



Cross Disciplines Seminar  
February 26, 2025  
10:30 AM

Lecture Hall (00.187) at BioZentrum I, Hanns-Dieter-Hüsch-Weg 15, 55128 Mainz

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# Mechanisms of Tumor Heterogeneity

## Insights from Mathematical Models

Tumor heterogeneity can be viewed as a collective phenomenon emerging from interactions between normal and malignant cells. As such, it can be studied using agent-based mathematical models, such as cellular automata. In this talk, I will present examples of these models to analyze the emergence of genotypic and phenotypic heterogeneity due to cellular interactions in growing tumors. I will also explore how these models help in understanding tumor evolution, clonal selection, and the role of the microenvironment in shaping heterogeneity.



### References

1. S. Syga, H. P. Jain, M. Krellner, H. Hatzikirou, A. Deutsch: Evolution of phenotypic plasticity leads to tumor heterogeneity with implications for therapy. *PLOS Comp. Biol.*, 2024
2. A. Deutsch, J. M. Nava-Sedeño, S. Syga, H. Hatzikirou. BIO-LGCA: a cellular automaton modelling class for analysing collective cell migration. *PLOS Comp. Biol.* (6), 2021
3. O. Ilina, P.G. ..., A. Deutsch, P. Friedl. Cell-cell adhesion and 3D matrix confinement determine jamming transitions in breast cancer invasion. *Nature Cell Biology* 1103-1115, 2020
4. A. Deutsch, S. Dormann. Cellular automaton modeling of biological pattern formation: characterization, examples, and analysis. Birkhauser, Basel, 2018

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