

pubs.acs.org/JPCB

On Kinetic Constraints That Catalysis Imposes on Elementary Processes

Yann Sakref, Maitane Muñoz-Basagoiti, Zorana Zeravcic, and Olivier Rivoire*



ABSTRACT: Catalysis, the acceleration of product formation by a substance that is left unchanged, typically results from multiple elementary processes, including diffusion of the reactants toward the catalyst, chemical steps, and release of the products. While efforts to design catalysts are often focused on accelerating the chemical reaction on the catalyst, catalysis is a global property of the catalytic cycle that involves all processes. These are controlled by both intrinsic parameters such as the composition and shape of the catalyst and extrinsic parameters such as the concentration of the chemical species at play. We examine here the conditions that catalysis imposes on the different steps of a reaction cycle and the respective role of intrinsic and extrinsic parameters of the system on the emergence of catalysis by using an approach based on first-passage times. We illustrate this approach for various decompositions of a catalytic cycle into elementary steps, including non-Markovian decompositions, which are useful when the presence and nature of intermediate states are a priori unknown. Our examples cover different types of reactions and clarify the constraints on elementary steps and the impact of species concentrations on catalysis.

1. INTRODUCTION

Catalysts are substances that accelerate the completion of chemical reactions without being consumed during the process. They have long been studied in chemistry for their role in the industrial production of chemical products and in biology for their role in the metabolism and regulation of living processes.¹ A newer context for catalysis is provided by developments in supramolecular chemistry, DNA nanotechnology, and soft-matter physics²⁻⁴ which yield alternative chemistries where the size, shape, valence, strength, and specificity of interaction of the molecules can be tuned.^{5–8} In these chemistries, catalysts can be designed with control over many parameters, which provides experimental models to study catalysis and, more generally, new contexts to engineer properties inspired by biology.

Motivated by these perspectives, we have recently taken up the task of computationally designing catalysts in the context of chemistry where the "atoms" consist of spherical colloidal particles that interact via pairwise potentials.⁹ In this chemistry, the interaction potentials are isotropic with a common shape and interaction range but with possibly different depths.

Different atoms (A, B,....) can be designed to have interactions with specific strengths (ϵ_{AA} , ϵ_{AB} , ϵ_{BB} ,). We considered catalyzing the dissociation of a dimer made of two bound particles of type A by a catalyst consisting of particles of another type B, i.e., catalyzing the reaction $A_2 \rightarrow 2A$. This problem led us to ask several general questions: how to assess the presence of catalysis? How to score the performance of a catalyst? To which extent do these questions depend on the extrinsic conditions under which catalysis is analyzed, including the volume of the reaction vessel and the presence of multiple substrates or multiple catalysts? Is it sufficient to analyze a system consisting of a single substrate and a single catalyst, as it is computationally most convenient to do, or could catalysis

Received: July 10, 2023 October 2, 2023 **Revised:** Accepted: November 16, 2023 Published: December 13, 2023



Article



arise only when sufficiently many molecules are present? What extrinsic conditions are most favorable to catalysis?

More specific questions arise when studying the different steps that catalysis typically requires, including binding of the substrate(s) to the catalyst, reaction(s) in the presence of the catalyst, and release of the product(s).¹⁰ Some of these steps can have a simple dependence on the design parameters. For instance, the final step of the catalyzed reaction that we studied in ref 9 is the release of a single particle A attached to a single particle B of the catalyst which depends on the interaction strength between A and B (ϵ_{AB}) but not on other features of the catalyst. Since this final release must occur faster than the spontaneous reaction for catalysis to take place, the interactions between A and B must be weaker than the interaction between two A, $\epsilon_{\rm AB}$ < $\epsilon_{\rm AA}$. It would be desirable to extend this reasoning to obtain bounds on the design parameters, but this opens up several additional questions: is it always necessary for a forward step along a catalytic cycle to be faster than a spontaneous reaction? What if we do not have the knowledge of each intermediate state, as is typically the case when starting to investigate different designs?

Several of these questions have been previously addressed.¹¹⁻¹⁵ Previous approaches, however, do not provide straightforward and consistent answers to all of the above questions. For instance, a well-accepted quantification of catalytic activity is the turnover number, defined as the number of substrates that a catalyst converts per unit of time.¹³ One limitation of the turnover number is that it does not refer to the spontaneous reaction. As a consequence, it cannot reveal if catalysis, understood as an acceleration relative to the spontaneous reaction, is indeed taking place. This is not an issue when studying substances such as enzymes, which are unambiguously accelerating reactions but can be an issue when designing catalysts in new contexts.⁹ The first designs are indeed likely to have limited activities, in which case, establishing that catalysis is taking place is critical before considering any improvement.

Beyond the turnover number, enzymes are typically characterized by two distinct quantities, the catalytic rate constant k_{cat} and the Michaelis constant K_{M} .¹¹ Taken together, they account for the dependence of the reaction rate on substrate concentration [S] by describing the rate of product formation per enzyme as $k_{cat}[S]/(K_M + [S])$, a relation known as Michaelis–Menten kinetics.¹¹ By accounting for the concentration of substrates, this relationship partly accounts for extrinsic parameters. In this case, a measure of catalytic efficiency relative to the spontaneous reaction has been proposed,¹² but Michaelis–Menten kinetics rely on several assumptions that prevent its general applications:¹⁶ the substrate must be in excess relative to the catalyst, and catalysis must involve a series of states with rates of transition justifying that an intermediate complex is at a quasi-steady state. These assumptions cannot be made, in general.

These limitations motivated us to develop our own criterion for defining and quantifying catalysis, a criterion from which necessary conditions for elementary steps can be derived. Our approach goes beyond assumptions that may be justified in particular cases but are not guaranteed when considering more general catalysts: (i) the spontaneous reaction must be accounted for, as catalysis refers to its acceleration;¹⁷ (ii) the substrate does not necessarily have to be in excess to the catalyst, as it is for instance of interest when studying some biochemical reactions^{18–20} or autocatalysis;²¹ (iii) the kinetics must not be imposed to cover both enzymes that follow Michaelis–Menten kinetics and heterogeneous catalysts that do not,²² and more generally not to restrain a priori the design of new catalysts. In what follows, we present our approach and show how it addresses the different questions that we raised through analysis of several examples.

2. METHODS

2.1. Formal Definition of Catalysis. We assume a thermal bath at a constant temperature and a reaction vessel of a fixed volume. A spontaneous reaction is a thermally induced transition between two states that differ in their composition, i.e., where some molecules $S_1,..., S_n$ (the reactants, collectively denoted S) are transformed into some other molecules $P_1,..., P_m$ (the products, collectively denoted P). We define the time of the spontaneous reaction to be the mean time $T_{S \rightarrow P}$ to reach a final state P from an initial one S. For instance, in the simple case of a single reactant S that converts into a product P at a given rate k_0 , the initial state is S, the final state is P, and the spontaneous reaction time is $T_{S \rightarrow P} = 1/k_0$.

The spontaneous reaction time is compared to the average time $T_{C+S \rightarrow C+P}$ to complete the reaction in the presence of an additional molecule C in the reaction vessel. We consider that C catalyzes the transformation of the substrate into a product if the reaction is on average faster in its presence, that is, if $T_{C+S \rightarrow C+P} < T_{S \rightarrow P}$ and if C is unchanged in the process. This excludes initial and final states in which C is not present or is interacting with other molecules, including substrate or product molecules. To define and quantify catalysis, we therefore propose to use the ratio $T_{S \rightarrow P}/T_{C+S \rightarrow C+P}$ both to assess the presence of catalysis, through $T_{S \rightarrow P}/T_{C+S \rightarrow C+P} > 1$ and to quantify its efficiency. This quantitative criterion comparing two average times is consistent with the definition of catalysis given by the International Union of Pure and Applied Chemistry (IUPAC), according to which a catalyst is "a substance that increases the rate of a reaction".²³ Beyond the average time to complete a reaction, it may be of interest to consider the complete distribution of completion times, particularly when the most probable time differs from the mean time, as illustrated in Section 3.2.1, where we compare catalysis with either one or two substrates. When considering systems with a large number of particles, however, the mean time is expected to correspond to the most likely time.

2.2. Decompositions in Elementary Steps. As it is essential to distinguish configurations in which substrates and products are bound to the catalyst from those where they are unbound, the simplest decomposition of the catalytic process is into a cycle where configurations are partitioned into just three states: an initial state where the substrates S and the catalyst C are unbound, an intermediate state where they are interacting, and a final state where the products P are not bound to the catalyst. For example, a model widely used in enzymology and which serves as a basis to derive Michaelis–Menten kinetics²⁴ is that of a Markov model with an intermediate state CS that is reached from an unbound state C + S at a rate k_1 and left either back to C + S at a rate k_{-1} or toward C + P at a rate k_2 , as represented by

$$C + S \underset{k_{-1}}{\overset{k_1}{\leftrightarrow}} CS \xrightarrow{k_2} C + P$$
(1)

Decomposing a catalytic process as a Markov chain rests on a major assumption of separation of time scales between the time to transition between states and the time spent in the states, with local equilibration within each state so that the memory of previous states is lost. Under this assumption, the transitions are quantified by rates, corresponding to exponential distributions of the dwelling times within each state. However, such an assumption is not necessarily valid. First, binding and unbinding events involving diffusion are described as Markov processes under a mean-field approximation that may not be justified. In particular, the mass action kinetics that underlies the definition of a single state C + S breaks down in low dimensions.²⁵ Even in three dimensions, rates may fail to describe transitions between intermediate states when they involve the restricted diffusion of part of a molecule. Second, the number and nature of intermediate states may be unknown.9

A more general framework is that of semi-Markov processes, also known as Markov renewal processes.²⁶ The only essential assumption of semi-Markov processes is a separation of time scales to define states with dwelling times within states that are not necessarily exponentially distributed. Each state *i* has a distribution of dwelling times $\mathbb{P}(\tau_i)$ in addition to probabilities $p_{i \rightarrow i}$ to transition to other states $j \neq i$ once this dwelling time is over. We will develop our formalism in this broader context. In contrast to decomposition into a Markov process, the decomposition of a catalytic cycle into a semi-Markov process is not unique and, as such, may be more or less informative. In particular, unknown states of a Markov process may be collected in different ways in fewer states to define a coarsegrained semi-Markov process. Below, we thus present a minimal decomposition with a single intermediate state, which we denote C. \cdot S, that allows us to give general conditions. The more states that are known, however, the more information, in the form of constraints, may be derived.

3. RESULTS

3.1. Single-Molecule Catalysis. To show how these informative constraints can be derived, we start by studying a unimolecular spontaneous reaction $S \rightarrow P$ that proceeds at a rate k_0 . For simplicity, we first consider a closed vessel containing a single substrate S and a single catalyst C. Applying our approach, we derive the necessary and sufficient kinetic conditions on the elementary processes of the cycles for C to catalyze the spontaneous reaction. The restrictions on the nature of the spontaneous reaction and the number of molecules will be lifted in the next sections.

3.1.1. Markovian Catalytic Cycles. Under the above assumptions, the most basic catalytic cycle comprises only one intermediate state denoted as CS, which is accessible from either the unbound states C + S or C + P and can also transition back to those states. Graphically, the cycle is represented aswith k_{+n} denoting the forward rate of an

$$C + S \xrightarrow{k_0} C + P$$

$$k_{-1} \xrightarrow{k_1} k_{-2}$$

$$CS$$

$$(2)$$

elementary reaction and k_{-n} its reverse rate. Given that we consider the mean first-passage time $T_{C+S\rightarrow C+P}$ from the initial state to the absorbing state C + P, the transitions with rates k_{-0} and k_{-2} can be ignored and an equivalent representation is

$$C + P \stackrel{k_0}{\leftarrow} C + S \stackrel{k_1}{\underset{k_{-1}}{\leftrightarrow}} CS \stackrel{k_2}{\rightarrow} C + P$$
(3)

where C + P is repeated on both sides. The mean first-passage time from C + S to C + P can then be written in terms of the elementary rates (see Supporting Information Section I) as

$$T_{C+S \to C+P} = \frac{k_1 + k_{-1} + k_2}{k_0 k_{-1} + k_0 k_2 + k_1 k_2}$$
(4)

or, to highlight constraints

 $T_{C+S \to C+P} = T_{S \to P} + \rho_{cat} (T_{cat} - T_{S \to P})$ (5)

with $T_{S \to P} = 1/k_0$, $T_{cat} = 1/k_2$ and

$$\rho_{\rm cat} = \frac{1}{1 + (1 + k_{-1}/k_2)k_0/k_1} \tag{6}$$

By our criterion, catalysis takes place if and only if $T_{C+S \rightarrow C+P}$ < $T_{S \rightarrow P}$, which, by eq 5, is the case if and only if $\rho_{cat} > 0$ and T_{cat} $< T_{S \rightarrow P}$. In terms of the elementary rates, this requires that (i) $k_1 > 0$, simply meaning that it is possible for the substrate to bind the catalyst, and (ii) $k_2 > k_0$, meaning that the transformation of the substrate into the product on the catalyst and its release from the catalyst are overall faster than the spontaneous reaction. In particular, the presence of catalysis is independent of k_{-1} and of the magnitude of k_1 , which only needs to be nonzero. These conditions are partly counterintuitive. Naively, one may expect that for catalysis to take place, every step forward process along the catalytic cycle must be faster than the spontaneous reaction. We find here that it is not the case, and that k_1 may take any value. This can be understood by noting that when a putative catalyst is present, the substrate has two pathways: it can either undergo the spontaneous reaction or proceed through the potential catalytic route. If the first pathway is taken, no delay is possibly incurred, and only if the second pathway is taken must we ensure that no additional time is spent relative to the spontaneous reaction $(k_2 > k_0)$. If this condition is not satisfied, the catalyst turns into an inhibitor that slows down the completion of the reaction. $\rho_{\rm cat}$ represents the probability of taking the catalytic pathway, and maybe also counterintuitively, this probability may be arbitrarily small, meaning that the reaction mostly occurs spontaneously, without preventing catalysis from occurring, although possibly only marginally.

However, although the presence of catalysis is independent of the values of k_1 and k_{-1} , the efficiency of catalysis certainly depends on these quantities. In particular, in the limit $k_1 \rightarrow 0$, catalysis becomes negligible $(T_{C+S\rightarrow C+P} \rightarrow T_{S\rightarrow P} = 1/k_0)$, while it is maximal in the limit $k_1 \rightarrow \infty$ $(T_{C+S\rightarrow C+P} \rightarrow T_{cat} = 1/k_2)$. In any case, the rates k_{-0} and k_{-2} that describe the rates at which state C + P is left are irrelevant, since we consider only the first time at which C + P is reached.

The calculation can easily be extended to cases where catalysis involves a series of N intermediate Markovian states of the form

$$C + P \stackrel{k_0}{\leftarrow} C + S \stackrel{k_1}{\underset{k_{-1}}{\rightleftharpoons}} CS_1 \stackrel{k_2}{\underset{k_{-2}}{\rightleftharpoons}} \dots \stackrel{k_N}{\underset{k_{-N}}{\rightleftharpoons}} CS_N \stackrel{k_{N+1}}{\longrightarrow} C + P$$
(7)

In this case, we may again write

$$T_{C+S \to C+P} = T_{S \to P} + \rho_{cat} (T_{cat} - T_{S \to P})$$
(8)

with

$$T_{\text{cat}} = \sum_{n=0}^{N-1} \sum_{i=1}^{N-n} \left(\prod_{j=1}^{n} \frac{k_{-(i+j)}}{k_{i+j}} \right) \frac{1}{k_{i+n+1}}$$
(9)

and

$$\rho_{\rm cat} = \frac{1}{1 + (1 + \theta k_{-1}/k_2)k_0/k_1} \tag{10}$$

where the variables T_{cat} and θ are functions of all the rates $k_{\pm i}$ with i > 1, with the exception of the rate $k_{-(N+1)}$ that describes the rate at which the final state C + P is left. In particular, with one intermediate state (N = 1), $T_{\text{cat}} = 1/k_2$ and $\theta = 1$, leading back to eq 6. With two intermediate states (N = 2), $T_{\text{cat}} = 1/k_2 + 1/k_3 + k_{-2}/(k_2k_3)$ and $\theta = 1 + k_{-2}/k_3$; because of the reversibility of the transitions between states, T_{cat} is longer $T_{\text{cat}} > 1/k_2 + 1/k_3$; more generally, $T_{\text{cat}} > \sum_{i=2}^{N+1} 1/k_i$.

From eq 8, we obtain necessary conditions for catalysis that generalize those obtained with a single intermediate state: the first step must occur at a nonzero rate $(k_1 > 0)$ and the following steps must each be faster than the spontaneous reaction $(k_i > k_0$ for $i \ge 2)$. While the value of k_{-1} is again irrelevant for the definition of catalysis, this is not the case for the elementary rates k_{-i} for $2 \le i \le N$: the conditions $k_i > k_0$ for $i \ge 2$ are necessary, although not sufficient. The necessary and sufficient condition is that $T_{\text{cat}} < T_{\text{S} \rightarrow \text{P}}$ and $\rho_{\text{cat}} > 0$, where T_{cat} depend on the reverse rates k_{-i} for $2 \le i \le N$.

3.1.2. Identification of Favorable Conditions. Equation 9 indicates that setting to zero all reverse rates k_{-n} with n > 0 effectively reduces T_{cat} ; this reflects the fact that, for catalytic cycles with a single loop, it is always favorable to operate in conditions where the backward transitions are suppressed.⁹ Here we illustrate how this may be achieved by considering reactions that are irreversible or by removing products as they are made.

As an illustration of the benefit of irreversible reactions, consider, for instance, a catalytic process with two intermediate states, $C + P \leftarrow C + S \rightleftharpoons CS \rightleftharpoons CP \rightarrow C + P$. The condition for catalysis $T_{cat} < T_{S \rightarrow P}$ can be rewritten

$$\frac{1}{k_{\text{CS}\to\text{CP}}} + \frac{1}{k_{\text{CP}\to\text{C+P}}} + \frac{k_{\text{CP}\to\text{CS}}}{k_{\text{CS}\to\text{CP}}k_{\text{CP}\to\text{C+P}}} < \frac{1}{k_{\text{S}\to\text{P}}} \quad (11)$$

The first two terms correspond to conditions on the elementary forward rates, while the third one involves an additional constraint on the reverse reaction. This third constraint vanishes when the reverse reaction on the catalyst is irreversible, that is, when $k_{CP \rightarrow CS} = 0$, which is expected when the spontaneous reaction is itself irreversible, $k_{P \rightarrow S} = 0$.

When designing a catalyst for a reaction with a forward rate, $k_{S \rightarrow P}$, we may therefore first identify the necessary conditions to catalyze an irreversible reaction with the same forward rate, which is generally easier. Catalysts for the reversible reaction must indeed necessarily satisfy those conditions as well.

As an illustration of the benefit of product removal, consider reaction $S \rightarrow P_1 + P_2$ and catalytic scheme $C + P \leftarrow C + S \rightleftharpoons$ $CS \rightleftharpoons CP_1 + P_2 \rightarrow C + P_1 + P_2$. Here the backward rate $CP_1 + P_2 \rightarrow CS$ can effectively be eliminated by removing P_2 from the system as soon as it is formed. This also makes catalysis easier, and again, a catalyst under conditions where the products remain in the reaction vessel must be a catalyst under conditions where the products are immediately removed.

These simple considerations provide insight into the conditions that are most favorable to catalysis.

3.1.3. Beyond Markov Chains. Can we extend eq 5 to decompositions of catalytic processes where we lack specific information about the potential configurations that C and S might assume upon being close to each other or when these configurations do not constitute Markovian states, meaning we cannot assume they are independent of prior configurations? To do so, we introduce a transitional state C··S that defines a boundary between interacting and noninteracting substrate–catalyst pairs, with a catalytic process described by

 $C + P \leftarrow \{C + S\} \rightleftharpoons C \cdot \cdot S \rightarrow \{CS\} \rightarrow C + P$ (12)

Here C.S represents a state where C and S are just about to interact, while {CS} gathers all other configurations where they interact and $\{C + S\}$ those where they do not interact. Braces are introduced to indicate that these configurations do not generally define a "state," in the sense of a set of configurations that are equivalent as far as future transitions are concerned, as is the case when considering a Markov chain. The arrows indicate possible transitions and C + P denotes the final configurations where the product is formed and dissociated from the catalyst. In this setting, eq 5 still provides a decomposition of the mean first-passage time from an initial configuration $(C + S)_0$ to any of the final configurations C + Pbut with T_{cat} representing the mean time to reach C + P from C··S when taking the route via {CS} and with $\rho_{cat} = p_0 q / (1 - p_0) q / (1 - p_$ (1 - q)p), where p_0 is the probability to reach C··S from the initial configuration $(C + S)_0$, q the probability to go toward $\{CS\}$ rather than $\{C + S\}$ once in C··S and p the probability to come back to C··S given that the path toward $\{C + S\}$ was taken (Supporting Information II). Formally, the same necessary and sufficient conditions for catalysis apply, namely, $T_{\text{cat}} < T_{\text{S} \rightarrow \text{P}} \text{ and } \rho_{\text{cat}} > 0.$

This generalization provides an example of the necessary conditions for catalysis given a non-Markovian decomposition of the catalytic cycle. It also addresses cases where the mechanisms of catalysis are unknown since eq 12 makes no assumption on the presence of intermediate states. In practice, C··S consists of configurations where C and S are at a given distance just above their interaction range. Remarkably, we need not distinguish configurations within $\{C + S\}$ or $\{CS\}$. In particular, we do not need to distinguish unbound states as a function of the distance between C and S. Instead, all of the relevant information can be encapsulated into the probability and the mean time for the system to come back to C··S after leaving it (Supporting Information Section II). This decomposition can then be used to limit the analysis to a volume of the same dimension, where configurations in $\{C + S\}$ are effectively excluded. These configurations are, indeed, irrelevant to the computation of T_{cat} . This is for instance convenient in the context of molecular dynamics simulations where the time spent in $\{C + S\}$ is otherwise wasted.

It is also of interest to consider situations where additional information is available that allows for a more detailed description of catalysis than eq 12, although still not necessarily in the form of a Markov chain. For instance, catalysis may be known to follow a sequence of states $C + P \leftarrow C + S \rightleftharpoons CS_1 \rightleftharpoons$... $\rightleftharpoons CS_N \rightarrow C + P$ described by a semi-Markov chain where the time spent in each state C + S or CS_i is not necessarily described by an exponential distribution, as it is the case for

pubs.acs.org/JPCB

A- Catalysis of Dimerization	B- Catalysis of Dissociation	
$C + 2S \longleftarrow C + P$ $\uparrow \downarrow \qquad \qquad \uparrow \downarrow$	$C + S \overleftarrow{C} + 2P$ $\uparrow \downarrow \qquad \uparrow \overleftarrow{\downarrow}$	Kinetic conditions on forward rates
$C \cdot S + S \iff C \cdot P$	$C \cdot S \iff C \cdot P + P$	No condition
$ \begin{array}{ccc} \uparrow & & \uparrow \\ C \cdot S_2 & \longleftrightarrow & C:P \end{array} $	$ \begin{array}{ccc} \uparrow & & \uparrow \\ C:S & \longleftrightarrow & C \cdot P_2 \end{array} $	Faster than spontaneous reaction Fast "enough"

Figure 1. Examples of the decomposition of catalytic processes into graphs with multiple loops. (A) Catalysis of dimer formation. (B) Catalysis of dimer dissociation. The steps corresponding to a spontaneous reaction are indicated in blue, and each forward step that must be faster than the spontaneous reaction is indicated in red. Steps indicated in orange must also be fast enough, although not necessarily faster than the spontaneous reaction.

Markov chains. In this case, we may generalize the necessary (but not sufficient) conditions obtained for Markov chains $(k_i > k_0 \text{ for } i \ge 2)$ to $T_{CS_{i-1} \to CS_i \setminus CS_{i-2}} < T_{S \to P}$ for each $i \ge 2$, where $CS_{i-1} \to CS_i \setminus CS_{i-2}$ indicates that we are considering the first passage from CS_{i-1} to CS_i when accessing CS_{i-2} is excluded (with $CS_0 = C + S$ when i = 2).

The previous extensions of eq 5 apply to a single substrate S, a single (candidate) catalyst C, and assume a spontaneous reaction $S \rightarrow P$ described by a single rate k_0 . It does not apply to spontaneous reactions with multiple steps without further assumptions. For instance, if the spontaneous reaction proceeds through intermediate states, no state C··S can generally be defined (Supporting Information II).

Difficulties also arise when considering multimolecular reactions, e.g., $S + S \rightarrow P$, as extending the definition of C·· S to these reactions is not straightforward. In practice, however, it may be possible to make additional assumptions. For example, $C + S + S \rightleftharpoons C \cdot S + S \rightleftharpoons CS_2$ may be described by a Markov chain and only $CS_2 \rightarrow \{CS_2\} \rightarrow C + P$ may require a non-Markovian description.

3.1.4. Application to Reactions with Bimolecular Substrates and Products. Conditions on elementary steps are more subtle when catalysis is not described by a cycle with a single loop but by a graph comprising several loops. This occurs, for instance, when considering the spontaneous reaction of the association of two monomers into a dimer or the dissociation of a dimer into two free monomers. In these cases, six states may typically be defined by the number of bonds formed between a catalyst C, the two monomers, and the dimer. As shown in Figure 1, these states are part of a graph that contains two loops. In the first loop, a spontaneous reaction can occur, while only one particle of the substrate is bound to the catalyst, whereas the second loop describes the catalyzed reaction.

Consider for illustration the case of the dissociation reaction (Figure 1B) which we studied previously in the context of colloidal particles.⁹ Assuming that the mean time for a spontaneous reaction is exponentially distributed (or, even more simply, that the entire process is Markovian), we can apply eq 5 on both loops to compute the mean time to form the product in the presence of the catalyst, $T_{C+S\to C+2P}$ (Supporting Information Section III). The necessary and sufficient conditions for catalysis take their simplest simple form if the product—the monomers in that case of a dissociation reaction—is removed upon forming. Let indeed ρ_0 be the probability of reaching C + 2P from C + S through C·S rather than through the direct transition C·S \rightarrow C·P + P.

We may again write $T_{C+S\rightarrow 2P} = T_{S\rightarrow 2P} + \rho_{cat}(T_{cat} - T_{S\rightarrow 2P})$ with t h is t i m e $\rho_{cat} = \rho_0 \rho_1$ and $T_{cat} = T_{C:S\rightarrow C.P+P\setminus C.S} + \rho_1^{-1}T_{C\cdot P\rightarrow C+P\setminus C.S}$. Necessary and sufficient conditions are therefore $\rho_0 > 0$, $\rho_1 > 0$, and $T_{cat} < T_{S\rightarrow P}$. The latter implies the following necessary conditions: $T_{C:S\rightarrow C\cdot P_2\setminus C.S} < T_{S\rightarrow P}$, $T_{C\cdot P_2\rightarrow C\cdot P+P\setminus C\cdot S} < T_{S\rightarrow P}$ and $T_{C\cdot P+P\rightarrow C+2P} < T_{S\rightarrow P}$. Note, however, that $T_{C+S\rightarrow C\cdot S} < T_{S\rightarrow P}$ is not a necessary condition, nor is $T_{C\cdot S\rightarrow C:S\setminus C+S} < T_{S\rightarrow P}$ but while catalysis can occur for arbitrary values of $T_{C+S\rightarrow C\cdot S}$, the transition $C\cdot S \rightarrow C:S$ is constrained as it enters ρ_1 which is itself part of T_{cat} . This case applies to the problem of catalyzing the dissociation of a dimer outlined in the Introduction.⁹ It provides another example where a forward step along a catalytic path needs not be faster than the spontaneous reaction, here arising from the fact that the catalytic process is decomposed into a graph with several loops.

3.2. Beyond Single Molecules. We have considered so far an initial situation with a single catalyst and a single substrate. This condition is particularly interesting to design catalysts as it has the lowest number of degrees of freedom, facilitating both calculations of first-passage times and numerical simulations. This raises, however, the question of whether this is necessarily the most favorable condition for catalysis. We provide an example showing that it is not necessarily the case: a molecule C can accelerate the formation of the product in the presence of multiple substrates when it does not in the presence of a single one. We will see that for unimolecular reactions, such conclusion can be directly drawn from studying a single catalyst and a single substrate provided we go beyond mean first-passage times $T_{cat} = \mathbb{E}[t_{cat}]$ to consider the full distribution of first-passage times $\mathbb{P}[t_{cat}]$. Moreover, the presence of additional molecules, either substrates, catalysts, or products, may also turn a catalyst into an inhibitor. As before, in this section, we assume the temperature and volume to be fixed so that the reaction rate constants do not change with time, and increasing the number of molecules increases their concentration.

3.2.1. Necessary Conditions for Catalysis with Multiple S and C. We first consider the substrate to be unimolecular and the reaction to proceed in a single step. For an initial condition consisting of n_s substrates S, n_c molecules C, and no product P, a generic decomposition of the catalytic cycle into elementary processes is represented in Figure 2. As there are multiple substrates and catalysts in the system, one substrate can be transformed into a product while other substrates are either free or interacting with catalysts. Thus, it is of interest to relax our original definition of catalysis and consider the mean time to form the first free product irrespectively of the other



Figure 2. Possible decomposition of the catalytic cycle for an unimolecular reaction starting with n_s substrates S and n_c molecules C. In the extended definition of catalysis presented in the main text, the final state is any of the states on the right. Blue horizontal arrows represent spontaneous reactions, with the first one being the fastest of them all (with n_s substrates). Red diagonal arrows represent catalytic reactions.

molecules, i.e., $T_{n_cC+n_sS\to X+P}$, where X represents all molecules other than a product P. These are all of the states in the right column of Figure 2. As this encompasses the case where all the molecules C are not interacting with other molecules, as per our original definition, the necessary conditions that we shall obtain under this extended definition also apply to the original one.

The possible final states on the right side of Figure 2 are reached through either a spontaneous reaction (horizontal blue arrows) or a catalyzed reaction (diagonal red arrows). Assuming the spontaneous reactions to be independent of each other, the mean time at which the first of n_s spontaneous r e a c t i o n s i s c o m p l e t e d i s

 $T_{n_s S \to P+(n_s-1)S} = \mathbb{E}[\min(t_{S \to P}^{(1)}, \dots, t_{S \to P}^{(n_s)})]$ where $t_{S \to P}^{(q)}$ denotes a random variable for the time of completion of the spontaneous reaction in the presence of a single substrate, such that $T_{S \to P} = \mathbb{E}[t_{S \to P}^{(q)}]$ for any q. A necessary condition for catalysis is that one of the transitions from the left column to the right column in Figure 2 is, on average, faster than the spontaneous reaction with n_s substrates (top blue arrow). Formally, this condition implies that there must exist an $r \geq 1$ for which

$$\mathbb{E}[\min(t_{S \to p}^{(1)}, \ldots, t_{S \to p}^{(n_s - r)}, t_{cat}^{(1)}, \ldots, t_{cat}^{(r)})] \\ < \mathbb{E}[\min(t_{S \to p}^{(1)}, \ldots, t_{S \to p}^{(n_s)})]$$
(13)

where $t_{cat}^{(q)}$ is the random variable whose mean is $T_{cat} = \mathbb{E}[t_{cat}^{(q)}]$. This necessary criterion involves only the distributions of $t_{S \rightarrow P}$ and t_{cat} defined for a single substrate and a single catalyst, respectively. However, as transitions within the first column of Figure 2 must also be considered, it is not a sufficient condition for catalysis.

The distributions $t_{S \rightarrow P}$ and t_{cat} are not necessarily exponential; if they are, however, the necessary condition eq 13 reduces to $((n_s - r)T_{S \to P}^{-1} + rT_{cat}^{-1})^{-1} < (n_s T_{S \to P}^{-1})^{-1}$, i.e., $T_{\text{cat}} < T_{\text{S} \rightarrow \text{P}}$, the necessary condition for catalysis in the presence of a single substrate. On the other hand, if the distributions are not exponential, a molecule C may not be a catalyst in a single copy with a single substrate but becomes one when in n_c copies in the presence of n_s substrates. An illustration is provided by the Markovian catalytic cycle represented in Figure 3. In this example, we have $\mathbb{E}[t_{cat}] > \mathbb{E}[t_{S \to P}]$ due to the presence of an inactive out-ofcycle intermediate of the cycle that can trap the substrate S with a small probability but for a long time. However, when the system is initialized with two substrates, we find $\mathbb{E}[\min(t_{\text{cat}}, t_{S \to P})] < \mathbb{E}[\min(t_{S \to P}^{(1)}, t_{S \to P}^{(2)})]$. This is because either the reaction proceeds very quickly through the catalyst or, if it does not, with one of the two substrates binding to the catalyst and getting trapped in the out-of-cycle intermediate because



Figure 3. Example of a molecule C that catalyzes the formation of a product P starting with two substrates S but not with a single substrate. (A) We consider a situation where $k_3^- \ll k_3^+ \ll k_2^+$ so that once a molecule C has bound a substrate S, it has a small chance to trap the substrate in the long-lived out-of-cycle intermediate C₁S. The mean time to form a product when the initial state contains only one substrate is longer in the presence of C than that without it, $\mathbb{E}[t_{s\to P}] \prec \mathbb{E}[t_{cat}]$, where $t_{s\to P}$ is the random variable for the time to complete the spontaneous reaction without catalyst and t_{cat} for reaching C + P from CS when a back transition to C + S is excluded. (B) On the contrary, it is faster to form a product in the presence of C if the system is initialized with two substrates, $\mathbb{E}[\min(t_{cat}, t_{s\to P})] \prec \mathbb{E}[\min(t_{S\to P}^{(1)}, t_{S\to P}^{(2)})]$. (C) In this case, the probability density $\mathbb{P}[t_{cat}]$ is the sum of two exponentials, one controlled by k_2^+ that dominates at small times and the other controlled by k_3^- that dominates at long times (Supporting Information IV). As a consequence, the necessary condition for catalysis is fulfilled with two substrate molecules, but not with one. The graph illustrates the case where $k_2^+/k_0^+ = 10$, $k_3^+/k_0^+ = 10^{-1}$ and $k_3^-/k_0^+ = 10^{-3}$.



Figure 4. Illustration of the effect of having multiple substrates or multiple catalysts in the reaction vessel on the efficiency of catalysis. We consider four types of spontaneous reactions: with a unimolecular substrate and product (green and red with no out-of-cycle intermediate or with one, respectively), with a bimolecular product (yellow), and with a bimolecular substrate (blue). (A) Catalytic efficiency with one substrate ($n_s = 1$) and an increasing number of catalysts n_c . With unimolecular substrates, if C is initially a catalyst, the catalytic efficiency increases with the number of catalysts. For reactions with bimolecular products and/or substrates, adding more catalysts can be detrimental and eventually transforms a catalyst into an inhibitor ($T_{S \rightarrow P}/T_{n_c C+S \rightarrow n_c C+P} < 1$). (B) Catalytic efficiency with one catalyst ($n_c = 1$) and an increasing number of substrates n_s . Here, we extend our original definition and compute the mean time to make the first product irrespective of whether catalysts are still interacting with other substrates, $T_{n_c C+n_s S \rightarrow X+P}$, where X represents all molecules other than a product P. For all four types of reactions, in the limit of a large number of substrates, the products most likely arise through a spontaneous reaction. The values of the reaction rates are detailed in Supporting Information V.

the other substrate can still spontaneously react. This is therefore an example where the reaction is on average faster with C in the reaction vessel when there are two substrates compared to when a single substrate is present.

3.2.2. Efficiency of Catalysis with Multiple S, Multiple C, or Multiple P. How the presence of multiple substrates, catalysts, or products affects catalysis also depends on the nature of the spontaneous reaction, as we illustrate here with three types of spontaneous reactions: with a unimolecular substrate and product, with a bimolecular product, and with a bimolecular substrate. In order to develop intuition, we vary successively the number of substrates, the number of catalysts, or the number of products individually while keeping the other two fixed. In this case, it is also easy to analytically compute the mean time to form the first product (see an example of formal derivation in Supporting Information V). We discuss the transformation of multiple substrates by multiple catalysts in the next section.

We start by considering the effect of increasing the number n_c of catalysts in the presence of a single substrate, $n_s = 1$. Examples of the mean time to form the first product are shown in Figure 4A (see details in Supporting Information V). For unimolecular reactions, adding more catalysts in the reaction vessel increases the probability that the substrate meets a catalyst before spontaneously reacting. Thus, if C is a catalyst when $n_c = 1$, then the catalytic efficiency increases with n_c . while if it is an inhibitor, then the efficiency decreases (Figure 4A). This is unlike the case of reactions with multimolecular products (e.g., $S \rightarrow P + P$) or/and multimolecular substrates (e.g., $S + S \rightarrow P$) where catalytic efficiency eventually decreases for large enough n_{c} up to a point where any catalyst is turned into an inhibitor. When the product is multimolecular, an excess of catalysts indeed causes any released molecule of the product (P) to bind to a catalyst, which hinders the joint release of all product's molecules (2P). With multimolecular substrates, an excess of catalysts causes the different substrates (2S) to bind to different catalysts, which hinders their joint interaction with the same catalyst. As illustrated in Figure 4A,

the dependence of catalytic activity on the number n_c of catalysts can be nonmonotonic as a small increase in the number of catalysts may favor the association of the substrate to a catalyst.

The inhibitory effects of large concentrations of catalysts are mitigated when the binding of different substrates and products to the catalyst is sequential (Supporting Information V). Notably, when considering a single substrate $(n_s = 1)$, for none of the four reactions studied in Figure 4 does a molecule C that is not a catalyst in a single copy $(n_c = 1)$ become one when present in multiple copies $(n_c > 1)$, see Supporting Information V). This justifies studying a single catalyst in these cases.

Varying the number n_s of substrates in the presence of a single catalyst ($n_c = 1$) can also lead to nonmonotonic effects. In the limit of a large number of substrates, the products most likely arise through a spontaneous reaction, and the catalytic efficiency is marginal (Figure 4B). Catalytic efficiency may vary nonmonotonically with the number of substrates for different reasons; for example, because of a nonexponential distribution of t_{cat} (as in Figure 3), or because the spontaneous reaction scales nonlinearly with the number of substrates (as for reactions with bimolecular substrates). We note that for none of the reactions without an out-of-cycle intermediate studied in Figure 4 does a molecule C that is not a catalyst with a single substrate ($n_s > 1$, see Supporting Information V). This justifies studying a single substrate in these cases.

In the presence of multiple products, extra states must be considered in which molecules C are bound to product molecules. Consistent with the general intuition that such product inhibition can only be detrimental, catalysis occurs in the presence of multiple products only if it occurs with no products (Supporting Information VI). An initial absence of products is therefore always most favorable for catalysis and, as we noted earlier, it is also advantageous to remove products as they are produced.



Figure 5. (A) Extension of Figure 2 beyond the first product, represented here with $n_c = n_s = 3$ for clarity. The spontaneous reaction is assumed to be irreversible, and the products are removed upon forming. (B) Gillespie simulations: mean time to form n_s products from n_s substrates and n_c catalysts. We recover the same dependency on the number of substrates and catalysts as in Figure 4 for the case of unimolecular reactions: as more substrate molecules need to be transformed, the catalytic efficiency eventually decreases, and adding more catalysts is always beneficial.

3.2.3. Catalysis of Multiple Substrates. We have focused so far on the time to make the first product, but the same principles apply when considering the time to make the first $n_p > 1$ products out of n_s substrates that are initially present. Indeed, representing the catalytic cycle as in Figure 5A, we can derive a series of necessary conditions of the form of eq 13 (Supporting Information VII). We find also the same patterns of dependency on the number of substrates and catalysts: as more substrate molecules need to be transformed, the catalytic efficiency eventually decreases, and for unimolecular reactions, adding more catalysts is always beneficial, as illustrated in Figure 5B.

3.3. Relation to Michaelis–Menten Kinetics. Mean first-passage times are known to provide an alternative derivation of Michaelis–Menten kinetics in enzymology,^{27,28} which is useful to analyze single-molecule experiments²⁹ or to understand theoretically what controls enzyme.³⁰ We explain here how the mean first-passage time $T_{\rm MM}$ defined in this context is related to but different from the mean first-passage time $T_{C+S\to C+P}$ introduced in this work.

This is most simply explained in the case where catalysis is described by a Markov chain with a single intermediate state CS, as considered in Section 1. In enzymology, this scheme is usually studied under the assumptions that the spontaneous reaction is negligible, in which case it is written $C + S \rightleftharpoons_{k-1}^{k_1} CS \xrightarrow{k_2} C + P$. The concentration [S] of substrates

is also assumed to be much larger than the concentration [C] of catalysts. The focus is then on the rate of product formation given by

$$\nu_{\rm MM} = \frac{\rm d[P]}{\rm dt} = \frac{k_{\rm cat}[S][C]}{K_{\rm M} + [S]}$$
(14)

with $k_{cat} = k_2$ and $K_M = (k_{-1} + k_2)/\overline{k}_1$. Here \overline{k}_1 is the secondorder rate constant of the transition $C + S \rightarrow CS$. v_{MM} can also be obtained as a mean first-passage time T_{MM} from C + S to C+ P as $v_{MM} = [C]/T_{MM}$ when viewing $C + S \rightleftharpoons CS \rightarrow C + P$ as describing the conversion into a product of one of the many available substrates by one particular enzyme.²⁷ In this context, k_1 is a pseudo-first-order rate constant giving the rate per catalyst at which the complex CS is produced from C + S, related to \overline{k}_1 by $k_1 = \overline{k}_1[S]$. In our approach, we take the point of view of a substrate rather than the point of view of a catalyst, which allows us to make a comparison with the spontaneous reaction. In this context, $k_1 = \overline{k_1}/V$ if considering a single catalyst, or $k_1 = \overline{k_1}[C]$ if considering many more catalysts than substrates. Here, $\overline{k_1}$ is the same second-order rate as above and V denotes the volume of the reaction vessel.

Comparing the expressions for $k_{\rm cat}$ and $K_{\rm M}$ in enzymology and those of $T_{\rm cat}$ and $\rho_{\rm cat}$ in our approach, we verify that $k_{\rm cat} = 1/T_{\rm cat}$ and that the probability of encountering $\rho_{\rm cat}$ is closely related to $k_{\rm cat}/K_{\rm M}$. Equation 6 indeed indicates that $\rho_{\rm cat}^{-1} - 1 = [(k_{-1} + k_2)/k_1](k_0/k_2) = (K_{\rm M}/k_{\rm cat})(k_0/V)$. To make a more general connection between the two approaches, we can rewrite eq 5 as

$$\frac{T_{S \to P}}{T_{C+S \to C+P}} = 1 + \frac{T_{S \to P}/T_{cat} - 1}{1 + (\rho_{cat}^{-1} - 1)T_{S \to P}/T_{cat}}$$
(15)

Assuming the spontaneous reaction to be negligible, $T_{\rm S \rightarrow P} = 1/k_0 \ll T_{\rm cat} = 1/k_2$, and using the expression for $\rho_{\rm cat}$, we obtain

$$\frac{1}{T_{C+S\to C+P}} \simeq \frac{k_2}{1 + (k_{-1} + k_2)/k_1} = \frac{k_{cat}}{1 + K_M V}$$
(16)

when considering a single substrate in the presence of a single catalyst, or $1/T_{C+S \rightarrow C+P} = k_{cat}[C]/(K_M + [C])$ when considering a single substrate in the presence of many catalysts.

Since $T_{cat} = 1/k_{cat}$ our criterion for catalysis $T_{cat} < T_{S \rightarrow P}$ reads $k_{cat} > k_0$ in the language of Michaelis–Menten kinetics. The ratio k_{cat}/k_0 used by Wolfenden and collaborators to compare a catalyzed reaction to a spontaneous one¹² therefore corresponds to $T_{S \rightarrow P}/T_{C+S \rightarrow C+P}$ in the saturation limit where $\rho_{cat} \rightarrow 1$ or $K_M \rightarrow 0$.

4. CONCLUSIONS

Motivated by the computational design of a minimal catalyst in the realm of colloids with programmable interactions,⁹ we studied several questions related to the kinetics of catalysis. First, we addressed the question of assessing the presence of catalysis, for which we propose to compare the mean firstpassage times from substrate(s) to product(s) in the absence and in the presence of the candidate catalyst: a substance is a catalyst if the reaction is faster on average, formally $T_{C+S \rightarrow C+P} <$ $T_{S \rightarrow P}$. Second, we addressed the question of scoring the performance of the catalyst, for which we propose to consider the ratio $T_{S \rightarrow P}/T_{C+S \rightarrow C+P}$ which, in the presence of catalysis, must be larger than one, that is, $T_{S \rightarrow P}/T_{C+S \rightarrow C+P} > 1$. These quantities depend not only on the intrinsic properties of the catalyst but also on the extrinsic conditions under which catalysis is analyzed. In particular, we illustrated in several examples how it depends on the presence of multiple substrates, products, or/and catalysts. We showed that it is not sufficient to analyze a single substrate and a single catalyst, although notable exceptions exist, including unimolecular single-step reactions. We also identified conditions that are more favorable to catalysis: if a substance is not a catalyst under these conditions, it is not a catalyst under other conditions. They include considering an irreversible spontaneous reaction with an equivalent forward rate or removing every product as soon as it is formed.

taneous reaction $S \xrightarrow{k_0} P$, it is required that the forward rates k_i satisfy $k_i > k_0$ for $i \ge 2$ but no such constraint applies to k_1 . Constraints on forward rates can take more relaxed forms when the elementary steps are not organized into a single cycle but instead in a graph with multiple loops. This happens, for instance, when the spontaneous reaction involves multiple substrates or multiple products. Informative necessary conditions for catalysis can also be derived when the decomposition of the catalytic cycle is not described by a Markov chain, although not without restrictive assumptions.

While motivated by the design of catalysts for experiments in soft-matter physics, our approach involves only the kinetics of catalysis and therefore has a broader scope. As we have shown, it is closely related to, although different from, quantifications of catalysis used in enzymology.^{27,28} The main difference is that we focus on the fate of a substrate rather than on the fate of a catalyst. This point of view is required to account for the spontaneous reaction, even in the presence of a catalyst. While the spontaneous reaction is typically negligible in the context of enzymes, a reference to the spontaneous reaction is essential both to define catalysis and to study it in conditions where catalytic efficiency is either poor or unknown, as it is for instance the case in experiments to design nonenzymatic autocatalysts.³¹ Our approach should therefore find applications to the design and study of catalysts beyond our original case study.

ASSOCIATED CONTENT

1 Supporting Information

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/acs.jpcb.3c04627.

Markovian catalytic cycles: derivation of the mean time to form a product in the presence of a putative catalyst and as per the Markovian catalytic cycle (eq 2), general criterion for catalysis: derivation of eq 5 for semiMarkovian cycles, catalysis for dimer dissociation: derivation of the mean time for a substrate to dissociate in the presence of a putative catalyst and as per the cycle represented in Figure 1B, example of a catalytic cycle with an out-of-cycle intermediate: derivation of the mean time to form a product when considering that the cycle contains an out-of-cycle intermediate as represented in Figure 3, conditions for catalysis in the presence of multiple catalysts or multiple substrates: explanation on how having multiple catalysts or multiple substrates change the catalytic cycle and may affect the relative efficiency of a molecule to act as a catalyst, catalysis in the presence of multiple products: explanation on how having multiple products changes the catalytic cycle and may affect the relative efficiency of a molecule to act as a catalyst, and catalysis beyond the first product: formalization of necessary and sufficient conditions for a molecule to act as an catalyst when considering multiple substrates (PDF)

AUTHOR INFORMATION

Corresponding Author

Olivier Rivoire – Gulliver UMR CNRS 7083, ESPCI Paris, 75005 Paris, France; orcid.org/0000-0002-1820-1324; Email: olivier.rivoire@espci.fr

Authors

- Yann Sakref Gulliver UMR CNRS 7083, ESPCI Paris, 75005 Paris, France
- Maitane Muñoz-Basagoiti Gulliver UMR CNRS 7083, ESPCI Paris, 75005 Paris, France; Institute of Science and Technology Austria, 3400 Klosterneuburg, Austria
- Zorana Zeravcic Gulliver UMR CNRS 7083, ESPCI Paris, 75005 Paris, France

Complete contact information is available at: https://pubs.acs.org/10.1021/acs.jpcb.3c04627

Notes

The authors declare no competing financial interest.

ACKNOWLEDGMENTS

We acknowledge funding from ANR-22-CE06-0037-02. This work has received funding from the European Unions Horizon 2020 research and innovation program under the Marie Sklodowska-Curie grant agreement No. 754387.

REFERENCES

(1) Chorkendorff, I.; Niemantsverdriet, J. W. Concepts of Modern Catalysis and Kinetics, 2010; Wiley-VCH; pp 349–376.

(2) Zhang, Y.; McMullen, A.; Pontani, L. L.; He, X.; Sha, R.; Seeman, N. C.; Brujic, J.; Chaikin, P. M. Sequential self-assembly of DNA functionalized droplets. *Nat. Commun.* **2017**, *8* (1), 21–27.

(3) Zhuo, R.; Zhou, F.; He, X.; Sha, R.; Seeman, N. C.; Chaikin, P. M. Litters of self-replicating origami cross-tiles. *Proc. Natl. Acad. Sci.* U.S.A. **2019**, *116* (6), 1952–1957.

(4) Rossi-Gendron, C.; El Fakih, F.; Bourdon, L.; Nakazawa, K.; Finkel, J.; Triomphe, N.; Chocron, L.; Endo, M.; Sugiyama, H.; Bellot, G.; et al. Isothermal self-assembly of multicomponent and evolutive DNA nanostructures. *Nat. Nanotechnol.* **2023**, *18*, 1311.

(5) Feng, L.; Pontani, L.-L.; Dreyfus, R.; Chaikin, P.; Brujic, J. Specificity, flexibility and valence of DNA bonds guide emulsion architecture. *Soft Mater.* **2013**, *9* (41), 9816.

(6) Seeman, N. C.; Sleiman, H. F. DNA nanotechnology. *Nat. Rev. Mater.* **2017**, 3 (1), 17068. (7) Hueckel, T.; Hocky, G. M.; Sacanna, S. Total synthesis of colloidal matter. *Nat. Rev. Mater.* **2021**, *6* (11), 1053–1069.

(8) Niu, R.; Du, C. X.; Esposito, E.; Ng, J.; Brenner, M. P.; McEuen, P. L.; Cohen, I. Magnetic handshake materials as a scale-invariant platform for programmed self-assembly. *Proc. Natl. Acad. Sci. U.S.A.* **2019**, *116* (49), 24402–24407.

(9) Muñoz-Basagoiti, M.; Rivoire, O.; Zeravcic, Z. Computational design of a minimal catalyst using colloidal particles with programmable interactions. *Soft Matter* **2023**, *19* (21), 3933–3939.

(10) Lehn, J.-M. Supramolecular Chemistry: Concepts and Perspectives; Wiley, 1995.

(11) Cornish-Bowden, A. Fundamentals of Enzyme Kinetics; Elsevier, 1979.

(12) Wolfenden, R.; Snider, M. J. The depth of chemical time and the power of enzymes as catalysts. *Acc. Chem. Res.* **2001**, *34* (12), 938–945.

(13) Kozuch, S.; Martin, J. M. L. turning over definitions in catalytic cycles. *ACS Catal.* **2012**, *2* (12), 2787–2794.

(14) Vojvodic, A.; Nørskov, J. K. New design paradigm for heterogeneous catalysts. *Natl. Sci. Rev.* **2015**, *2* (2), 140–143.

(15) Grima, R.; Leier, A. Exact product formation rates for stochastic enzyme kinetics. *J. Phys. Chem. B* **2017**, *121* (1), 13–23.

(16) Srinivasan, B. A guide to the michaelis-menten equation: steady state and beyond. *FEBS J.* **2022**, 289 (20), 6086-6098.

(17) McNaught, A. D.; Wilkinson, A. Compendium of Chemical Terminology; Blackwell Science Oxford, 1997; Vol. 1669.

(18) Shacter, E.; Chock, P. B.; Stadtman, E. R. Energy consumption in a cyclic phosphorylation/dephosphorylation cascade. *J. Biol. Chem.* **1984**, 259 (19), 12260–12264.

(19) Luby-Phelps, K. Cytoarchitecture and physical properties of cytoplasm: volume, viscosity, diffusion, intracellular surface area. *Int. Rev. Cytol.* **1999**, *192*, 189–221.

(20) Bispo, J. A. C.; Bonafe, C. F. S.; Koblitz, M. G. B.; Silva, C. G. S.; de Souza, A. R. Substrate and enzyme concentration dependence of the henri-michaelis-menten model probed by numerical simulation. *J. Math. Chem.* **2013**, *51* (1), 144–152.

(21) Hanopolskyi, A. I.; Smaliak, V. A.; Novichkov, A. I.; Semenov, S. N. Autocatalysis: Kinetics, mechanisms and design. *ChemSusChem* **2020**, *3*, No. e2000026.

(22) Laidler, K. J.; Meiser, J. H. *Physical Chemistry*; The Benjamin Cummings Publishing Company, 1982.

(23) Laidler, K. J. A glossary of terms used in chemical kinetics, including reaction dynamics (IUPAC Recommendations 1996). *Pure Appl. Chem.* **1996**, *68* (1), 149–192.

(24) Srinivasan, B. A guide to the michaelis-menten equation: steady state and beyond. *FEBS J.* **2021**, *289*, 6086-6098.

(25) Savageau, M. A. Michaelis-menten mechanism reconsidered: implications of fractal kinetics. J. Theor. Biol. 1995, 176 (1), 115–124.
(26) Karlin, S. A First Course in Stochastic Processes; Academic Press, 2014.

(27) Ninio, J. Alternative to the steady-state method: derivation of reaction rates from first-passage times and pathway probabilities. *Proc. Natl. Acad. Sci. U.S.A.* **1987**, *84* (3), 663–667.

(28) Qian, H. Cooperativity and specificity in enzyme kinetics: a single-molecule time-based perspective. *Biophys. J.* **2008**, 95 (1), 10–17.

(29) Min, W.; English, B. P.; Luo, G.; Cherayil, B. J.; Kou, S. C.; Xie, X. S. Fluctuating enzymes: lessons from single-molecule studies. *Acc. Chem. Res.* **2005**, 38 (12), 923–931.

(30) Banerjee, K.; Kolomeisky, A. B.; Igoshin, O. A. Accuracy of substrate selection by enzymes is controlled by kinetic discrimination. *J. Phys. Chem. Lett.* **2017**, *8* (7), 1552–1556.

(31) von Kiedrowski, G. Minimal replicator theory i: Parabolic versus exponential growth. *Bioorganic Chemistry Frontiers*; Springer, 1993; Vol. 3.