

Piecewise Deterministic Markov Processes and Bacterial Growth

Bertrand Cloez, Benoîte de Saporta , **Nathalie Krell** ,Tristan Roget,
Emeline Schmisser

Abstract

This presentation will focus on the study of piecewise deterministic Markov process (PDMP) models used to describe dynamic phenomena involving occasional random events. In the first part, I will present work done in collaboration with Emeline Schmisser [6] on the nonparametric estimation of the jump rate of a deterministic PDMP based on observations of the embedded Markov chain (Z_k) . We construct an adaptive estimator of the jump rate λ and establish an L^2 norm risk bound as well as a quasi-minimax result.

In a second part, I will focus on a bacterial growth model based on a PDMP-type multi-type branching process. Each cell grows exponentially at an individual rate, then divides at a rate depending on its size. This work extends a model introduced by Doumic, Hoffmann, Krell, and Robert [2] by integrating two cell types according to pole position. We show that the process is well defined, satisfies a “many-to-one” formula, and that its average empirical measure converges to a growth-fragmentation equation with multiple state variables. I will conclude with perspectives on estimating the division rate in this framework, in collaboration with Benoîte de Saporta, Bertrand Cloez, and Tristan Roget [1]. To this end, I will rely on a proceedings paper [4] in which I constructed a two-type branching process to model bacterial growth.

References

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- [4] N. Krell 2025 *Branching processes and bacterial growth*. To appear Proceedings IWBPA24.
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