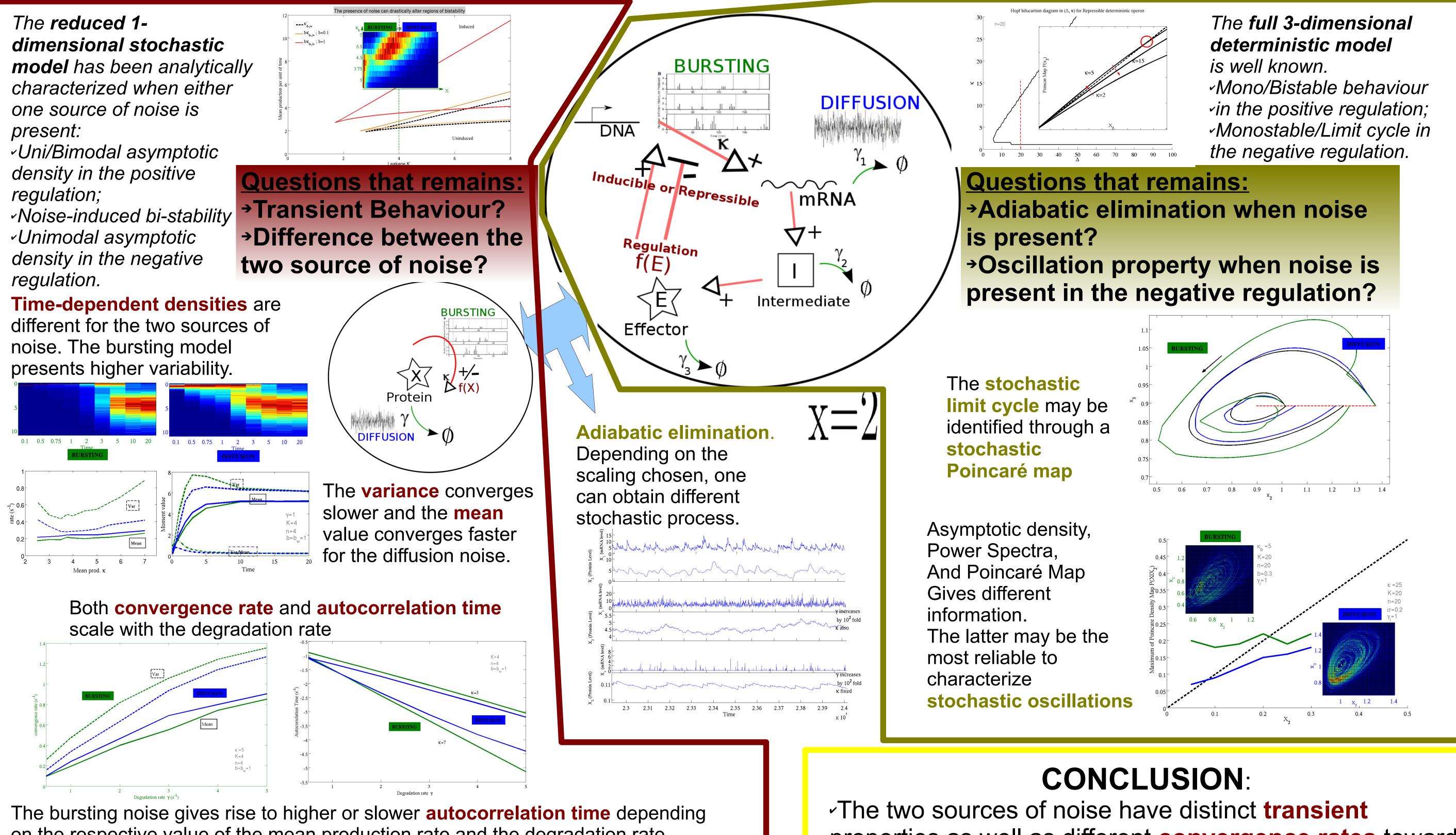
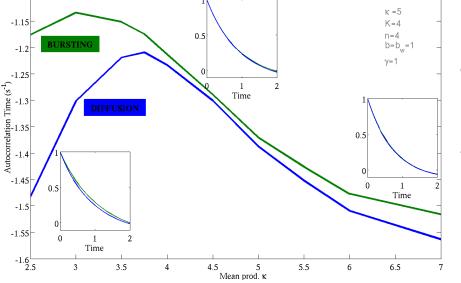
Getting to understand more and more a (simple?) model of stochastic gene expression Self-Regulation White noise Two different sources of noise

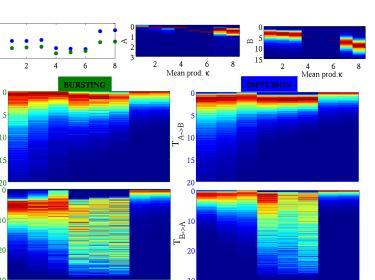
Using Sample paths simulation



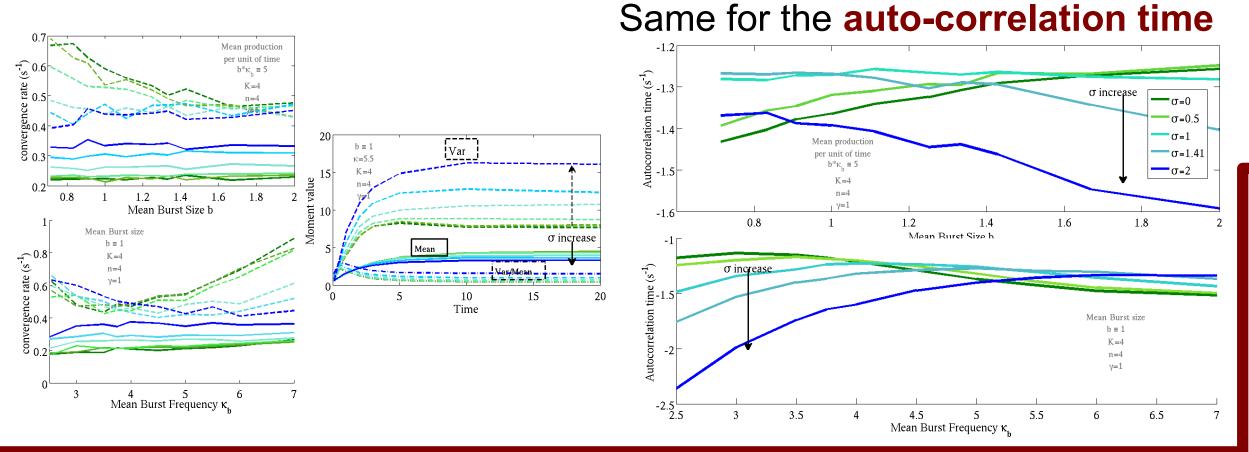
on the respective value of the mean production rate and the degradation rate.



The diffusion model **switch** more an faster, specifically due to a non-refractory time in the bursting model while going from a high to a low value (the bursting noise is asymmetric)



When both noise are present, the strength of the noise governs mainly the **convergence rate** of the **mean**. For the **variance**, we can observe non-monotonic behaviour and more complex dependencies of the two source of noise.



The two sources of noise have distinct transient properties as well as different convergence rates toward equilibrium; Memory and switch property also differs;

- Noisy and bistable regime may occur transiently;
- Key scaling parameters has been identified;
- Oscillations and limit cycle property is being characterized.
- *Interplay between the two source of noise is still mysterious;
- *Interpretation in terms of biology: possible interpretation as Intrinsic and Extrinsic noise?
- *Adiabatic reduction not fully understood. Will some properties such bistable regime be transmitted? *Analytical results....

References:

Molecular distributions in gene regulatory dynamics.

Mackey MC, Tyran-Kamińska M, Yvinec R. J Theor Biol. 2011 274(1):84-96

On the Poincaré—Hill cycle map of rotational random walk: locating the stochastic limit cycle in a reversible Schnakenberg model, Vellela M. and Qian H., Proc. R. Soc. 2010 vol. 466 no. 2115 771-78