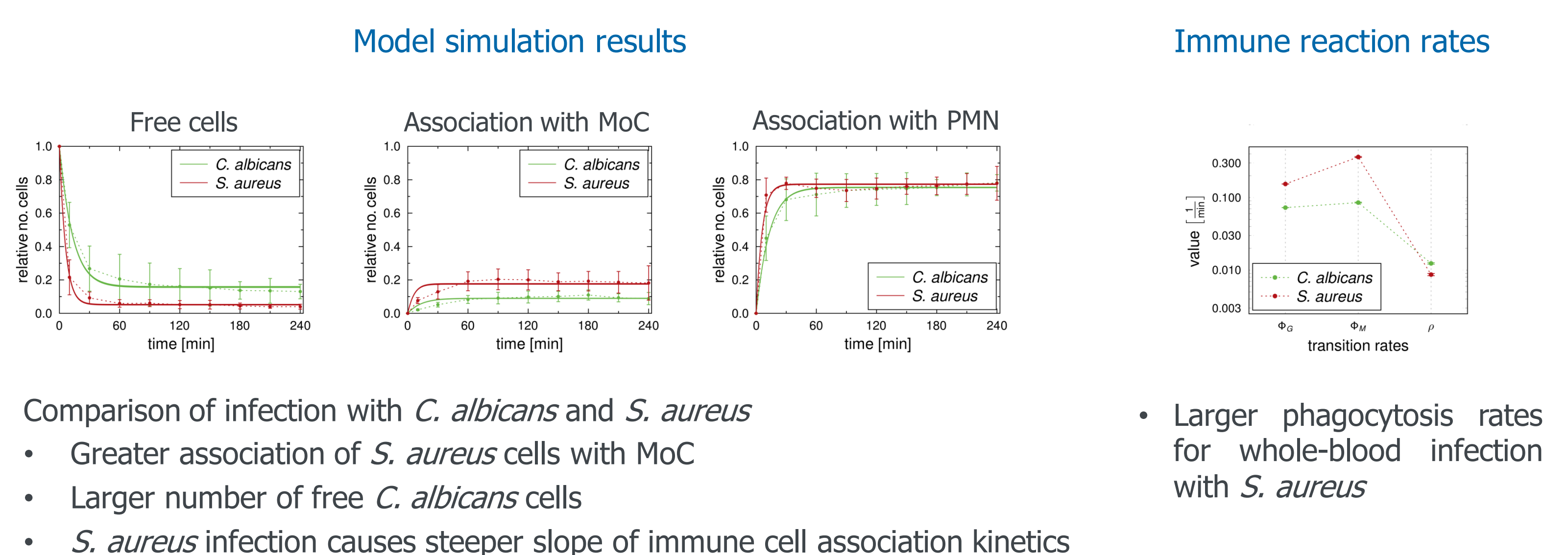


Results

Analysing pathogen association and immune activation in blood from healthy volunteers

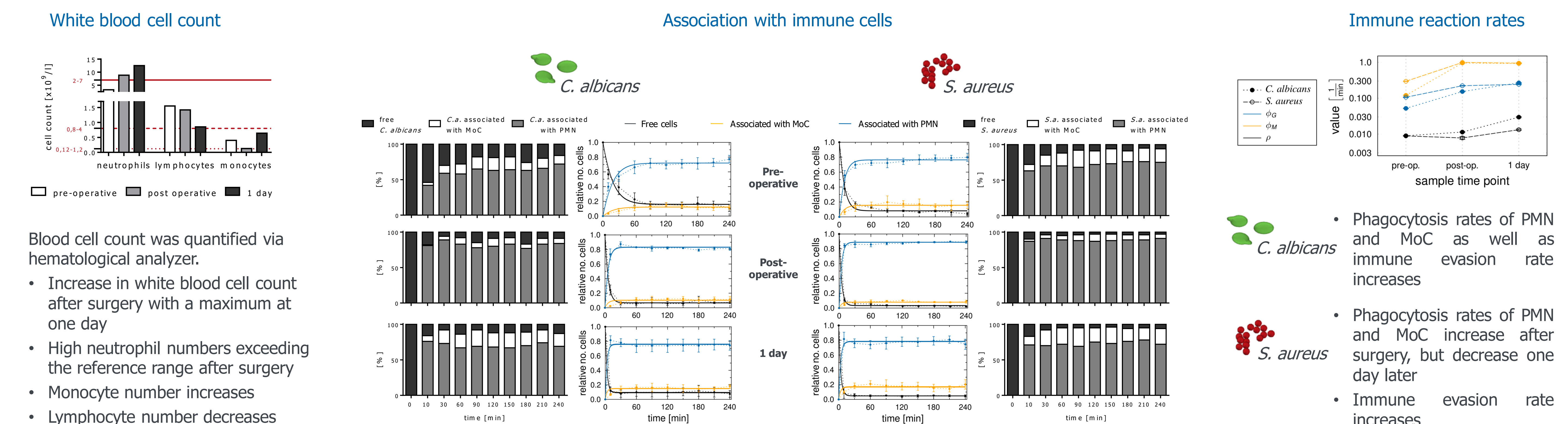


Quantification of immune effector mechanisms by biomathematical analyses



Analysing pathogen association and innate immune activation in blood from HLM patients

- Within a pilot study, blood samples of 3 patients that underwent cardiac surgery with extracorporeal circulation (heart-lung machine, HLM) were analysed
- Blood samples were obtained before cardiac surgery (pre-operative), immediately after surgery (post-operative), and 1 day after admission to ICU



Conclusions

Once optimized, analyses of blood samples from sepsis patients and patients who have survived severe sepsis will follow. This will allow identifying patterns of the dysregulated immune homeostasis providing functional classifiers for the differentiation of sepsis patients, thereby forming a basis for future patient stratification.