Traditional kinetic models	Monte-Carlo Markov Chain	Kinetic Monte-Carlo models	References	Annexes
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Kinetic Monte-Carlo models

Ruben Staub

ENS de Lyon

Friday 18, 2018

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- Derivations
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Unimolecular decomposition				



Figure: Reactant is adsorbed on an unoccupied adsorption site, independently of local coverage.

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Unimolecular decomposition				



Figure: Adsorbed reactant at an adsorption site does not feel the influence of other adsorbed species.

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Unimolecular decomposition				



Figure: Reactant is decomposed and rapidly desorbed.

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Unimolecular decomposition				

Unimolecular decomposition

Reaction considered:

$$A(g) + S_{free} \xrightarrow[k_{-1}]{k_1} A(ads) \xrightarrow{k_2} Products(g) + S_{free}$$

Assuming elementary steps, the total reaction rate r is:

$$r = -\frac{dP_{\rm A}}{dt} = k_2[{\rm A}({\rm ads})] = k_2\theta[{\rm S}]$$
(1)

where θ is the surface coverage (i.e. fraction of sites occupied), and [S] is the surface concentration of adsorption sites

Applying a steady-state approximation to A (ads), we get:

$$r = \frac{k_1 k_2 P_{\rm A}[{\rm S}]}{k_1 P_{\rm A} + k_{-1} + k_2}$$
(2)

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Unimolecular decomposition				

Limitations

This approach has multiple limitations/approximations:

- Only unimolecular decomposition is considered
- Steady-state approximation of adsorbed species
- All catalytic sites are equivalent
- Adsorption sites can only be occupied by the reactant
- Each site can be occupied by at most one adsorbate
- The adsorption and decomposition/desorption steps are considered elementary

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- No spatial correlation/lateral interactions
- Kinetic constants are independent of coverage

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Langmuir-Hinshelwood mode				



Figure: Reactants are adsorbed on unoccupied adsorption sites, independently of local coverage.

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Langmuir-Hinshelwood model				



Figure: Adsorbed reactants at an adsorption site does not feel the influence of other adsorbed species, and diffuse freely over the surface.

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Langmuir-Hinshelwood mode				



Figure: Bimolecular reaction occurs and product is rapidly desorbed.

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Langmuir-Hinshelwood model				

Reaction rate

$$\begin{array}{l} \mathsf{A}(\mathsf{g}) + \mathsf{S}_{\mathsf{free}} \xrightarrow[k_{-1}]{k_{-1}} \mathsf{A}(\mathsf{ads}) \\ \texttt{considered:} \qquad \mathsf{B}(\mathsf{g}) + \mathsf{S}_{\mathsf{free}} \xrightarrow[k_{-2}]{k_{-2}} \mathsf{B}(\mathsf{ads}) \\ \\ \mathsf{A}(\mathsf{ads}) + \mathsf{B}(\mathsf{ads}) \xrightarrow[k_{-3}]{k_{-3}} \mathsf{AB}(\mathsf{g}) \end{array}$$

Reactions considered

Assuming elementary steps, the total reaction rate r is:

$$r = -\frac{dP_{\rm A}}{dt} = -\frac{dP_{\rm B}}{dt} = k_3 \theta_{\rm A} \theta_{\rm B} [\rm S]^2 \tag{3}$$

Applying a steady-state approximation to A(ads) and B(ads), and considering the rate determining step being the reaction/diffusion, we get:

$$r = \frac{K_{\rm A}K_{\rm B}P_{\rm A}P_{\rm B}[{\rm S}]^2}{(1 + K_{\rm A}P_{\rm A} + K_{\rm B}P_{\rm B})^2} \tag{4}$$

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Langmuir-Hinshelwood model				

Limitations

This approach also has multiple limitations/approximations:

- Steady-state approximation of adsorbed species
- Reaction/diffusion between A(ads) and B(ads) is the rate determining step
- All catalytic sites are equivalent
- Adsorption sites can only be occupied by the reactants
- Each site can be occupied by at most one adsorbate
- The adsorption and decomposition/desorption steps are considered elementary
- No spatial correlation/lateral interactions
- Kinetic constants are independent of coverage

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Traditional kinetic models	Monte-Carlo Markov Chain	Kinetic Monte-Carlo models	References	Annexes
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Metropolis-Hastings Monte-C	arlo			

 Multiple MCMC (Monte-Carlo Markov Chain) methods for generating samples from a probability distribution P(S).

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Traditional kinetic models	Monte-Carlo Markov Chain	Kinetic Monte-Carlo models	References	Annexes
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Metropolis-Hastings Monte-C	arlo			

 Multiple MCMC (Monte-Carlo Markov Chain) methods for generating samples from a probability distribution P(S).

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Based on random walk onto a Markov chain, with initial proposal distribution $g(S_i \rightarrow S_j)$.

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Traditional kinetic models	Monte-Carlo Markov Chain	Kinetic Monte-Carlo models	References	Annexes
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Metropolis-Hastings Monte-C	arlo			

- Multiple MCMC (Monte-Carlo Markov Chain) methods for generating samples from a probability distribution P(S).
- Based on random walk onto a Markov chain, with initial proposal distribution $g(S_i \rightarrow S_j)$.
- Reaches asymptotically a stationary distribution $\pi(S)$, such that $\forall S, \pi(S) = P(S)$.

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Metropolis-Hastings Monte-C	arlo			

- Multiple MCMC (Monte-Carlo Markov Chain) methods for generating samples from a probability distribution P(S).
- Based on random walk onto a Markov chain, with initial proposal distribution $g(S_i \to S_j)$.
- Reaches asymptotically a stationary distribution π(S), such that ∀S, π(S) = P(S).
- Requires only the computation of an estimator W(S) proportional to P(S), over all possible S.

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Metropolis-Hastings Monte-C	arlo			



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Traditional kinetic models	Monte-Carlo Markov Chain	Kinetic Monte-Carlo models	References	Annexes		
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Metropolis- Hastings Monte-Carlo						

 A next candidate S_j is selected from the current state S_i using the proposal distribution g.

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MC models			

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Traditional kinetic models	Monte-Carlo Markov Chain	Kinetic Monte-Carlo models	References	Annexes		
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Metropolis- Hastings Monte-Carlo						

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KMC models

- A next candidate S_j is selected from the current state S_i using the proposal distribution g.
- An acceptance probability is computed as:

$$A(\mathcal{S}_i, \mathcal{S}_j) = \min\left(\frac{W(\mathcal{S}_j)g(\mathcal{S}_i \to \mathcal{S}_j)}{W(\mathcal{S}_i)g(\mathcal{S}_j \to \mathcal{S}_i)}, 1\right)$$

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Traditional kinetic models	Monte-Carlo Markov Chain	Kinetic Monte-Carlo models	References	Annexes		
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Metropolis- Hastings Monte-Carlo						

- A next candidate S_j is selected from the current state S_i using the proposal distribution g.
- An acceptance probability is computed as:

$$A(\mathcal{S}_i, \mathcal{S}_j) = \min\left(\frac{W(\mathcal{S}_j)g(\mathcal{S}_i \to \mathcal{S}_j)}{W(\mathcal{S}_i)g(\mathcal{S}_j \to \mathcal{S}_i)}, 1\right)$$

In practice the targeted probability distribution P(S) is usually a Boltzmann distribution. No need for partition function, as $W(S_i) = e^{\frac{-S_i}{kT}} \propto P(S_i)$ is enough.

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Traditional kinetic models	Monte-Carlo Markov Chain	Kinetic Monte-Carlo models	References	Annexes		
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Metropolis- Hastings Monte-Carlo						

- A next candidate S_j is selected from the current state S_i using the proposal distribution g.
- An acceptance probability is computed as:

$$A(\mathcal{S}_i, \mathcal{S}_j) = \min\left(\frac{W(\mathcal{S}_j)g(\mathcal{S}_i \to \mathcal{S}_j)}{W(\mathcal{S}_i)g(\mathcal{S}_j \to \mathcal{S}_i)}, 1\right)$$

- In practice the targeted probability distribution P(S) is usually a Boltzmann distribution. No need for partition function, as $W(S_i) = e^{\frac{-G_i}{kT}} \propto P(S_i)$ is enough.
- Pick random number $x \in [0, 1]$, candidate S_j accepted iff:

$$x \leq \min\left(e^{rac{-(G_j-G_i)}{kT}} imes rac{g(\mathcal{S}_i o \mathcal{S}_j)}{g(\mathcal{S}_j o \mathcal{S}_i)}, \ 1
ight)$$

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Metropolis-Hastings Monte-Carlo						



- With such acceptance criterion, asymptotic behaviour is mathematically proven¹.
- Sampling at the thermodynamic equilibrium.
- No time defined, no temporal evolution, no kinetics.
- Need to introduce kinetic elements.

 ¹Proof is straightforward, starting from detailed balance condition.
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From Metropolis Monte-Carlo	to Kinetic Monte-Carlo			

Theoretical background

Master equation²:

$$\frac{dP(\mathcal{S}_i, t)}{dt} = \sum_{j \neq i} \left(R(\mathcal{S}_j \to \mathcal{S}_i) P(\mathcal{S}_j, t) - R(\mathcal{S}_i \to \mathcal{S}_j) P(\mathcal{S}_i, t) \right)$$

- Note: In the Metropolis-Hastings MC, a stronger version of the master equation (with $\forall i$, $\frac{dP(S_i,t)}{dt} = 0$) was satisfied: detailed balance.
- Need to define states {S₁, S₂,...}, transitions S_i → S_j, and associated transition rates R(S_i → S_j)

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■ Analytical solution cannot be computed for real systems ⇒ numerical methods are required

²Can be proven from first principles

Traditional kinetic models	Monte-Carlo Markov Chain 000 0	Kinetic Monte-Carlo models ●000	References 0	Annexes
Algorithms overview				

Standard algorithm

- Generate all $N(S_i)$ possible transitions (i.e. elementary steps) $S_i \rightarrow S_j$ from current state S_i .
- 2 Choose a transition $\mathcal{S}_i \to \mathcal{S}_j$ at random.
- 3 Retrieve associated transition rate $R(S_i \to S_j)$, and compute $R_{\max}(S_i) = \max_{\substack{j \neq i}} (R(S_i \to S_j)).$
- 4 Pick a random number $r_1 \in [0, 1]$. Apply transition $S_i \to S_j$ iff $r_1 \leq \frac{R(S_i \to S_j)}{R_{\max}(S_i)}$.
- **5** If transition accepted, pick a random number $r_2 \in (0, 1]$, and increase time by $\Delta t = \frac{-\ln(r_2)}{N(S_i)R_{\max}(S_i)}$.
 - Otherwise, transition is rejected. (If most transition rates are negligible compared to $R_{max} \Rightarrow most$ attempts rejected).

6 Repeat.

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Algorithms overview				

First Reaction Method

- I Generate all possible transitions (i.e. elementary steps) S_i → S_j from current state S_i, and retrieve associated transition rate R(S_i → S_j)
- 2 For each possible transition $S_i \to S_j$, pick random number $r_j \in (0, 1]$ and compute associated time $\tau_j = \frac{-\ln(r_j)}{R(S_i \to S_i)}$.
- **3** Select transition $S_i \to S_j$ with lowest τ_j .
- **4** Apply selected transition, and increase time by τ_j .
- 5 Repeat (previous computations might be stored, and updated).

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Algorithms overview				

Variable Step Size Method / N-fold way / BLK algorithm

- **I** Generate and store in a list all $N(S_i)$ possible transitions (i.e. elementary steps) $S_i \rightarrow S_j$ from current state S_i , and retrieve associated transition rate $R(S_i \rightarrow S_j)$.
- 2 Compute the cumulative function $R_c(S_i, j) = \sum_{k \neq i}^{J} R(S_i \to S_k)$, and define $R_{tot}(S_i) = R_c(S_i, N(S_i))$.
- 3 Pick random number $r_1 \in (0, 1]$, and select transition $S_i \to S_j$ such that $R_c(S_i, j-1) < r_1 R_{tot}(S_i) \le R_c(S_i, j)$.
- Apply selected transition, pick a random number $r_2 \in (0, 1]$, and increase time by $\Delta t = \frac{-\ln(r_2)}{R_{tot}(S_i)}$.
- **5** Update list of stored possible positions (add new transitions, update rates, delete transitions no longer possible).
- 6 Repeat.

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Algorithms overview				

BLK algorithm flowchart



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Determining the kinetic rates				

Transition state theory

• How to compute transition rates $R(S_i \rightarrow S_j)$?

Common model, transition state theory:

$$k_{\text{TST}} = \kappa \frac{k_B T}{h} \exp\left(-\frac{\Delta G_{ij}^{\ddagger}}{k_B T}\right)$$

In practice, we define:

$$R(\mathcal{S}_i \to \mathcal{S}_j) = k_{\text{TST}} = \kappa \frac{Q_{ij}^{\ddagger}}{Q_i} \frac{k_B T}{h} \exp\left(-\frac{\Delta E_{ij}^{\ddagger}}{k_B T}\right)$$

where Q_{ij}^{\ddagger} and Q_i are the partition functions of the activated complex and current state S_i , and ΔE_{ij}^{\ddagger} is the energy barrier for the elementary step $S_i \rightarrow S_i$.

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Determining the kinetic rates				

Lateral interactions

- How to compute ΔE[‡]_{ij}? Depends on the composition and configuration of S_i and S_j
- Cluster expansion techniques:

$$\Delta E_{ij} = E_j - E_i = (E_{j,0} + L_j) - (E_{i,0} + L_i) = \Delta E_{ij,0} + L_{ij}$$

where $E_{i,0}$ is the total energy of state S_i as if the entities were infinitely separated; and L_i are the lateral interactions present in state S_i .

Activation energies are calculated with linear interpolation:

$$\Delta E_{ij}^{\ddagger} = \Delta E_{ij,0}^{\ddagger} + \omega \cdot L_{ij}$$

where $\omega \in [0, 1]$ is a proximity factor expressing how reactant- or product-like the transition state is.

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- Derivations
- Discussions

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Derivations				

Unimolecular decomposition: derivations

Using steady-state approximation to A(ads):

$$\frac{d[A(ads)]}{dt} = 0 = k_1 P_A[S_{free}] - (k_{-1} + k_2)[A(ads)]$$
(5)

$$\Leftrightarrow \quad \mathbf{0} = k_1(1-\theta)P_{\mathbf{A}}[\mathbf{S}] - (k_{-1}+k_2)\theta[\mathbf{S}] \qquad (6)$$

$$\Leftrightarrow \quad \theta = \frac{k_1 P_{\rm A}}{k_1 P_{\rm A} + k_{-1} + k_2} \tag{7}$$

Injecting (7) into (1):

$$r = k_2 \theta[S] = \frac{k_1 k_2 [A][S]}{k_1 [A] + k_{-1} + k_2}$$
(8)

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Derivations				

Langmuir-Hinshelwood model: derivations (part 1)

Using steady-state approximation to A(ads) and B(ads):

$$\frac{d[\mathrm{A(ads)}]}{dt} = 0 = k_1 P_{\mathrm{A}}[\mathrm{S}_{\mathrm{free}}] - k_{-1}[\mathrm{A(ads)}] - k_3 \theta_{\mathrm{A}} \theta_{\mathrm{B}}[\mathrm{S}]^2$$
(9)

$$\Leftrightarrow \quad \mathbf{0} = k_1 P_{\mathrm{A}}[\mathrm{S}_{\mathrm{free}}] - k_{-1} \theta_{\mathrm{A}}[\mathrm{S}] - k_3 \theta_{\mathrm{A}} \theta_{\mathrm{B}}[\mathrm{S}]^2 \tag{10}$$

$$\Leftrightarrow \quad \mathbf{0} = k_1 (1 - \theta_{\mathrm{A}} - \theta_{\mathrm{B}}) P_{\mathrm{A}}[\mathrm{S}] - (k_{-1} + k_3 \theta_{\mathrm{B}}[\mathrm{S}]) \theta_{\mathrm{A}}[\mathrm{S}] \qquad (11)$$

$$\Leftrightarrow \quad 0 = k_1 (1 - \theta_{\rm A} - \theta_{\rm B}) P_{\rm A} - (k_{-1} + k_3 \theta_{\rm B}[{\rm S}]) \theta_{\rm A} \tag{12}$$

$$\frac{d[\mathrm{B(ads)}]}{dt} = 0 = k_2 P_{\mathrm{B}}[\mathrm{S}_{\mathrm{free}}] - k_{-2}[\mathrm{B(ads)}] - k_3 \theta_{\mathrm{A}} \theta_{\mathrm{B}}[\mathrm{S}]^2$$
(13)

$$\Leftrightarrow \quad \mathbf{0} = k_2 P_{\mathrm{B}}[\mathrm{S}_{\mathrm{free}}] - k_{-2} \theta_{\mathrm{B}}[\mathrm{S}] - k_3 \theta_{\mathrm{A}} \theta_{\mathrm{B}}[\mathrm{S}]^2 \tag{14}$$

$$\Leftrightarrow \quad 0 = k_2 (1 - \theta_{\mathrm{A}} - \theta_{\mathrm{B}}) P_{\mathrm{B}}[\mathrm{S}] - (k_{-2} + k_3 \theta_{\mathrm{A}}[\mathrm{S}]) \theta_{\mathrm{B}}[\mathrm{S}] \qquad (15)$$

$$\Leftrightarrow \quad 0 = k_2(1 - \theta_{\rm A} - \theta_{\rm B})P_{\rm B} - (k_{-2} + k_3\theta_{\rm A}[{\rm S}])\theta_{\rm B} \tag{16}$$

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Derivations				

Langmuir-Hinshelwood model: derivations (part 2)

Assuming that the rate determining step is the reaction between A(ads) and B(ads):

$$k_{-2} \gg k_3 \theta_{\rm A}[{\rm S}]$$
(17)
$$k_{-1} \gg k_3 \theta_{\rm B}[{\rm S}]$$
(18)

Therefore, we obtain (defining $K_{\rm A} = \frac{k_1}{k_{-1}}$ and $K_{\rm B} = \frac{k_2}{k_{-2}}$):

$$\frac{d[\mathrm{A(ads)}]}{dt} = 0 = k_1(1 - \theta_{\mathrm{A}} - \theta_{\mathrm{B}})P_{\mathrm{A}} - k_{-1}\theta_{\mathrm{A}}$$
(19)

$$\Leftrightarrow \quad \mathbf{0} = \mathcal{K}_{\mathrm{A}}(1 - \theta_{\mathrm{A}} - \theta_{\mathrm{B}})\mathcal{P}_{\mathrm{A}} - \theta_{\mathrm{A}} \tag{20}$$

$$\frac{d[\mathrm{B(ads)}]}{dt} = 0 = k_2(1 - \theta_\mathrm{A} - \theta_\mathrm{B})P_\mathrm{B} - k_{-2}\theta_\mathrm{B} \qquad (21)$$

$$\Leftrightarrow \quad \mathbf{0} = \mathcal{K}_{\mathrm{B}}(1 - \theta_{\mathrm{A}} - \theta_{\mathrm{B}}) \mathcal{P}_{\mathrm{B}} - \theta_{\mathrm{B}} \tag{22}$$

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Langmuir-Hinshelwood model: derivations (part 3)

Ratioing (22) by (20), we get:

$$\frac{\theta_{\rm B}}{\theta_{\rm A}} = \frac{K_{\rm B}P_{\rm B}}{K_{\rm A}P_{\rm A}} \tag{23}$$

Injecting (23) into (20) and (22):

$$\theta_{A} = \frac{K_{A}P_{A}}{1 + K_{A}P_{A} + K_{B}P_{B}}$$
(24)
$$\theta_{B} = \frac{K_{B}P_{B}}{1 + K_{A}P_{A} + K_{B}P_{B}}$$
(25)

Leading finally to:

$$r = k_3 \theta_A \theta_B [S]^2 = \frac{K_A K_B P_A P_B [S]^2}{(1 + K_A P_A + K_B P_B)^2}$$
(26)

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Time step justification

Let us assume state S_i is accepted at time $t = t_i$, and define $\tau = t - t_i \ge 0$. So that $P(S_i, 0) = 1$.

$$P(S_i, \tau + \Delta \tau) = P(S_i, \tau) \overline{(1 - \frac{R_{tot}(S_i)\Delta \tau}{(any-transition \ proba)}})}$$
(27)

$$\Leftrightarrow \quad \frac{dP(\mathcal{S}_i,\tau)}{d\tau} = -P(\mathcal{S}_i,\tau)R_{\rm tot}(\mathcal{S}_i) \tag{28}$$

$$\Rightarrow P(\mathcal{S}_i, \tau) = e^{-R_{\text{tot}}(\mathcal{S}_i)\tau}$$
(29)

Probability p to time τ meaningful conversion:

$$\tau = -\frac{\ln(p)}{R_{\rm tot}(S_i)} \tag{30}$$

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