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Modeling the host-pathogen interactions of macrophages and Candida albicans using game theory and dynamic optimization
Sybille Dühring (Jena/Germany)

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Dimensionality of motion but not receptor morphology governs affinity of receptor-ligand binding as revealed by molecular agent-based models
Teresa Lehnert (Jena/Germany)

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Iron redistribution after Candida albicans infection in the murine kidney
Theresia Conrad (Jena/Germany)

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Investigating Candida albicans resistance in whole-blood assays by virtual infection models using parallelized parameter estimation
Maria Praußе (Jena/Germany)

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Simulation of the dynamics of primary immunodeficiencies in CD4+ T-cells
Gabriel Teku (Lund/Sweden)

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Michael Weber (Jena/Germany)

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Carolin Zitzmann (Greifswald/Germany)

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Nelly F. Mostajo Berrospi (Jena/Germany)

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Jennifer Scheidel (Frankfurt am Main/Germany)

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Ralf Takors (Stuttgart/Germany)

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Genome-wide gene regulatory network in the opportunistic human pathogenic fungi Aspergillus fumigatus
Silvia Gerber (Jena/Germany)

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Sabrina Ellenberger (Jena/Germany)

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Polynomial parametric modeling of synthetic transcriptional circuits in S. cerevisiae
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Christoph Schmal (Berlin/Germany)

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Andreas Kremling (Garching/Germany)

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David van Niekerk (Stellenbosch/South Africa)

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Rikard Johansson (Linköping/Sweden)

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Shahid Khan (Berkeley/United States)

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Kiril S. Zaytsev (Moscow/Russian Federation)

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Dipali Singh (Oxford/Great Britain)

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