

Linear Eyring Plots Conceal a Change in the Rate-Limiting Step in an Enzyme Reaction

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S Supporting Information

ABSTRACT: The temperature dependence of psychrophilic and mesophilic (*R*)-3-hydroxybutyrate dehydrogenase steady-state rates yields nonlinear and linear Eyring plots, respectively. Solvent viscosity effects and multiple- and single-turnover pre-steady-state kinetics demonstrate that while product release is rate-limiting at high temperatures for the psychrophilic enzyme, either interconversion between enzyme–substrate and enzyme–product complexes or a step prior to it limits the rate at low temperatures. Unexpectedly, a similar change in the rate-limiting step is observed with the mesophilic enzyme, where a step prior to chemistry becomes rate-limiting at low temperatures. This observation may have implications for past and future interpretations of temperature–rate profiles.

Temperature–rate profiles are powerful tools for gaining insight into thermodynamic activation parameters of chemical reactions.¹ In enzyme-catalyzed reactions, analysis of the temperature-dependent behavior of rate constants is widely used to uncover catalytic properties among psychrophilic, mesophilic, and thermophilic enzymes² and to probe protein dynamics and hydrogen tunneling³ and even allosteric regulation.^{4,5} Plotting $\ln(k/T)$ versus $1/T$ (Eyring plot) often results in a straight line, and fitting data to eq 1 yields the activation enthalpy and entropy, assuming they are constant over the temperature range and recrossing is negligible.^{1,6,7} However, nonlinear Eyring plots have been reported,⁸ and a recent and elegant hypothesis for interpreting them invokes a role for activation heat capacity in enzyme catalysis upon fitting data to eq 2.^{9,10} In eqs 1 and 2, k is the rate constant; k_B , h , and R are the Boltzmann, Planck, and gas constants, respectively; ΔH^\ddagger and ΔS^\ddagger are the activation enthalpy and entropy, respectively; T is the temperature; T_0 is a reference temperature; $\Delta H^\ddagger_{T_0}$ and $\Delta S^\ddagger_{T_0}$ are ΔH^\ddagger and ΔS^\ddagger at T_0 , respectively; and ΔC_p^\ddagger is the activation heat capacity.

$$\ln \frac{k}{T} = \ln \frac{k_B}{h} - \frac{\Delta H^\ddagger}{RT} + \frac{\Delta S^\ddagger}{R} \quad (1)$$

$$\ln \frac{k}{T} = \ln \frac{k_B}{h} - \frac{\Delta H^\ddagger_{T_0} + C_p^\ddagger(T - T_0)}{RT} + \frac{\Delta S^\ddagger_{T_0} + C_p^\ddagger \ln(T/T_0)}{R} \quad (2)$$

Nonlinear Eyring plots without protein denaturation could also arise from a switch in the rate-limiting step at a given temperature, which must be tested.^{2,8} Conversely, linear Eyring plots are often assumed to reflect the same rate-limiting step at all temperatures.^{2,6,9,11,12} Here we show that linear Eyring plots may accommodate a change in the rate-limiting step.

We cloned and expressed the bacterial (*R*)-3-hydroxybutyrate dehydrogenase (HBDH)-encoding gene from mesophilic *Acinetobacter baumannii*¹³ and from psychrophilic *Psychrobacter arcticus*¹⁴ and purified the recombinant mesophilic (*Ab*HBDH) and psychrophilic (*Pa*HBDH) enzymes (Figure S1). HBDH (EC 1.1.1.30) catalyzes the NADH-dependent reduction of acetoacetate to (*R*)-3-hydroxybutyrate, with 3-oxovalerate being an alternative substrate turned over more slowly by the enzyme (Scheme S1).¹⁵

HBDH is part of the short-chain dehydrogenase/reductase (SDR) superfamily, one of the largest known protein groups.¹⁶ SDR proteins share a low level of sequence identity but highly conserved three-dimensional architecture and catalytic residues.¹⁷ HBDH has applications in asymmetric synthesis of precursors of pharmaceuticals and monomeric constituents of biodegradable polyhydroxyalkanoates.^{18,19}

We measured saturation curves for *Ab*HBDH and *Pa*HBDH at various temperatures (283–330 K for *Ab*HBDH and 283–318 K for *Pa*HBDH) (Figure S2 and Tables S1–S4). Differential scanning fluorimetry demonstrated no denaturation occurs in this temperature range (Figure S3 and Table S5). Apparent steady-state catalytic constants (k_{cat}) were used to construct Eyring plots for each enzyme (Figure 1). Fitting all data with both eqs 1 and 2 resulted in best-fit lines that are linear for *Ab*HBDH but nonlinear for *Pa*HBDH. Accordingly, a

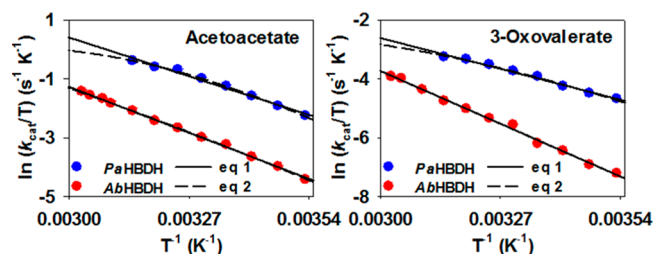


Figure 1. Eyring plots for *Pa*HBDH and *Ab*HBDH with acetoacetate and 3-oxovalerate as substrates.

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significant negative ΔC_p^\ddagger is predicted for the psychrophilic enzyme but not for the mesophilic one (Table S6).

A negative ΔC_p^\ddagger for k_{cat} could suggest a decrease in the number of vibrational modes as the Michaelis complex progresses toward the transition state.^{9,11,12,20} However, enzymatic rates often contain contributions from several steps, and a negative ΔC_p^\ddagger can report on vibrational modes associated with chemical^{16,20} and physical steps.²

To test if diffusional steps limit the steady-state rate in *Pa*HBDH and *Ab*HBDH reactions at 298 K, we determined solvent viscosity effects by measuring saturation curves at different concentrations of glycerol (Figure S4 and Tables S7–S10), and plots of k_{cat} ratios versus relative viscosity²¹ are shown in Figure 2. Fitting data to eq S5 yielded slopes of 0.43

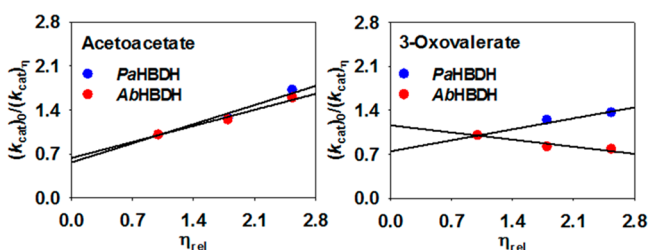


Figure 2. Solvent viscosity effects on k_{cat} for *Pa*HBDH and *Ab*HBDH with acetoacetate and 3-oxovalerate as substrates. Lines are fits of data to eq S5.

± 0.04 and 0.36 ± 0.02 for *Pa*HBDH and *Ab*HBDH, respectively, indicating product release is significantly rate-limiting with acetoacetate as the substrate.^{21,22} With 3-oxovalerate as the substrate, slopes decrease to 0.25 ± 0.02 for *Pa*HBDH, consistent with only a modest product release contribution to the rate-limiting step, and -0.16 ± 0.02 for *Ab*HBDH, showing no contribution from diffusional steps to k_{cat} . Diffusional steps from substrate binding do not limit reaction rates for either enzyme (Figure S4), and 5% PEG-8000 as a macroviscogen had no effect on saturation curves (Figure S5), pointing to the effects of glycerol resulting from an increase in solvent microviscosity.^{21,23}

The solvent viscosity effects on k_{cat} suggest that the curvatures in the Eyring plots and the consequent negative ΔC_p^\ddagger calculated for *Pa*HBDH may report on distinct steps when acetoacetate and 3-oxovalerate are used as substrates. To probe further the nature of the rate-limiting step for this enzyme, we employed multiple-turnover pre-steady-state kinetics to characterize the approach to the steady state²⁴ at 283, 298, and 318 K (Figure 3A). No burst in substrate consumption is observed with 3-oxovalerate as the substrate. These results support the conclusion from solvent viscosity effects at 298 K that k_{cat} is not limited by the product release rate with 3-oxovalerate as the substrate.

When acetoacetate is used as the substrate, even though a burst in substrate depletion at 298 and 318 K cannot be directly observed, it can be inferred²⁵ (Figure 3A). The concentration of co-substrate NADH at time zero in the presence of *Pa*HBDH and acetoacetate is significantly offset from its value in the control without acetoacetate (Figure S6), suggesting at least one turnover has happened within the 0.9 ms dead time of the stopped-flow spectrophotometer. This suggests a step after chemistry, likely product release, is rate-limiting at 298 and 318 K, in agreement with the significant solvent viscosity effect on k_{cat} at 298 K. At 283 K, however,

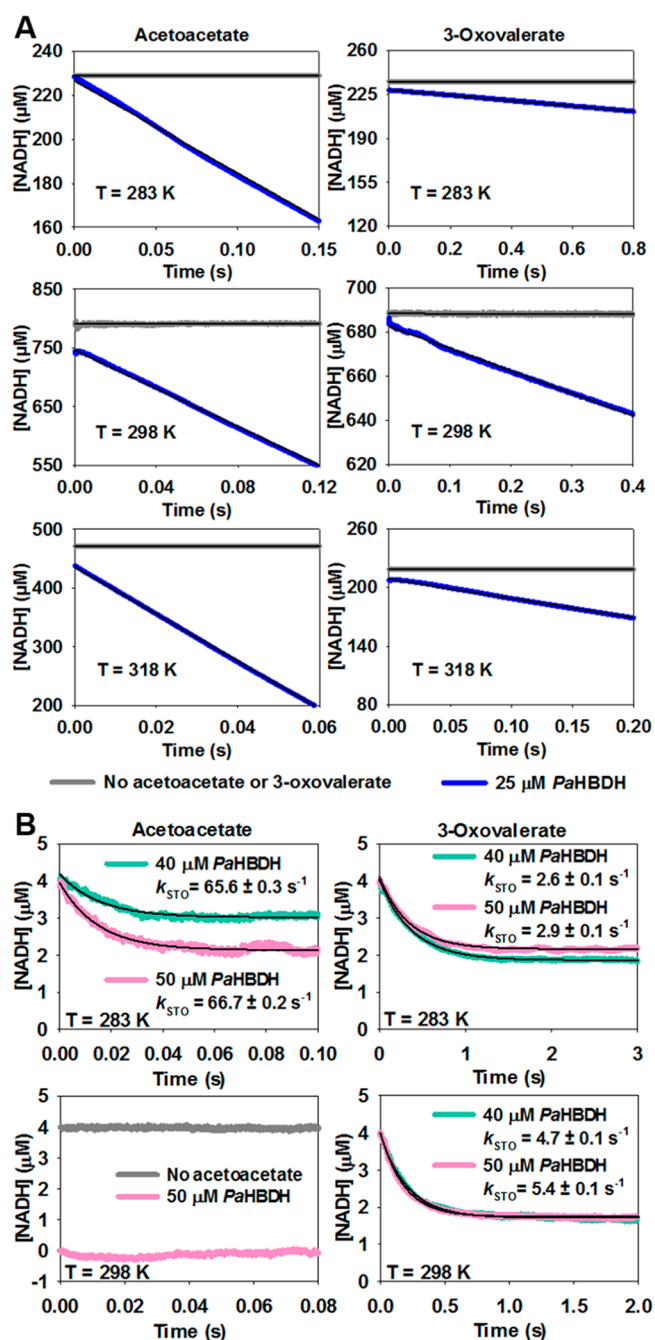


Figure 3. Rapid kinetics for *Pa*HBDH with acetoacetate and 3-oxovalerate as substrates. (A) Multiple-turnover pre-steady-state kinetics. Black lines are fits of data to eq S3. (B) Single-turnover pre-steady-state kinetics. Black lines are fits of data to eq S4.

there is no burst (Figure 3A and Figure S6), indicating a change in the rate-limiting step at low temperatures.

The hypotheses regarding rate-limiting steps in the *Pa*HBDH reaction at different temperatures were tested further by single-turnover pre-steady-state kinetics at 283 and 298 K with NADH as the limiting reagent. *Pa*HBDH single-turnover rate constants (k_{STO}) were independent of enzyme concentration (Figure 3B), demonstrating NADH is saturated with *Pa*HBDH and k_{STO} is unimolecular.²⁴ With 3-oxovalerate, k_{STO} and k_{cat} values are similar at the same temperatures (Table S11). Along with the absence of a pre-steady-state burst and a negligible solvent viscosity effect on k_{cat} with this substrate, this

result indicates that interconversion between *Pa*HBDH-NADH-3-oxoalate and *Pa*HBDH-NAD⁺-3-hydroxyvalerate is rate-limiting. With acetoacetate as the substrate, if the hypothesis emerging from multiple-turnover pre-steady-state kinetics and solvent viscosity effects holds, k_{STO} should be measurable at 283 K but should not be possible to monitor at 298 K as it takes place within the dead time of the instrument. This is exactly what is observed (Figure 3B and Table S11), with k_{STO} being too fast to be detected at 298 K although NADH is consumed. At 283 K, k_{STO} is 2.2-fold higher than k_{cat} , which, combined with the absence of a burst, indicates a step prior to acetoacetate reduction is rate-limiting at this temperature.

Whereas there is precedent for a change in the rate-limiting step in nonlinear Eyring plots of enzyme reactions,²⁶ *Pa*HBDH Eyring plots are linear in the range of 283–308 K (Figure S7), and the switch in the rate-limiting step occurs between 283 and 298 K. This indicates a temperature-dependent shift in the rate-limiting step giving rise to a linear Eyring plot.

To assess whether this unusual behavior is displayed by the mesophilic enzyme, we carried out rapid kinetics experiments with *Ab*HBDH, whose Eyring plots are linear throughout the experimental temperature range. Remarkably, multiple-turnover pre-steady-state kinetics at 283, 298, and 328 K (Figure 4A) show a change in the rate-limiting step with both substrates. A burst in substrate consumption at 298 and 328 K with acetoacetate and at 328 K with 3-oxoalate is too fast to be directly observed but is clearly inferred²⁵ from the offset in NADH concentration at time zero in the presence and absence of acetoacetate or 3-oxoalate (Figure S6), indicating the first turnover occurs within the dead time of the stopped-flow spectrophotometer. Conversely, there is no burst at 283 K with either substrate or at 298 K with 3-oxoalate (Figure 4A and Figure S6). The results at 298 K are in strict agreement with the conclusions drawn from solvent viscosity effects for both substrates.

As with *Pa*HBDH, we performed single-turnover pre-steady-state kinetics experiments at 283 and 298 K, with NADH as the limiting reagent, to examine further the hypothesis of a change in the rate-limiting step in *Ab*HBDH catalysis at different temperatures. *Ab*HBDH k_{STO} values were independent of enzyme concentration (Figure 4B), demonstrating NADH is saturated with *Ab*HBDH and k_{STO} is unimolecular. The k_{STO} for acetoacetate reduction can be measured at 283 K, but it is too fast to be monitored at 298 K, even though NADH is consumed. This is in accordance with the results from rapid kinetics under multiple-turnover conditions. The k_{STO} for 3-oxoalate reduction can be measured at 283 and 298 K, as predicted by the absence of a burst at these temperatures. The affinity between *Ab*HBDH and its substrates decreases severely at high temperatures, which prevented single-turnover experiments at 328 K (the enzyme could not be concentrated enough to ensure sufficient binding to NADH). Combined, these results demonstrate a shift in the rate-limiting step between 283 and 298 K for *Ab*HBDH-catalyzed reduction of acetoacetate and between 298 and 328 K for reduction of 3-oxoalate despite linear Eyring plots with both substrates.

The k_{STO} s for *Ab*HBDH-catalyzed reduction of 3-oxoalate are 10- and 7-fold higher than k_{cat} at 283 and 298 K, respectively, and the k_{STO} for acetoacetate reduction at 283 K is 8-fold higher than the corresponding k_{cat} (Table S12). Therefore, unlike the *Pa*HBDH reaction, a step preceding chemistry limits the *Ab*HBDH reaction rate with 3-oxoalate

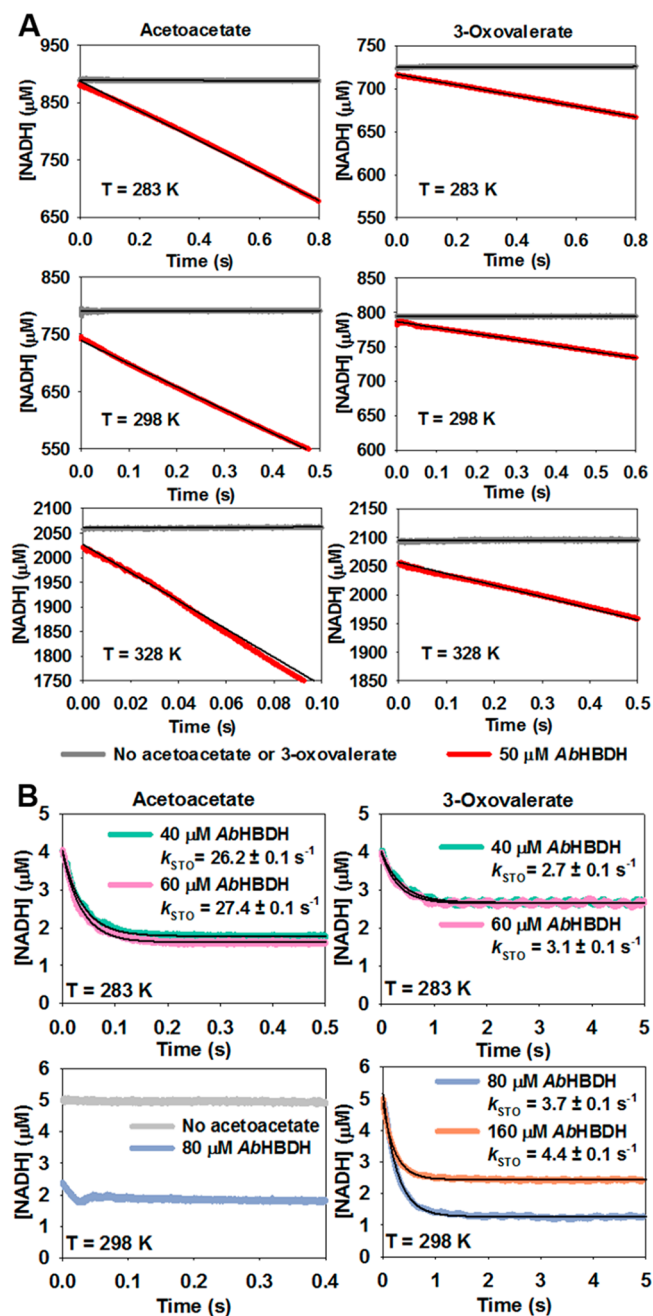


Figure 4. Rapid kinetics for *Ab*HBDH with acetoacetate and 3-oxoalate as substrates. (A) Multiple-turnover pre-steady-state kinetics. Black lines are fits of data to eq S3. (B) Single-turnover pre-steady-state kinetics. Black lines are fits of data to eq S4.

at 283 and 298 K and with acetoacetate at 283 K. Distinct rate-limiting steps between enzyme orthologues have been reported for other members of the SDR superfamily.^{27,28}

Temperature–rate profiles are invaluable for the investigation of enzymatic mechanisms, encompassing probing quantum mechanical tunneling,³ enzyme dynamics,^{10,12} allostery,^{4,5} and calculation of thermodynamic parameters of activation.^{6,9–12} They are also instrumental for studying the evolution of modern enzymes from primitive catalysts that arose in a hot Earth.^{2,29,30}

A temperature-dependent change in the rate-limiting step could produce a linear Eyring plot for k_{cat} if the ratio of Eyring plot slopes for the microscopic rate constant(s) determining

each step is relatively small. This scenario is demonstrated (Supporting Information) from simulated data (Figure S8) for a hypothetical two-step mechanism (Scheme S2) and may explain the results with AbHBDH. This highlights the importance of confirming whether the same rate-limiting step holds throughout the experimental temperature range before further interpretation of both nonlinear and linear Eyring plots of macroscopic rate constants.

■ ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acs.biochem.8b01099.

Materials and Methods, Schemes S1 and S2, Figures S1–S8, a discussion of simulated Eyring plots, and Tables S1–S12 (PDF)

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Notes

The authors declare no competing financial interest.

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■ ABBREVIATIONS

HBDH, (R)-3-hydroxybutyrate dehydrogenase; SDR, short-chain dehydrogenase/reductase; ΔC_p^\ddagger , activation heat capacity; k_{cat} , steady-state catalytic constant; k_{STO} , single-turnover rate constant.

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