# Taming Highly Enolizable Aldehydes via Enzyme Catalysis for Enantiocomplementary Construction of $\beta$ -Hydroxyphosphonates

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**Cite This:** J. Am. Chem. Soc. 2025, 147, 3102–3109

anions). Unlike NHC-mediated reactions that yield complex





mixtures of multiple adducts, our enzymatic process selectively produces biologically active  $\beta$ -hydroxy phosphonates with high yields (up to 95%) and excellent enantioselectivities (up to 99% ee). The products can be obtained on gram scales and exhibit rich reactivity for downstream transformations to afford diverse molecules. *Pf*BAL (or its mutant A28G) and *Pa*BAL enzymes serve as enantiocomplementary pairs, enabling the synthesis of both product configurations. Mechanistic studies proved that the entrance directions of the active cavities of these two enzyme pairs were distinct, leading to acyl anions formed from these two enzyme pairs attacking 2-phosphonate aldehydes from different orientations.

# INTRODUCTION

Chiral  $\beta$ -hydroxyphosphonates are unique moieties with a large presence in medicines, agrochemicals, and natural products (Figure 1A).<sup>1</sup> For example, phosphonothrixin, a natural product isolated from Saccharothrix, contains a  $\beta$ -hydroxyphosphonate moiety and exhibits herbicidal activity.<sup>1a</sup> Valinophos, a phosphonopeptide natural product, exhibits important antibiotic activity.<sup>Ib</sup> Fosfomycin is a widely used antibiotic for treating bacterial infections.<sup>1c</sup> Therefore, the development of an efficient and enantioselective strategy for the construction of  $\beta$ -hydroxyphosphonates is of great interest. Traditional synthetic methods rely on the lipase- or organocatalyst-mediated kinetic resolution of racemic alcohols or esters and afford products with a less than 50% yield (Figure 1B, left).<sup>2</sup> Although reductase or metal-catalyzed reduction of ketophosphonates or vinylphosphonates has been developed, these methods suffer from lengthy routes for substrate preparation (Figure 1B, right).<sup>3</sup> A more attractive approach, based on retrosynthetic analysis, is C-C bond coupling of highly enolizable 2-phosphonate aldehydes with C-nucleophiles (or addition of C-nucleophiles to highly enolizable 2phosphonate aldehydes) as it uses readily available materials as substrates and offers easily tunable product structures (Figure 1B, down). However, compared with widely explored simple alkyl or aryl aldehydes,<sup>4</sup> taming highly enolizable aldehydes (introducing a strong electron-withdrawing group, such as -PO<sub>3</sub>R, -COOR, -NO<sub>2</sub>, -SO<sub>2</sub>R, etc., to the  $\alpha$ -position of aldehyde moieties) for catalytic asymmetric C-C coupling remains an elusive challenge and has been scarcely explored

(Figure 1C).<sup>5</sup> The main difficulties are probably attributed to the following factors: (i) the deactivation of common nucleophiles by acidic H at the  $\alpha$ -position of aldehyde moieties;<sup>6</sup> (ii) undesirable aldol-type side reactions in the reaction system;<sup>7</sup> (iii) limited catalytic asymmetric strategies;<sup>8</sup> and (iv) the inherent instability of highly enolizable aldehydes.

Herein, we report a ThDP-dependent enzyme-enabled C-C bond coupling reaction between highly enolizable 2phosphonate aldehydes and nucleophiles, producing biologically active  $\beta$ -hydroxy phosphonates with high yields and excellent chemo- and enantioselectivities (Figure 1D). In this reaction, C-nucleophiles (acyl anions) are catalytically generated in situ from the polarity inversion of another aldehyde, which is mild, dynamically reversible,9 and compatible with the acidic H-containing 2-phosphonate aldehydes. In addition, the protein cavity of the enzyme catalyst provides subtle noncovalent interactions and molecular recognition, ensuring high reactivity and excellent chemo-/ enantioselectivity. Although ThDP-dependent enzymes are well-known for catalyzing benzoin-type C-C couplings between two simple aldehydes,<sup>10</sup> their application to highly enolizable aldehydes remains underexplored despite offering

Received:August 29, 2024Revised:December 11, 2024Accepted:January 9, 2025Published:January 17, 2025





Figure 1. Chiral  $\beta$ -hydroxyphosphonate-containing molecules and their synthetic strategies.

access to hard-to-reach products and facilitating the concise synthesis of complex natural products or medicines.<sup>11</sup> In addition, the synthesis of two mirror-image molecules with ThDP-dependent enzymes is very difficult, owing to the lack of enantiocomplementary enzyme pairs. In 2014, Pohl and coworkers engineered ApPDC to produce (S)-benzoins, complementing the (R)-benzoins from natural enzymes, but the substrates were limited to aromatic aldehydes.<sup>12</sup> Moreover, only a few substrates have been reported for enantiocomplementary cross-coupling between aromatic aldehydes and simple alkyl aldehydes, typically with low to moderate yields and ee.<sup>10a,13</sup> In this work, PfBAL and PaBAL were demonstrated to be enantiocomplementary ThDP-dependent enzyme pairs, yielding  $\beta$ -hydroxyphosphonates in two configurations. Mechanistic studies proved that the active cavities of the two proteins had distinct entrance directions, leading to acyl anions attacking 2-phosphonate aldehydes in different orientations.

#### **RESULTS AND DISCUSSION**

Reaction Development. We initiated our studies by investigating a panel of structurally diverse ThDP-dependent enzymes (cell-free lysates; for phylogenetic analysis, see Figure S2 in the SI) for their ability to catalyze the benzoin-type coupling reaction of 1a or 1m with the 2-phosphonate acetaldehyde 2a. Key results are summarized in Table 1 (see Table S7 in the SI for details). Benzaldehyde lyase from Pseudomonas fluorescens (PfBAL, PDB ID: 3D7K) furnished the (S)-selective  $\beta$ -hydroxyphosphonate 3a with 90% yield and 98% ee (entry 10). However, the yield dropped sharply when using the electron-withdrawing-group-substituted 1m (entry

Table 1. Optimization of the Reaction Conditions<sup>a</sup>

;	X=H, X=F,	H EtO EtO 1a 2a	EtO EtO 2a EtO EtO EtO EtO EtO EtO EtO EtO EtO EtO			
			$1a \rightarrow 3a$		$1m \rightarrow 3m$	
	entry <sup>a</sup>	enzyme	yield (%) <sup>b</sup>	R/S ratio <sup>b</sup>	yield (%) <sup>b</sup>	R/S ratio <sup>b</sup>
	1	EcMend	trace		0	
	2	<i>Tp</i> BFD	trace		0	
	3	KdcA	0		0	
	4	SsPDC	0		0	
	5	EcTK	0		0	
	6	ScTK	0		0	
	7	SeAAS	0		0	
	8	SsBAL	10	1:99	trace	
	9	SuBAL	7	1:99	0	
	10	<i>Pf</i> BAL	90	1:99	10	1:99
	11	PaBAL	48	96:4	19	95:5
	12	PaBAL (cell) <sup>c</sup>	62	96:4	31	95:5
	13	PaBAL (cell) <sup>d</sup>	85	96:4	60	95:5

<sup>a</sup>Reaction conditions: 1a or 1m (0.02 mmol), 2a (0.04 mmol), cellfree extract (50 mg/mL, 400 µL), ThDP (0.15 mM), MgSO<sub>4</sub> (2.5 mM), KPB buffer (50 mM, pH 7.25), DMSO (20% v/v), 1000 rpm, 20 °C, 24 h. <sup>b</sup>Yield and R/S ratio were determined by chiral HPLC. <sup>c</sup>pH 7.25, whole cell (50 mg/mL). <sup>d</sup>pH 8.0, whole cell (100 mg/mL).

10). Surprisingly, benzaldehyde lyase from Polymorphobacter arshaanensis  $(PaBAL)^{14}$  yielded 3a and 3m with the opposite (R)-selective configuration (entry 11), enabling the enantio-



Figure 2. (a) Molecular docking of the ThDP-bound 3m precursor. (b) Screening of mutants. (c) Specific activity of *Pf*BAL and A28G.



(b) One-step transformations of (S)-3a



Figure 3. Gram-scale reaction and product transformation.

complementary synthesis of  $\beta$ -hydroxyphosphonates. The yields of **3a** and **3m** were further improved using whole-cell *Pa*BAL and increasing the loading of the enzyme catalyst (entry 13). Subsequently, we focused on engineering *Pf*BAL to enhance the reactivity of substrate **1m** through site saturation mutagenesis (SSM). Molecular docking of the ThDP-bound **3m** precursor with *Pf*BAL identified four key amino acid residues (Y397, L112, Q113, and A28) located within 4 Å of the ThDP-bound **3m** precursor as hot-spots for SSM (Figure 2a). However, no positive mutants were identified when we performed SSM at residues Y397, L112, and Q113, with most mutants exhibiting no catalytic activity (see Tables S11 and S12 in the SI). Surprisingly, the next round of SSM at residue

A28 resulted in notable yield improvement using the mutant A28G, producing the desired 3m in 95% yield and 99% ee (Figure 2b; see Table S11 in the SI for details). Molecular dockings revealed that the attack distance of the 1m-bound acyl anion intermediate to substrate 2a in mutant A28G was shorter than that in wild-type PfBAL (4.4 vs 5.8 Å). This may be because the replacement of Ala with Gly at position A28 creates more space for substrate conformational rotation, resulting in a more efficient catalytic conformation (see Figure S3 in the SI). Testing the reactivity of mutant A28G with substrate 1a also achieved excellent yield and enantioselectivity (93% yield, 99% ee) (Figure 2b). Subsequent specific activity analysis revealed that the mutant A28G showed slightly higher specific activity than PfBAL toward 1a and a notable improvement of specific activity toward 1m (Figure 2c), aligning with the yield results. Although four (cross) benzoin products could potentially form, the only byproduct observed was the self-benzoin product of arylaldehydes, with its yield decreased to below 5% (see Tables S7 and S11 in the SI).

Reaction Scope. With the enantiocomplementary enzyme pairs in hand, we then investigated their catalytic scope for the benzoin-type C-C coupling reaction of highly enolizable aldehydes on a preparative scale (Table 2). The results of PfBAL A28G catalysis showed that methyl substituents at the para-, meta-, and ortho-positions of aryl aldehydes were well tolerated, furnishing the corresponding (S)-selective products (3b to 3d) with 32-95% yields and excellent ee values. However, the ortho substituent proved to be unreactive for the wild-type PfBAL and PaBAL. Strong electron-donating groups at the phenyl ring of aryl aldehydes, such as hydroxyl or methoxy group, were also well compatible with PfBAL A28G, wild-type PfBAL, and PaBAL (except para-hydroxyl), providing products 3e-3g with excellent yields and ee values. Disubstituted aldehyde afforded the product 3h with an increased yield for PfBAL A28G compared with wild-type PfBAL. Furthermore, the larger substrate 2-naphthaldehyde failed to yield the corresponding product with wild-type PfBAL, while PfBAL A28G produced 3i with 85% yield and 97% ee, attributed to the expansion of the active pocket in PfBAL A28G. Both disubstituted aldehyde and 2-naphthaldehyde were suitable substrates for wild-type PaBAL, yielding (R)-selective products (3h and 3i, respectively). Although wild-type PfBAL showed good compatibility with various electron-withdrawing groups (such as Br, Cl, and F) at the meta position of aryl aldehydes (3j to 3l), sharply decreased yields were observed for para-substituted electron-withdrawing groups (3m to 3q). However, the mutant A28G was compatible with both meta- and para-substituted electronwithdrawing groups (3j to 3q). Substrates with various electron-withdrawing groups were also suitable for PaBAL (3i to 3q) and showed a better result for substrates with parasubstituted electron-withdrawing groups (3m to 3q) compared to wild-type PfBAL and its mutant A28G. The investigation of furan and thiofuran heterocycles (3r to 3t) showed that wildtype PfBAL displayed higher reactivities than PfBAL A28G and PaBAL. Notably, cyano- and pyridine-containing substrates were unreactive under the current reaction conditions (not shown). Next, we tested the compatibility of the other phosphonate substrates. Diisopropoxyphosphate substrates were also well accepted in this benzoin-type coupling process by both mutant A28G and wild-type PfBAL, producing (S)selective products (**3u** to **3w**). *Pa*BAL also yielded (*S*)-selective products (3u to 3w), but with low ee values. We imagine that

Table 2. Substrate Scope<sup>a</sup>



<sup>*a*</sup>Reaction condition: 1 (0.15 mmol), 2 (0.3 mmol), cell-free extract for *Pf*BAL and *Pf*BAL A28G (50 mg/mL, 4 mL), whole cell for *Pa*BAL (100 mg/mL, 4 mL), ThDP (0.15 mM), MgSO<sub>4</sub> (2.5 mM), KPB buffer (50 mM, pH 7.25 for *Pf*BAL and *Pf*BAL 28G; pH 8.0 for *Pa*BAL), DMSO (20% v/v), 1000 rpm, 20 °C, 24 h; isolate yield. <sup>*b*</sup>2a (2.5 equiv). <sup>*c*</sup>Whole-cell *Pa*BAL (150 mg/mL).

protein engineering could be used to further invert the enantioselectivity for 3u to 3w (see Figure S9 in the SI) and should be applied to phosphate substrates with extended carbon chains (3x and 3y).

**Synthetic Applications.** To demonstrate the utility of the current method, gram-scale reactions and product transformations were conducted (Figure 3). Both the wild-type PfBAL and its mutant A28G successfully catalyzed the benzoin-type coupling reaction of highly enolizable aldehydes on gram scales, affording the desired products in 80% yield

(1.5 g) for 3a and 85% yield (1.45 g) for 3m (Figure 3a). Furthermore, the enantioenriched phosphonate product 3a was easily converted to various derivatives through a one-step reaction (Figure 3b). For instance, the reduction and Grignard reaction of 3a provided vicinal diol-containing phosphonates 4a and 4b. The ketone unit of 3a was efficiently transformed into imine, producing 4c with hydroxyl imine. The phosphonate product 3a also reacted with the Wittig reagent to form alkene 4d without the loss of ee. In addition, 3a was easily acylated and oxidized, affording the corresponding



<sup>a</sup>Reaction conditions: **1a** (0.15 mmol), **2a** (0.3 mmol), NHC (20 mol %), DIPEA (1.0 equiv) in THF (0.75 mL) at rt for 24 h. <sup>b</sup>Conversion was determined by GC. <sup>c</sup>Yields were determined by GC. <sup>d</sup>ee of **3a** was determined by chiral HPLC; ee of **3a**' and **3ab** was not measured.

products (4e and 4f) in good yields. Finally, the elimination of the hydroxyl group in product 3a was realized by reacting 3a with TsCl in the presence of a base. Notably, all of the products derived from 3a via a one-step reaction could serve as versatile synthetic precursors for further transformations, such

as the reduction of imine **4c** or various functionalizations of alkenes **4d** and **4g**.

Comparison with NHC Organic Catalysis. Achieving chemoselective cross-benzoin reactions between two different aldehydes is a classic challenge in synthetic chemistry owing to the presence of three undesirable competing reactions (two homobenzoins and one cross-benzoin).<sup>15</sup> This problem is further complicated when highly enolizable aldehydes are involved as the substrates because of a series of potential side reactions, such as self-/cross-benzoin, self-/cross-aldol reactions, iterative aldol reactions, and the Stetter reaction of aldol products. To demonstrate the unique ability of enzymes to control the chemical selectivity in cross-benzoin reactions involving highly enolizable aldehydes, we conducted the same reaction using different NHC organic catalysts for comparison (Table 3). As speculated, commonly used chiral NHC catalysts A-F failed to achieve chemoselective cross-benzoin reactions (Table 3). In addition, insufficient conversion, low yields of benzoin-related products, and unsatisfactory stereoselectivities further limited the development of chemoselective crossbenzoin reactions involving highly enolizable aldehydes in NHC organic catalysis. Self- or cross-aldol reactions also appeared in all reaction systems along with other complicated byproducts. These results indicate that avoiding or minimizing competing pathways in cross-benzoin reactions with highly enolizable aldehydes remains a daunting challenge in NHC organic catalysis. In contrast, our enzymatic approach gives unparalleled chemoselective control for this reaction with simultaneous control of the stereochemistry.

**Mechanistic Study.** To understand the reaction pathway, mechanistic experiments were conducted (Figure 4). Reaction process experiments showed that the benzoin 3a' (*R* configuration, > 99% ee) formed within 1 h under *Pf*BAL and its mutant A28G catalysis, with *Pf*BAL exhibiting better



Figure 4. Mechanism-related experiments.

https://doi.org/10.1021/jacs.4c11957 J. Am. Chem. Soc. 2025, 147, 3102-3109



**Figure 5.** Docking and MD simulations for clarifying the origins of enantioselective complementarity. (a) Active pocket of PfBAL; (b) active pocket of PaBAL; (c) MD simulation for PfBAL; and (d) MD simulation for PaBAL. (e) Relative frequency distribution of the distance between the oxygen atom of **2a** and the hydrogen of the G398 residue. **1a**-bonded acyl anion intermediate, **2a**, and the residue are shown as sticks.

reactivity for producing benzoin 3a' (Figure 4a,i,ii). Upon prolonging the reaction time, 3a' was gradually converted into the desired product 3a under the catalysis of PfBAL or its mutant A28G while maintaining a high level of stereoselectivity. Specific activity analysis demonstrated that the reactivity of natural benzaldehyde was higher than that of unnatural 2-phosphonate aldehyde for PfBAL and PfBAL A28G catalysis (see Table S14 in the SI). These results indicated that the formation of the desired product 3a through PfBAL and its mutant A28G catalysis involves a dynamically reversible process of benzoin 3a' formation. Further investigations showed that racemic benzoin or (R)-benzoin could also be used as the starting material in our new reaction catalyzed by PfBAL or PfBAL A28G (see Table S15 in the SI). In contrast, no benzoin 3a' was detected during the formation of the desired product 3a under PaBAL catalysis (Figure 4a,iii). Next, 3a was separately exposed to PfBAL, PfBAL A28G, or PaBAL, and benzaldehyde 1a was detected after 24 h in all experiments (Figure 4b). These results prove that the formation of 3a and the nucleophilic acyl anion intermediate are also dynamically reversible processes. The dynamically reversible feature of the in situ-generated acyl anion makes it an ideal nucleophile compatible with acidic H-containing 2phosphonate aldehydes. Built upon these experiments (Figure

4a,b), a postulated reaction pathway is briefly illustrated in Figure 4c.

To clarify the origins of enantioselective complementarity, AlphaFold 2 was employed to predict the structure of PaBAL. We found that the overall structure of PaBAL is strikingly similar to that of PfBAL (Figures S5 and S6 in the SI) despite PaBAL sharing 41% sequence identity with PfBAL based on amino acid sequence alignment (Figure S4 in the SI). Interestingly, the entrance direction of the active cavities in the two proteins is distinct (Figure 5a,b), which may contribute to the different configurations of products. Next, 100 ns molecular dynamic (MD) simulations were conducted using AutoDock vina<sup>16</sup> and Amber24. We found that the acyl anion intermediate in PfBAL attacks the Si face of 2a, yielding the S configuration products (Figure 5c). Further analysis of the amino acid residues in the active pocket of PfBAL showed that four residues (G394, A395, L396, and L399) prevented **2a** from reaching the bottom of the acyl anion intermediate to produce the R configuration products. In contrast, the Re face of 2a is attacked by an acyl anion intermediate in PaBAL to form R configuration products (Figure 5d). It is obvious that there is not enough space on the top of the acyl anion intermediate to accommodate 2a (to produce the S configuration products) in PaBAL due to the appearance of H32, G33, G34, H35, and T79. Notably, a weak hydrogen bond between the G398 residue and 2a was observed in PaBAL (Figure 5d,e), which also controls the orientation of the aldehyde group in 2a. These results are consistent with our experimental observations.

# CONCLUSIONS

In summary, we have developed a novel C-C coupling system for the enantiocomplementary synthesis of chiral  $\beta$ -hydroxyphosphonates. This enzyme-mediated transformation is hard to achieve with small-molecule NHC catalysts owing to the complex mixture of multiple adducts. Diverse control experiments and mechanism studies clarified the reaction pathway and the source of the enantioselective complementarity of *Pf*BAL. The chiral  $\beta$ -hydroxyphosphonate products can be obtained on gram scales and exhibit rich reactivities for further downstream transformations, creating additional opportunities for the development of novel C–P drugs.<sup>1</sup> Further studies on the bioactivities of these chiral phosphonates for agricultural and medical applications, as well as the application of this method to the total synthesis of complex natural products, are in progress in our laboratories. Our study offers a new perspective on the C-C coupling of highly enolizable aldehydes with C-nucleophiles (or the addition of Cnucleophiles to highly enolizable aldehydes) and will encourage the further development of new systems to tame highly enolizable aldehydes for asymmetric synthesis.

#### ASSOCIATED CONTENT

#### **Supporting Information**

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/jacs.4c11957.

Experimental methods; optimization data; protein engineering; details of docking and MD simulations; chemical characterization; and NMR spectra (PDF)

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### Notes

The authors declare no competing financial interest.

# ACKNOWLEDGMENTS

The authors acknowledge the National Natural Science Foundation of China (32301231, 32121005), the East China University of Science and Technology for startup funding (YF0142210), the Fundamental Research Funds for the Central Universities (JKF01231815), the National Key Research and Development Program of China (2020YFA0907200, 2020YFA0907800, 2022YFC2303100, and 2022YFC2303104), the Open Project Funding of the State Key Laboratory of Bioreactor Engineering, and the 111 Project (B18022).

# ABBREVIATIONS

MD, molecular dynamics; NHC, *N*-heterocyclic carbene; ThDP, thiamine diphosphate

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