

Scientific Programme

International Conference on Mathematical Methods and Models (Biomath 2023), 18-23 June 2023, Pomorie, Bulgaria

08:00 – 08:50	Registration.	SYS Lecture: Maya Mincheva, Biodynamics of reaction networks
09:00 – 09:20	Opening	
		Chair: Roumen Anguelov
09:20 – 10:05	Keynote: Irina Doytchinova	<i>Novel acetylcholinesterase inhibitors developed by structure-based drug design in DDBL@MUS</i>
10:05 – 10:50	Keynote: Andreas Deutsch	<i>Mechanisms of cancer invasion and progression: insights from cellular automaton models</i>
10:50 – 11:10	Tea/Coffee	
		Chair: Andreas Deutsch
11:10 – 11:35	Miroslaw Lachowicz	<i>Self-organization or disorder phenomena in biology</i>
11:35 – 12:00	Svetoslav Nikolov	<i>A mathematical model for compression of poroviscoelastic biological material-like articular cartilage</i>
12:00 – 12:25	Ryszard Rudnicki	<i>Some aspects of mathematical modelling of cell cycle</i>
12:25 – 12:50	Katarzyna Pichór	<i>A general model of immune status</i>
12:50 – 14:00	Lunch	
		Chair: Irina Doytchinova
14:00 – 14:25	Mariyana Atanasova	<i>In silico screening of natural compounds and discovery of novel acetylcholinesterase inhibitors</i>
14:25 – 14:50	Ivan Dimitrov	<i>PreDQ - a software tool for peptide binding prediction to HLA-DQ2 and/or HLA-DQ8</i>
14:50 – 15:15	Anjana Ghelani	<i>Genome variation analysis and strategic clustering to sub-lineage of double mutant strain B.1.617 of SARS-CoV-2 t</i>
15:15 – 15:40	Anelia Horvath	<i>Identification and analysis of cell-specific expressed genetic variants from scRNA-seq data</i>
15:40 – 16:05	Julien Vincent	<i>Modelling horizontal gene transfer of plasmid-mediated resistance in biofilms</i>
16:05 – 16:30	Tea/Coffee	
		Chair: Ami Radunskaya
16:30 – 16:55	Maria C. A. Leite	<i>Modeling the control of breathing using a Boolean framework</i>
16:55 – 17:20	Nikolet Doneva	<i>In silico immunogenicity prediction of viral proteins</i>
17:20 – 17:45	Monika J. Piotrowska	<i>Analysis of a basic mathematical model of CAR-T cell therapy for glioblastoma</i>
17:45 – 18:10	Urszula Foryś	<i>Periodic treatment in a mathematical model of CAR-T cell therapy for glioblastoma</i>
19:30 – 20:30	Welcoming Cocktail (Lobby of the hotel)	

Tuesday	20 June 2023	
		Chair: Yves Dumont
09:00 – 09:45	Keynote: Ami Radunskaya	Mathematical challenges in triggered drug delivery: getting the right dose to the right place at the right time
09:45 – 10:30	Keynote: Somdatta Sinha	Modelling cellular processes: from single cells to collective behaviour
10:30 – 11:00	Tea/Coffee (Bring all posters to the Vista Hall, where the tea/coffee is)	
		Chair: Michael Chapwanya
11:00 – 11:25	Yves Dumont	About sterile insect control strategies in a two patches system
11:25 – 11:50	Daiver Cardona-Salgado	Optimal control approach for establishing Wolbachia in wild population of Aedes aegypti mosquitoes
11:50 – 12:15	Jacek Banasiak	Impact of demography on the dynamics of malaria
12:15 – 12:40	Ivric Valaire Yatat-Djeumen	About sterile females contamination and residual fertility in a mosquito control program using the Sterile Insect Technique. Impact on Dengue control.
12:40 – 12:50	Conference Photo	
12:50 – 14:00	Lunch (All posters to be displayed)	
		Chair: Hristo Kojouharov
14:00 – 14:25	Stanislav Sotirov	Assessment of human proteins for potential tumour immunogenicity by in silico models
14:25 – 14:50	Marek Bodnar	Mathematical model of CAR-T therapy taking into account cells targeting off-tumour antigens
14:50 – 15:15	Ivanka Stamova	On the qualitative analysis of fractional-order impulsive gene regulatory networks
15:15 – 15:40	Charlise Basson &	Mathematical modelling for CTCE-9908 (a CXCR4 inhibitor) on B16 F10 melanoma cell proliferation: Part I and Part II
15:40 – 16:05	Avulundiah Edwin Phiri	
16:05 – 16:30	Tea/Coffee	
		Chair: Jean Lubuma
16:30 – 16:55	Hong-Ming Yin	On an optimal control problem arising from the tumor treatment
16:55 – 17:20	Mmatlou S. Kubyana	Adaptive strategies destabilise the rock-paper-scissors game but increase the eco-evolutionary performance
17:20 – 18:20	Poster Session (posters to remain displayed until 16:30 on 21 June)	
18:30 – 19:30	Open ISC Meeting	SYS Lecture: Ami Radunskaya How to get the most out of a model

Posters

Yusuke Asai	<i>Evaluation of the effectiveness of health countermeasures against infectious disease arrival times</i>
R. Anguelov, C. Basson, A. E. Phiri	<i>Mathematical modelling for CTCE-9908 (a CXCR4 inhibitor) on B16 F10 melanoma cell proliferation</i>
Milen Borisov	<i>On the construction of amiloid growth models</i>
Daiver Cardona-Salgado	<i>Modeling the introduction of sterilizing treatment for tuberculosis in low- and middle-income countries</i>
Eunjin Eom	<i>Construction of a new infectious disease model using the time delay</i>
Slavi Georgiev	<i>Dynamical analysis combined with parameters identification for a model of infection in honeybee colonies with social immunity</i>
Ivan M. Ivanov	<i>Ultrasound experimental model for knee joint intra-articular movements</i>
Elena V. Nikolova	<i>Analytical and numerical study of a diffusive predator-prey model incorporating an Allee effect</i>
Peter Rashkov	<i>Time-scale separation in models for dengue fever</i>
Fatima Sapundzhi	<i>The impact of molecular docking programs and virtual screening in modern drug design</i>
Stoyan Stoytchev	<i>Estimation of the parameters of a mathematical model for articular cartilage compression</i>
Ivan Trenchev	<i>Integrating Mixed Reality Technologies in Genomic Data Visualization and Analysis for Bioinformatics Research</i>
Iryna Zabaikina	<i>Quantifying noise modulation from coupling of stochastic expression to cellular growth: an analytical approach</i>

Wednesday	21 June 2023	
		Chair: Greg Rempala
09:00 – 09:30	Invited: Manuel Molina	<i>A stochastic model to mathematically describe the dynamics of long-lived raptor species</i>
09:30 – 10:00	Invited: Miguel González	<i>Controlled branching processes as models for logistic population growth</i>
10:00 – 10:30	Invited: Inés del Puerto	<i>Multitype controlled branching process as a model for progenitor cell populations</i>
10:30 – 11:00	Tea/Coffee	
	Session A	Chair: Alf Gerisch
11:00 – 11:25	María Vela-Pérez	<i>Modeling the COVID-19 incorporating variants and vaccines</i>
11:25 – 11:50	Arsène Jaures Ouemba Tassé	<i>Influence of the co-dynamics Ebola-COVID-19 in the population</i>
11:50 – 12:15	Slavi Georgiev	<i>Optimal coefficient restoration for COVID-19 epidemic modelling</i>
12:15 – 12:40	Fawaz K. Alalhareth	<i>Analysis and an NSFD method of a model of bacterial competition in the presence of a plasmid</i>
	Session B	Chair: Stoyan Stoytchev
11:00 – 11:25	Claudia Timofte	<i>Asymptotic analysis for diffusion problems in thin periodic media</i>
11:25 – 11:50	Zirhumanana Balike	<i>Mathematical modeling of trace-metals precipitation in biofilms</i>
11:50 – 12:15	Ahmed Fathi	<i>Upscaling a mixed-culture biofilm model in homogeneous porous media via multiscale asymptotics approach</i>
12:15 – 12:40	Andrey Vassilev	<i>The hybrid Gompertz distribution - derivation, characterization and estimation (a reaction network treatment)</i>
12:40 – 14:00	Lunch	
		Chair: Jacek Banasiak
14:00 – 14:25	Petra Macková	<i>Distributional solutions of nonlinear diffusion equations with a moving Dirac source term</i>
14:25 – 14:50	Ekaterina Auer	<i>Predicting Hereditary BRCA1/2 Mutations Using Publicly Available Data</i>
14:50 – 15:15	Maya Mincheva	<i>Identifying parameter values for oscillations in reaction networks</i>
15:15 – 15:40	Sara Hamis	<i>Spatial cumulant models enable spatially informed treatment strategies in theoretical cancer systems</i>
15:40 – 16:05	Jean Lubuma	<i>On reliable numerical methods for some real-life differential equation models with singularities</i>
16:05 – 16:30	Tea/Coffee	
	Section A	Chair: Manuel Molina
16:30 – 16:55	Nikolay Yanev	<i>Nonhomogeneous multitype Markov branching stochastic processes as models of cell population dynamics</i>
16:55 – 17:20	Boseung Choi	<i>Statistical model for identification of the relationship between SARS-CoV-2 prevalence and wastewater concentration with vaccination and delta variant mutation</i>
17:20 – 17:45	Marta Tyran-Kamińska	<i>A stochastic model of seasonal savanna</i>
	Section B	Chair: Claudia Timofte
16:30 – 16:55	Nadezhda Zh. Bunzarova	<i>On an integrable version of the generalized totally asymmetric simple exclusion process on open chains</i>
16:55 – 17:20	Schehrazad Selmane	<i>Demographic and epidemiological characteristics of animal bites in the North West Health Region, Algeria</i>
17:20 – 17:45	Yibeltal Adane Terefe	<i>Impact of self-protection measures to reduce antibiotic resistant gonorrhoea infection</i>
18:30 – 23:00	Official Dinner	

Thursday	22 June 2023	
08:30 – 15:30	Excursion	
15:30 – 16:00	Tea/Coffee	
		Chair: Mirosław Lachowicz
16:00 – 16:25	Dariusz Wrzosek	<i>Pursuit-evasion dynamics in predator-prey models</i>
16:25 – 16:50	Pravin Dudhagara	<i>Response surface designs and R package based modelling for predictive capabilities, sensitivity and validation of model for improving the bacterial growth, laccase production, and textile dye decolorization alongside a detoxification study</i>
16:50 – 17:15	Morelyn Sigauke	<i>Analysis of the transmission dynamics of pneumonia disease in a developing country: the effect of environment on pneumonia transmission, significance of hospital and community based care</i>
17:15 – 17:40	Michael Chapwanya	<i>Pattern formation in the Holling-Tanner predator-prey model with prey-taxis</i>
17:40 – 18:05	Hristo V. Kojouharov	<i>Higher-order modified nonstandard finite difference methods for autonomous dynamical systems</i>

Friday	23 June 2018	
		Chair: Somdatta Sinha
09:00 – 09:45	Keynote: Alf Gerisch	<i>Exploring the spatial tissue environment: non-local modelling, applications, numerical challenges</i>
09:45 – 10:30	Keynote: Greg Rempala	<i>Modeling biological systems using dynamical survival models. Lessons from the recent pandemic.</i>
10:30 – 11:00	Tea/Coffee	
		Chair: Peter Rashkov
11:00 – 11:25	Jian-Jun Shu	<i>Fat-tail test of regulatory DNA sequence</i>
11:25 – 11:50	Nadia Antonova	<i>On the analysis of hemorheological data with mathematical models</i>
11:50 – 12:15	Svetoslav Markov	<i>The SEI epidemiological model generates natural classes of biological functions: analysis using reaction network theory</i>
12:15 – 14:00	Lunch	