## 33rd M. Smoluchowski Symposium on Statistical Physics

Thursday 03 December 2020 - Friday 04 December 2020



## **Book of Abstracts**

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## **Conference Agenda**

Thursday, 03/12/2020	
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09:00 - 09:30 CET	<b>Kim Sneppen</b> Super-spreaders in the Corona Epidemics
09:30 - 10:00 CET	<b>Adam Kleczkowski</b> Using stochastic models to describe the coronavirus epidemic
10:00 - 10:30 CET	<b>Thierry Dauxois</b> Energy cascade in internal wave attractors
10:30 - 11:00 CET	<b>Gleb Oshanin</b> Spectral content of a single trajectory

12:00 - 12:30 CET	<b>Eli Barkai</b> Packets of diffusing particles exhibit universal exponential tails
12:30 - 12:50 CET	<b>Adrian Baule</b> Loopy Lévy flights enhance tracer diffusion in active suspensions
12:50 - 13:10 CET	<b>Alexandre Guillet</b> Extreme-Value Statistics of Stochastic Transport Processes
13:10 - 13:30 CET	<b>Aleksander Stanislavsky</b> <i>Confined random motion: What is statistics? Gaussian or</i> <i>Laplacian?</i>
13:30 - 13:50 CET	<b>Horacio S. Wio</b> Variational Approach to KPZ: Fluctuation Theorems, Large Deviation Function, and Probability Distribution Function

16:00 - 16:30 CET	<b>Sidney Redner</b> <i>Optimization and Growth in First-Passage Resetting</i>
16:30 - 17:00 CET	<b>Shlomi Reuveni</b> <i>Restart: The Physics Of Starting Anew</i>

17:00 - 17:20 CET	<b>Martin Bier</b> Lévy Noise, Time-Reversal Symmetry, Nonequilibrium Stochastic Thermodynamics, and Bak's Sandpile
17:20 - 17:40 CET	<b>Jerzy Gorecki</b> <i>Simple mathematical models for self-propelled motion on the</i> <i>water surface</i>
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09:50 - 10:10 CET	<b>Baruch Meerson</b> Geometrical Optics of Constrained Brownian Motion
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13:00 - 13:20 CET	<b>David Kessler</b> Biological Networks Regulating Cell Fate Choice Are Minimally Frustrated	
13:20 -13:40 CET	<b>Andreas Dechant</b> <i>Continuous time-reversal and the thermodynamic uncertainty</i> <i>relation</i>	
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## Presentations

#### Thursday 03/12, 09:00 - 09:30 CET

## **Super-spreaders in the Corona Epidemics**

Kim Sneppen<sup>1</sup>

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Recently a powerful example of a replicating nano-machinery entered our society. In principle it's just a normal disease, that one attempt to model with 3 or 4 simple coupled equations with 2 important parameters: a timescale, and a replication factor (the famous  $R_0$ ). And then one try to guess how changes in society changes  $R_0$  and perhaps adopt some more or less strong lock-down measures. However, this virus have more "personality" than that. It behaves different in different persons, and persons behave differently. Presumably only a few of us infect a lot, while most does not infect so much. This assumption is supported by the observation that couples living together only infect each other with about 15 percent probability, indicating that most infected people are not really infectious. I will discuss this and other aspects of Covid-19 in perspective of models that describe heterogeneous individuals in a society. In particular we suggest that limiting superspreading opportunities is a cost effective strategy to mitigate Covid-19.

## Presentations (continued)

#### Thursday 03/12, 09:30 - 10:00 CET

### Using stochastic models to describe the coronavirus epidemic

A. Kleczkowski<sup>1</sup>, Jonathan Wells, Chris Robertson<sup>1</sup>, Vincent Marmara<sup>2</sup> and Alan Yeung

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In December 2019, a novel new strain of coronavirus (COVID-19) was identified as it spread through China and subsequently throughout the world, resulting in a pandemic. As the health systems became overwhelmed, a need arose for real-time surveillance and modelling to identify the breaking points caused by increased numbers of infections and hospitalised individuals. We combined surveillance data for Scotland, detailing new COVID-19 cases, deaths, hospitalisations and critical care beds, with a modified stochastic Susceptible Exposed Infected Recovered Hospitalised Critical Dead (SEIR-HCD) model. The model parameters are adjusted daily, to fit the real-world data, using a Bayesian statistical particle filtering technique. The estimation technique uses an acceptance/rejection method combined with correlated multi-dimensional diffusion and drift model to effectively search the parameter space. We show close agreement between data and model output and discuss how changes in parameters reflect the changing transmission dynamics of the virus and severity of the disease.

## Presentations (continued)

#### Thursday 03/12, 10:00 - 10:30 CET

### Energy cascade in internal wave attractors

Thierry Dauxois<sup>1</sup>

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Internal gravity waves play a primary role in geophysical fluids : they contribute significantly to mixing in the ocean and they redistribute energy and momentum in the middle atmosphere. In addition to their very interesting and very unusual theoretical properties, these waves are linked to one of the important questions in the dynamics of the oceans: the cascade of mechanical energy in the abyss and its contribution to mixing. Combining the physics of waves, dynamical systems theory and oceanography, I will discuss a unique self-consistent experimental and numerical setup that models a cascade of triadic interactions transferring energy from large-scale monochromatic input to multi-scale internal wave motion. I will also provide explicit evidence of a wave turbulence framework for internal waves.

## Presentations (continued)

#### Thursday 03/12, 10:30 - 11:00 CET

## Spectral content of a single trajectory

Gleb Oshanin<sup>1</sup>

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In this presentation I will overview a recent progress in the theoretical, numerical and experimental analyses of spectral densities of individual random trajectories of a finite length. On example of a broad class of anomalous diffusions - the so-called fractional Brownian motion, I will demonstrate that one may calculate analytically the full probability density function of such random functionals, parametrised by a frequency and a finite observation time, and extract a very meaningful information on the evolution of a process under study. In particular, I will show that the large frequency behaviour of the coefficient of variation of this distribution provides a robust criterion of anomalous diffusion, which analytical prediction is validated by a comparison with experiments on dynamics in live cells and in agarose hydrogels, and also by extensive numerical simulations. If time permits, I will also discuss the peculiarities of spectra in out-of-equilibrium systems, as exemplified by a Brownian Gyrator model, and also of the so-called active Brownian motion.

## Presentations (continued)

#### Thursday 03/12, 12:00 - 12:30 CET

# Packets of diffusing particles exhibit universal exponential tails

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Brownian motion is a Gaussian process described by the central limit theorem. How- ever, exponential decays of the positional probability density function P(X, t) of packets of spreading random walkers, were observed in numerous situations that include glasses, live cells, and bacteria suspensions. We show that such exponential behavior is generally valid in a large class of problems of transport in random media. By extending the large deviations approach for a continuous time random walk, we uncover a general universal behavior for the decay of the density [1]. It is found that fluctuations in the number of steps of the random walker, performed at finite time, lead to exponential decay (with logarithmic corrections) of P(X,t). This universal behavior also holds for short times, a fact that makes experimental observations readily achievable. Time permitting we then formulate the hitchhiker model where interacting molecules form aggregates, that lead to fluctuations in the diffusion field, and a many body mechanism for the exponential tails [2].

#### References

[1] E. Barkai and Stas Burov, Packets of diffusing particles exhibit universal exponential tails, Phys. Rev. Lett. **124**, 060603 (2020).

[2] M. Hidalgo-Soria, and E. Barkai, The Hitchhiker model for Laplace diffusion processes in the cell environment, Phys. Rev. E **102**, 012109 (2020).

## Presentations (continued)

#### Thursday 03/12, 12:30 - 12:50 CET

## Loopy Lévy flights enhance tracer diffusion in active suspensions

Kiyoshi Kanazawa<sup>1,2</sup>, Tomohiko Sano<sup>3,4</sup>, Andrea Cairoli<sup>5,6,7</sup> and Adrian Baule<sup>6</sup>

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Brownian motion is widely used as a model of diffusion in equilibrium media throughout the physical, chemical and biological sciences. However, many real-world systems are intrinsically out of equilibrium owing to energy-dissipating active processes underlying their mechanical and dynamical features. The diffusion process followed by a passive tracer in prototypical active media, such as suspensions of active colloids or swimming microorganisms, differs considerably from Brownian motion, as revealed by a greatly enhanced diffusion coefficient and non-Gaussian statistics of the tracer displacements. Although these characteristic features have been extensively observed experimentally, there is so far no comprehensive theory explaining how they emerge from the microscopic dynamics of the system. Here we develop a theoretical framework to model the hydrodynamic interactions between the tracer and the active swimmers, which show s that the tracer follows a non-Markovian coloured Poisson process that accounts for all empirical observations [1]. The theory predicts a long-lived Lévy flight regime of the loopy tracer motion with a non-monotonic crossover between two different power-law exponents. The duration of this regime can be tuned by the swimmer density, suggesting that the optimal foraging strategy of swimming microorganisms might depend crucially on their density in order to exploit the Lévy flights of nutrients. Our framework can be applied to address important theoretical questions, such as the thermodynamics of active systems, and practical ones, such as the interaction of swimming microorganisms with nutrients and other small particles (for example, degraded plastic) and the design of artificial nanoscale machines.

[1] K. Kanazawa, T. Sano, A. Cairoli, A. Baule; Nature 579, 364–367 (2020)

## Presentations (continued)

#### Thursday 03/12, 12:50 - 13:10 CET

## **Extreme-Value Statistics of Stochastic Transport Processes**

Alexandre Guillet<sup>1</sup>, Edgar Roldan<sup>1</sup> and Frank Jülicher<sup>1</sup>

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We derive exact expressions for the finite-time statistics of extrema (maximum and minimum) of the spatial displacement and the fluctuating entropy flow of biased random walks. Our approach captures key features of extreme events in molecular motor motion along linear filaments. For one-dimensional biased random walks, we derive exact results which tighten bounds for entropy production extrema obtained with martingale theory and reveal a symmetry between the distribution of the maxima and minima of entropy production. Furthermore, we show that the relaxation spectrum of the full generating function, and hence of any moment, of the finite-time extrema distributions can be written in terms of the Marchenko-Pastur distribution of random-matrix theory. Using this result, we obtain efficient estimates for the extreme-value statistics of stochastic transport processes from the eigenvalue distributions of suitable Wishart and Laguerre random matrices. We confirm our results with numerical simulations of stochastic models of molecular motors.

## Presentations (continued)

#### Thursday 03/12, 13:10 - 13:30 CET

## Confined random motion: What is statistics? Gaussian or Laplacian?

Aleksander Stanislavsky<sup>1</sup>

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In biophysics, the single-particle tracking (SPT) is broadly used to quantify the kinetics of fluorescently-labelled molecules/proteins in live cells. This powerful tool in microscopy permits ones to resolve modes of motion of individual molecules for better understanding their role in such systems. The motion is not limited to pure (Brownian motion) diffusion, but directed, confined, anomalous diffusion are observed too. The study of their effects in live cells is of great interest.

Based on strict analytical calculations and simulations, we show that the confined motion in presence of Brownian motion in complex systems may have two different statistics . One of them has the normal distribution, whereas another, surprisingly, is under the Laplace distribution with fatter tails than the normal case (with thin tails always). Using the experimental data obtained from a recent SPT study on two particle types, G-protein coupled receptors and the G proteins with which the receptors interact, we have detected the normal and Laplace confinements. It is interesting that the contribution of the Laplace confinement is different for G proteins and receptors.

The results are an important step to experimental cell probing and paves the way for a new strategy acting on the processes occurring in live cells through changing the confined motion of molecules into live cells, for example, by drugs. This analytic treatment can be used for different models, thus opening a new direction towards understanding the role of confinement in complex systems.

## Presentations (continued)

#### Thursday 03/12, 13:30 - 13:50 CET

## Variational Approach to KPZ: Fluctuation Theorems, Large Deviation Function, and Probability Distribution Function

<u>Horacio S. Wio</u><sup>1</sup>, Miguel A. Rodriguez<sup>2</sup>, Rafael Gallego<sup>3</sup>, Roberto R. Deza<sup>4</sup> and Jorge A. Revelli<sup>5</sup>

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The KPZ equation is - as known - a stochastic field theory that describes the kinetic roughening of surfaces and interfaces, plus many other nonequilibrium processes that belong to its universality class. For  $\lambda \neq 0$ , the systems described are intrinsically unstable and "lack of stationary probability distribution". This can be readily inferred from the structure of the functional from which the deterministic KPZ equation stems in a variational formulation [1]. In fact, the time behavior of the aforementioned functional resembles that of a particle in a gravitational field [2]. A path-integral scheme has been set up in [3] that allows obtaining detailed and integral fluctuation theorems - as well as a Large Deviation Function for entropy production for the KPZ equation, "regardless of the substrate dimensionality" [4]. After reviewing these issues, we close by discussing an explicit expression of the probability distribution for the KPZ system.

[1] H.S. Wio, "Variational formulation for the KPZ and related kinetic equations", Int. J. Bif. Chaos 19, (2009) 2813-2821.

[2] H.S. Wio, M.A.Rodriguez, R.Gallego, J.A.Revelli, A. Ales and R.R.Deza, "d-Dimensional KPZ Equation as a Stochastic Gradient Flow in an Evolving Landscape: Interpretation and Time Evolution of Its Generating Functional", Front.Phys.4 (2017): 52.

[3] H.S. Wio, R.R. Deza and J.A. Revelli, "Fluctuation theorems and large deviation functions in systems not featuring a steady state", J. Stat. Mech. (2020) 024009.

[4] H.S. Wio, M.A. Rodriguez and R. Gallego, "Variational approach to KPZ: Fluctuation theorems and large deviation function for entropy production", Chaos 30, (2020) 073107.

## Presentations (continued)

#### Thursday 03/12, 16:00 - 16:30 CET

## **Optimization and Growth in First-Passage Resetting**

Sidney Redner<sup>1</sup>, Benjamin De Bruyne<sup>2</sup> and Julien Randon-Furling<sup>3</sup>

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We combine resetting and first-passage to define "first-passage resetting", where a random walk is reset to a fixed position due to a first-passage event of the walk itself. On the infinite half-line, first-passage resetting of isotropic diffusion is non-stationary, in which the number of resetting events grows with time as  $t^{1/2}$ . We calculate the resulting spatial probability distribution of the particle, and also obtain this distribution by a path decomposition approach. In a finite interval, we define an first-passage-resetting optimization problem that is motivated by reliability theory. Here, the goal is to operate a mechanical system close to its maximum capacity without experiencing too many breakdowns. When a breakdown occurs, the system is reset to its minimal operating point. We define and optimize an objective function that maximizes the reward (being close to maximum operation) minus a penalty for each breakdown. Finally, we study a first-passage-driven domain growth dynamics in which its boundary recedes by a specified amount when a diffusing particle reaches the boundary, after which resetting occurs. We find a wide range of dynamical behaviors for the domain growth rate in the interval and the semi-infinite line.

## Presentations (continued)

#### Thursday 03/12, 16:30 - 17:00 CET

## **Restart: The Physics Of Starting Anew**

Shlomi Reuveni<sup>1</sup>

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Stopping a process in its midst—only to start it all over again—may prolong, leave unchanged, or even shorten the time taken for its completion. Among these three possibilities, the latter is particularly interesting as it suggests that restart can be used to expedite the completion of complex processes involving strong elements of chance. This turned out to be important in computer science where restart drastically improves performance of randomized algorithms, but is not less relevant to many physical, chemical, and biological processes where restart plays a central role. In this talk, I will provide an introduction to the theory of restart phenomena and review some of its applications in statistical, chemical, and biological physics.

#### References

- [1] S. Reuveni, Phys. Rev. Lett. 116, 170601 (2016).
- [2] A. Pal & S. Reuveni, Phys. Rev. Lett. 118, 030603 (2017).
- [3] A. Pal, I. Eliazar & S. Reuveni, Phys. Rev. Lett. 122, 020602 (2019).
- [4] S. Ray, D. Mondal & S. Reuveni, J. Phys. A. 52, 255002 (2019).
- [5] S. Ray & S. Reuveni, J. Chem. Phys. 152, 234110 (2020).
- [6] O. T. Friedman, A. Pal, A. Sekhon, S. Reuveni & Y. Roichman, J. Phys. Chem. Lett. 11, 7350 (2020).
- [7] A. Pal, Ł. Kuśmierz & S. Reuveni, Phys. Rev. Research 2, 043174 (2020).

## Presentations (continued)

#### Thursday 03/12, 17:00 - 17:20 CET

## Lévy Noise, Time-Reversal Symmetry, Nonequilibrium Stochastic Thermodynamics, and Bak's Sandpile

Steven Yuvan<sup>1</sup> and Martin Bier<sup>1</sup>

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The Brownian noise in nonequilibrium systems commonly contains more outliers. In that case the noise is best described with a Levy distribution. Many systems in which there are fluctuations around a steady-state throughput can be modeled as a Levy-noise-subjected particle in a parabolic potential. We consider an overdamped Brownian particle in a parabolic potential. If the noise is Levy, time-reversal symmetry for the particle's trajectory is violated. We formulate a parameter, r, to express and detect this violation. With solar flare data it is shown how r can be readily obtained and next used to obtain a good estimate of the stability index,  $\alpha$ , of the underlying noise. Self-organized-criticality, i.e. the famous avalanching sandpiles of Per Bak et al, can also be modeled with a Levy-noise-subjected particle on a potential. Preliminary results from such an approach are shown.

## Presentations (continued)

#### Thursday 03/12, 17:20 - 17:40 CET

## Simple mathematical models for self-propelled motion on the water surface

Jerzy Gorecki<sup>1</sup>, Hiroyuki Kitahata<sup>2</sup>, Satoshi Nakata<sup>3</sup> and Yuki Koyano<sup>3</sup>

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Self-propelled motion on the water surface has fascinated scientists for almost two hundred years. There are many solid substances, such as camphor, camphene,

and phenanthroline, which can develop to the water surface as a molecular layer, evaporate to the air phase and continuously dissolve into the water phase. This surface layer decreases the water surface tension and this decrease is local and time-dependent. Inhomogeneities in the surface concentration, resulting from fluctuations in release and evaporation, can break the symmetry around an object and drive its motion. The mathematical description of the self-propelled motion is complex because it should include object location, generated hydrodynamic flows, and dissipation of surface-active molecules. The simplest mathematical models consider the object position together with the concentration of surface-active molecules described by a reaction-diffusion equation in which the effective diffusion coefficient includes a hydrodynamic effect. However, there are many interesting examples of systems where the motion of a self-propelled object is strongly coupled with generated hydrodynamic flows and in such cases, the model mentioned above fails. We introduce another simple mathematical model that describes the time evolution of a self-propelled object on a liquid surface using such variables as the object location, the surface concentration of active molecules, and the hydrodynamic surface flow. The model is applied to simulate the time evolution of a rotor composed of a polygonal plate with camphor pills at its corners. We have qualitatively reproduced results of experiments, in which the inversions of rotational direction under periodic stop- and releaseoperations were investigated. The model correctly describes the probability of the inversion as a function of the duration of the phase when the rotor is stopped. Moreover, the model allows us to introduce the rotor asymmetry unavoidable in real experiments and study its influence on the studied phenomenon. Our numerical simulations have revealed that the probability of the inversion of rotational direction is determined by the competition among the transport of the camphor molecules by the flow, the intrinsic asymmetry of the rotor, and the noise amplitude.

## Presentations (continued)

#### Friday 04/12, 09:00 - 09:30 CET

## Galilean invariance for stochastic diffusive dynamics

Andrea Cairoli<sup>1</sup>, Rainer Klages<sup>2</sup> and Adrian Baule<sup>2</sup>

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Galilean invariance is a cornerstone of classical mechanics. It states that the equations of motion are the same in different inertial frames meaning they do not change under a Galilean transformation. Inertial frames, in turn, are reference frames describing closed systems where the frame-internal physics is not affected by frame-external forces. The description of real world systems, however, usually requires coarse-grained models integrating complex internal and external interactions indistinguishably as friction and stochastic forces, which intrinsically violates Galilean invariance. Starting from the Kac-Zwanzig Hamiltonian for a tracer particle in a heat bath of harmonic oscillators generating Brownian motion, we show how Galilean invariance is broken during the coarse graining procedure when deriving stochastic Langevin dynamics. We argue that traces of Galilean invariance survive for stochastic dynamics yielding a set of alternative rules, which we call weak Galilean invariance.

[1] A.Cairoli, R.Klages, A.Baule, Weak Galilean invariance as a selection principle for coarse-grained diffusive models, PNAS 115, 5714 (2018)

## Presentations (continued)

Friday 04/12, 09:30 - 09:50 CET

## **Anomalous Diffusion and Generalized Cattaneo Equation**

Katarzyna Gorska<sup>1</sup>

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The generalized Cattaneo equation (GTE) involving memory effect is introduced by smearing the time derivatives. Consistency conditions which the smearing functions obey restrict freedom in their choice. The proposed scheme goes beyond the approach based on using fractional derivatives. I present the conditions under which solutions of the GTE can be recognized as probability distribution, i.e. are normalizable and nonnegative on their domain.

## Presentations (continued)

#### Friday 04/12, 09:50 - 10:10 CET

## **Geometrical Optics of Constrained Brownian Motion**

Baruch Meerson<sup>1</sup>

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The optimal fluctuation method - essentially geometrical optics - gives a valuable insight into large deviations of constrained Brownian motion, and it achieves this purpose by simple means. I will illustrate these points by revisiting the Airy distribution - the probability distribution of the area under a Brownian excursion - and considering some additional statistics of Brownian excursions.

The geometrical optics immediately gives the large-area tail of the Airy distribution. It also predicts the position distribution of a Brownian excursion at an intermediate time, conditioned on a large area. Finally, it gives the area distribution on a sub-interval of Brownian excursion. The last two distributions exhibit dynamical phase transitions which have a simple geometric origin.

## Presentations (continued)

Friday 04/12, 10:10 - 10:30 CET

## Effective Langevin equations leading to large deviation function of time-averaged velocity for a nonequilibrium Rayleigh piston

Masato Itami<sup>1</sup>, Yohei Nakayama<sup>2</sup>, Naoko Nakagawa<sup>3</sup> and Shin-ichi Sasa<sup>4</sup>

- <sup>1</sup> Nagoya University
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- <sup>3</sup> Ibaraki University
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We study fluctuating dynamics of a freely movable piston that separates an infinite cylinder into two regions filled with ideal gas particles at the same pressure but different temperatures. To investigate statistical properties of the time-averaged velocity of the piston in the long-time limit, we perturbatively calculate the large deviation function of the time-averaged velocity. Then, we derive an infinite number of effective Langevin equations yielding the same large deviation function as in the original model. Finally, we provide two possibilities for uniquely determining the form of the effective model.

## Presentations (continued)

Friday 04/12, 10:30 - 10:50 CET

## Random walks with asymmetric time delays

Jacek Miękisz<sup>1</sup> and Kamil Łopuszański<sup>1</sup>

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We study simple random-walk models with asymmetric time delays. Probability of a walker to move to the right or to the left depends on a difference between two state-dependent functions evaluated at states of the walker at two different times in the past. This might be seen as a model of a discrete replicator dynamics with strategy-dependent time delays. We assume hyperbolic-tangent fitness functions and to obtain analytical results we approximate them by step functions.

We observe a novel behavior. Namely, the mean position of the walker depends on time delays. This is a joint effect of stochasticity and time delays present in the system. In the deterministic version of a hyperbolic-tangent model, there appear symmetric cycles around a stationary point so the mean position of the walker stays the same. In stochastic versions without time delays, the expected value of the position of the walker is given by the stationary point.

If one interprets a position of the walker as a fraction of the population with a given strategy, then our results show that this fraction is a decreasing function of a delay, in fact a linear one for small delays. Moreover, bigger is the region with the (almost) unbiased random walk, smaller is the fraction of the population of the strategy with a bigger time delay.

## Presentations (continued)

Friday 04/12, 12:00 - 12:20 CET

### Spontaneous symmetry breaking of active phase in coevolving nonlinear voter model

<u>Arkadiusz Jędrzejewski</u><sup>1</sup>, Joanna Toruniewska<sup>2</sup>, Krzysztof Suchecki<sup>2</sup>, Oleg Zaikin<sup>3</sup> and Janusz Hołyst<sup>2</sup>

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A feedback loop between the network topology and dynamical processes that occur between nodes is common in real-world networks. The topology impacts the evolution of node states, which in turn influence the way the structure itself is modified. This feedback is a signature of networks that are called adaptive or coevolutionary. Adaptive networks are especially relevant for social systems, where they can model phenomena such as the emergence of consensus and polarization, opinion formation, or group fragmentation. These coevolutionary models rely on two basic mechanisms. One accounts for the changes in the node states, whereas the other for the link rewiring. Both of them may be implemented in various ways. The voter model, as a minimalist model of opinion formation process, provides the basis for the evolution of state variables in many adaptive networks that represent social interactions. Being analytically tractable, it has played a fundamental role in understanding the process of network fragmentation. This work extends the study in this area by the analysis of one of the nonlinear extensions of the coevolving voter model.

In the analyzed model, each node in the network represents a voter and can be in one of two states that correspond to different opinions shared by the voters. A voter disagreeing with its neighbor's opinion may either adopt it or rewire its link to another randomly chosen voter with any opinion. The system is studied by means of the pair approximation in which a distinction between the average degrees of nodes in different states is made. This approach allows us to identify two dynamically active phases: a symmetric and an asymmetric one. The asymmetric active phase, in contrast to the symmetric one, is characterized by different numbers of nodes in the opposite states that coexist in the network. The pair approximation predicts the possibility of spontaneous symmetry breaking, which leads to a continuous phase transition between the symmetric active phases. In this case, the absorbing transition occurs between the asymmetric active and the absorbing phases after the spontaneous symmetry breaking. Discontinuous phase transitions and hysteresis loops between both active phases

## Presentations (continued)

are also possible. Interestingly, the asymmetric active phase is not displayed by the model where the rewiring occurs only to voters sharing the same opinion, studied by other authors. Our results are backed up by Monte Carlo simulations.

During the talk, we will compare both the versions of the model, and we will see how a seemingly small difference in the link rewiring dynamics leads to profound differences in the phase diagrams exhibited by the models.

## Presentations (continued)

Friday 04/12, 12:20 - 12:40 CET

## The q-voter model with attractive and repulsive interactions and independence on random graphs

Andrzej Krawiecki<sup>1</sup>

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The *q*-voter model with both attractive (roughly speaking, ferromagnetic-like) and repulsive (antiferromagnetic-like) interactions on random graphs is investigated. In this model the agent, represented by a two-state spin located in a node of a graph, with probability 1 - p changes his/her opinion under the influence of a clique of *q* randomly chosen neighbors and with probability *p* acts independently and changes opinion randomly. In the former case the agent changes opinion if opinions of all selected neighbors interacting with him/her attractively via the attached edges of the graph ("friends") are opposite and simultaneously opinions of all selected neighbors interacting with him/her repulsively ("disliked persons") are the same as the agent's one. The parameter *p* measures the level of stochastic noise in the model. For  $q \ge 4$  the model on graphs with large mean degree of nodes exhibits first-order ferromagnetic transition with decreasing *p*, with a clearly visible hysteresis loop. The width of this loop decreases with increasing fraction of the repulsive interactions and the transition can eventually become second-order. For q < 4 the transition is always second-order. An extension of the pair approximation taking into account presence of the repulsive interactions predicts quantitatively well results of Monte Carlo simulations of the model in a broad range of parameters.

## Presentations (continued)

Friday 04/12, 12:40 - 13:00 CET

## Modeling Echo Chambers and Polarization Dynamics in Social Networks

Fabian Baumann<sup>1</sup>, Philipp Lorenz-Spreen<sup>2</sup>, Igor Sokolov<sup>1</sup> and Michele Starnini<sup>3</sup>

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Echo chambers and opinion polarization recently quantified in several sociopolitical contexts and across different social media raise concerns on their potential impact on the spread of misinformation and on the openness of debates. Despite increasing efforts, the dynamics leading to the emergence of these phenomena remain unclear. We propose a model that introduces the dynamics of radicalization as a reinforcing mechanism driving the evolution to extreme opinions from moderate initial conditions. Inspired by empirical findings on social interaction dynamics, we consider agents characterized by heterogeneous activities and homophily. We show that the transition between a global consensus and emerging radicalized states is mostly governed by social influence and by the controversialness of the topic discussed. Compared with empirical data of polarized debates on Twitter, the model qualitatively reproduces the observed relation between users' engagement and opinions, as well as opinion segregation in the interaction network. Our findings shed light on the mechanisms that may lie at the core of the emergence of echo chambers and polarization in social media.

## Presentations (continued)

Friday 04/12, 13:00 - 13:20 CET

## Biological Networks Regulating Cell Fate Choice Are Minimally Frustrated

David Kessler<sup>1</sup>, Shubham Tripathi<sup>2</sup> and Herbert Levine<sup>2</sup>

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Characterization of the differences between biological and random networks can reveal the design principles that enable the robust realization of crucial biological functions including the establishment of different cell types. Previous studies, focusing on identifying topological features that are present in biological networks but not in random networks, have, however, provided few functional insights. We use a Boolean modeling framework and ideas from the spin glass literature to identify functional differences between five real biological networks and random networks with similar topological features. We show that minimal frustration is a fundamental property that allows biological networks to robustly establish cell types and regulate cell fate choice, and that this property can emerge in complex networks via Darwinian evolution. The study also provides clues regarding how the regulation of cell fate choice can go awry in a disease like cancer and lead to the emergence of aberrant cell types.

## Presentations (continued)

Friday 04/12, 13:20 - 13:40 CET

## Continuous time-reversal and the thermodynamic uncertainty relation

Andreas Dechant<sup>1</sup> and Shin-ichi Sasa<sup>1</sup>

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We introduce a continuous time-reversal operation which connects the time-forward and timereversed trajectories in the steady state of an irreversible Markovian dynamics via a continuous family of stochastic dynamics. This continuous time-reversal allows us to derive a tighter version of the thermodynamic uncertainty relation (TUR) involving observables evaluated in different physical systems. Moreover, the family of dynamics realizing the continuous time-reversal contains an equilibrium dynamics halfway between the time-forward and time-reversed dynamics. We show that this equilibrium dynamics, together with an appropriate choice of the observable, turns the inequality in the TUR into an equality. We demonstrate our findings for the example of a two-dimensional rotational flow and discuss the role of the observable close to and far from equilibrium.

## Presentations (continued)

#### Friday 04/12, 13:40 - 14:00 CET

## The role of ergotropy in the quantum thermodynamics

Marcin Łobejko1

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Ergotropy is a state function of a density matrix which physical interpretation is the optimal work that can be extracted through the arbitrary unitary channel. The concept naturally appears in frameworks with implicit work reservoirs (e.g. external fields modeled by time-dependent Hamiltonians), where, in particular, the process of charging and discharging of so-called quantum batteries is studied. We reveal that the same quantity appears in autonomous systems with an idealized model of the work reservoir - a quantum weight. Despite similarities between those two approaches, we reveal a fundamental difference, namely the emergence of the locked energy in coherences, i.e. the quantum part of the state that contributes to ergotropy but cannot be extracted as a work. Furthermore, we prove the relation between the ergotropy and free energy, where the former can be interpreted as a generalization of the latter for systems coupled to finite-size heat baths, such that in the thermodynamic limit the total ergotropy of the system and the heat bath approaches the free energy. As a consequence, we derive the second kind of locked energy, due to the finite-size of the bath, which is given by the difference of free energy between the global passive state and the corresponding equilibrium state.

## Presentations (continued)

Friday 04/12, 16:00 - 16:30 CET

## Thermodynamics in biology: From molecular motors to metabolism

Massimiliano Esposito<sup>1</sup>

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I will describe recent progress in nonequilibrium thermodynamics based on stochastic thermodynamics and chemical reaction network theory. I will discuss the relevance of these results for biology and the many open challenges to be addressed.

## Presentations (continued)

Friday 04/12, 16:30 - 17:00 CET

## On zero-delay synchronization in a network of timed automata modeling cardiac tissue

Danuta Makowiec<sup>1</sup>

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A network of timed automata (NTA), inspired by Greenberg-Hastings cellular automata, can efficiently and faithfully reflects the work of the real pacemaker [1]. In this system, each automaton cyclically switches between three states of a certain length of time: from F firing state of length f, to R refractory state of length r and then to A activity state of length a, what closely follows the phenomenologically known oscillations of myocytes. As a results of interactions between neighboring cells, the self-organization of the oscillating units to a common oscillation emerges which then leads to forming a signal of initiating a contraction of the whole heart. Thus NTA establishes an appropriate platform to model a natural human pacemaker, in which we gain insight into conditions and stages of the synchronization process [2].

The visible effect of synchronization in NTA is the excitation wave, i.e. the emergence of steady move of a cluster formed by automata staying in F state.

The synchronization can be of frequency locking type, when some automata force the neighboring automata to faster switch the state A to F than it is expected by intrinsic automaton's cycle, or of phase-locking type, when neighboring automata exhibit the common difference in oscillatory phase. Thus, the frequency-locking synchronizations result in the fast rhythms. The visible effect of such synchronization is a spiral origin of the excitation wave. The phase-locking synchronization occurs when neighboring automata are one step late in the advancement of the cyclic transitions of intrinsic time periods. A visible effect of such synchronization is propagation of the excitation wave with rhythm comparable to the period of oscillating automata.

The strength of synchronization is due to the reduction of uncorrelated degrees of freedom to a collective mode of operation, which then enables the long-term ordering and coordination of biological processes.

The emergence of zero-delay phase-locking synchronization, the marching squad synchronization, has been observed in NTA when in response to the F state of neighbors, the length of the refractory phase r was increased [1].

On the other hand, in NTA system in which the vagal regulation was modeled as increase in the length a of the activity state in response to the vagal activity, such synchronization is also observed but only when the activity of vagal system often changes [2].

#### Presentations (continued)

The mechanisms behind synchronizing without any delay in spatially distributed systems are not clear and debated. The presentation will be devoted to analyze possible mechanisms leading to the marching squad synchronization in the NTA system modeling the pacemaker.

[1] D. Makowiec "Pacemeaker rhythm through networks of pacemaker automata - a review" Acta Physica Polonica B Proceedings Supplement, 7(2014) 347

[2] D. Makowiec, W. Miklaszewski, J. Wdowczyk, A. Lawniczak "From cellular automata model of vagal control of human right atrium to heart beats patterns" Physica D, to appear, (2020)

#### Presentations (continued)

Friday 04/12, 17:00 - 17:20 CET

# Protein noise floor in gene expression: How cell growth and division contribute to it?

Jakub Jędrak<sup>1</sup> and <u>Anna Ochab-Marcinek<sup>1</sup></u>

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Gene expression in cells is a stochastic process. Experiments have shown that noise in protein levels does not decrease to zero as mean gene expression increases. The origins of the noise floor are still debated. The goal of our study was to check how several basic mechanisms affect the noise floor level. These mechanisms are: Cell-cycle dependent gene expression, translational bursting, random protein partitioning between daughter cells at division, and cell cycle desynchronization resulting in cell-cycle age distribution within the population. Our model gives analytical predictions of the existence of the noise floor and it semi-quantitatively reproduces the shapes of the experimental noise vs. protein concentration plots. We show what additional experimental information is needed for the mean-noise fitting to be unambiguous.

[1] J. Jędrak, A. Ochab-Marcinek, Contributions to the 'noise floor' in gene expression in a population of dividing cells, Scientific Reports, 10, 13533 (2020)

#### Presentations (continued)

#### Friday 04/12, 17:20 - 17:40 CET

#### Symmetries and asymmetries of times to complete a chemical cycle

Artem Ryabov<sup>1</sup>, David Voráč<sup>1</sup>, Dominik Lips<sup>2</sup> and Philipp Maass<sup>2</sup>

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Measurements of durations of nonequilibrium stochastic processes provide valuable information on underlying microscopic kinetics and energetics. Theories for corresponding experiments to date are well-developed for single-particle systems only. Little is known for interacting systems in nonequilibrium environments. We introduce and discuss a basic model for cycle processes interacting with an environment that can be out of thermodynamic equilibrium. We find a surprising richness of cycle time variations with environmental conditions. This manifests itself in unequal cycle times in forward and backward cycle directions, speeding up of backward cycles by interactions, and dynamical phase transitions, where cycle times become multimodal functions of a bias. The model allows us to relate these effects to specific microscopic mechanisms, which can be helpful for interpreting experiments.

#### Presentations (continued)

Friday 04/12, 17:40 - 18:00 CET

# Diffusion limitations and translocation barriers in atomically thin biomimetic pores

Michael Zwolak<sup>1</sup> and Subin Sahu<sup>2</sup>

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Ionic transport in nano- to sub-nano-scale pores is highly dependent on translocation barriers and potential wells. These features in the free-energy landscape are primarily the result of ion dehydration and electrostatic interactions. For pores in atomically thin membranes, such as graphene, other factors come into play. Ion dynamics both inside and outside the geometric volume of the pore can be critical in determining the transport properties of the channel due to several commensurate length scales, such as the effective membrane thickness, radii of the first and the second hydration layers, pore radius, and Debye length. In particular, for biomimetic pores, such as the graphene crown ether we examine, there are regimes where transport is highly sensitive to the pore size due to the interplay of dehydration and interaction with pore charge. Picometer changes in the size, e.g., due to a minute strain, can lead to a large change in conductance. Outside of these regimes, the small pore size itself gives a large resistance even when electrostatic factors and dehydration compensate each other to give a relatively flat - e.g., near barrierless - free energy landscape. The permeability, though, can still be large and ions will translocate rapidly after they arrive within the capture radius of the pore. This, in turn, leads to diffusion and drift effects dominating the conductance. The current thus plateaus and becomes effectively independent of pore free energy characteristics. Measurement of this effect will give an estimate of the magnitude of kinetically-limiting features and experimentally constrain the local electromechanical conditions.

#### Posters

## 

# Detection of interaction and energy exchange with invisible partners in localized Brownian dumbbells

Alexander S. Serov<sup>1</sup>, Christian L. Vestergaard<sup>1</sup> and Jean-Baptiste Masson<sup>1</sup>

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When recording trajectories of biomolecules in crowded media, a question of fundamental interest is whether a tracked molecule interacts with other molecules or structures invisible to the observer. One also wants to know whether energy exchange occurs during such interactions since out-of-equilibrium interactions can be a sign of a specific biological function. Addressing these questions using recorded single-particle trajectories is, in general, an ill-posed problem. In this work, we demonstrate how and when possible interactions and energy transfer between the particles can be detected in an idealized system consisting of two linked localized Brownian particles, where only one of the two trajectories is recorded.

To these means, we designed a Bayesian test for whether the observed trajectory corresponds to a single particle or 2 linked particles. The test leverages Bayesian statistics to compare the evidence for that the input trajectory's power spectrum is generated by an independent particle or a particle with a partner. Extensive numerical simulations allowed us to establish parameter regions, within which such partner detection is possible. As expected, the test is most sensitive to the link strength and the ratio of particle diffusivities. As a byproduct, we also obtained estimates of the localization strength and diffusivities of both particles.

For systems where a difference in diffusivities can be interpreted as a difference in local temperatures, we further extended our approach to determine whether the physical system is out of equilibrium. Using numerical simulations, we identified the parameter range where the distinction can be reliably made from recorded data.

Despite the simplicity of the model, the application of the method to real experimental trajectories may help identify the presence of an interaction partner, the interaction direction, and the presence of potential energy fluxes.

#### Posters (continued)

2

# Statistical physics of inhomogeneous transport equations: first passage to the space-dependent diffusion

<u>Roman Belousov</u><sup>1</sup>, Edgar Roldán<sup>1</sup>, Ricardo Franklin Mergarejo<sup>1</sup>, Ali Hassanali<sup>1</sup> and Muhammad Nawaz Qaisrani<sup>2</sup>

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As the science and modern technologies are seizing the realm of nanoscale systems, characterization of composite materials with solid-fluid and fluid-fluid interfaces has become a problem of broad interdisciplinary interest. Properties of such complex systems, which vary in space over microscopic scales, shape important processes in engineering and biology. Estimation of the inhomogeneous—space-dependent—transport coefficients that characterize these processes, e.g. the diffusion coefficient or the heat conductivity, is however a challenging task. Moreover, two alternative formulations of the inhomogeneous transport equations exist in the literature. Using the theory of statistical physics, in my talk I will show that the two formulations, often regarded as distinct models, are in fact equivalent. In particular, one of them conveniently links the mass diffusion equation with statistics of molecules' first-passage events, which provide an efficient inference technique for the space-dependent diffusion coefficient.

## Posters (continued)

3

### Kramers-like problem for underdamped Levy flights

Karol Capała<sup>1</sup> and Bartłomiej Dybiec<sup>1</sup>

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The noise driven motion in a bistable potential acts as the archetypal model of various physical phenomena. In the weak noise limit, for the overdamped particle driven by a non-equilibrium,  $\alpha$ -stable noise the ratio of forward and backward transition rates depends only on the width of the potential barrier separating both minima. The poster presents analytical and numerical results showing that in the regime of full dynamics the ratio of transition rates depends both on widths and heights of the potential barrier separating minima of the double-well potential.

K. Capała and B. Dybiec, *Underdamped, anomalous kinetics in double-well potentials*, Phys. Rev. E **102**, 052123 (2020).

## Posters (continued)

4

#### Exploring shape space for densest random sequential adsorption packing

Konrad Kozubek<sup>1</sup>, Michał Cieśla<sup>1</sup>, Piotr Kubala<sup>1</sup> and Adrian Baule<sup>2</sup>

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Random sequential adsorption of various shapes built of disks is studied to determine the shape, which follows to densest random packing. Using the evolutionary algorithm to sample the space of shapes, we found that independently of the number of used disks, the optimal shape tends to a triangle with rounded corners. Thus, we run the same algorithm to study packings built of rounded polygons. Both these approaches indicate that the shape building the densest RSA packings is the rounded triangle, for which the packing fraction is  $0.600608 \pm 0.000017$ , which is the highest known value so far.

## Posters (continued)

## 5

#### **Photoluminescence of Complex Systems**

Ambra Lattanzi<sup>1,2</sup>, Giuseppe Dattoli<sup>2</sup> and Giuseppe Baldacchini<sup>2</sup>

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The photoluminescence degradation of the thermally evaporated Alq3 thin films can be described by a four components model based on the Kohlrausch-Williams-Watt (KWW) function. This model improved the agreement between experimental data and the theoretical fit with respect to the four components model based on "pure" exponential functions. In fact, the negative bump in PL intensity decay observed in the reference not-annealed sample is perfectly fitted, and the last part of the time resolved spectroscopy improved its adherence to experimental data both in annealed and not-annealed samples.

A material frame of reference has been introduced in order to investigate the physical nature of the KWW function in relaxation processes and at the same time, explain the peculiar experimental features emerged in the time-spectral PL behaviour. The introduction of a nonlinear time variable, named material clock or material time, overcomes the difficulties carried by the anomalous behaviour considering the process from a bare point of view: the relaxation behaves ideally whereas the material frame of reference is going to be stretched or compressed. This allows to model the PL degradation dynamics as a damped harmonic oscillator. Once the physics has been understood, it is possible to restore the laboratory frame of reference and define the reduced mass, a time-depending function that unlocks the physical meaning of the experimental peculiarities observed in the PL emission.

The results are framed in the context of physical-chemical reactions in order to emphasize how the KWW function is a sum of sub-processes. The physical meaning of these subprocesses can be related to internal and environmental agents interactions under defined physical conditions. These insights highlight the usefulness of the proper mathematical procedures and properties, such as the monotonicity and the complete monotonicity, for investigating the PL emission of this organometallic molecule. Moreover, this method is also promising for describing the photoluminescent processes of similar organic molecules both for basic research and optoelectronic applications.

## Posters (continued)

6

#### **Temperature-dependent Smoluchowski equations**

Nikolai Brilliantov<sup>1</sup>, Alexander Osinsky<sup>1</sup>, Arno Formella<sup>2</sup> and Thorsten Poeschel<sup>3</sup>

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- <sup>2</sup> Department of Computer Science, University of Vigo
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We report a new class of Smoluchowski-like equations for ballistically aggregating particles in a space-uniform system. They naturally emerge when the derivation of the aggregation equations is performed starting from the Boltzmann kinetic equation for the mass-velocity distribution functions [1]. Then the system of equations for the zero-order moments of the distribution functions - the concentrations of aggregates of different size, corresponds to the conventional Smoluchowski equations. The aggregation rate coefficients of these equations depend on the second moments of the distribution function - partial temperatures of the aggregates. In this way, we obtain two coupled systems of equations - for the concentrations and temperatures [1]. We show that, depending on the aggregation probability, these equations provide two main types of scaling. The first one, when all collisions are aggregative leads to Smoluchowski-type evolution with a gradual decrease in temperature. The second one, when only a small fraction of collisions are aggregative, results in a stunning effect of the aggregation with temperature growth. We show that the temperature increase may be stable and perpetual until the system remains large enough for the notion of the temperature to persist. We also estimate the system parameters (restitution coefficient, aggregation probability), which can lead to this type of scaling. The theoretical results are confirmed by the numerical simulations.

[1] N. Brilliantov, T. Poeschel, and A. Formella, Increasing temperature of cooling granular gases, Nature Communications, 9 (2018) 797.

## Posters (continued)

## 7

## Energetics of critical oscillators in active bacterial baths

Ashwin Gopal<sup>1</sup>, Edgar Roldan<sup>2</sup> and Stefano Ruffo<sup>3</sup>

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We investigate the nonequilibrium energetics near a critical point of a non-linear driven oscillator immersed in an active bacterial bath. At the critical point, we reveal a scaling exponent of the average power  $\langle \dot{W} \rangle \sim (D_{\rm a}/\tau)^{1/4}$  where  $D_{\rm a}$  is the effective diffusivity and  $\tau$  the correlation time of the bacterial bath described by Gaussian colored noise. Other features that we investigate are the average stationary power and the variance of the work both below and above the saddle-node bifurcation. Above the bifurcation, the average power attains an optimal, minimum value for finite  $\tau$  that is below its zero-temperature limit. Furthermore, we reveal a finite-time uncertainty relation for active matter, which leads to values of the Fano factor of the work that can be below  $2k_{\rm B}T_{\rm eff}$ , with  $T_{\rm eff}$  the effective temperature of the oscillator in the bacterial bath. We analyze different Markovian approximations to describe the nonequilibrium stationary state of the system. Finally, we illustrate our results in the experimental context by considering the example of driven colloidal particles in periodic optical potentials within a E. Coll bacterial bath.

### Posters (continued)

## 8

#### Electronic relaxation in solution: Exact solution of multistate problems in time domain

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Electronic relaxation in solution is an interesting phenomenon from experimental as well as theoretical point of view. This diffusion controlled process has been theoretically modeled by using Smoluchowski equation with a position dependent sink term. The problem has been solved for different potentials and sink functions. Single state problem where Smoluchowski equation for harmonic potential with a Dirac delta sink is the most general model which has been solved analytically in Laplace domain. Due to the complexity of equation, there is no time domain solution have been reported. In this work, we will give the exact analytical solution for this problem in time domain. We have solved one dimensional Smoluchowski equation for harmonic potential with a sink of ultra-short width. This problem has been solved for single state as well as two state model exactly in time domain. We have used Green's function method to solve the equation and calculated survival probability. For two state model, survival probability for both the states have been calculated explicitly. We have developed a general method to solve single state as well as coupled two state model which can be used to solve many similar problems.

#### Posters (continued)

## 9

# Search efficiency of (discrete) fractional Brownian motion in a random distribution of targets

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Efficiency of search for randomly distributed targets is a prominent problem in many branches of the sciences. For the stochastic process of Lévy walks, a specific range of optimal efficiencies was suggested under variation of search intrinsic and extrinsic environmental parameters. We study fractional Brownian motion as a search process, which under parameter variation generates all three basic types of diffusion, from sub- to normal to superdiffusion. In contrast to Lévy walks, fractional Brownian motion defines a Gaussian stochastic process with power law memory yielding anti-persistent, respectively persistent motion. Computer simulations of this search process in a uniformly random distribution of targets show that maximising search efficiencies sensitively depends on the definition of efficiency, the variation of both intrinsic and extrinsic parameters, the perception of targets, the type of targets, whether to detect only one or many of them, and the choice of boundary conditions. We find that different search scenarios favour different modes of motion for optimising search success, defying a universality across all search situations. Some of our results are explained by a simple analytical model. Having demonstrated that search by fractional Brownian motion is a truly complex process, we propose an over-arching conceptual framework based on classifying different search scenarios. This approach incorporates search optimisation by Lévy walks as a special case.

### Posters (continued)

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#### Efficiency of energy harvesting out of colored Lévy fluctuations, by a harmonic piezoelectric transducer

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Based on studies where a linear piezoelectric energy harvester is subjected to external random mechanical excitations modeled through Gaussian noises (white and colored), it was decided to propose a more general colored noise of the Levy  $\alpha$  - stable type. Analytical, exact and approximate results and numerical results are presented on the characteristics of such excitations and the electrical power that can be extracted from them.

#### Posters (continued)

## 11

# Exploring "inverse stochastic resonance" and nonstandard stochastic resonance with information-theoretic tools

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Some pacemaker neurons show a depression of their mean firing rate for intermediate noise amplitudes, which reminds the response enhancement known as "stochastic resonance" (SR). Even though its analogy with SR goes no further, this phenomenon has been termed "inverse stochastic resonance" (ISR). In a recent work, the ISR phenomenon -as well as its close relative "nonstandard SR" (or more properly, noise-induced activity amplification, NIAA)have been shown to stem from the weak-noise quenching of the initial distribution, in bistable regimes where the metastable state has a larger attraction basin than the global minimum. To further illustrate the mechanism whereby the ISR and NIAA phenomena take place, we revisit the one-variable model that led to that conclusion, exploring it with information-theoretic tools.

## Posters (continued)

## (12)

#### Phase transitions in the q-voter model with generalized anticonformity

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Recent empirical studies provide evidence that so-called social hysteresis [1] is present in animal [1-3] as well as in human societies [4-6], which would suggest that (at least some) phase transitions observed in real social systems are discontinuous. It occurs that discontinuous phase transitions are not that typical in models of opinion dynamics. Within several versions of the *q*-voter model [7], belonging to the class of the binary-state dynamics [8, 9], only continuous phase transitions has been observed, including the original *q*-voter model [10] or the *q*-voter model with anticonformity [11]. However, the change of transition from continuous to discontinuous (for q > 5) has been reported for the *q*-voter model with independence [11].

In [12] we introduce a q-voter model with generalized anticonformity. Previously it was assumed that the size of the unanimous group of influence needed for both conformity and anticonformity is equal [11]. We abandon this unjustified assumption and introduce a generalized model, in which the size of the influence group needed for conformity  $q_c$  and the size of the group needed for anticonformity  $q_a$  are independent variables and in general  $q_c \neq q_a$ .

We consider a system of N voters that form vertexes of an arbitrary network. Each of them is characterized by the dynamical binary variable  $S_i(t) = \pm 1, i = 1, ..., N$  which, in case of social systems, can be interpreted as an opinion on a given subject (yes/no, agree/disagree) at a given time t. In each elementary time step we randomly choose one agent that will reconsider its opinion. With probability p the chosen voter behaves like an anticonformist, whereas with complementary probability 1 - p like a conformist. In any case we randomly choose a group of influence from the nearest neighbours of the agent without repetitions. The size of the group depends on the voter's response to social pressure ( $q_a$  for anticonformity,  $q_c$  for conformity). For  $q_a = q_c = q$  the model reduces to the original q-voter model with anticonformity introduced in [11]. If the group of influence is unanimous, the voter is influenced by the group and adapts to it (in case of conformity) or rebels against it (in case of anticonformity).

### Posters (continued)

We analyse the model on a complete graph using linear stability analysis, numerical methods and Landau's theory. We calculate the analytical formulas for the lower spinodal and the tricritical point for which the phase transition changes from continuous to discontinuous. It has occurred that the generalized model displays both continuous and discontinuous phase transitions depending on the sizes of the groups of influence needed for conformity  $q_c$  and anticonformity  $q_a$ . If the parameter  $q_c$  is sufficiently larger than  $q_a$ , the type of the phase transition changes to discontinuous.

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## Posters (continued)

## (13)

#### Numerical solution of temperature-dependent Smoluchowski equations

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In this talk, we provide a numerical study of the recently developed generalised temperaturedependent Smoluchowski equations. To solve the new complex system, we adapt and improve the low-rank approach of solving large ODE systems. This allows us to quickly find approximate solutions for generalised systems of Smoluchowski equations even when the collision kernels change during aggregation. Our results confirm the analytical predictions of the temperature-dependent scaling, including analytically obtained scaling parameters. We also use a special type of Monte-Carlo simulations (temperature-dependent Monte-Carlo) to plot a phase diagram for different values of aggregation probability. We observe both temperature decrease and temperature increase scaling, as well as find more interesting behaviours at small system times.

## Posters (continued)

## (14)

# Combinatorial aspects of the scattering on the Dirac delta potential

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We study a system where a particle scatter on the periodic Dirac delta potential. The problem is canonical i.e. it is considered during typical quantum mechanics course. There are several approaches to solve the Schrodinger equation for such a systems [1, 2]. However most of solutions rely on different matrix method. We would like to introduce new, combinatorial method by writing system of recursive equations. Based off this solution we write transmission and reflected coefficients of a wave functions of a particle, which is the common way to describe Quantum tunnelling.

The most interesting conclusion from the introduced technique: we reveal combinatorial structure of the typical quantum mechanics problem and write transmission and reflected coefficients using binomial coefficients as the number of combinations with repetition.

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## Posters (continued)

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### Cyclic Kuramoto models and Byzantine attack

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Cyclic, one-dimensional Kuramoto models with various types of interactions are reviewed. Ruch models are less resilient to Byzantine perturbations than a classic solution would suggest.

### Posters (continued)

(16)

# The study of structure influence on diffusion across alginate membranes filled with magnetite

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The combination of polymer and inorganic substance in hybrid membrane results in favourable selectivity and permeability, and consequently such membrane becomes a promising alternative to conventional membrane materials. The resulting membranes show improved chemical, mechanical, and thermal stability and hydrophilic-hydrophobic balance. The objective of this research is to determine whether there is a relation between membrane morphology, which is characterized by different parameters, and the diffusive transport in the membrane. We expect that different amount of magnetite particles in alginate matrix cross-linked by different agents influence on structure and morphology properties and also affects the diffusion and transport properties. In this work, we investigate the morphology of cross-sections of the hybrid alginate membranes filled with various amount of magnetite  $(Fe_3O_4)$  and crosslinked using four different agents: calcium chloride, phosphoric acid, glutaraldehyde and citric acid. A key aspect of showing structure - diffusion relationship can be the study of the simulation of particle motion in a membrane environment, and in our case, it is a simulation of a random walk on the structures of hybrid alginate membranes. For a better understanding of the problem, we model structures of two-dimensional heterogenic membranes which resemble real structures and then simulate random walk on them. The prototype structures of hybrid polymeric membranes are created with the desired quantity, size and distribution of obstacles, which corresponds to the given amount of magnetite in the hybrid alginate membrane. Generated membranes possessing specific parameters are comparable to the real hybrid alginate membranes filled with magnetite and give a real chance to find the relation between diffusion and structure properties. This observation may support a better understanding of structure influence on mass transport through polymer materials.

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#### Posters (continued)

## (17)

#### Diffusion-controlled reactions: Extension of time-dependent Smoluchowski's rate coefficient to reactions in media with relaxation

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Processes involving Brownian motion of small reactants toward much larger static particles (sinks) and subsequent trapping of these reactants by the sinks are very commonly encountered in both nature and in artificial media. Theoretically these processes are described by so-called trapping model of the irreversible bulk diffusion-controlled reactions. To avoid difficulties of classical diffusion theory the hyperbolic diffusion approximation is often used [1, 2]. Within the scope of the telegrapher equation the density distribution of reactants  $\rho(r, t)$  about a spherical sink of radius R obeys the initial boundary-value problem with Smoluchowski's condition on the reaction surface [1]. The primary task for the theory of diffusion-controlled reactions is the calculation of the reaction rate coefficient:

$$k(t) = k_S \frac{R}{\tau_d} \int_0 t \exp\left(-\frac{t-\zeta}{\tau_d}\right) \left.\frac{\partial \rho(r,\zeta)}{\partial r}\right|_{r \to R+} d\zeta.$$
(1)

Hereafter  $k_S = 4\pi RD$  is the Smoluchowski steady state rate coefficient;  $\tau_d$  and D are the relaxation time for the gradients of density distribution of reactants and diffusion coefficient, respectively. We solved the posed problem exactly and, using expression (1), for the time-dependent reaction rate coefficient obtained

$$k(t) = k_S \left\{ \left[ 1 - \exp\left(-\frac{t}{\tau_d}\right) \right] + \frac{R}{\sqrt{\tau_d D}} \left[ \exp\left(-\frac{t}{2\tau_d}\right) I_0\left(\frac{t}{2\tau_d}\right) - \exp\left(-\frac{t}{\tau_d}\right) \right] \right\},\tag{2}$$

where  $I_0(z)$  is the first kind modified Bessel function of order zero. It is important to note that the last term in expression (1) was erroneously omitted in the relevant formula derived by Rice (see expression (289) in page 330 of book [1]). Formula (2) generalizes the classical Smoluchowski result to short time values, which is important for reactions occurring in media with relaxation [2].

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## Posters (continued)

## (18)

# Finite element modelling of atomic force indentation of an animal cell

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We present a model of atomic force microscope indentation measurements using the finite element method. The focus is set on a thorough representation of the complex structure of an animal cell. Crucial constituent is the cell cortex - a stiff layer of cytoplasmic proteins present on the inner side of the cell membrane. It plays a vital role in the mechanical interactions between cells. In our model, the cell cortex is modelled by a three-dimensional solid characterized primarily by its bending stiffness. This approach allows us to interpret the measurements of the mechanical properties of the cells, such as elasticity. During the simulations, we probe a broad range of parameters defining cell properties and experimental conditions. Finally, we derive a simple and closed-form formula that approximates the simulated results with satisfactory accuracy. Our formula is as easy to use as Hertz's function in order to extract cell properties from the measurement, with additional consideration of the cell inner structure.

### Posters (continued)

## (19)

# Steady oscillations in kinetic model of aggregation process with collisional fragmentation

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In this work, we study a kinetic model of aggregation process with collisional fragmentation with use of two efficient implementations of numerical methods: direct simulation Monte Carlo and finite-difference scheme exploiting the low-rank matrix representations of the utilized kinetic coefficients. We concentrate our efforts on the analysis of the solutions for a particular class of non-local aggregation kernels

$$K_{i,j} = i^a j^{-a} + i^{-a} j^a,$$

with multiplicative expression for the fragmentation rates  $F_{i,j} = \lambda K_{i,j}$  with  $0 < \lambda \ll 1$ . For a > 0.5 and  $\lambda < \lambda_c$  never-ending collective oscillations of the aggregates' concentrations take place [1].

The main contribution of this work is cross-validation of our previous observations with the utilization of the well-known stochastic acceptance-rejection method [2] and its modification to an accounting of the fragmentation events.

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#### Posters (continued)

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# Experimental study of the energy flux between two NESS thermostats

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We address the question of energy transport in out-of-equilibrium systems. The experiment consists in two coupled granular gas Non-Equilibrium Steady State (NESS) heat baths, in which 2 cm-scale rotors are imbedded. These Brownian-like mobiles are electro-mechanically coupled with a resistor R, allowing energy to flow between them. The mean flux is non-zero if the temperatures of the baths differ. The resistor R allows measurement of the flux and the temperatures in each bath simultaneously. Varying R, we show that, in the limit of non-dissipative coupling ( $R \rightarrow 0$ ), the heat flux obey the 'eXchange Fluctuation Theorem' (XFT), in a form proposed by Jarzynski and Wojcik in 2004, for the fluctuations of the flux between finite size equilibrium heat baths. This result, here obtained for granular gas thermostats experiment, is nevertheless representative of a large class of NESS systems.

## Posters (continued)

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# Determination of psychotic behaviour using a network of chemical oscillators

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Schizophrenia is the most common form of psychotic behaviour where patients experiences hallucination, dillusion or chaotic speech. Schizophrenia is difficult to detect and easily go undetected for years. Here we propose the idea of detecting schizophrenia by a network of interacting chemical oscillators. We optimized a classifier based on six interacting oscillator using genetic algorithm and obtained 82% accuracy of schizophrenia detection on a selected training dataset.

Keywords: Schizophrenia, EEG signal, chemical computing, oscillatory network, Oregonator model, genetic optimization

### Posters (continued)

(22)

#### Force spectroscopy in the study of endometrial diseases

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The endometrium is the tissue lining the uterus cavity. The most important role of this tissue is enabling the embryo to implant and providing it with good conditions for growth and development. According to many literature reports, cell and tissue mechanical properties have a significant role in many disease states. Changes in cell properties such as elasticity are observed in the case of various cancers or blood diseases.

The main technique for cell and tissue mechanical properties analysis is Force Spectroscopy. The force curves obtained during indentation reflect the interaction between the surface of the sample surface and the measuring probe. The course of these curves allows the determination of such properties as adhesion and Young modulus.

In this poster, we present results of research aimed at checking the relationship between Young's modulus of endometrial tissue and women's receptivity.

## Posters (continued)

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#### Moments of the Van Hove dynamic scattering law

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Studying the dynamics of a system at the atomic level provides important information about the behaviour of the system. For example, studying the dynamics of enzymes allows us to understand their biological function. Among the available techniques, studying the dynamics of a system by means of neutron scattering is of exceptional meaning, as neutrons scatter at the nuclei themselves. Van Hove's functions  $G_s(r,t)$  and  $G_d(r,t)$ , as well its Fourier transforms  $I_s(k,t)$  and  $I_d(k,t)$  contain all accessible information on the dynamics of the system. In order to find the function  $I_s(k,t)$  for small times we use its lowest time derivatives taken at the time t = 0. These are also the moments of the scattering law  $S_s(k,\omega)$ . Moments of the odd orders are equal to zero, what results from the invariance of the mechanics equations with respect to the time inversion. We give even moments up to the tenth order.

## Posters (continued)

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# Generalised 'Arcsine' laws for run-and-tumble particle in one dimension

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The 'Arcsine' laws of Brownian particles in one dimension describe distributions of three quantities: the time  $t_m$  to reach maximum position, the time  $t_r$  spent on the positive side and the time  $t_l$  of the last visit to the origin. Interestingly, the cumulative distribution of all three quantities are the same and given by Arcsine function. In this paper, we study distribution of these three times  $t_m$ ,  $t_r$  and  $t_l$  in the context of single run-and-tumble particle in one dimension, which is a simple non-Markovian process. We compute exact distributions of these three quantities for arbitrary time and find that all three distributions have a delta function part and a non-delta function part. Interestingly, we find that the distributions of  $t_m$  and  $t_r$  are identical (reminiscent of the Brownian particle case) when the initial velocities of the particle are chosen with equal probability. On the other hand, for  $t_l$ , only the non-delta function part is the same as the other two. In addition, we find explicit expressions of the joint distributions of the maximum displacement and the time at which this maxima occurs.

## Posters (continued)

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#### Express your scientific track record in just three numbers

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The growing popularity of bibliometric indexes goes hand in hand with their critique by those who claim that scientist's impact cannot be reduced to a single number. Some even believe that such a complex reality fails to submit to any quantitative description. We argue that neither of the two controversial extremes is true. With our new agent-based model (doi.org/10.1073/pnas.2001064117) we can describe the emergence of citation curves very accurately.

In the model we assumed that some citations are distributed according to the rich get richer rule while some others are allocated totally at random. This yields a very accurate model that is governed by merely three easily interpretable parameters: productivity, total impact, and rho, which measures preferential vs. random rules ratio.

#### Posters (continued)

## (26)

#### Nanoscale Lubrication in Model Biosystems as Rationalized in Terms of Fractons and Spectral-Mechanical Properties of Networked Biopolymers in Solutions

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A concept of a biopolimer network immersed within an aqueous solution as addressed in terms of flexibility vs. mechanical stability criterion has been proposed. It is based on a transmission of correlated wave of (hydrogen) ions emerging from breaking in a massive way the hydrogen bonds between the biopolymers, such as hyaluronan, and their non-ideal agueous solution's surroundings. Based on the argumentation presented in a paper by Reuveni et al. [1] it has been demonstrated that there exists a clear connection between the  $\ln(N)$  (a natural logarithm of the biopolymer length N) and an inverse of a difference between two major contributions of this Landau-Peierls instability type paradigm. Providing that the so-called Alexander-Orbach conjecture for the oscillating biopolymeric system applies [2,3] one of the contributions is of mechanical nature, with an exponent q represented by 1/(2-3q) whereas the other appears to be a surface-to-volume characteristic exponent, attaining preferentially a value of ca. 2/3 for a three-dimensional adjacent (articulating) space. It has been shown in a numerical way that for N of the order of milion(s) biopolymer's residues, for example for hyaluronan equivalent to its molecular weight of  $10^6 - 6 \times 10^6$  Daltons, a measure of the best viscoelastic efficiency for the hydronan, there exists an equality  $\ln(N) = b/[1/(2-3g)-2/3]$ that, for example, for  $N = 10^6$  gives the value of mechanical exponent q close to 1/3, yielding according to [3,4], an excellent passage of the (hydrogen) ions' wave derived from a breakage of the adjacent hydrogen bonds in the biopolymer-solution system of interest, provided that the constant b, according to [1], can be taken at b = 4.5. (In general, for the exponent g > 1/6holds.) The first results seem to be promising when thoroughly rationalizing nanoscale frictionlubrication properties of biopolymer-solution articulating/confined subspaces exposed to very small nano-Newton loading conditions. For another thermomechanical scenario describing phase-transition and relaxation kinetics of a biopolymeric system, see [5].

## Posters (continued)

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## Posters (continued)

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#### The effect of substrate roughness on random sequential adsorption packing properties

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Adsorption on smooth and atomistically flat surface is well understood, while still very few works concern adsorption on rough and heterogeneous surfaces. In this study, we investigate the properties of packings of balls obtained via random sequential adsorption (RSA) on a sinusoidal substrate. Our main goal is to observe surface inhomogeneities on a scale smaller than particles' size. We study how various characteristics, such as packing fraction, packing growth kinetics, available surface function and 2-particle density correlation function depend on the wavenumber and the amplitude of a sine-wave-shaped surface. These quantities indeed allow us to discern very minute surface impurities from the packing alone.

## Posters (continued)

## (28)

# Controlling particle currents with evaporation and resetting from an interval

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We investigate the Brownian diffusion of particles in one spatial dimension and in the presence of finite regions within which particles can either evaporate or be reset to a given location. For open boundary conditions, we highlight the appearance of a Brownian yet non-Gaussian diffusion: at long times, the particle distribution is non-Gaussian but its variance grows linearly in time. Moreover, we show that the effective diffusion coefficient of the particles in such systems is bounded from below by  $(1-2/\pi)$  times their bare diffusion coefficient. For periodic boundary conditions, i.e., for diffusion on a ring with resetting, we demonstrate a "gauge invariance" of the spatial particle distribution for different choices of the resetting probability currents, in both stationary and non-stationary regimes. Finally, we apply our findings to a stochastic biophysical model for the motion of RNA polymerases during transcriptional pauses, deriving analytically the distribution of the length of cleaved RNA transcripts and the efficiency of RNA cleavage in backtrack recovery.