Jakub Pękalski College of Inter-Faculty Individual Studies in Mathematics and Natural Sciences, University of Warsaw E-mail: japek@wp.pl Savas Tay Department of Biosystems Science and Engineering Swiss Federal Institute of Technology ETH Zürich E-mail: savas.tay@bsse.ethz.ch Tomasz Lipniacki Division of Modeling in Biology and Medicine Institute of Fundamental Technological Research Polish Academy of Sciences E-mail: tlipnia@ippt.gov.pl

## Stochastic switching of NF- $\kappa B$ system

NF- $\kappa$ B is a key transcription factor controlling immune responses such as inflammation, proliferation and apoptosis. Its regulatory system is tightly controlled by the numerous feedback loops. We pursue our earlier studies [1, 2] by considering not only two negative loops mediated by NF- $\kappa$ B inducible inhibitors I $\kappa$ B $\alpha$  and A20, which assures the oscillatory responses of NF- $\kappa$ B, but also a positive feedback loop mediated by the NF- $\kappa$ B inducible cytokine TNF $\alpha$ . This loop is negligible in many cell lines, but may become, as suggested by our study, dominant in immune cells like monocytes or macrophages that have a high level of TNF $\alpha$  expression.

## References

- T. Lipniacki, K. Puszynski, P. Paszek, A. R. Brasier, M. Kimmel, Single TNFα trimers mediating NF-κB activation: Stochastic robustness of NF-κB signaling, BMC Bioinformatics 8 (2007), 376.
- [2] S. Tay, J. Hughey, T. Lee, T. Lipniacki, M. Covert, S. Quake, Single-cell NF-κB dynamics reveal digital activation and analogue information processing, Nature 466 (2010), 267–271.