

# IbSB-12

## Causal inference analysis of live cell imaged confrontation assays with agent-based modelling

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### Causal inference analysis of live cell imaged confrontation assays with agent-based modelling

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#### Introduction

Here we combine agent-based models (ABMs) with Dynamical Bayesian Networks (DBNs) [1] to analyse the cell track data in confrontation assays. DBNs can in principle describe the heterogeneity that is often observed in cell tracks [2] is caused by previous interactions or natural variability and even find causal interactions between different observables. Limitations are the amount of tracks normally available for each time point in live cell imaging, that the cells disappear from the field of view and that interactions occur because of the spatial distribution of the system rather than internal cell states. Therefore, we here use simulated data of a confrontation assay containing neutrophils and fungal cells based on an agent-based model [3]. In this model we can introduce causal relationships and remove them to investigate how these are represented in the DBN. We give an example how DBNs can identify a causal relationship between immune cell-fungal touching and the probability of individual immune cells to phagocytose fungal cells. We especially look at the possible dependencies that are introduced by spatial dynamics compared to the intrinsic causal effects that we introduce in the ABM.

$t$ : discrete time index with range  $t \in [0, t_{max}]$   
 $T_t$ : number of fungal cell touchings per immune cell until time  $t$   
 $F_t$ : additional touching event occurring between  $t$  and  $t + 1$   
 $P_t$ : number of phagocytosis events per immune cell until time  $t$   
 $\hat{P}$ : additional phagocytosis event occurring between  $t$  and  $t + 1$

#### Agent-based model and simulations

- 2D environment
- Immune cells
- Fungal cells
- 500 cells of each type
- Simulated for 2h
- Time step: 0.01s
- C++ implementation

Provides track files with:

- Immune cell positions
- Immune cell - fungal cell interactions
- Phagocytosis events

Control simulation:

- All immune cells are activated at the start of the simulation
- Phagocytosis occurs throughout the simulation

Causal simulation:

- Immune cell activation necessary for phagocytosis
- No immune cells are activated at the start of the simulation
- At least one touching event necessary for immune cell activation
- Multiple touchings increase activation probability

#### Network inference

##### Model selection

- The dependence of  $F$  and  $P$  on  $T_t, P_t$  and  $\hat{P}$  is determined by logistic regression
- The terms  $T_t$  and  $P_t$  are intrinsic causal factors, specific for each immune cell
- The term  $\hat{P}$  is an extrinsic factor that captures changes on the population level
- All possible linear combinations of the factors are tried
- Model selection is made based on the Bayesian information criterion (BIC)
- Testing the method by generating non-spatial data according to:

$$T_{t+1} = T_t + F_t, \quad P_{t+1} = P_t + \hat{P}_t$$
$$p(F = 1) = S_1(0.0071 + 0.031T_t - 0.0004P_t)$$
$$p(\hat{P} = 1) = S_2(0.052 - 0.048P_t + 0.00917T_t)$$

##### Inferred networks from ABM simulations

Best fitting network, ABM control:  $p(F = 1) = S_1(0.0071 + 0.031T_t - 0.0004P_t)$

Best fitting network, ABM causal:  $p(F = 1) = S_1(0.0027 + 0.020T_t - 0.00031P_t)$

##### Summary

- DBNs can be used to identify intrinsic and extrinsic causal effects
- Spatial dynamics may introduce network connections that are not intended
- Realistic ABM simulations are necessary to create control simulations

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References  
[1] Ebenau et al 2015, *Aliment Pharmacol Ther* 2015; 41:158-169.  
[2] Malmgren et al 2015, *PLoS ONE* 10(12):e0182006.  
[3] Lehnert et al 2015, *PLoS ONE* 10(12):e0182006.

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