

Sensitivity analysis for the Chemical Master Equation

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ANR project OPT-MC

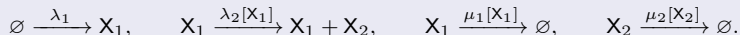
(OPTimal control of Markov Chains & OPTogenetics control of Microbial Communities)

between DISCO¹ team and LIFEWARE² team



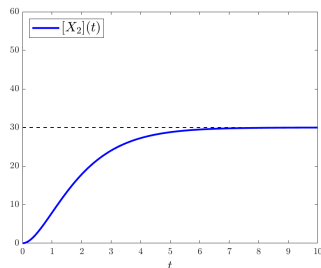
- The Chemical Master Equation (CME)
- A toy example of optimal control problem for the CME
- Sensitivity analysis for the CME

Simplified protein-production model ($X_1 = \text{mRNA}$ & $X_2 = \text{protein}$)

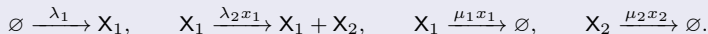


Chemical kinetic equation (deterministic point of view)

$$\frac{d}{dt}[X_1] = \lambda_1 - \mu_1[X_1], \quad \frac{d}{dt}[X_2] = \lambda_2[X_1] - \mu_2[X_2].$$

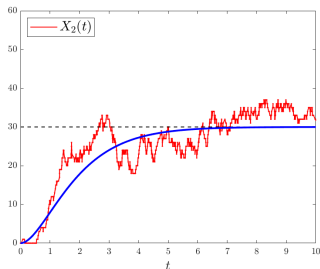


Simplified protein-production model ($X_1 = \text{mRNA}$ & $X_2 = \text{protein}$)



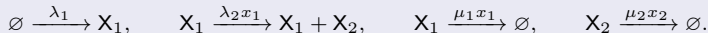
Chemical Master Equation (stochastic point of view)

- ▷ We denote by $X(t) = (X_1(t), X_2(t))$ the state of the system at time t .
 - ▷ We assume that the instants at which reactions occur are random, so that $X(t)$ is a **continuous-time Markov chain** on \mathbb{N}^2 .
 - ▷ Each **reaction** R_r is defined by its **propensity** $a_r(x)$ and its **stoichiometric vector** ν_r .
- ↪ If the system is in state x , the probability that R_r fires in time Δt is equal to $a_r(x)\Delta t$.
- Ex: For the 2nd reaction, we have $a_2(x) = \lambda_2 x_1$ and $\nu_2 = (0, 1)$.



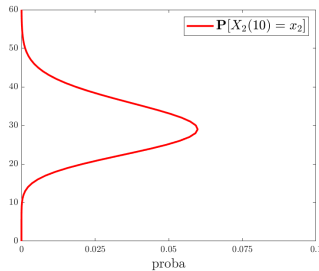
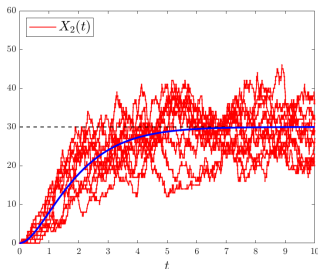
Introduction

Simplified protein-production model ($X_1 = \text{mRNA} \ \& \ X_2 = \text{protein}$)



Chemical Master Equation (stochastic point of view)

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Chemical Master Equation

- ▷ The law of $X(t)$ satisfies a Kolmogorov equation, called the Chemical Master Equation (CME) in this context,

$$\frac{d}{dt}\mathbb{P}[X(t) = x] = \sum_{r=1}^4 \left(a_r(x - \nu_r) \mathbb{P}[X(t) = x - \nu_r] - a_r(x) \mathbb{P}[X(t) = x] \right), \quad x \in \mathbb{N}^2.$$

↪ Thus, the CME is a **countable collection of coupled ODEs** indexed by $\Omega = \mathbb{N}^2$.

General CME

For a set of R reactions involving d species, we set $\Omega = \mathbb{N}^d$ and $p(t, x) = \mathbb{P}[X(t) = x]$, then

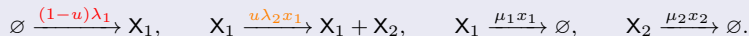
$$\dot{p}(t, x) = \sum_{r=1}^R \left(a_r(x - \nu_r) p(t, x - \nu_r) - a_r(x) p(t, x) \right), \quad x \in \Omega.$$

In what follows, we rewrite the CME in matrix form as

$$\dot{p}(t) = p(t) \mathbf{Q} \quad \text{and} \quad p(0) = p_0,$$

with $p(t)$ the vector of the probabilities $p(t, x)$ and \mathbf{Q} the matrix of the propensities $a_r(x)$.

Simplified protein-production model (X_1 = mRNA & X_2 = protein)



- ▷ The **control** u (e.g. a light stimulus) acts globally on the system to:
 - **inhibit the growth of mRNAs** X_1 .
 - **strengthen the production of proteins** X_2 .
- ▷ We may want to maximize the expected number of proteins X_2 at time T , i.e. solve

$$\max_u \varphi(u), \quad \text{with} \quad \varphi(u) = \sum_{x \in \Omega} x_2 p_u(T, x),$$

and where p_u is solution of the controlled CME

$$\dot{p}(t) = p(t)Q_u \quad \text{and} \quad p(0) = p_0.$$

- ↪ To derive the **optimality conditions** for this problem, we need to perform a **sensitivity analysis** of the CME w.r.t. the matrix Q .

CME with source term

In what follows, we denote by $P[\mathbf{Q}, p_0, f]$ the solution of the CME

$$\dot{p}(t) = p(t)\mathbf{Q} + f(t) \quad \text{and} \quad p(0) = p_0.$$

Existence and uniqueness

- ▷ Depending on the growth of $a_r(x)$ as $\langle x \rangle \rightarrow +\infty$, $P[\mathbf{Q}, p_0, f]$ might not be unique.
 \leadsto If $a_r(x)$ grows at most linearly as $\langle x \rangle \rightarrow +\infty$, then $P[\mathbf{Q}, p_0, f]$ is unique.
- ▷ In any case, we can work with the so-called minimal solution.
 \leadsto The constructive definition of the minimal solution can be used to get energy estimates.

Minimal solution

- ▷ Let $(\omega_n)_{n \in \mathbb{N}}$ be any sequence of finite subsets of Ω s.t. $\omega_n \uparrow \Omega$ as $n \rightarrow +\infty$.
- ▷ Then, the minimal solution $P[\mathbf{Q}, p_0, f]$ is defined by

$$P[\mathbf{Q}, p_0, f] = \lim_{n \rightarrow +\infty} \uparrow P_n[\mathbf{Q}, p_0, f],$$

where $P_n[\mathbf{Q}, p_0, f]$ is the solution of the restricted CME

$$\dot{p}(t) = p(t)\mathbf{Q}|_{\omega_n} + f(t)|_{\omega_n} \quad \text{and} \quad p(0) = p_0|_{\omega_n}.$$

Functional setting

- ▷ Since the solution of the CME is a **probability mass function**, it naturally belongs to the space ℓ^1 of vectors $p \in \mathbb{R}^\Omega$ s.t. $\sum_{x \in \Omega} p(x) < +\infty$.
- ▷ Moreover, we set:
 - ℓ_1^1 the subspace of $p \in \ell^1$ with **finite expected value**, i.e. s.t. $\sum_{x \in \Omega} x_i p(x) < +\infty$;
 - ℓ_2^1 the subspace of $p \in \ell^1$ with **finite variance**, i.e. s.t. $\sum_{x \in \Omega} x_i^2 p(x) < +\infty$;
 - etc...

Energy estimates

Let $P[\mathbf{Q}, p_0, f]$ be the minimal solution of the CME

$$\dot{p}(t) = p(t)\mathbf{Q} + f(t) \quad \text{and} \quad p(0) = p_0.$$

For $m \in \mathbb{N}$, there exists $C_m \geq 0$ s.t.

$$\|P[\mathbf{Q}, p_0, f](t)\|_{\ell_m^1} \leq C_m \left(\|p_0\|_{\ell_m^1} + \|f\|_{L^1(\ell_m^1)} \right), \quad t \in [0, T].$$

Sensitivity w.r.t. \mathbf{Q}

- ▷ Let $\mathbf{Q}, \overline{\mathbf{Q}}$ be CME matrices and $p_0 \in \ell_1^1$.
- ▷ We denote $p = P[\mathbf{Q}, p_0, 0]$ and $\bar{p} = P[\overline{\mathbf{Q}}, p_0, 0]$. We then have

$$p - \bar{p} = P[\overline{\mathbf{Q}}, 0, p(\mathbf{Q} - \overline{\mathbf{Q}})]$$

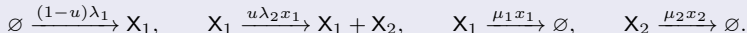
and there exists $C \geq 0$ s.t.

$$\|p(t) - \bar{p}(t)\|_{\ell^1} \leq C \|p_0\|_{\ell_1^1} \|\mathbf{Q} - \overline{\mathbf{Q}}\|_{\ell_1^1, \ell^1}, \quad t \in [0, T].$$

- ⚠ Since the propensities $a_r(x)$ can grow linearly as $\langle x \rangle \rightarrow +\infty$, \mathbf{Q} and $\overline{\mathbf{Q}}$ are only bounded from ℓ_1^1 to ℓ^1 .

Sensitivity w.r.t. u

Simplified protein-production model ($X_1 = \text{mRNA}$ & $X_2 = \text{protein}$)



▷ Here, the CME matrix is of the form $\mathbf{Q}_u = \mathbf{Q}_0 + u \delta \mathbf{Q}$, where $\delta \mathbf{Q} = \mathbf{Q}_1 - \mathbf{Q}_0$.

Sensitivity w.r.t. u

▷ Let $u, \bar{u} \in L^\infty(0, T; [0, 1])$ and $p_0 \in \ell_2^1$.

▷ We denote $p_u = \mathbf{P}[\mathbf{Q}_u, p_0, 0]$ and $p_{\bar{u}} = \mathbf{P}[\mathbf{Q}_{\bar{u}}, p_0, 0]$. We then have

$$p_u - p_{\bar{u}} = \mathbf{P}[\mathbf{Q}_{\bar{u}}, 0, (u - \bar{u})p_{\bar{u}}\delta\mathbf{Q}] + \mathbf{P}[\mathbf{Q}_{\bar{u}}, 0, (u - \bar{u})(p_u - p_{\bar{u}})\delta\mathbf{Q}].$$

▷ The 1st-order term $\mathbf{P}[\mathbf{Q}_{\bar{u}}, 0, (u - \bar{u})p_{\bar{u}}\delta\mathbf{Q}]$ is linear w.r.t. $u - \bar{u}$.

▷ For the 2nd-order term, there exists $C' \geq 0$ s.t.

$$\left\| \mathbf{P}[\mathbf{Q}_{\bar{u}}, 0, (u - \bar{u})(p_u - p_{\bar{u}})\delta\mathbf{Q}](t) \right\|_{\ell_1} \leq C' \|p_0\|_{\ell_2^1} \|u - \bar{u}\|_{L^\infty}^2, \quad t \in [0, T].$$

↪ For the cost $\varphi(u) = \sum_{x \in \Omega} x_2 p_u(T, x)$, we thus have

$$\varphi(u) - \varphi(\bar{u}) = \sum_{x \in \Omega} x_2 \mathbf{P}[\mathbf{Q}_{\bar{u}}, 0, (u - \bar{u})p_{\bar{u}}\delta\mathbf{Q}](T, x) + o(\|u - \bar{u}\|_{L^\infty}).$$

▷ Summary:

- We consider a toy example of **optimal control problem** for the CME.
- We use the notion of **minimal solution** to perform a **sensitivity analysis** for the CME.
- This analysis leads to a **Taylor expansion of the cost** w.r.t. the control and ultimately to **optimality conditions**.

▷ Perspectives:

- Study relevant simple biological models.
~ grower/producer model, bursty protein production, ...
- Solve these optimal control problems numerically.

Thank you for your attention