

# The role of neutrophil-derived EVs in promoting *Candida albicans* escape from phagocytosis

Anastasia Solomatina<sup>1</sup>, Kerstin Hünigler-Ast<sup>2,3</sup>, Jennifer Patitz<sup>3</sup>, Natalie Nieuwenhuizen<sup>3</sup>, Yann Bachelot<sup>1,4</sup>, Carl-Magnus Svensson<sup>1</sup>, Ann-Kathrin Zimmermann<sup>6</sup>, Matthew Blango<sup>5</sup>, Thomas Krüger<sup>6</sup>, Olaf Kniemeyer<sup>6</sup>, Axel A. Brakhage<sup>4,6</sup>, Oliver Kurzai<sup>2,3</sup>, Marc Thilo Figge<sup>1,4</sup>

<sup>1</sup> Applied Systems Biology, Leibniz Institute for Natural Product Research and Infection Biology, Hans Knöll Institute (HKI), Jena, Germany

<sup>2</sup> Fungal Septomics, Leibniz Institute for Natural Product Research and Infection Biology, Hans Knöll Institute (HKI), Jena, Germany

<sup>3</sup> Institute for Hygiene and Microbiology, University of Würzburg, Würzburg, Germany

<sup>4</sup> Institute of Microbiology, Faculty of Biological Sciences, Friedrich Schiller University, Jena, Germany

<sup>5</sup> RNA Biology of Fungal Infections, Leibniz Institute for Natural Product Research and Infection Biology, Hans Knöll Institute (HKI), Jena, Germany

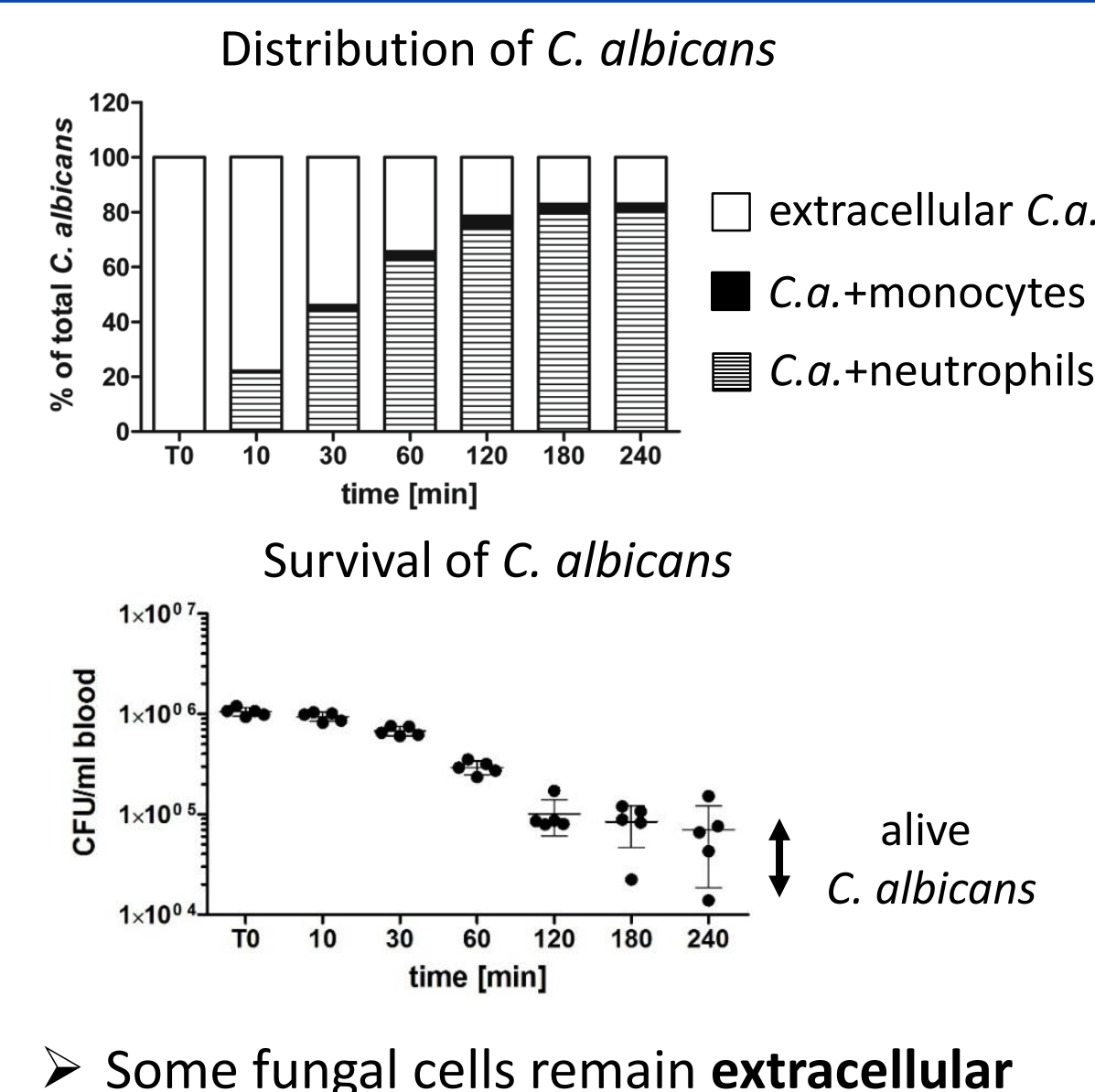
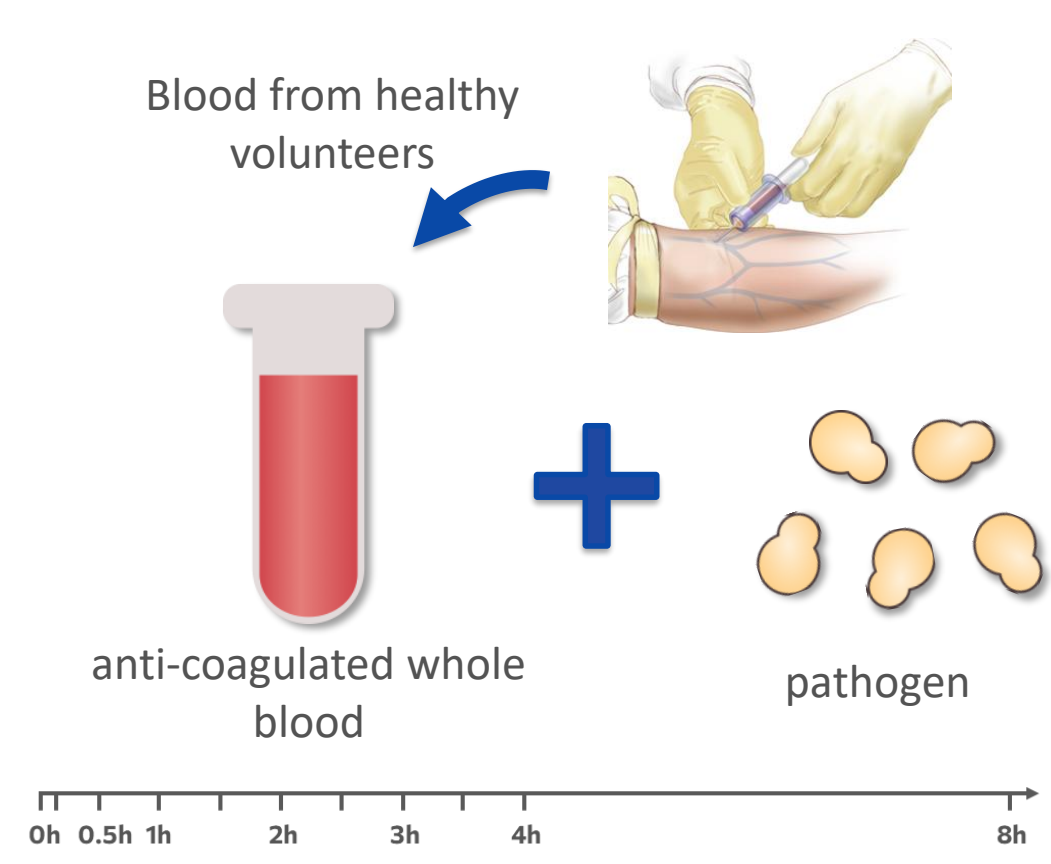
<sup>6</sup> Molecular and Applied Microbiology, Leibniz Institute for Natural Product Research and Infection Biology, Hans Knöll Institute (HKI), Jena, Germany

## Abstract

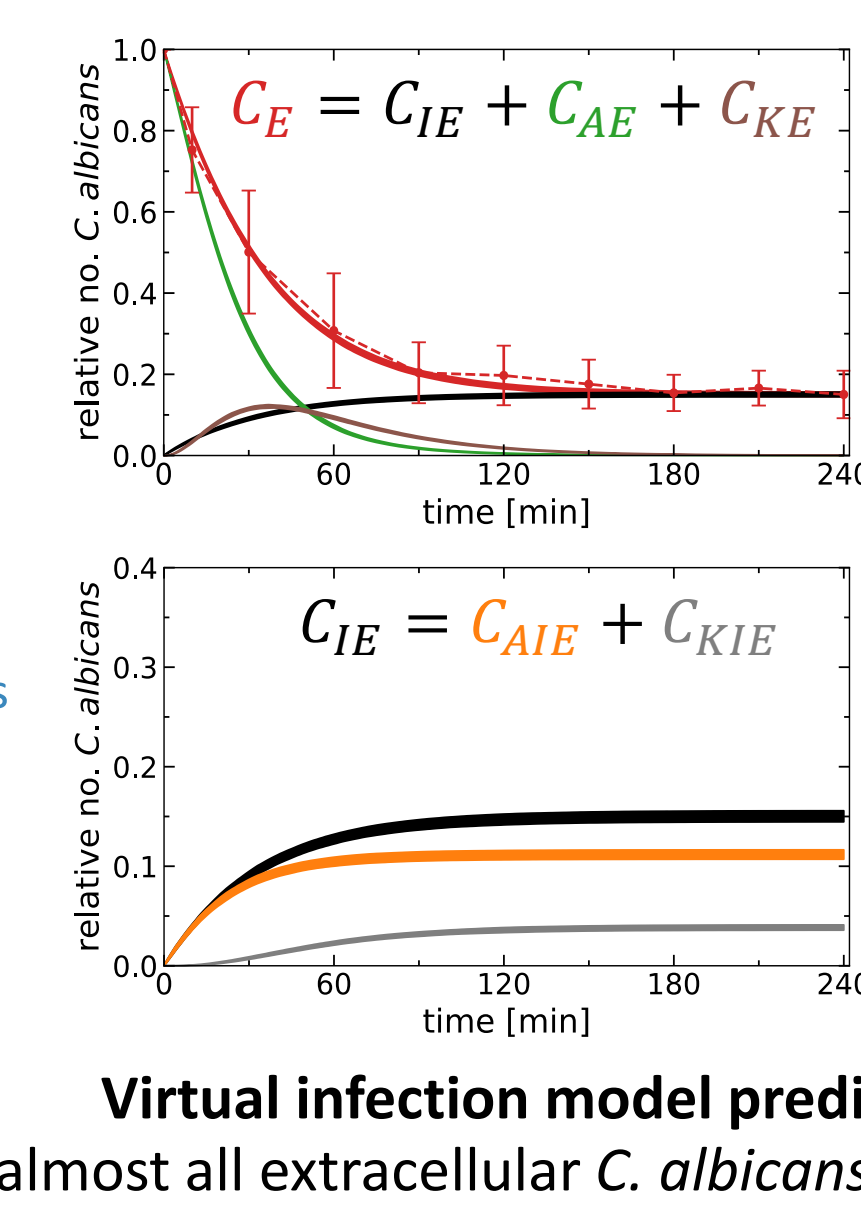
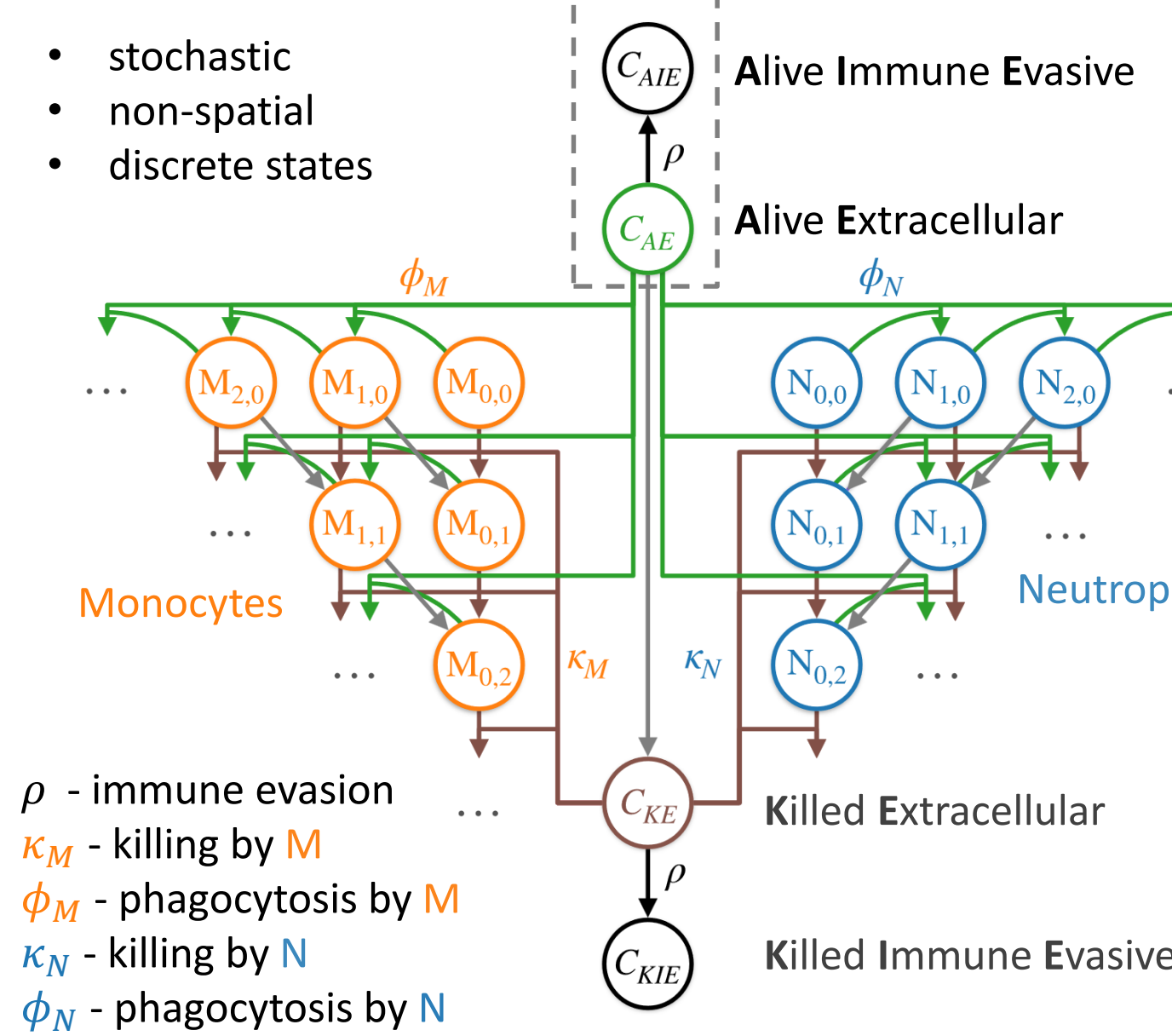
- Opportunistic fungus *Candida albicans* is one of the leading causes of **bloodstream infections**, especially in immunocompromised patients
- A substantial proportion of fungal cells remain **extracellular** in the human whole-blood infection assay, indicating effective **immune evasion strategies**
- Extracellular *C. albicans* cells acquire **host-cell molecules** through interactions with **neutrophil-derived extracellular vesicles (EVs)**
- Mathematical modeling** is used to **unravel the mechanism** how EV-decoration of the pathogen surface modulates its interaction dynamics with the host
- EV-decorated *C. albicans*** are **phagocytosed about 60% less efficiently** than cells lacking host-derived surface features
- These results uncover a previously **unrecognized role of neutrophil-derived EVs in promoting immune evasion** of *C. albicans* by modulating its susceptibility to phagocytosis

## Human whole-blood infection assay

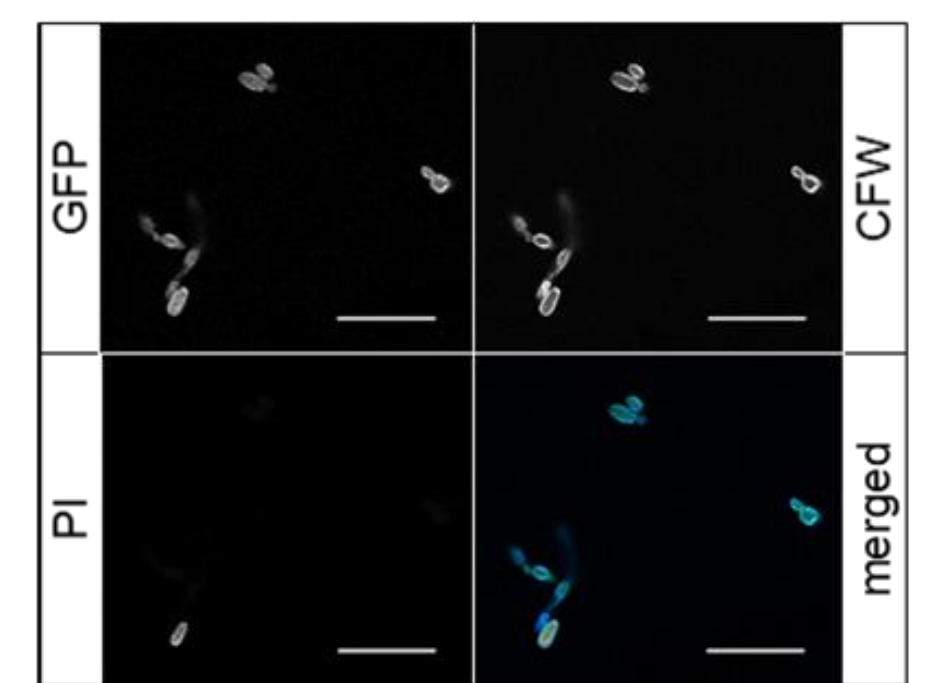
- mimics the *in vivo* scenario of the innate immune response<sup>[1]</sup>
- allows monitoring **host-pathogen interactions**
- many variables are accessible to **direct experimental quantification**



## State-based model<sup>[1,2]</sup>

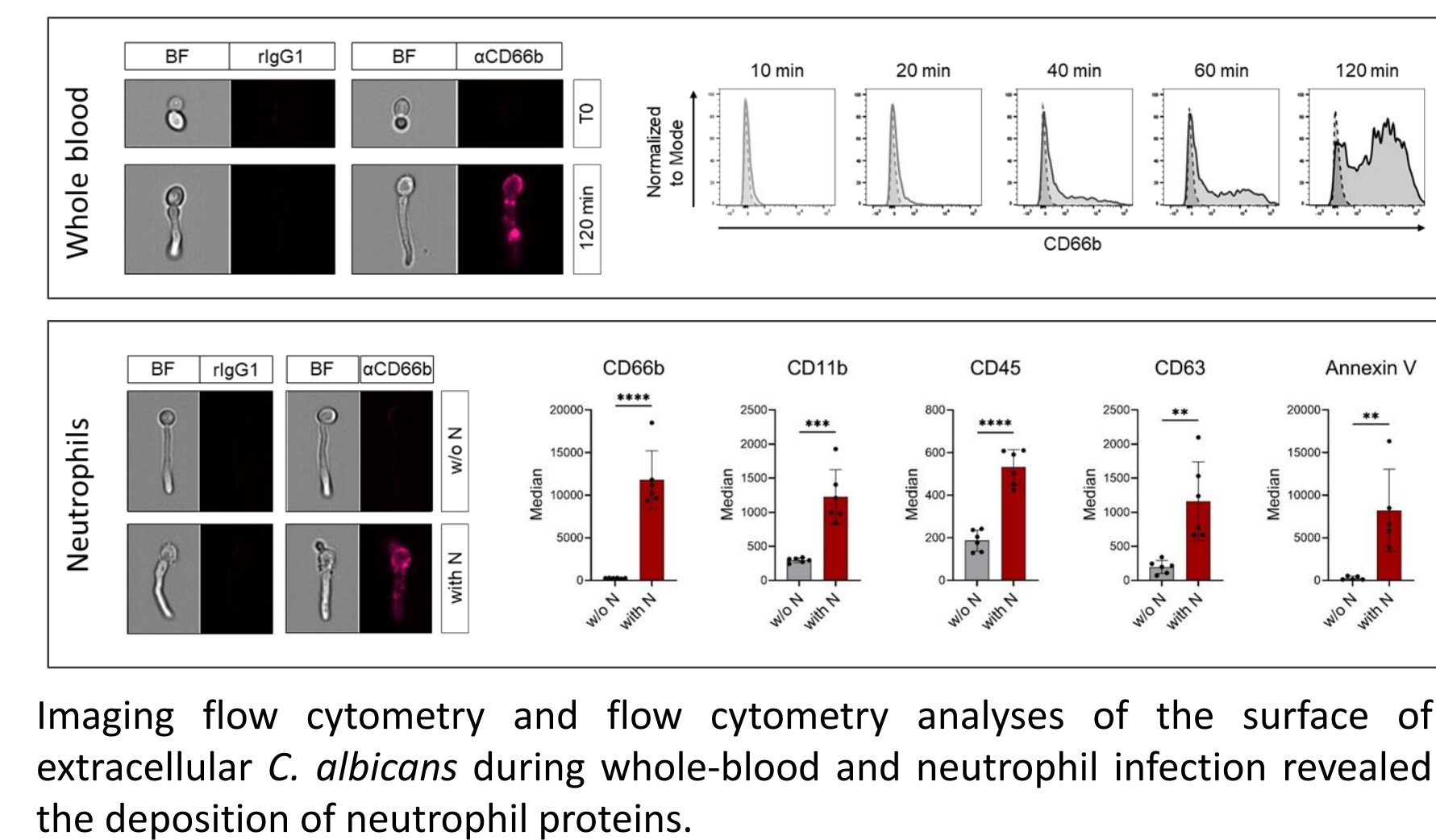


### Experimental validation:



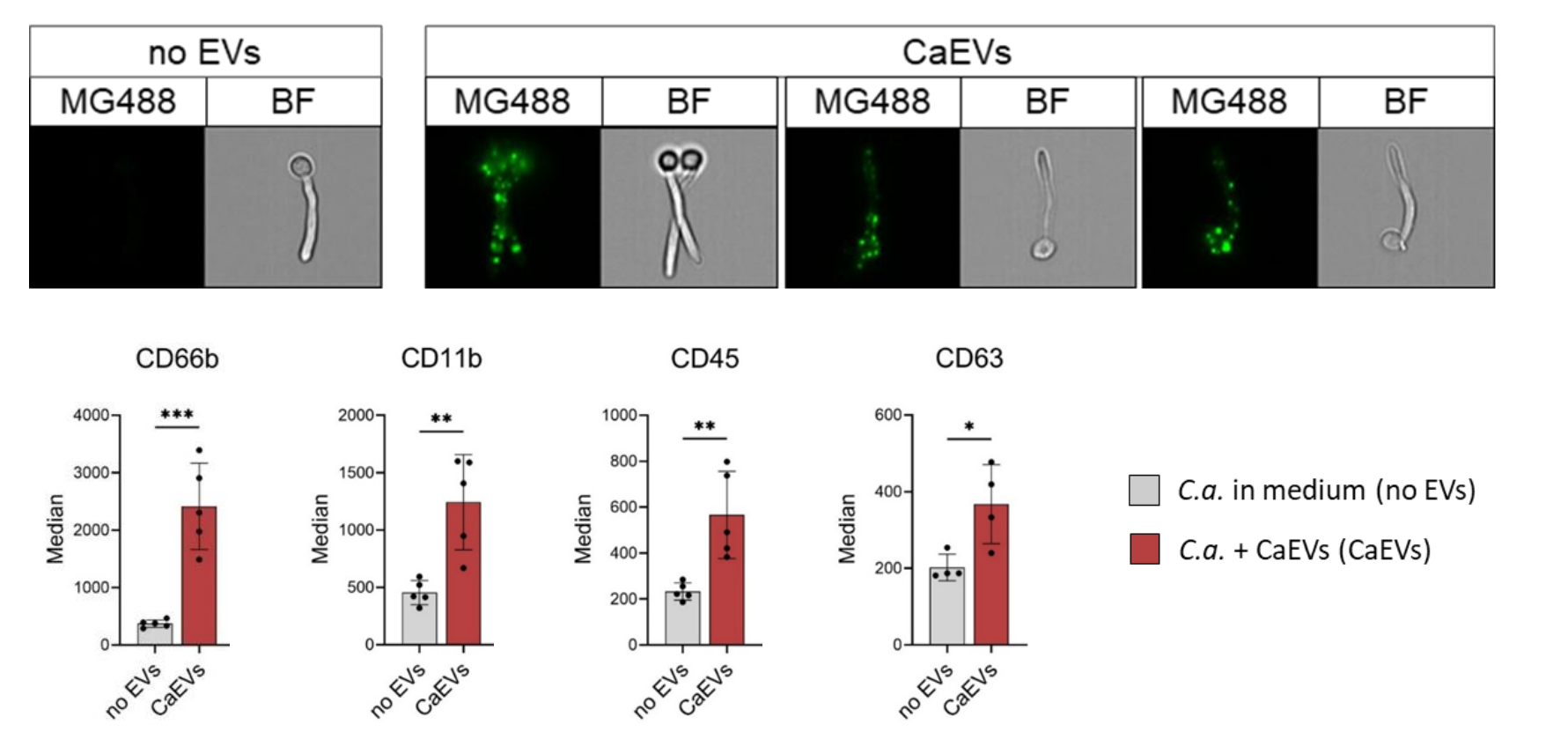
## Experimental characterization of EVs

### Neutrophils coat *C. albicans* with host molecules



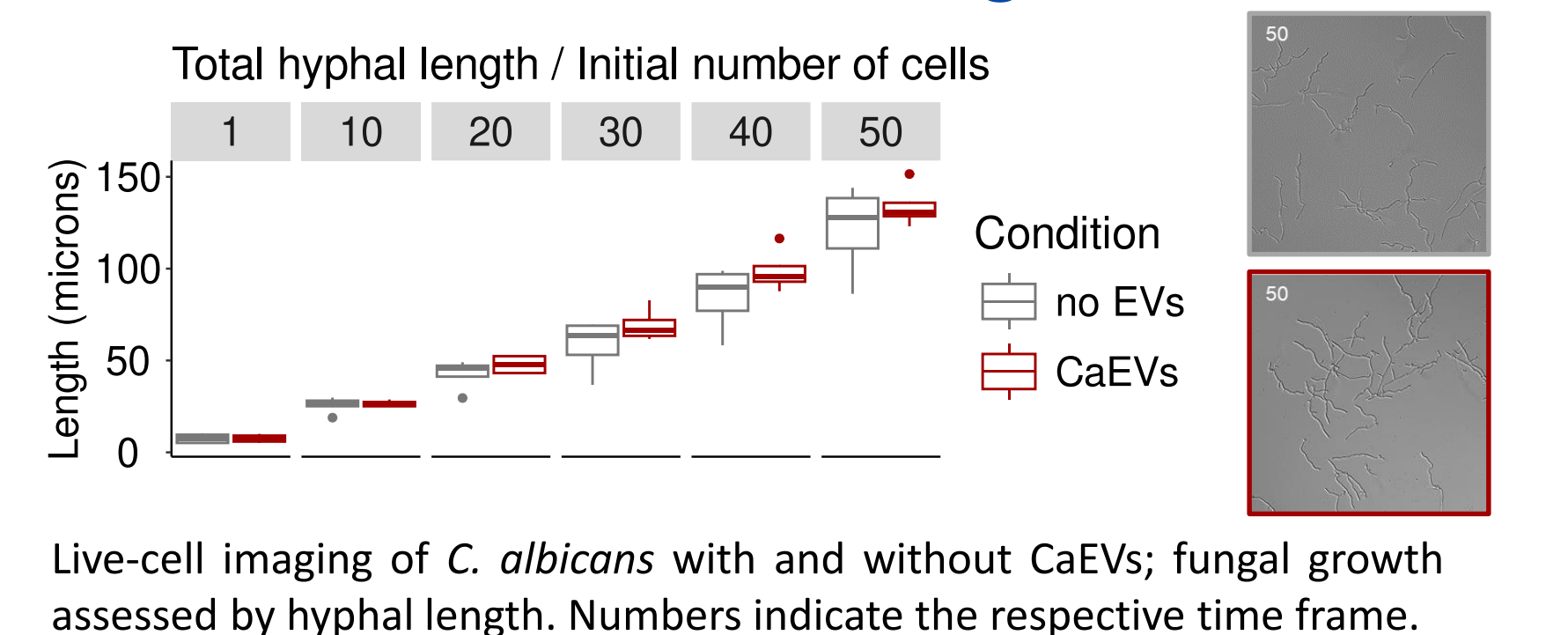
Imaging flow cytometry and flow cytometry analyses of the surface of extracellular *C. albicans* during whole-blood and neutrophil infection revealed the deposition of neutrophil proteins.

### Isolated EVs associate with *C. albicans*

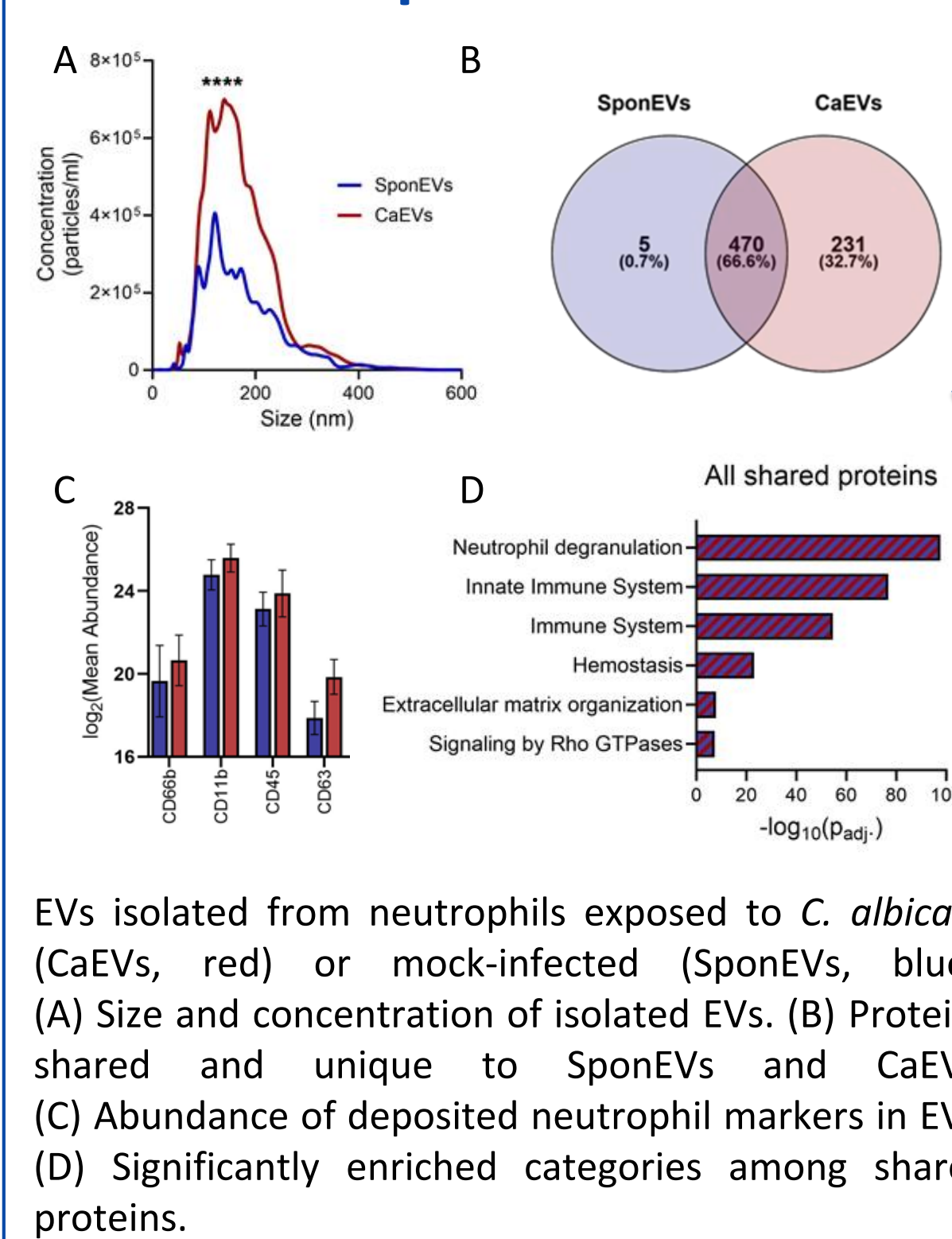


Neutrophil-derived EVs bind to *C. albicans* and mediate the transfer of neutrophil surface markers, as demonstrated by flow cytometry.

### EVs do not affect *C. albicans* growth

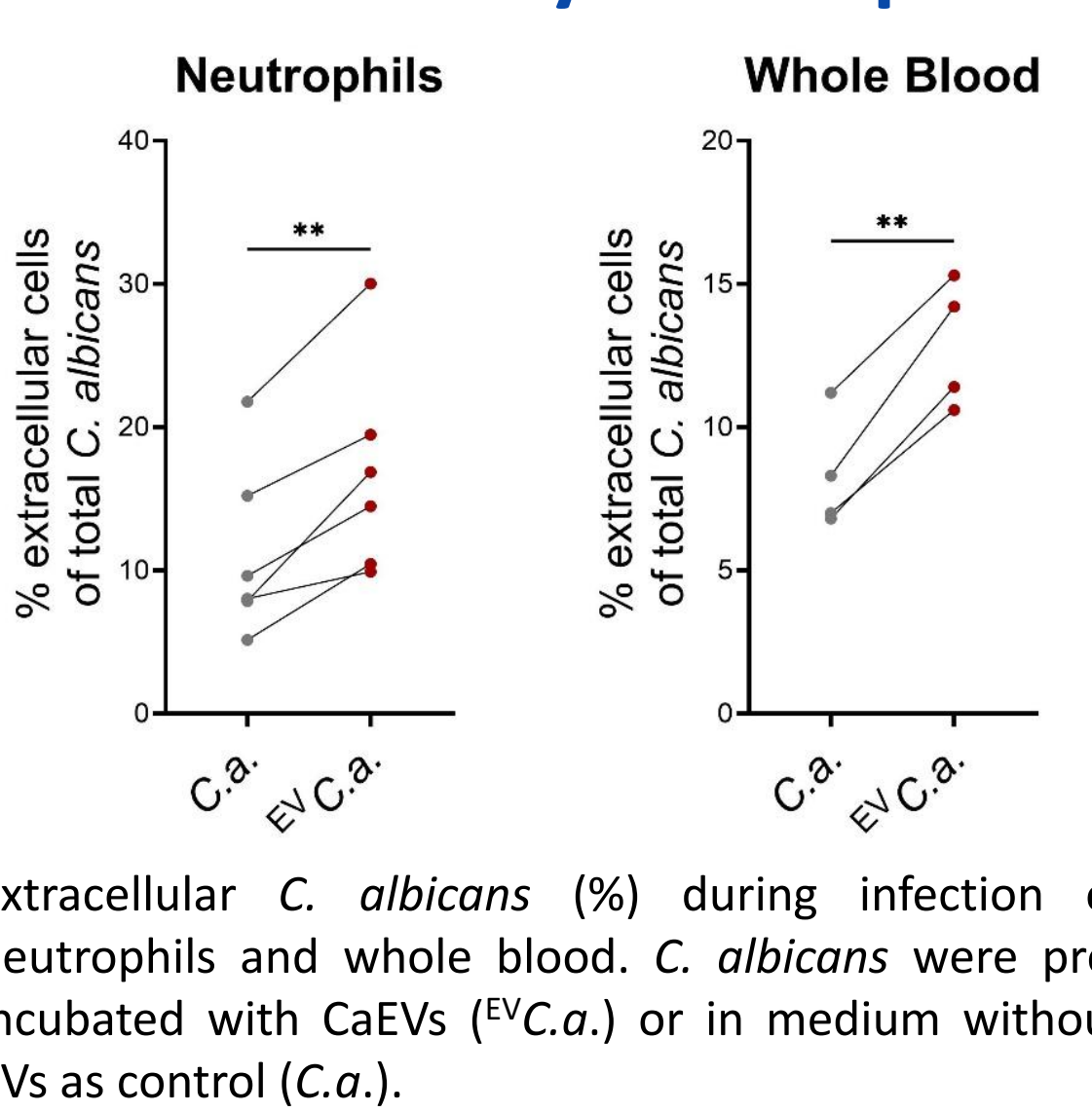


### Characterization of neutrophil-derived EVs



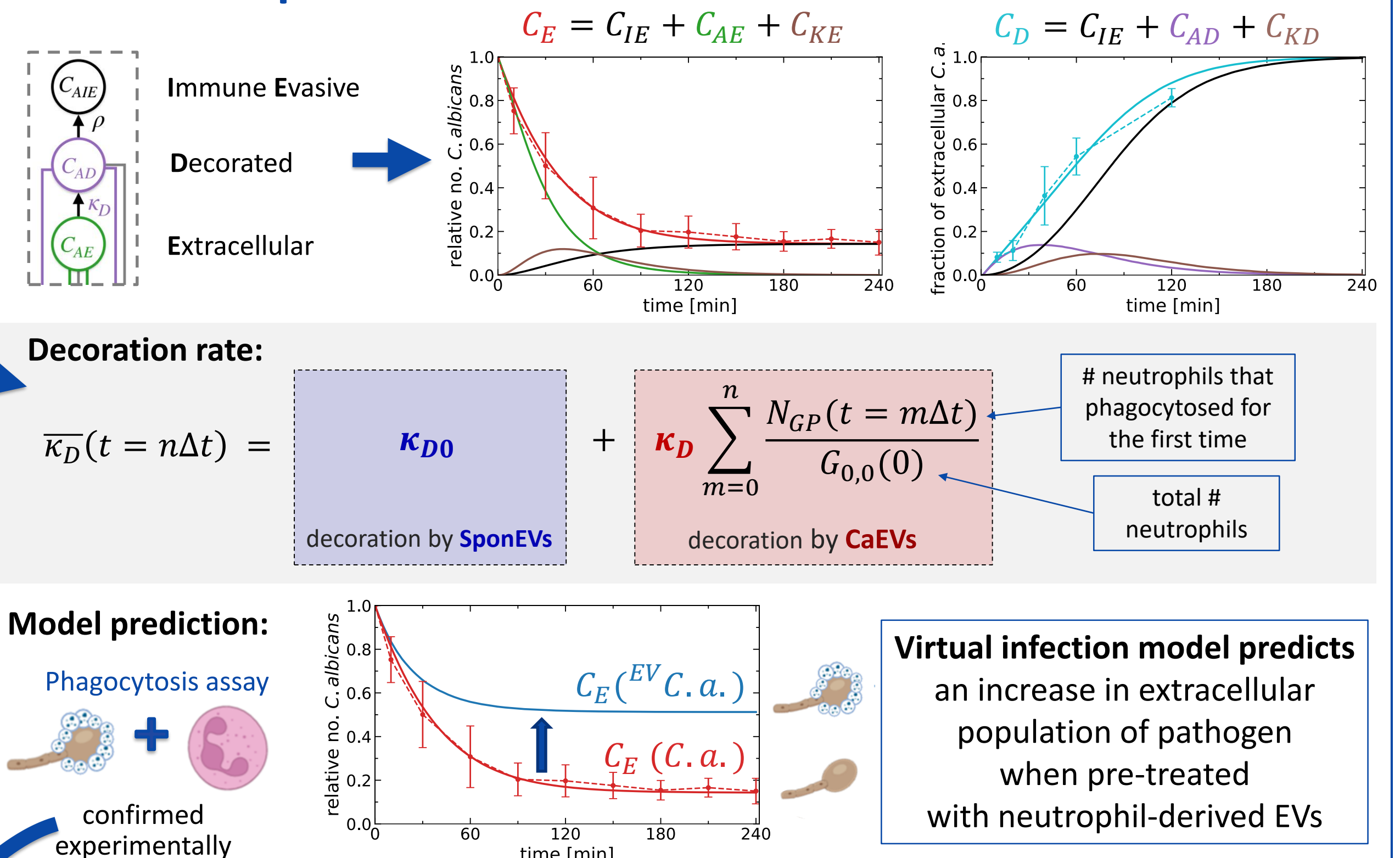
EVs isolated from neutrophils exposed to *C. albicans* (CaEVs, red) or mock-infected (SponEVs, blue). (A) Size and concentration of isolated EVs. (B) Proteins shared and unique to SponEVs and CaEVs. (C) Abundance of deposited neutrophil markers in EVs. (D) Significantly enriched categories among shared proteins.

### EVs reduce phagocytosis of *C. albicans* by neutrophils

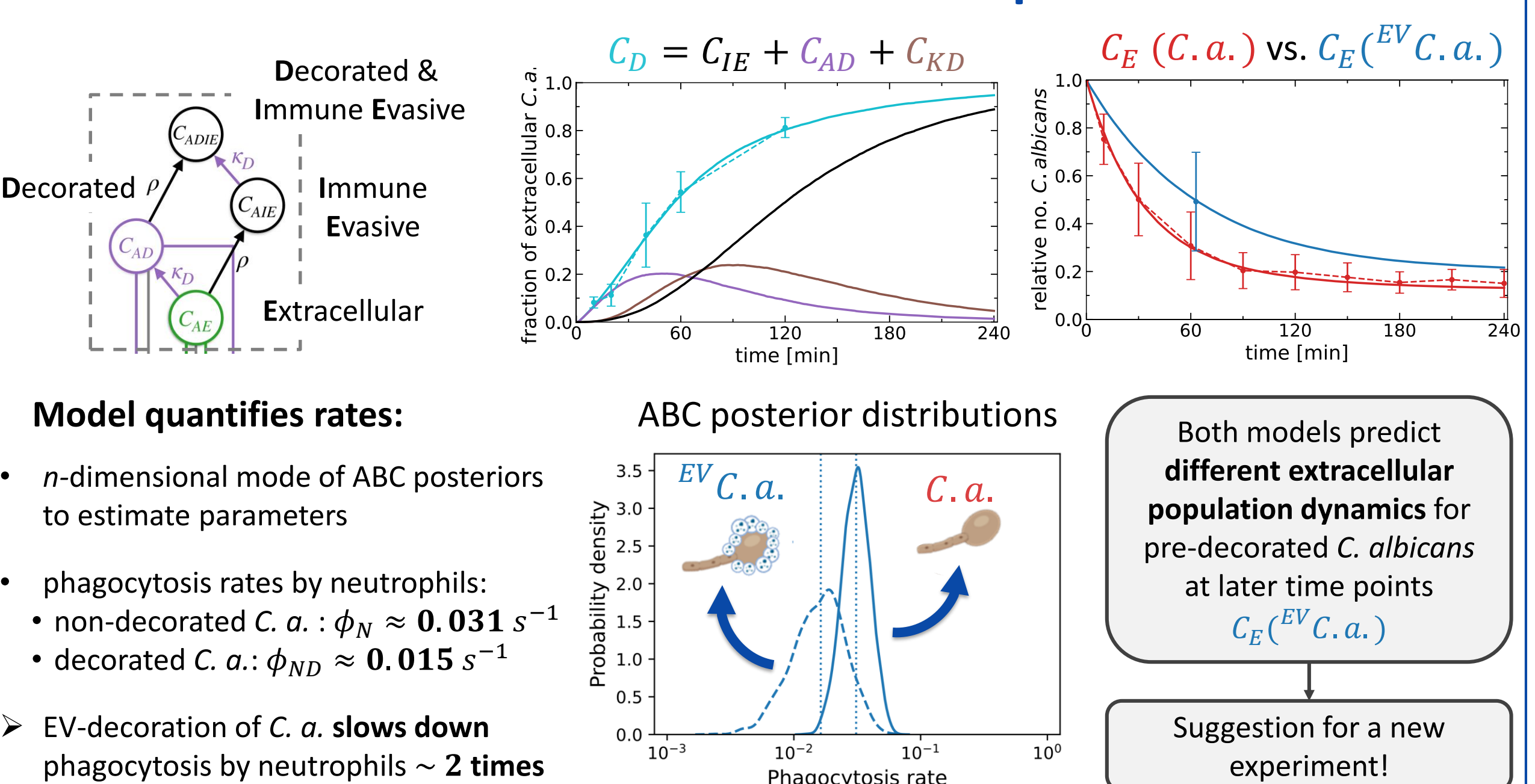


## Hypothesis-driven virtual models

### Decoration precedes immune evasion



### Decoration and immune evasion are independent



## Conclusion and outlook

- Neutrophil-derived EVs** are released upon *C. albicans* exposure and bind to fungal cells
- EV binding does not affect fungal growth
- Mathematical modeling predicts that **EV binding reduces phagocytosis efficiency** of neutrophils; this result was confirmed experimentally
- Mathematical modeling does not exclude an **alternative pathway** for immune evasion **without prior decoration**
- Quantification shows that **decorated pathogens are phagocytosed ~2x slower**
- To distinguish between competing hypotheses, we suggest an **additional experiment**

anastasia.solomatina@leibniz-hki.de

kerstin.huenniger@leibniz-hki.de

www.leibniz-hki.de

### References

- [1] Hünigler et al. 2014. PLOS Comput Biol. 10(2)  
[2] Lehnert et al. 2015. Front Microbiol. 6:608