Modeling and Verification of Probabilistic Systems

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http://moves.rwth-aachen.de/teaching/ws-1516/movep15/

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Overview



2 Continuous-time Markov chains



The advance of time in DTMCs

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Continuous random variables

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- X is continuously distributed if there exists a function f(x) such that:

$$F_x(d) = Pr\{X \leqslant d\} = \int_{-\infty}^d f(x) \ dx$$
 for each real number d

where f satisfies: $f(x) \ge 0$ for all x and $\int_{-\infty}^{\infty} f(x) dx = 1$

- \triangleright $F_X(d)$ is the (cumulative) probability distribution function
- f(x) is the probability density function

Density of exponential distribution

The density of an *exponentially distributed* r.v. Y with rate $\lambda \in \mathbb{R}_{>0}$ is:

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► Variance $Var[Y] = \int_0^\infty (x - E[X])^2 \lambda \cdot e^{-\lambda \cdot x} dx = \frac{1}{\lambda^2}$

Exponential pdf and cdf



The higher λ , the faster the cdf approaches 1.

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- Yield a maximal entropy if only the mean is known

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8/30

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Proof of 2. : By contraposition, using the total law of probability.

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Minimum closure theorem for several exponentially distributed r.v.'s

For independent, exponentially distributed random variables X_1, X_2, \ldots, X_n with rates $\lambda_1, \lambda_2, \ldots, \lambda_n \in \mathbb{R}_{>0}$ the r.v. $\min(X_1, X_2, \ldots, X_n)$ is exponentially distributed with rate $\sum_{0 \le i \le n} \lambda_i$, i.e.,:

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Proof:

Generalization of the proof for the case of two exponential distributions.

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Overview

Negative exponential distribution





Continuous-time Markov chains

- labeled transition systems augmented with rates
- discrete state space
- continuous time steps
- delays exponentially distributed

Suited to modelling

- reliability models
- control systems
- queueing networks
- biological pathways
- chemical reactions
- ▶ ...

Continuous-time Markov chain

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Interpretation

• residence time in state s is exponentially distributed with rate r(s).

Continuous-time Markov chain

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- $(S, \mathbf{P}, \iota_{\text{init}}, AP, L)$ is a DTMC, and
- $r: S \to \mathbb{R}_{>0}$, the exit-rate function

Interpretation

- residence time in state s is exponentially distributed with rate r(s).
- phrased alternatively, the average residence time of state s is $\frac{1}{r(s)}$.
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Interpretation

- ▶ residence time in state *s* is exponentially distributed with rate *r*(*s*).
- phrased alternatively, the average residence time of state s is $\frac{1}{r(s)}$.
- ► thus, the higher the rate r(s), the shorter the average residence time in s.

Example



r(s) = 25, r(t) = 4, r(u) = 2 and r(v) = 100

Example: a classical perspective



We use $(S, \mathbf{P}, \mathbf{r}, \iota_{\text{init}}, AP, L)$ and $(S, \mathbf{R}, \iota_{\text{init}}, AP, L)$ interchangeably.



CTMC semantics

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$$Pr\{X_{s_0, s_2} \leqslant X_{s_0, s_1} \cap X_{s_0, s_2} \leqslant X_{s_0, s_3}\} = \\ R(s_0, s_2)$$

$$\overline{\mathbf{R}(s_0, s_1) + \mathbf{R}(s_0, s_2) + \mathbf{R}(s_0, s_3)}$$

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► Probability of staying at most t time in s_0 is: $Pr\{\min(X_{s_0,s_1}, X_{s_0,s_2}, X_{s_0,s_2}) \leq t\}$

20/30

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Simple CTMC example

Modelling a queue of jobs

- initially the queue is empty
- > jobs arrive with rate 3/2 (i. e., mean inter-arrival time is 2/3)
- ▶ jobs are served with rate 3 (i. e., mean service time is 1/3)
- maximum size of the queue is 3
- ▶ state space $S = \{s_i | 0 \leq i \leq 3\}$ where s_i indicates *i* jobs in queue.



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State-to-state timed transition probability

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Proof:

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Continuous-time Markov chains

Enzyme-catalysed substrate conversion

Kinetics

Main article: Enzyme kinetics



Enzyme kinetics is the investigation of how enzymes bind substrates and turn them into products. The rate data used in kinetic analyses are commonly obtained from enzyme assays, where since the 90s, the dynamics of many enzymes are studied on the level of individual molecules.

In 1902 Victor Henri^[57] proposed a quantitative theory of enzyme kinetics, but his experimental data were not useful because the significance of the hydrogen ion concentration was not yet appreciated. After Peter Lauritz Sorensen had defined the logarithmic pH-scale and introduced the concept of buffering in 1909^[58] the German chemist Leonor Michaelis and his Canadian postdoc Maud Leonora Menten repeated Henri's experiments and confirmed his equation which is referred to as Henri-Michaelis-Menten kinetics (termed also Michaelis-Menten kinetics).^[59] Their work was further developed by G. E. Briggs and J. B. S. Haldane, who derived kinetic

equations that are still widely considered today a starting point in solving enzymatic activity.[60]

The major contribution of Henri was to think of enzyme reactions in two stages. In the first, the substrate binds reversibly to the enzyme, forming the enzyme-substrate complex. This is sometimes called the Michaelis complex. The enzyme then catalyzes the chemical step in the reaction and releases the product. Note that the simple Michaelis Menten mechanism for the enzymatic activity is considered today a basic idea, where many examples show that the enzymatic activity involves structural dynamics. This is incorporated in the enzymatic mechanism while introducing several Michaelis Menten pathways that are connected with fluctuating rates ^{[44][45][46]}. Nevertheless, there is a mathematical relation connecting the behavior obtained from the basic Michaelis Menten mechanism (that was indeed proved correct in many experiments) with the generalized Michaelis Menten mechanisms involving dynamics and activity; ^[61] this means that the measured activity of enzymes on the level of many enzymes may be explained with the simple Michaelis. Menten equation, yet, the actual activity of enzymes is richer and involves structural dynamics.

Source: wikipedia (June 2011)

[edit]

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This process is a continuous-time Markov chain.

Enzyme-catalyzed substrate conversion as a CTMC



Transitions:
$$E + S \xrightarrow{1}{\stackrel{\frown}{=}} C \xrightarrow{0.001} E + P$$

e.g., $(x_E, x_S, x_C, x_P) \xrightarrow{0.001 \cdot x_C} (x_E + 1, x_S, x_C - 1, x_P + 1)$ for $x_C > 0$

Markovian queueing networks

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Joost-Pieter Katoen
Overview

Negative exponential distribution

Continuous-time Markov chains



Main points

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- A CTMC is a DTMC where state residence times are exponentially distributed.
- CTMC semantics distinguishes between enabledness and taking a transition.
- CTMCs are frequently used as semantical model for high-level formalisms.