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PERSPECTIVE

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1. Introduction

Oxidation is an essential reaction for chemical synthesis. Historically, oxidation reactions (also on an industrial scale) utilize high-valent Cr- and Mn-salts, generating enormous amounts of environmentally questionable wastes. In the last decades, the research focus has shifted towards more acceptable stoichiometric oxidants, such as O_2 or H_2O_2 .¹ Next to organocatalytic² and transition-metal-catalyzed³ approaches, biocatalytic oxidation methods are also gaining interest, and a broad range of enzymes are available nowadays to perform selective oxidation reactions under very mild reaction conditions (Fig. 1).

Mechanistically, biocatalytic oxidation reactions fall either into the 'oxyfunctionalizing' or the 'dehydrogenative' branch.⁴ Oxyfunctionalizing reactions have been traditionally catalyzed by so-called monooxygenases. These enzymes reductively activate O_2 of which one O-atom is inserted into the starting material while the other O-atom is used to form H_2O as a final product. More recently, 'unspecific peroxygenases' (UPOs) are receiving increasing interest as alternatives to monooxygenases. UPOs, in contrast to monooxygenases, directly utilize partially reduced oxygen (in the form of hydrogen peroxide, H_2O_2).⁵ Dehydrogenative oxidation reactions typically imply the oxidation of alcohols or amines to the corresponding ketones (imines), also aldehyde oxidations to the corresponding carboxylic acids fall into this category. The reducing equivalents

Light-driven biocatalytic oxidation

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Enzymes are the catalyst of choice for highly selective reactions, offering nature-inspired approaches for sustainable chemical synthesis. Oxidative enzymes (*e.g.*, monooxygenases, peroxygenases, oxidases, or dehydrogenases) catalyze a variety of enantioselective oxyfunctionalization and dehydrogenation reactions under mild conditions. To sustain the catalytic cycles of these enzymes, constant supply with or withdrawal of reducing equivalents (electrons) is required. Being redox by nature, photocatalysis appears a 'natural choice' to accomplish the electron-relay role, and many photoenzymatic oxidation reactions have been developed in the past years. In this contribution, we critically summarize the current developments in photoredoxbiocatalysis, highlight some promising concepts but also discuss the current limitations.

liberated in these transformations are transferred either to nicotinamide cofactors (dehydrogenases) or to molecular oxygen (oxidases).⁴

As redox catalysts, all above-mentioned enzymes rely on the supply with redox equivalents to maintain their catalytic cycles. Counter-intuitively (at first sight) monooxygenases necessitate supply with reducing equivalents, which are—more or less directly—delivered to the monooxygenases' active sites from the reduced nicotinamide cofactors [NAD(P)H].⁶ Considering the overall four electron reduction of O_2 catalyzed by mono-oxygenases and the fact that only two electrons stem from the starting material, this dependence on reducing equivalents, however, becomes clear. Established enzymatic methods to *in situ* regenerate the reduced nicotinamide cofactors comprise, amongst others, formate dehydrogenase-catalyzed oxidation of formic acid or glucose dehydrogenase-catalyzed oxidation of glucose.

Envisioning the use of clean solar power as energy source, photocatalysis has emerged as an alternative to the abovementioned enzymatic regeneration systems.7-10 The promise of photocatalytic regeneration systems lies with access to a broader range of stoichiometric sources or acceptors of redox equivalents and simplified reaction schemes. Many photocatalysts-such as organic dyes, metal nanoparticles, and semiconductor quantum dots-have been tested to activate redox enzymes via the transfer of photoinduced electrons. In these systems, sacrificial electron donors [e.g., triethanolamine (TEOA), ethylenediaminetetraacetic acid (EDTA), ascorbic acid, or even water] have been used to recycle a cofactor (or a mediator) to sustain the biocatalytic reaction cycle. The reducing equivalents are transferred either directly to the oxidoreductase's active site or indirectly via the natural nicotinamide cofactors.

Alternatively, in recent years, so-called peroxyzymes (*i.e.*, enzymes using H_2O_2 as a stoichiometric oxidizing agent) have



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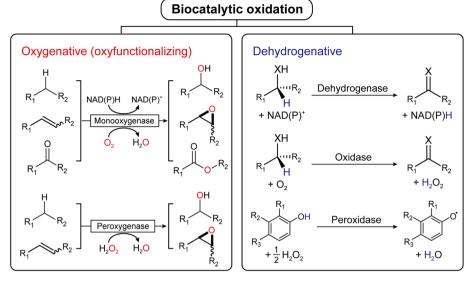


Fig. 1 Biocatalytic oxidation is categorized into either oxygenative (oxyfunctionalizing) or dehydrogenative reactions.

gained considerable interest in the biocatalysis community. Particularly, peroxygenases [E.C. 1.11.2.] (https:// www.iubmb.qmul.ac.uk/enzyme/EC1/11/2/) are promising alternatives to well-known P450 monooxygenases because of the significantly simpler catalytic cycle. Nevertheless, the poor robustness of heme-enzymes against H_2O_2 necessitates its provision in controlled amounts to balance stability and reactivity. This offers enormous possibilities for photocatalysis to solve this issue.

Finally, $NAD(P)^+$ -dependent oxidation reactions are also an attractive target for photocatalysis as simple photocatalytic NAD(P)H oxidation reactions are at hand.

More detailed information about the enzymes discussed in this contribution (*e.g.*, genbank/Uniprot ID, heterologous expression conditions, and other technical details) have been summarized in Table 1.

2. Light-driven monooxygenase catalysis

Monooxygenases catalyze the insertion of one oxygen atom (O) from molecular oxygen (O₂) into an organic molecule. For photochemical activation of a monooxygenase, photoinduced electrons should be transferred directly or indirectly to the enzyme's active site (Fig. 2). During direct transfer of photoinduced electrons, a photocatalyst and a monooxygenase having a redox prosthetic group (*e.g.*, heme, flavin), interact in a relationship between an electron donor and an acceptor, respectively. The excited state of the photocatalyst is quenched by the oxidation of a sacrificial electron donor (*e.g.*, EDTA, TEOA) to sustain the turnover of the photocatalyst.

In 2007, Hollmann *et al.* devised and implemented experimentally a light-driven catalytic scheme for flavin-dependent Baeyer–Villiger monooxygenase (BVMO, *e.g.* PAMO-P3)²⁹ (Fig. 2a). Photochemical reduction of free flavin molecules (a photocatalyst) using EDTA produces reduced ones, which then transfer electrons to the FAD cofactor bound to the PAMO-P3 active site. Overall, visible light promotes direct reductive regeneration of the PAMO-P3-bound FAD; thus, the scheme does not require costly NADPH (or a complicated NADPH regeneration system). Here, the light-driven platform of PAMO-P3 for Baeyer–Villiger reactions of ketone substrates showed 48–93% conversion of ketone substrates with excellent enantiose-lectivity, which were comparable to conventional NADPH-based process (albeit at reduced catalytic rates). However, already then two major limitations had been pointed out: oxidative uncoupling of the regeneration reaction from enzymatic O₂-activation and the slow electron transfer between free and enzyme-cofactors.³⁰

A whole-cell-based, light-driven P450 platform has been devised using eosin Y (EY) as a photocatalyst and EDTA as an electron donor³¹ (Fig. 2b). EY could easily enter into the *Escherichia coli* cytoplasm and then bind to the P450's heme domain. The catalytic reaction of P450 was mediated *via* the direct transfer of light-induced electrons from the photo-activated EY to the heme domain of P450 under visible light. Photochemical activation of the P450 catalysis has been successfully demonstrated using many variants of CYP102A1 and human P450s for the bioconversion of various substrates. A possible contribution of peroxygenase activity in the whole-cell P450 photocatalysis however cannot be excluded, which needs further investigation.

Besides, several other light-driven whole-cell systems have been reported: (1) Rieske oxygenase using 5(6)-carboxyeosin as a photosensitizer and MES buffer as electron donor;³² (2) photoautotrophic cyanobacterium *Synechocystis* expressing the heterologous P450 (ref. 33); (3) BVMO reactions in metabolically engineered cyanobacteria;³⁴ (4) microbial photosynthesis works for O₂ generation for biocatalytic oxyfunctionalizations.³⁵

In addition to the photosensitizing dyes (*e.g.*, flavin,^{12,13} EY^{31}), ruthenium (Ru)³⁶ complexes have been used as

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 Table 1
 Detailed information about representative oxidative enzymes discussed in this article

Representative enzyme Genbank ID (Uniprot ID) Phenylacetone CP000088 (Q47PU3) Pnenylacetone CP000088 (Q47PU3) nonoxygenase (PAMO-P3 variant) from bacterium Thermobifida fusca CP100058 (Q47PU3) Thermobifida fusca CP100058 (Q47PU3) Thermobifida fusca CP100058 (Q47PU3) Thermobifida fusca CP1000058 (Q47PU3) CYP1021 (Personants) WP_03450556 (P14779.2) Thuman CYP2E1 NM_000773 (P05181.1) Human CYP2E1 NM_000773 (P05181.1) Human CYP2E1 NM_000773 (P05181.1) Pacterium Bacilius subitis NM_000773 (P05181.1) Pacterium Bacilius subitis SU02100 (03140.1) Pacterium Costridium NM_000773 (P051821.1) Pacterium Costridium SU02100 (03140.1) Pacterium Costridium SU02100 (03140.1) AccUP from mushroom Hms72457 (B9W4V6) AccUP from mushroom AccUP (P04963) AccUP from mushroom AccP1741821 AccUP from mushroom AccP1741821 AccUP from mushroom AccP1745666 AccUP from mushroom <th>Classification of</th> <th></th> <th></th> <th></th> <th></th>	Classification of				
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Muman CYP2E1NM_000773 (P05181.1)Human CYP2E1NM_000773 (P05181.1)Human CYP2E1NM_000773 (P05181.1)CYP152A2 (P450c1A) from bacterium Bacillus subilis str. 168BSU02100 (031440.1)bacterium Bacillus subilis str. 168CYP152A2 (P450c1A) from bacterium Costridium acetobupticum ATCG 824WP_198687844 (E9NSU2)Peorgalicoccus sp. 8456WP_198687844 (E9NSU2)Peorgalicoccus sp. 8456FM872457 (B9W4V6)AacUPO from mushroom Agroybe acgrita (wild-type, PaDa-1 mutant, SoLo mutant)NP_103557336.1 (G2RGE5)DA9E from fungus Thielavia paretriaXP_003657336.1 (G2RGE5)DA9E from fungus Thielavia fungoNP_625478.1 (Q9RJY2)OA9E from fungus Thielavia fungoNP_625478.1 (Q9RJY2)DA9E from fungus Thielavia fungoNP_625478.1 (Q9RJY2)OA9E from fungus Thielavia fungoNP_625478.1 (Q9RJY2)DA0F from fungus Thielavia fundoNP_625478.1 (Q9RJY2)DA0F from fungus Thielavia fungoNP_625478.1 (Q9RJY2)DA0F from fungus Thielavia fundoNP_625478.1 (Q9RJY2)DA0F from fungus Thielavia fundoNP_625478.1 (Q9RJY2)DA0F from fungus Thielavia fundoNP_625478.1 (Q9RJY2)DA	P450	CYP102A1 (heme domain) from bacterium <i>Bacillus</i> moottorium	1	<i>E. coli</i> XL1-blue cells, Luria–Bertani (LB) medium, ampicillin (100 µg mL ⁻¹), 0.4 mM IPTG (isopropyl β -D-1-thiogalactopyranoside), and 0.5 mM, δ -ALA (δ -aminolenulinio aciel) 30 °C for 6 h	12
CYP152A1 (P450 _{has}) from bacterium <i>Bacilius subilis</i> subsp. Subtilis str. 168 corp152A2 (P430c) from bacterium <i>Clostridium</i> <i>Clostridium</i> <i>Clostridium</i> 		Human CYP2E1	$NM_{-}000773$ (P05181.1)	E. coli DH5 α cells, LB medium with 20 g glucose per L, ampicillin (100 µg mL ⁻¹), 1 mM IPTG, and 0.5 mM δ -ALA, 30 °C for 48 h	13 and 14
CYP132A2 (P450 _{c1A}) from CAC3330 (Q97DZ0) bacterium <i>Clostridium cetobuylicum</i> ATCC 834 WP_198687844 (E9NSU2) bacterium <i>Clostridium MP_198687844</i> (E9NSU2) <i>cetobuylicum</i> ATCC 834 WP_198687844 (E9NSU2) <i>cetobuylicum</i> ATCC 834 WP_198687844 (E9NSU2) <i>forgalifoccus</i> sp. 8456 FM872457 (B9W4V6) <i>Agropye acgerita</i> (wild-type, MP_198687844 (E9NSU2) <i>Agropye acgerita</i> (wild-type, RM872457 (B9W4V6) <i>Agropye acgerita</i> (wild-type, RA9411 (P00331) <i>Ipmo from fungus Thielavia</i> CA_900343105.1 <i>terrestris</i> (copper saturated TAA9 <i>Abbl from fungus Thielavia</i> GCA_900343		CYP152A1 (P450 _{bsb}) from bacterium <i>Bacillus subtilis</i> subsp. Subtilis str. 168	BSU02100 (O31440.1)	$E.~coli^{'}$ BL21 (DE3) cells, LB medium, kanamycin (30 $\mu g~m L^{-1}$), 1 mM IPTG, 30 °C for 4 h	15
CYP152.1.1 from bacterium WP_19687844 (E9NSU2) Jeotgalicoccus sp. 8456 FM872457 (B9W4V6) Jeotgalicoccus sp. 8456 FM872457 (B9W4V6) Agroopbe acgeria (wild-type, PaDa-I mutant, SoLo mutant) FM872457 (B9W4V6) Option Caldariomyces X04486 (P04963) <i>funago</i> XP_108657336.1 (G2RG5) D AA9E from fungus Thielavia XP_003657336.1 (G2RG5) LPMO from Streptomyces NP_625478.1 (Q9RJY2) coelicolor (ScA10C) IPMO from Streptomyces LPMO from fungus Thielavia GCA_900343105.1 drogenase Alcohol dehydrogenase (ADH) AA34411 (P00331) from Saccharomyces cerevisiae GCA_900343105.1 from Saccharomyces cerevisiae GCA_90033105.1 from Saccharomyces cerevisiae GCA_9003310 from Saccharomyces cerevisiae GCA_9003310 from Saccharomyces cerevisiae GCA_9003310 <		CYP152A2 (P450 _{CLA}) from bacterium <i>Clostridium</i> acetoburvlicum ATCC 824	CAC3330 (Q97DZ0)	$E\ coli$ BL21 (DE3) cells, LB medium, kanamycin (30 µg mL^{-1}), 1 mM IPTG, 30 °C for 4 h	15
AccordFM872457 (B9W4V6)Agrophe acgerita (wild-type, PaDa-1 mutant, SoLo mutant)FM872457 (B9W4V6)Agrophe acgerita (wild-type, PaDa-1 mutant, SoLo mutant)FM872457 (B9W4V6)GfuUPO from CaldariomycesX04486 (P04963)fumagoAA9E from fungus ThielaviaXP_003657336.1 (G2RGE5)furnestris (TtAA9E)NP_625478.1 (Q9RJY2)coelicolor (ScAA10C)RCA_900343105.1terrestris (copper saturated TtAA9)AA34411 (P00331)drogenaseAlcohol dehydrogenase (ADH)AA334411 (P00331)from Saccharomyces cerevisiae (ScADH)BU64864 (P00327)ADH from horse liver (HLADH)M64864 (P00327)Formate dehydrogenase from Candida boidinii (CbFDH)M454879		CYP152L1 from bacterium <i>leotealicoccus</i> sp. 8456	WP_198687844 (E9NSU2)	<i>E coli</i> C43 cells, TB medium, kanamycin (50 μg mL ⁻¹). 0.1 mM IPTG, 1 mM δ-ALA. 30 °C for 20 h	16
PaDa-1 mutant, SoLo mutant)PaDa-1 mutant, SoLo mutant)CfuUPO from CaldariomycesJumagoMA9E from fungus ThielaviaA9E from fungus ThielaviaAPE from fungus ThielaviaLPMO from StreptomycesLPMO from Streptomycescoelicolor (ScAA10C)LPMO from fungus ThielaviaCoelicolor (ScAA10C)LPMO from fungus ThielaviaCoelicolor (ScAA10C)LPMO from fungus ThielaviaGCA_900343105.1terrestris (copper saturated TtAA9)TRAA9)IogenaseAlcohol dehydrogenase (ADH)AA34411 (P00331)from Saccharomyces cerevisiae (ScADH)ADH from Thermus sp. ATN1FU681191 (B2ZRE3)form sate dehydrogenase from Candida boidmii (CbFDH)KM454879 	UPO	AgeUPO from mushroom Agrocybe aegerita (wild-type,	FM872457 (B9W4V6)	Saccharomyces cerevisiae, minimal medium, chloramphenicol (25 $\mu g~m L^{-1}$), 25 °C for 72 h	17 and 18
CfuUPO from CaldariomycesX04486 (P04963)fumagoAA9E from fungus ThielaviaXP_003657336.1 (G2RGE5)AA9E from fungus ThielaviaXP_003657336.1 (G2RGE5)AA9E from fungus ThielaviaNP_625478.1 (Q9RJY2)LPMO from StreptomycesNP_625478.1 (Q9RJY2)coelicolor (ScAA10C)IPMO from fungus ThielaviaLPMO from fungus ThielaviaGCA_900343105.1terrestris (copper saturated TtAA9)AA34411 (P00331)from Saccharomyces cerevisiae (ScADH)AA34411 (P00331)from Saccharomyces cerevisiae (ScADH)M64864 (P00327)ADH from horse liver (HLADH) 		PaDa-I mutant, SoLo mutant)		<i>Pichia pastoris</i> , basal salts medium, <i>P</i> TM1 trace salts (4.35 mL L^{-1}) and antifoam 204 (0.25 mL L^{-1}), methanol (0.5%, v/v), 25 °C	
AA9E from fungus ThielaviaXP_003657336.1 (G2RGE5)terrestris (TtAA9E)NP_625478.1 (Q9RJY2)terrestris (TtAA9E)NP_625478.1 (Q9RJY2)LPMO from StreptomycesNP_625478.1 (Q9RJY2)coelicolor (ScAA10C)LPMO from fungus ThielawiaLPMO from fungus ThielawiaGCA_900343105.1terrestris (copper saturated TtA9)AA34411 (P00331)from Saccharomyces cerevisiae (ScADH)AA34411 (P00331)from Saccharomyces cerevisiae (ScADH)MA434411 (P00331)ADH from Thermus sp. ATN1EU681191 (B2ZRE3)ADH from horse liver (HLADH) Formate dehydrogenase from Candida boidinii (CbFDH)M64864 (P00327)		CfuUPO from Caldariomyces fumago	X04486 (P04963)	<i>Cfu</i> UPO is obtained by fermentation of <i>C. fumago</i> on synthetic medium, filtration of the mycelium, and subsequent chromatography. The strain was obtained from the International Mycological Institute, Surrey, UK	19
LPMO from StreptomycesNP_655478.1 (Q9RJY2)coelicolor (ScAA10C)LPMO from fungus ThielaviaGCA_900343105.1LPMO from fungus ThielaviaGCA_900343105.1terrestris (copper saturated TtAA9)AA34411 (P00331)Alcohol dehydrogenase (ADH)AAA34411 (P00331)from Saccharomyces cerevisiae (ScADH)BU681191 (B2ZRE3)ADH from Thermus sp. ATN1EU681191 (B2ZRE3)ADH from horse liver (HLADH)M64864 (P00327)Formate dehydrogenase from Candida boidinii (CbFDH)KM454879	LPMO	AA9E from fungus <i>Thielavia</i> terrestris (TtAA9E)	XP_003657336.1 (G2RGE5)	The preculture and production culture in <i>P. pastoris</i> SMD1168H were both done in <i>YPD</i> medium. The strain was grown for production for 4 days. Purified <i>T. terrestris</i> LPMO (<i>TtLPMO9E</i> , previously <i>TtGH61E</i>) was donated from Novozwes AS (<i>Denmark</i>).	20
LPMO from fungusThielaviaGCA_900343105.1terrestris (copper saturated TtAA9)TtAA9)GCA_900343105.1Alcohol dehydrogenase (ADH)AAA34411 (P00331)from Saccharomyces cerevisiae (ScADH)GCA_91191 (B22RE3)ADH from Thermus sp. ATN1EU681191 (B22RE3)ADH from horse liver (HLADH)M64864 (P00327)Formate dehydrogenase from Candida boidinii (CbFDH)KM454879		LPMO from <i>Streptomyces</i> coelicolor (ScAA10C)	NP_625478.1 (Q9RJY2)	Fresh colonies were incounted in LB-ampicillin (50 μ g mL ⁻¹) media and grown at 37 °C (Gel22) for 20 h	21 and 22
Alcohol dehydrogenase (ADH)AAA34411 (P00331)from Saccharomyces cerevisiae(ScADH)(ScADH)EU681191 (B2ZRE3)ADH from Thermus sp. ATN1EU681191 (B2ZRE3)(TADH)M64864 (P00327)ADH from horse liver (HLADH)M64864 (P00327)Formate dehydrogenase fromKM454879Candida boidinii (CbFDH)KM454879		LPMO from fungus <i>Thielavia</i> <i>terrestris</i> (copper saturated <i>Tt</i> AA9)	GCA_900343105.1	Thielavia terrestris LPMO (TtAA9) was provided as a broth from Novozymes A/S (Denmark). The TtAA9 was saturated with Cu(i) chloride under anaerobic conditions and on ice for 2 h, filtered, purified by chromatorraphy, and concentrated by ultra-filtration	23
m <i>Thermus</i> sp. ATN1 EU681191 (B2ZRE3) m horse liver (HLADH) M64864 (P00327) dehydrogenase from KM454879 <i>boidinii</i> (CbFDH)	Dehydrogenase	Alcohol dehydrogenase (ADH) from <i>Saccharomyces cerevisiae</i> (ScADH)	AAA34411 (P00331)	<i>ScADH</i> gene was transformed into <i>S. cerevisiae</i> 302–21#2 strain that does not produce active alcohol dehydrogenase, and cultivated in YEPD medium and antimvcin A (1 $\mu m mL^{-1}$) at 30 °C for 24 h	24 and 25
M64864 (P00327) KM454879		(TADH from <i>Thermus</i> sp. ATN1 (TADH)	EU681191 (B2ZRE3)	TADH was produced by transforming pASZ2 (PET) into <i>E. coli</i> BL21 (DE3) pLySS cells and cultivation of the recombinant <i>E. coli</i> using the autoinduction system	26
		ADH from horse liver (HLADH) Formate dehydrogenase from <i>Candida boidinii</i> (CbFDH)	M64864 (P00327) KM454879	Commercially available CbFDH plasmids were transformed into <i>E. coli</i> BL21 (DE3) pLysS cells and grown in LB medium with ampicillin (100 μ g mL ⁻¹) and 1 mM IPTG at 37 °C	26 27 and 28

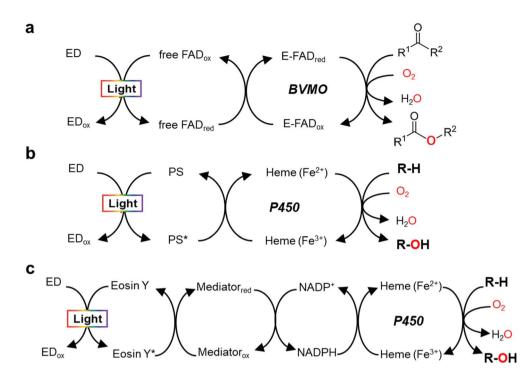


Fig. 2 Light-driven monooxygenase catalysis *via* direct activation by electron transfer to active site (heme or flavin) (a and b) and indirect route by photoregeneration of NAD(P)H (c). (a) Light-driven direct electron transfer to enzyme-bound FAD (E-FAD) of the Baeyer–Villiger mono-oxygenase (BVMO, *e.g.* PAMO) using a flavin as a photosensitizer (PS) and EDTA as an sacrificial electron donor (ED).²⁹ (b) Direct electron transfer from activated PS [*e.g.*, eosin Y,³¹ flavin,^{12,13} Ru(II)³⁶], which are made by light and ED (TEOA, EDTA, or diethyldithiocarbamate), to P450 heme. (c) The catalytic turnover of P450 is indirectly accomplished by photochemical reduction of NADPH through a cascaded electron delivery in a photocatalytic system having an ED (TEOA), a PS (eosin Y), and an electron mediator M [Cp*Rh(bpy)H₂O]^{2+,40}

a photocatalyst to reduce P450 heme iron (Fig. 2b). Ru(II) has been widely applied for single electron transfer in metalcontaining hydrogenases and photochemical enzymatic reduction of small molecules by nitrogenases.37 The Cheruzel group attached a Ru(II)-diimine complex specifically close to the enzyme active site of CYP102A1 via an engineered mutant (L407C) for direct electron transfer from photoreduced Ru(I) to the P450 heme iron using diethyldithiocarbamate as a sacrificial electron donor.36 The Ru(II)-CYP102A1 platform catalyzed light-driven reaction of lauric acid hydroxylation, with an initial reaction rate of 125 min^{-1} and a total turnover number (TTN = maximum product concentration/enzyme concentration) of 935.36 Furthermore, the Ru(II)-CYP102A1 platform was also applied to the hydroxylation reaction of 10-undecenoic acid³⁸ to produce (R)-9-hydroxy-10-undecenoic acid and the O-dealkylation reaction of nitrophenolate-based substrates as chromogenic substrates.39

In 2019, Le *et al.* reported that a flavin/EDTA/light platform can support the catalytic activities of CYP102A1 heme domain through direct delivery of photoexcited electrons to the P450 heme iron and indirect H_2O_2 generation for peroxygenase activity as well¹² (Fig. 2b). Without the requirement of reductase and NADPH, the presence of only light and EDTA mediates the direct electron transfer from reduced flavins to the heme iron. The reduced heme can drive the hydroxylation of the organic substrate (R–H). On the other hand, the photoreduced flavin reacts quickly with O_2 to produce H_2O_2 . Hence, again, a contribution of the H_2O_2 shunt pathway cannot be fully excluded.¹² For human CYP2E1, in contrast, the flavin/EDTA/light system works only *via* direct transfer of light-induced electrons to the P450 heme iron, not by the H_2O_2 -mediated peroxide shunt.¹³

Indirect transfer of photoinduced electrons to monooxygenases has been extensively accomplished through photochemical reduction of biological cofactors [e.g., NAD(P)H, FADH₂]. Here, the redox mediators function as a freely diffusing, remote communicator between a photocatalyst and a monooxygenase. For example, photochemical regeneration of NADPH for CYP102A1 (Y51F/F87A variant)-catalyzed O-dealkylation reaction was performed using EY as a photosensitizing dye, TEOA as an electron donor, and [Cp*Rh(bpy)H₂O] as a selective NAD(P)H regeneration catalyst (Fig. 2c). The P450 sustainably maintained its catalytic turnover with continuous photoregeneration of NADPH. Thus, visible light-driven recycling of NADPH provides a new route for P450 monooxygenase catalysis.40 Later on, natural sunlight-driven platform for P450 catalysis has been demonstrated at preparative scale using P450s immobilized on poly(3hydroxybutyrate) [P(3-HB)] scaffold. Through photochemical regeneration of NADPH using EY as a photocatalyst, the P450-P(3HB) complex could successfully catalyze O-dealkylation reactions. In addition, a P450-catalyzed reaction was accomplished under natural sunlight for four consecutive days using a solartracking module.41 This study hints at practical applicability of natural sunlight for driving photobiocatalysis, which can be replaced by LED light during night.

Interestingly, ferredoxin (Fd)-mediated transfer of electrons in photosystem I (PSI) has been adopted for indirect transfer of photoinduced electrons to P450 heme. Jensen and coworkers demonstrated that the heme domain of plant CYP79A1, which is involved in tyrosine oxidation, can be photochemically reduced by the PSI using Fd as a mediator.^{42,43} Upon photoactivation, the Fd-mediated electron transfer chain readily provides the required electrons to support the P450 catalysis.

In addition to abiotic photochemical and photoelectrochemical approaches, cyanobacteria have been reported to activate oxidative redox enzymes (*e.g.*, BVMO,³⁴ P450,³³ alkane monooxygenase⁴⁴) by producing O₂ gas and reducing equivalents.

3. Photobiocatalytic oxyfunctionalization *via* H₂O₂-dependent pathway

 $\rm H_2O_2$ -Dependent pathways are recently gaining increasing attention—instead of the above-mentioned reductive regeneration of the enzymes' active sites—for photobiocatalytic oxy-functionalization. The catalytically active species in monooxygenases contain or are derived from an intermediate comprising activated oxygen in the oxidation state of $\rm H_2O_2$.⁴ The latter has been derived from molecular oxygen *via* reductive

activation within the enzyme active site *via* more or less complex and vulnerable mechanisms.⁶ Therefore, simplifying the catalytic mechanism and reaction schemes using already reduced oxygen in form of H_2O_2 is a very appealing approach. As H_2O_2 , however, also acts as a strong oxidative inactivator of many enzymes, its stoichiometric application is not feasible and photochemical *in situ* generation methods providing H_2O_2 in appropriate rates represent a promising approach.

Most prominent examples for H_2O_2 -utilizing enzymes are the heme-dependent P450 monooxygenases and peroxygenases as well as the Cu-dependent lytic polysaccharide monooxygenases (LPMOs). Further enzymes such as hydrolases following the perhydrolase pathway,^{45,46} vanadium-dependent haloperoxidases⁴⁷ or artificial peroxyzymes⁴⁸ are worth mentioning but have not been used together with a photochemical H_2O_2 generation system yet.

3.1 Unspecific peroxygenases (UPOs)

UPOs catalyze a wide range of selective oxyfunctionalization reactions on inert C–H and C==C bonds. The catalytic cycle of peroxygenases represents the well-known H_2O_2 -shunt shortcut on P450 monooxygenases (Fig. 3). P450 monooxygenases are mostly rather inefficient in utilizing the H_2O_2 -shunt pathway, which has been assigned to the lack of a catalytic base

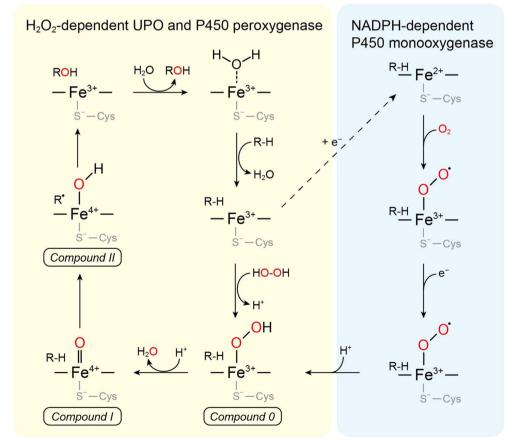


Fig. 3 Comparison between the mechanism of P450s and UPOs for C-hydroxylation. The left cycle describes the H_2O_2 -supported "peroxide shunt" shared by UPOs and P450 peroxygenases and the right one is the "classical" NADPH-dependent P450 monooxygenase route. Both cycles share compounds 0 (ferric peroxo-complex), I (oxo-ferryl cation radical complex), and II (ferryl hydroxide complex).

Perspective

(glutamate) facilitating the conversion of Cpd-0 to Cpd-I. This catalytically relevant amino acid is present in peroxygenases but not in P450 monooxygenases⁴⁹ (Fig. 4).

The first reported peroxygenase stems from *Leptoxyphium fumago* (previously *Caldariomyces fumago*)⁵⁰ and found some popularity as oxyfunctionalization catalyst.⁵¹ However, poor activity and difficulties in the recombinant expression of this enzyme have limited its broad applicability in organic synthesis. This situation changed in 2004 with Hofrichter and coworkers reporting the UPO (initially termed as haloperoxidase) from *Agrocybe aegerita (Aae*UPO).⁵² Later, Alcalde and coworkers succeeded in its recombinant expression and engineering^{17,18} and more recently it's large-scale production.⁵³ These pioneering contributions have paved the way to the engineering and identification of new, promising peroxygenases.^{5,52} Even though only a handful of UPOs have so far been functionally expressed

and evaluated with respect to their catalytic properties, the synthetic potential is enormous.^{4,48,55}

3.2 Cytochrome P450 peroxygenases

P450 monooxygenases and peroxygenases are structurally and mechanistically related. Hence, in principle, the H_2O_2 -shunt pathway represents a doable alternative pathway to simplify the catalytic mechanism of P450 monooxygenases (Fig. 3). Unfortunately, however, most P450 monooxygenases exhibit very low activity with H_2O_2 (whereas the oxidative degradation of the catalytic heme moiety with H_2O_2 remains a challenge) resulting in comparably few catalytic cycles of P450 monooxygenases with H_2O_2 . Protein engineering principally offers a solution to this issue⁵⁶ which however yet remains to be fully exploited. The socalled P450 peroxygenases (bacterial CYP152 family) represent

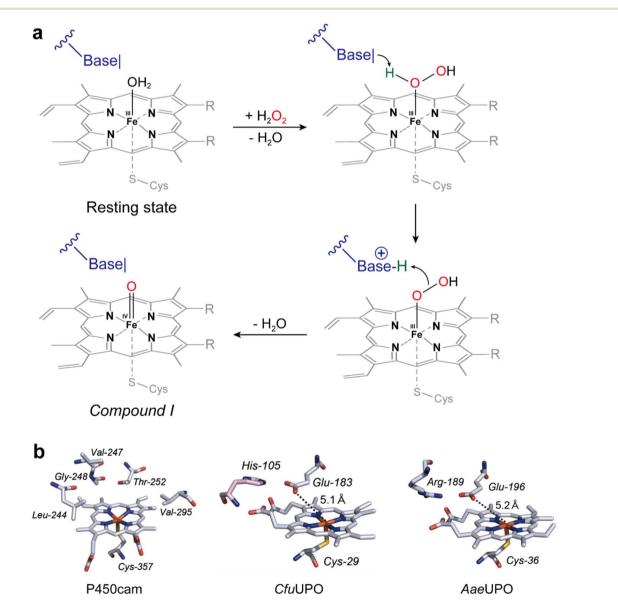


Fig. 4 (a) Acid-base-catalysis as key difference between P450 monooxygenases and peroxygenases. (b) Active-site architectures of P450cam (POB code 2CPP), *Cfu*CPO (PDB code 1CPO) and *Aae*UPO (PDB code 2YP1). Adapted from ref. 49 with permission from Springer.

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Table 2 LPMO catalysis driven by *in situ* photochemical H₂O₂ generation

Chemical Science

LPMO	Photosensitizer	Reductant of O ₂	Light source	Substrate	Product	Ref.
AA9E from Thielavia terrestris (TtAA9E)	Chlorophyllin (Chl) with ascorbic acid (AscA)	Photo-excited Chl	$150-200 \mu mol photons s^{-1} m ^{-2}$	Phosphoric acid-swollen cellulose (PASC), Avicel microcrystalline cellulose, lignin, xyloglucan	Diverse oligosacharides, gluconic acid, p-glucose, p- cellobiose, cello- oligosaccharides, xvlogluco-oligosaccharides	20
LPMO from Streptomyces coelicolor (ScAA10C)	Chl with AscA, (or V-TiO ₂)	Photo-excited Chl (or H ₂ O)	150 W, xenon lamp visible light $(I = 25\%$ Imax. 42 W cm ⁻²)	Crystalline cellulose (Avicel)	Áldonic acid, GlcGlc1A, (Glc) ₂ Glc1A	21 and 22
LPMO from <i>Thielavia</i> <i>terrestris</i> (copper saturated <i>Tt</i> AA9)	Chl with AscA	Photo-excited Chl	White LEDs (200 μ mol photons m ⁻² s ⁻¹)	Microcrystalline cellulose (Avicel) Phosphoric acid-swollen cellulose (PASC) Cellulose nanofibrils (CNF)	Gluconic acid (GlcA) GlcA GlcA	23

a subclass of the P450 monooxygenases, which in addition to the natural cycle also accept H_2O_2 .⁵⁷ Interestingly, these enzymes are all involved in fatty acid metabolism.

The first reported P450 peroxygenases—CYP152B1 and CYP152A2—hydroxylate fatty acids exclusively in α -position.^{15,58,59} In contrast, CYP152A1 shows a preference for the β -position of fatty acids.^{15,59,60} On the other hand, CYP152L isozymes mainly perform oxidative fatty acid decarboxylation, producing terminal alkenes, along with α - and β -hydroxylation reactions.^{61,62} CYP152N1 catalyzes oxidative decarboxylation reaction of fatty acids to produce one-carbon shorter fatty acids.⁶³ Thus, CYP152 peroxygenase reactions for alkene production are useful for the production of biofuels and fine chemicals.^{62,64}

Finally, also the decoy-molecule—dual function small molecule—approach is worth mentioning.⁶⁵ Using modified imidazoles as decoy molecules, Cong and coworkers successfully introduced H_2O_2 -reactivity with some 'conventional' P450 monooxygenases. The imidazole is believed to partially replace the missing catalytic bases present in peroxygenases.

3.3 Lytic polysaccharide monooxygenases

LPMOs are key enzymes that catalyze degradation and decomposition of recalcitrant polysaccharides (*e.g.*, cellulose, lignin, and chitin) by fungi and bacteria.^{66–68} Their active sites have a single copper ion, which is coordinated by an N-terminal histidine side chain and a further histidine residue to arrange the Cu-histidine brace.⁶⁹ LPMOs are widespread in nature and classified as seven sequence-distinct families (*i.e.*, 9–11 and 13–16) of the auxiliary activities according to the Carbohydrate Active enZymes (CAZy) database (https://www.cazy.org/Auxiliary-Activities.html).⁷⁰

Mechanistically, LPMOs catalyze the reductive activation of O_2 for oxidative cleavage of the glycosidic bonds of polysaccharides *via* hydroxylation at either the C1 or the C4 position, followed by the cleavage of the scissile glycosidic bond and the formation of aldonic acids or 4-keto sugars at the oxidized chain ends.⁶⁸ In addition to using O_2 as an oxidant, LPMOs can utilize H_2O_2 as a cosubstrate to perform peroxygenation reactions on polysaccharides, catalyzing similar hydroxylation reactions with O_2 , which then lead to the spontaneous bond cleavage of the polysaccharides.^{68,71} Details of the catalytic mechanism remain to be fully elucidated.⁶⁸

Nowadays, LPMOs are receiving a lot of attention because of their potential as an efficient decomposer of biomass.^{68,72,73} According to Bissaro *et al.*,⁷¹ LPMOs prefer H₂O₂ over O₂ (*via* reductive activation) for hydroxylation of microcrystalline α -cellulose substrate (*i.e.*, Avicel). The biocatalytic performance and stability could be controlled by regulating the supply of H₂O₂ in the absence of O₂, suggesting that LPMOs possess more efficient peroxygenase activity than the monooxygenase activity (Table 2).

4. Photocatalytic platforms for in situ H_2O_2 generation

 H_2O_2 represents a 'double-edged sword' enabling simplified reaction schemes on the one hand but also, on the other hand, inactivates enzymes. To balance both reactivities, *in situ* generation of H_2O_2 has proven suitable. Next to enzymatic,^{74,75} chemical^{76,77} and electrochemical⁷⁸⁻⁸⁰ H_2O_2 generation schemes, also photo(electro)catalytic systems have been developed. The light-driven methods require a proper photocatalyst that, when excited by light, oxidises an electron donor and transfers the reducing equivalents to oxygen, producing H_2O_2 . The steady-state H_2O_2 concentration is intrinsically in an equilibrium between photocatalytic generation and degradation.⁸¹ In this section, we discuss organic or inorganic photocatalytic platforms for light-driven H_2O_2 generation and biocatalytic oxyfunctionalization.

4.1 Flavins

Flavin molecules, such as flavin adenine mononucleotide (FMN), flavin adenine dinucleotide (FAD) and riboflavin, can produce H₂O₂ photochemically at the expense of electron donors and molecular oxygen. For example, FMN extracts an electron from EDTA after being activated by light, then the reduced FMN generates H2O2 via O2 reduction.82 The flavinbased in situ H₂O₂ generation strategy was first designed for driving CfuCPO-83 and AaeUPO-catalyzed84 oxyfunctionalization reactions (Table 3). Even though in these early works, the benefits of photochemical in situ H2O2 generation could be demonstrated, EDTA as sacrificial electron donor exhibits several disadvantages impeding it widespread use. Particularly, the degradation products, eventually formaldehyde and ethylene diamine, are not desirable from a practical and an environmental point of view. Other sacrificial electron donors such as ascorbic acid, formate, hydrazine or phosphite can substitute for EDTA but still need to be further explored.

Nevertheless, mostly due to its simplicity, the flavin/EDTA/ light system has also been applied for some P450 peroxygenase reactions including CYP152A1, CYP152A2, and CYP152L1 (Fig. 5). The P450 fatty acid peroxygenases hydroxylate (or decarboxylate) saturated fatty acids $(C_{12}-C_{20})^{101-103}$ (Fig. 5a and b) The catalysis of CYP152L1 using this photochemical method was also applied to the conversion of ω hydroxylated fatty acids (C_{12} , C_{15} , C_{16}), obtained from palm kernel oil, into the corresponding alkenols¹⁰⁴ (Fig. 5c). In the flavin/EDTA/light system, a peroxygenase activity of bacterial CYP102A1 heme domain were confirmed to catalyse the hydroxylation of 4-nitrophenol and lauric acid¹² (Fig. 5d and e).

Next to waste generation, another drawback of flavins as photocatalysts is the rather narrow wavelength spectrum of flavins, only partially covering the spectrum of sunlight. In view of future sunlight-driven processes Willot *et al.* explored wavelength-complementary photosensitizers such as FMN (yellow), phenosafranine (green), and methylene blue (red) to utilize the different wavelengths of visible light more efficiently.⁹⁴ In these experiments, co-catalysis by formate dehydrogenase and the nicotinamide cofactors enabled replacement of EDTA by formate.

Choi *et al.* demonstrated photoelectrocatalytic formation of H_2O_2 driven by flavin-hybridized single-walled carbon nanotube (SWNT) photoelectrode¹⁰⁰ (Table 3). Lumichrome, a flavin derivative, was used as a photosensitizer. When flavin-

hybridized SWNTs were exposed to visible light, flavins allowed a significant anodic shift of the O₂ reduction potential. Because of flavin's high reactivity toward O₂, photoactivated flavins boost electron delivery from the SWNT cathode to O₂ for more efficient H₂O₂ production. The flavin-functionalized SWNT photoelectrode system enabled *Aae*UPO to hydroxylate ethylbenzene to (*R*)-1-phenylethanol at a TTN and STY of 123 900 (95% ee) and 1.0 (g L⁻¹ d⁻¹), respectively. However, when 2-phenoxypropionic acid was used as a substrate, the TTN and STY was only 5900 and 0.13 (g L⁻¹ d⁻¹), respectively.

4.2 Titanium dioxide

Zhang et al. reported the use of gold-loaded rutile titanium dioxide (Au-TiO₂) and methanol to drive AaeUPO-catalyzed oxyfunctionalization of various alkane substrates90 (Table 3 and Fig. 6a). Through the methanol-driven reductive O2 activation, Au-TiO₂ provided H₂O₂ in concentrations with which the enzyme remained active and stable. Later, the same authors, demonstrated that also water may serve as sacrificial electron donor.91 Compared to the use of methanol, reaction rates and robustness of the overall reaction were significantly impaired. This may be attributed to higher rate of the photocatalytic oxidation of alcohols compared to water. Furthermore, alcohols act as radical scavengers thereby prolonging the enzyme lifetime.92 The radical inactivation of the biocatalyst by surfaceborne by hydroxyl radicals (HO') can also be alleviated by spatial separation of the photo- and biocatalyst. Under optimized reaction conditions maximum TTN for the biocatalyst of 220 000 and space time yields (STY) of 2.6 g L^{-1} d⁻¹ together with photonic efficiencies of 13.6% were achieved.

4.3 Carbon nitrides

Carbon nitrides are known to catalyse light-induced water oxidation yielding O_2 and H_2 , they can also be used for the oxidation of alcohols to the corresponding carbonyl groups.¹⁰⁵ Instead of transferring electrons obtained from water- or alcohol oxidation to protons also O_2 can serve as electron acceptor eventually yielding H_2O_2 . Graphitic carbon nitride (g- C_3N_4) for example is a good alternative to the TiO₂-based photocatalysts⁹³ (Table 3 and Fig. 6a). Again, however, surfacebourne hydroxyl radicals impair the stability of the biocatalyst, which can be overcome by spatial separation of the bioand photo catalyst. Using different wavelengths can tune the redox potential of a carbon nitride photocatalyst (CN-OA-m) and thereby influence the selectivity of the photoenzymatic cascade.⁹⁸

4.4 Lignin

Lignin, a major component that accounts for 20–30% of plants, is a heterogeneous polymer mostly consisting of aromatic subunits. This year Kim *et al.* identified lignin as a multifunctional photocatalyst for such as *in situ* H_2O_2 formation through H_2O oxidation and O_2 reduction.⁹⁹ Particularly appealing is the intrinsic hydroxyl radial-scavenging activity enabling more robust photoenzymatic cascade reactions without the need for spatial separation of both catalysts (Fig. 6a). This very

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Table 3 Comparison of *in situ* H_2O_2 generation methods for ethylbenzene hydroxylation to (R)-1-phenylethanol by AaeUPO^a

Method	AaeUPO (concentration)	Reagents, light sources, and remarks	TTN (ee%)	STY (g L ⁻¹ day ⁻¹)	Ref.
Stoichiometric	WT (40 nM)	$ m H_2O_2$ (1 mM), 1 mM ethylbenzene	6000 (97)	1.8	84
addition	WT (600 nM)	t-BuooH (60 mM), 1 mM ethylbenzene	90 000 (95- 98)	40	85
Enzymatic	WT (100 nM)	AOx (60 nM), FDM (295 nM), FDH (1 μ M), NAD ⁺ (1.6 mM), 3HB6H (11 μ M), MeOH (5 mM), 50 mM ethylbenzene	468 500 (>99)	17.6	86
	PaDa-I (25 nM)	AOX (60 nM), MeOH (5 mM), 15 mM ethylbenzene FDH (250 nM), NAD ⁺ (0.5 mM), YqjM (2.67 µM), MeOH (200 mM),	291 500 (>99) 390 000 (>96)	11.0 5.2	86 87
	PaDa-I or SoLo	sodium formate (75 mM), 10 mM ethylbenzene SO (200 nM), calcium sulfite (100 mM), O ₂ , 50 mM ethylbenzene	30 800 (98)	19	88
	PaDa-I (4 µM)	Soluble hydroxylase (95 nM), H_2 , FMN (200 μ M), two-phase system: 0.75 mL ethylbenzene, 0.75 mL KPi buffer (50 mM, pH 7.5), 7 mL excess of headspace filled with gas (40% O_2 and 60% H_2)	11 250 (99)	3.2	88
Photochemical	WT (40 nM)	F ⁻¹⁵ FMN (50 μM), EDTA (1 mM), O ₂ , 1 mM ethylbenzene. Visible light (Philins 7748XHP 200W. white light bulb)	11 470 (>97)	3.5	84
	PaDa-I (150 nM)	V_{1} V_{2} V_{2} V_{1} V_{2} V_{2	79 000 (>98)	0.48	90
	PaDa-I (150 nM)	Rutile-Au-TiO ₂ (30 mg mL ⁻¹), H ₂ O, O ₂ , 10 mM ethylbenzene, visible light ($\lambda > 400$ nm) (Philips 7748XHP 150 W, white light bulb)	37 000 (98)	0.17	91
		Carbon nanodot (CND) (1 mg mL ^{-1}), FMN (0.1 mM), H ₂ O, O ₂ , 10 mM ethylbenzene, visible light (Philips 7748XHP 150 W, white light hulb)	100 000 (98)	2.4	
	WT (200 nM)	TiO ₂ (2 mg mL ⁻¹), MeOH (5%, v/v), O ₂ , 10 mM ethylbenzene, 365 nm LED (either M365LP1, Thorlabs or LEDMOD365.1050.V2, Omicron	220 000	2.6	92
	PaDa-I (100 nM)	Graphitic carbon nitride (g- C_3N_4) (5 mg mL ⁻¹), MeOH or formic acid (250 mM), 50 mM ethylbenzene, white light bulb (Osram 200 W light bulb)	60 000 (98)	0.59	93
	PaDa-I (100 nM)	Wavelength-complementary photosensitizers (5 µM FMN, 10 µM methylene blue, 10 µM phenosafranin), MeOH (200 mM), NAD ⁺ (0.4 mM), 4.8 µM FDH, O ₂ , 10 mM ethylbenzene, Commercial LEDs (24 W) or LIGHTINGCURE LC8 L9566 (Hamamatsu)	100 000 (>95)	7.3	94
	PaDa-I (100 nM)	Nitrogen-doped carbon nanodots (N-CNDs) (5 mg mL ⁻¹), O_2 , 50 mM ethylbenzene. white light bulb (Philips 7748XHP 150 W)	000 06	0.56	95
	PaDa-I (150 nM)	Au-TiO ₂ (5 mg mL ⁻¹), MeOH (250 mM), O ₂ , 60 mM ethylbenzene, LED (405 mm), SMBB405V-1100-02Z, 125 mA	372 000	7.8	96
	PaDa-I (100 nM)	Anthraquinone-2-sulfonate (0.5 mM), MeOH (40%), O ₂ , 50 mM ethylbenzene, visible light ($\lambda > 400$ mm) (white light bulb, Philips 7748XHP 150W)	\sim 150 000 (34)	0.46	97
	PaDa-I (25 nM)	Carbon nitride photocatalyst (CN-OA-m) (2 mg mL ^{-1}), MeOH (250 mM), O ₂ , 10 mM ethylbenzene, green light (528 nm) (Kessil PR160-525 LED)	284 000 (99)	0.87	86

Table 3 (Contd.)					
Method	AaeUPO (concentration)	Reagents, light sources, and remarks	TTN (ee%)	${ m STY}~({ m g} { m L}^{-1}~{ m day}^{-1})$	Ref.
	PaDa-I (50 nM)	Kraft lignin (or lignosulfate) (8 mg mL ⁻¹), O ₂ , 10 mM ethylbenzene, visible light ($\lambda > 400$ nm, photon flux: 1.74 µE cm ⁻² s ⁻¹), a xenon arc lamp (with an infrared water filter and 400 nm lonebass filter)	72 600 (81 000) (>99)	0.0819 (0.0914)	66
Photoelectro- chemical	PaDa-I (100 nM)	Flavin-SWNT (single walled carbon nanotubes) electrode, U, O ₂ , 100 mM ethylbenzene, U– power source and electrodes: WMPG100 potentiostat, 450 W Xe lamp with a 420 nm cut-off filter	123 900	1.0	100
Electrochemical	PaDa-I (50 nM)	U, O ₂ , acetone (3.3%, v/v), 139 mM ethylbenzene, U: -10 (TTN) or -30 (STY) mA cm ⁻² by multichannel potentiostat/galvanostat G300 TM (projected area $A_{normed} = 2$ cm ²)	400 000	25	78
Chemical	PaDa-I (15 U mL _{RM} ⁻¹)	Metal catalyst (0.1 mg mL _{RM} ⁻¹) of 0.5% Au–0.5% Pd/TiO ₂ , 10 mM ethylbenzene, under 2 bar of 80% H ₂ and 20% air from a high-pressure gas reservoir	25 900 (98)	1.5	76
^a Abbreviations: AaeUPO: U yield, AOx: alcohol oxidase hydroxylase from Rhodococ	JPO from Agrocybe aegerita, W from <i>Pichia pastoris</i> , FDM: fo <i>cus jostii</i> , YgJM: homologue o	^a Abbreviations: <i>Aae</i> UPO: UPO from <i>Agrocybe aegerita</i> , WT: wild-type, PaDa-I or SoLo: recombinant mutant of <i>Aae</i> UPO, TTN: total turnover number, ee: enantiometric excess (%), STY: space time yield, AOx: alcohol oxidase from <i>Pichia pastoris</i> , FDM: formaldehyde dismutase from <i>Pseudomonas putida</i> F61, FDH: formate dehydrogenase from <i>Candida boidinii</i> , 3HB6H: 3-hydroxybenzoat-6-hydroxylase from <i>Rhodococcus jostii</i> , YgjM: homologue of old yellow enzyme from <i>Bacillus subtilis</i> , SO: sulfite oxidases SO from <i>Arabidopsis thaliana</i> .	rnover number, ee: enar Irogenase from <i>Candida</i> bidopsis thaliana.	htiometric excess (%), STY: sp boidinii, 3HB6H: 3-hydroxybe	ace time enzoat-6-

promising combination will certainly yield interesting photoenzymatic reaction schemes.

4.5 Anthraquinone

Water-soluble anthraquinone 2-sulfonate (SAS) is a simple, robust, and promising organo-photocatalyst enabling simpler sacrificial electron donors than EDTA such as primary alcohols.^{97,106} This H_2O_2 generation photocatalyst has been demonstrated with peroxygenase- and haloperoxidase-catalyzed oxidation reactions⁹⁷ (Fig. 6b). The ability of SAS to also oxidize secondary alcohols to ketones enabled a truly chemoenzymatic reaction converting cyclohexane into cyclohexanone with the first step being peroxygenase-catalyzed and promoted by the H_2O_2 obtained from the second, SAS-photocatalytic cyclohexanol-oxidation step⁹⁷ (Fig. 6c).

4.6 γ -Ray-induced H₂O₂ formation

 γ -Rays induce (non-catalytic) water splitting and significant formation of H₂O₂ in aqueous media exposed to γ -irradiation.¹⁰⁷ This phenomenon can be exploited to promote peroxygenasecatalysed oxyfunctionalization reactions,¹⁰⁸ potentially opening up new possibilities of making use of nuclear wastes. For example, a proper dose rate of gamma radiation (12.9 Gy min⁻¹) could provide suitable H₂O₂ concentration (0.1 mM at steady state with an initial rate of 0.1 mM h⁻¹ for H₂O₂ formation) that enables significantly increased product formation while minimizing the inactivation of r*Aae*UPO.¹⁰⁸ Again, inactivation of the biocatalysts by the *in situ* formed reactive oxygen species represented a major impediment en route to preparative application, which may be overcome by spatial separation of the biocatalysts from the radiation source (*e.g.* in a flow chemistry setup).

5. Photoenzymatic oxyfunctionalization: current state of the art

The peroxygenase from *Agrocybe aegerita* (*Aae*UPO) represents the most frequently used representative of novel peroxygenases capable of selective oxyfunctionalization chemistry.^{48,54,75,109} Therefore, it is not very astonishing that it also dominates the following section of practical examples of photoenzymatic oxyfunctionalization reactions.

Table 3 and Fig. 7 summarize the current state of the art of photoenzymatic oxyfunctionalization reactions. Whereas possible, a particular focus was put on the TTN of the biocatalyst as this number also gives the reader an impression of the biocatalyst usage. Today, still enzymes are generally perceived as expensive, which *per se* is not true. The costs of enzyme production are subject to economy of scale which means that the price for the production of an enzyme exponentially decrease with the scale of its fermentation.¹¹⁰ If produced on small scale (*i.e.* <1 m³) production costs for an enzyme can easily surpass 10 000 \in kg⁻¹ or more. This is typically the case with enzymes commercialized by SMEs and general chemical vendors and probably motivates scientists for

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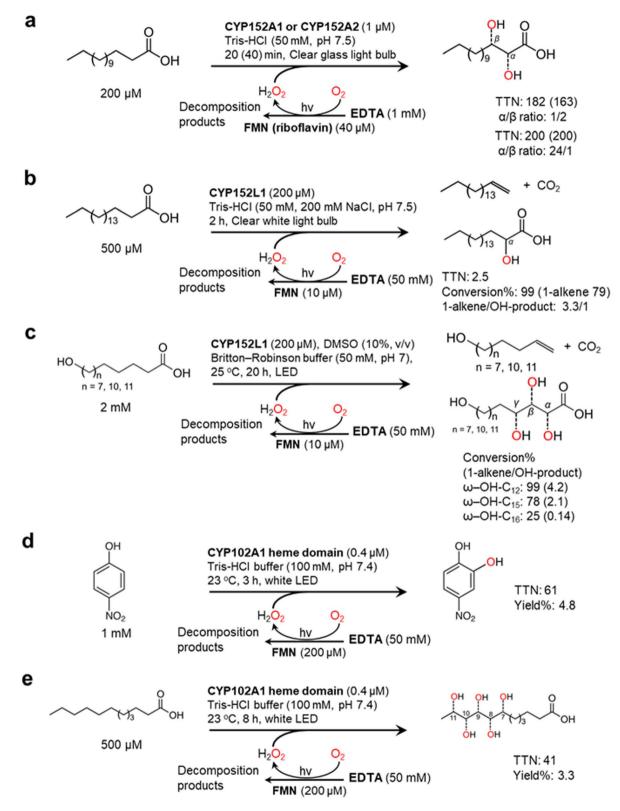


Fig. 5 Photochemically driving P450 peroxygenases using FMN as a photocatalyst and EDTA as an electron donor. Hydroxylation of fatty $acids^{101-103}$ (a and b) and ω -hydroxylated fatty $acids^{104}$ (c) is catalyzed by CYP152 peroxygenases. CYP102A1 heme domain also shows peroxygenase activity towards 4-nitrophenol (d) and lauric acid (e).¹²

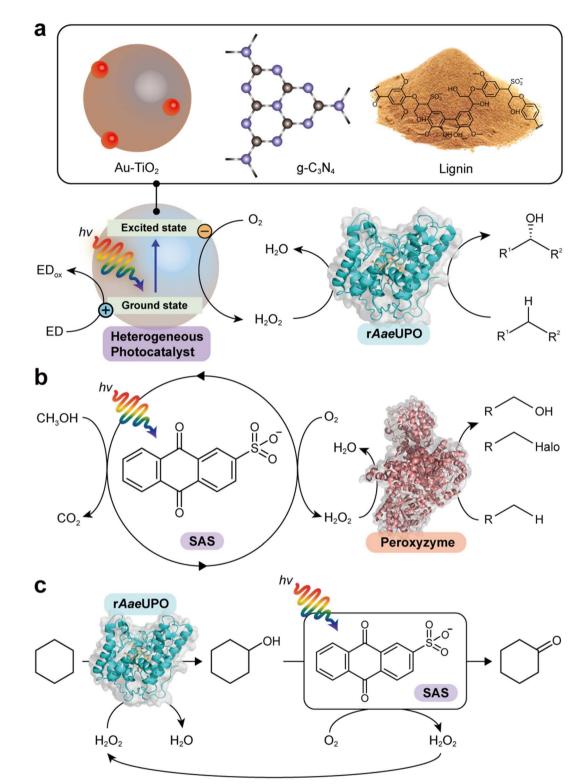


Fig. 6 Photocatalytic formation of H_2O_2 and selective oxidation reactions catalyzed by r*Aae*UPO and peroxyzyme. Representative photocatalyts include Au–TiO₂,⁹⁰ g-C₃N₄,⁹³ lignin⁹⁹ (a), and anthraquinone (SAS)^{97,106} (b). (c) Photocatalytic oxidation of cyclohexane to cyclohexanone. The SAS-catalyzed aerobic overoxidation of cyclohexanol to cyclohexanone produces H_2O_2 as a byproduct, which itself is used to support the r*Aae*UPO-catalyzed hydroxylation of cyclohexane.⁹⁷

statements like 'enzymes are expensive'. If, however, a biocatalyst is produced on large scale (*i.e.* \gg 10 m³) the production costs drop to values in the range of 250–1000 \in kg⁻¹, prices that are difficult to beat by traditional chemical catalysts (especially of rare transition metal catalysts or catalysts obtained from complex multi-step syntheses).

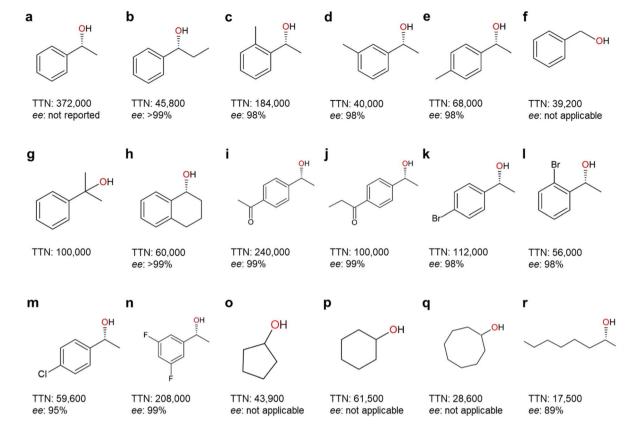


Fig. 7 Highest-record photobiocatalytic hydroxylation reactions of non-activated C-H bonds. All the photobiocatalysis were catalyzed by rAaeUPO (PaDa-I mutant) except (f) which was catalyzed by wild-type AaeUPO. TTN (nmol product per nmol enzyme) and enantioselectivity (ee%) for (*R*)-isoform were shown. References: (a);⁹⁶ (b), (m), (o)–(r);⁹⁰ (c)–(e), (g), (i)–(l), (n);⁹⁸ (f);⁸⁴ (h).⁹⁹

Furthermore, the cost contribution of an enzyme to the cost of the desired product directly correlate with its catalytic performance (or TTN). Hence, to attain a cost contribution of less than $10 \in \text{mol}_{\text{Product}}^{-1}$ (acceptable for the production of active pharmaceutical ingredients) enzymes produced at 250 or $10\ 000 \in \text{kg}_{\text{enzyme}}^{-1}$ would have to perform at least 1200 or $50\ 000$, respectively. Consequently, reaching a cost contribution of less than $0.1 \in \text{mol}_{\text{Product}}^{-1}$ requires TTN of more than $120\ 000\ or\ 5\ 000\ 000$, respectively. Of course, many other factors contribute to the final production costs of a chemical compound but the TTN at least gives a first orientation about the economic potential of an enzymatic process.

A range of photobiocatalytic oxyfunctionalization reactions, especially using *Aae*UPO, have been reported (Table 3). The reaction- and product scope attainable matches the scope of *Aae*UPO reactions reported using conventional *in situ* H_2O_2 generation systems and/or simple addition of H_2O_2 .⁵ The same is true concerning TNs reported for *Aae*UPO in these photobiocatalytic reactions that fall in the range of 17 500 to 372 000. In line with the above-discussed issues limitations of P450 monooxygenases following the H_2O_2 shunt pathway, the TNs observed with those enzymes typically fall back by one to two orders of magnitude behind those reported with peroxygenases (Fig. 5 and 7).

One common feature of most photoenzymatic reactions reported so far is the very low product concentrations, generally being in the lower millimolar range. Obviously, this is a result of the poor water solubility of the (mostly hydrophobic) starting materials of interest. Such low product titres, however, significantly impair their economic viability and also result in enormous amounts of waste water.¹¹¹ One possibility is to apply water-miscible organic cosolvents to increase the solubility of the hydrophobic reagents. Particularly, peroxygenases appear very suitable for such harsh reaction conditions as e.g. AaeUPO can withstand very high concentrations of typical organic cosolvents.112 Kara and coworkers recently reported g-C3N4adsorbed AaeUPO in a microaqueous reaction system comprised of cyclohexane as the main liquid phase.95 This approach, provided it is further optimized, may address this substrate-solubility issue at least for liquid starting materials. Another option to increase the overall reagent concentration is to make use of the so-called two-liquid-phase-system approach (2LPS).¹¹³ In the 2LPS approach, a nonpolar organic phase serves substrate reservoir and product sink as well with the reagents partitioning between the organic layer and the aqueous, catalyst-containing, aqueous phase. Phase transfer limitations, as often observed in such heterogeneous reaction media, can be alleviated by emulsion formation and thereby maximization of the interphase area as demonstrated by Churakova et al. in the photoenzymatic sulfoxidation thioanisol.114

6. Light-driven biocatalytic dehydrogenation

6.1 Photo(electro)chemical oxidation of NAD(P)H

NAD(P)⁺-dependent dehydrogenases catalyse a range of synthetically relevant oxidation reactions such as the regio- and/ or stereoselective oxidation of alcohols to the corresponding aldehydes or ketones. In the catalytic cycle NAD(P)⁺ is reduced to NAD(P)H. For economic reasons, the latter needs to be recycled back into its oxidised form. From a thermodynamic point-of-view, O₂ is an attractive electron acceptor because the redox potential of O₂ reduction [$E_{red}(O_2/O_2^{--}) = 0.33$ V vs.

normal hydrogen electrode (NHE), $E_{red}(O_2/H_2O_2) = 0.27 \text{ V} vs.$ NHE at pH 7]^{115,116} is more positive than that of NADH oxidation $[E_{ox}(NAD^+/NADH) = -0.32 \text{ V} vs.$ NHE at pH 7.0].¹¹⁷ Furthermore, only H_2O_2 (which can easily be dismutated into O_2 and H_2O) or H_2O are formed as by-products. NAD(P)H, however, is comparably stable against direct oxidation with O_2 necessitating catalytic acceleration of the reaction. Amongst the enzymatic methods, NAD(P)H oxidases have been established.¹¹⁸ Also nonenzymatic catalysts such as ABTS,^{119,120} quinoids,¹²¹ flavins^{26,122,123} or organic dyes¹²⁴ such as Congo Red²⁴ have been reported for the aerobic reoxidation of NAD(P)H. The sometimes rather sluggish NAD(P)H oxidation kinetics of these reagents can be accelerated by illumination.

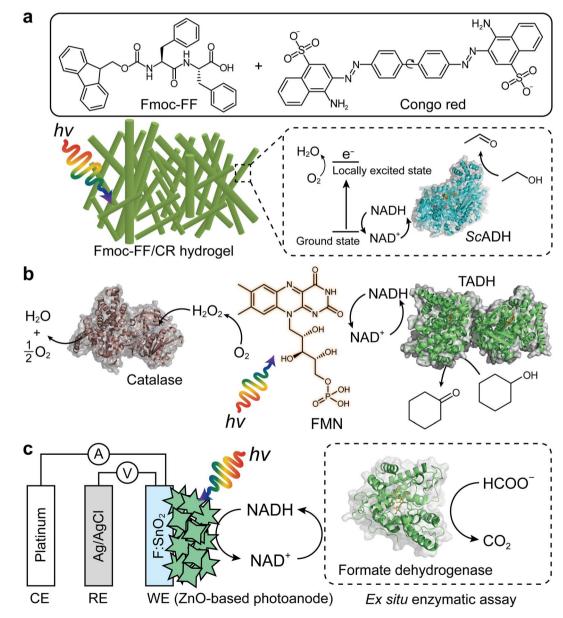


Fig. 8 Photo(electro)chemical regeneration of NAD(P)⁺ for biocatalytic oxidation reactions. (a) Fmoc-FF/CR-driven NAD⁺ recycling for biocatalytic oxidation of alcohol under visible light.²⁴ (b) Photoenzymatic synthesis of cyclohexanone through coupling of flavin mononucleotidesensitized regeneration of NAD⁺ and TADH-driven oxidation.²⁶ (c) ZnO-Based photoelectrochemical oxidation of NADH to NAD⁺.²⁷ The formation of NAD⁺ was confirmed by *ex situ* enzymatic assay. WE: working electrode. RE: reference electrode. CE: counter electrode.

6.2 Light-driven biocatalytic dehydrogenation reactions

The Park group reported a representative example for photobiocatalytic oxidation reactions,²⁴ in which β -sheet-rich amyloid nanofibers performed NAD⁺ regeneration in couple with enzymatic alcohol oxidation under an aerobic environment (Fig. 8a). Note that in contrast to many studies on photo(electro)catalytic NADH recycling,^{125,126} the reverse reaction (*i.e.*, NAD(P)⁺ production) has been rarely investigated despite the importance of the process. A solvatochromic Congo red (CR) was hybridized into an amyloid-derived peptide [e.g., Fmoc-diphenylalanine (Fmoc-FF)] through π - π interaction between aromatic functional groups of the Fmoc-FF and CR. This interaction made CR photocatalytic because the CR's donor-acceptor couple in the βsheet-rich nanostructure became highly planarized, which restrained nonradiative relaxation via CR's twisted intramolecular charge-transfer. As a result, CR performed O₂-fueled regeneration of NAD⁺ under visible light by (i) transferring CR's photoexcited electrons (in a locally excited state) to O_2 and (ii) delivering NADH's electrons to CR. The regenerated NAD⁺ activated alcohol dehydrogenase (ADH) from Saccharomyces cerevisiae (ScADH) to oxidize ethanol to acetaldehyde [ScADH's total turnover number (TTN_{SCADH}): 57 692, TTN_{NADH}: 3, TTN_{CR}: 30, yield: 60%].

Another example of photocatalyst-using O2 as an electron acceptor-are flavin molecules,26,123 such as flavin mononucleotide (FMN), flavin adenine dinucleotide, and riboflavin. The Hollmann group demonstrated the use of flavins to oxidize NAD(P)H under light conditions²⁶ (Fig. 8b). Flavin photoexcitation made flavin's redox potential more positive, facilitating its $NAD(P)^+$ regeneration in the presence of O₂. The flavinsensitised regeneration of NAD(P)⁺ activated ADH from Thermus sp. ATN1 (TADH) to catalyse the oxidation of cyclohexanone to cyclohexanol [TTN_{NADH}: 170, TTN_{FMN}: 340, TTN_{TADH}: 4359, FMN's turnover frequency (TOF_{FMN}): 38 h^{-1} , TOF_{TADH}: 498 h^{-1}]. In the photoenzymatic reaction, catalase was used to dismutate H_2O_2 to O_2 and H_2O . The photocatalytic NAD⁺ regeneration system was also applied to make ADH from horse liver (HLADH) catalyse the oxidation of diols e.g., 1,4-butanediol,²⁶ 1,5-pentanediol,²⁶ meso-3-methyl-1,5-pentanediol¹²³ to enantiopure lactones via hemiacetal intermediates [yield: 85-100%, TTN_{NADH}: 8.5-200, TTN_{FMN}: 85-400, TTN_{HLADH}: 1150].

One disadvantage of O_2 as terminal electron acceptor is its low solubility in aqueous media necessitating efficient external aeration. Photoelectrochemical (PEC) systems can overcome this issue because mechanistically (photo)anodes serve as electron acceptors.¹²⁷⁻¹³⁰ Ottone *et al.*²⁷ reported ZnO materials for PEC regeneration of NAD⁺ (Fig. 8c). Dual simulated solar irradiation (180 mW cm⁻²) and applied voltage (1.41 V *vs.* NHE at pH 7) on porous ZnO and flower-like ZnO (ZnOF) photoanodes drove the oxidation of NADH to NAD⁺ with conversions of 34.7% and 58.7%, respectively. An *ex situ* enzymatic assay using formate dehydrogenase confirmed that enzymatically active form of NAD⁺—formed by the ZnOF film—was 94%, not 100%. The minor loss should originate from dimerization and/ or fragmentation of NADH due to direct oxidation. Taken together, multiple photochemical and photoelectrochemical systems open up sustainable enzymatic oxidation reactions through light-powered $NAD(P)^+$ regeneration.

7. Conclusions and perspectives

Photobiocatalytic oxidation is an emerging synthetic route but still in infancy. For example, only a few cases of light-driven monooxygenase catalysis have been reported,12,13,29,31,36 and their catalytic activities are quite low (TTN < 1000). For operation of H₂O₂-accepting enzymes, light-driven in situ H₂O₂ generation is advantageous because it uses a clean light energy source and does not need any cofactors and additional enzymes. But the limited number of UPOs and P450 peroxygenases poses an obstacle to broader applications. Genome mining of new UPOs and heterologous expression in a suitable heterologous host have been tried, but the number of characterized UPOs remains limited. While the current target for genome mining mostly focused on fungal sources, its range should be expanded to other kingdoms, such as bacteria and plants, to obtain a diverse spectrum of UPOs. For directed evolution to achieve desired activity and stability, highthroughput screening (HTS) system should be applied to select mutants having the enhanced properties from the enzyme library. Information on 3D structures of diverse UPOs are essential for enzyme engineering, only a few structures of UPOs are available. On the other hand, the P450 peroxygenases (CYP152 family) catalyze only hydroxylation and decarboxylation reactions of fatty acids. More genome mining of new P450 peroxygenases is necessary for the expansion of substrate scope and their applications.

Light-driven reaction systems compete with more traditional approaches and today still the light-driven approaches appear to be less efficient compared to the latter. The direct photochemical regeneration of PAMO for example allowed for only a fraction of catalytic turnovers for the enzyme as compared to the reaction system utilizing NADPH together with an enzymatic regeneration system (96 TTNs as compared to 9471).^{29,131} Similarly, for the CYP152L1-catalyzed oxidative decarboxylation and hydroxylation of ω-OH lauric acid, the conversion achievable by a light-driven system (FMN/EDTA/light) was less than 10% of that by a secondary enzymatic system (CamAB/NADH/ FDH/formic acid).104 Obviously, direct photochemical regeneration systems need significant improvement in terms of activity and robustness. Particularly, the interaction of the non-natural redox partner with the enzyme active site has to be engineered. Furthermore, in case of O₂-dependent enzymes, the oxygen dilemma is still pending a promising solution.6 On the other hand, for ADH-catalyzed oxidation reactions, the aerobic regeneration of NAD⁺ by photochemical system (FMN/light) (TTN_{NAD} of 170)²⁶ appeared to be comparable in terms of cofactor recycling to the NADH oxidase system (TTN_{NAD} of 109).¹³² Since most reports on light-driven biocatalytic oxidation have focused so far on the demonstration of proof-of-concepts, further engineering studies on the optimization of reaction conditions should enhance their performances.

We should also consider technical issues for practical applications of photobioreactors. TiO₂-based photocatalysts

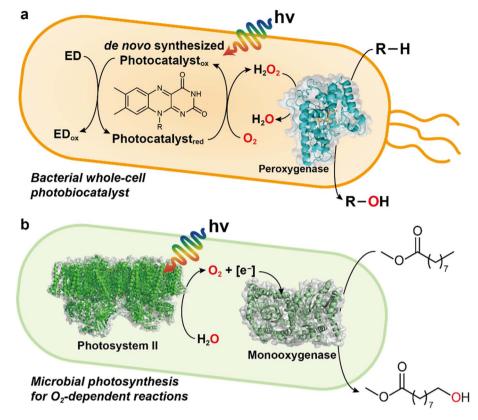


Fig. 9 Examples of light-driven biocatalytic oxidation in microorganisms. (a) Photoenzymatic, whole-cell biocatalyst expressing peroxygenase genes. *De novo* synthesized photocatalysts (*e.g.*, FMN, FAD) quickly react with O_2 to yield H_2O_2 in the presence of electron donor (ED) and light. The peroxygenase can hydroxylate the substrate (R–H) by using H_2O_2 as an oxygen donor. EDTA or TEOA can be used as an ED and other EDs such as methanol, formate, or simple amino acids may serve as alternatives. (b) Photosystem-assisted monooxygenase reaction. Photosystem II oxidizes water for producing O_2 and reducing equivalents (*e.g.*, NADH). These molecules are consumed for monooxygenase-driven catalysis (*e.g.*, oxyfunctionalization).

rely on a limited wavelength range (to near UV and blue light), thus leaving a significant spectrum of solar energy unexploited. In this context, a recent study using multiple wavelengthcomplementary photocatalysts provides insight into more efficient use of the polychromatic light energy.⁹⁴ Regarding the light sources, white light bulbs or light-emitting diodes (LEDs) have been widely used for photochemically driving several redox enzymes⁷ including UPOs (Table 3), P450 peroxygenases (Fig. 5), and dehydrogenases (Fig. 8). A versatile and advanced photoreactor platform is desired for HTS, preparative-scale batch reactions, and continuous processing, like the 'Photochemistry LED Illuminator' that identifies the optimal wavelength and power for specific photochemical transformation.⁹⁶

Whole cell-based approaches are attractive for the scale-up and industrial applications of light-driven biocatalytic oxidation (Fig. 9). Future efforts are needed for *de novo* synthesis of photocatalysts, peroxygenase engineering through rational design and/or directed evolution, and reaction optimization in a one-pot cascade process. The whole-cell platform should allow for high substrate concentration, consequently high productivities can be obtained. Therefore, engineering of strains by integration of desired peroxygenase genes into the genome will also accelerate the development of improved bacterial strains for massive production of desired chemicals.

Future research should focus on preparative oxyfunctionalization chemistry of peroxygenases. In a recent review, Dong et al.4 provided guidelines to assess the catalytic activity and productivity for the practical usefulness of enzymes for the production of desired products. According to the enzyme performance requirements of a given product category of pharma, fine chemical, speciality chemical, and bulk chemicals, required minimal productivity (TTN) for an enzyme is 4000, 26 666, 160 000, and 800 000, respectively.^{4,109} Meanwhile, minimally required STY (g L^{-1} d⁻¹) values for fine chemical production and pharmaceuticals are 2.4 and 0.024, respectively.133 Based on AaeUPO's performance by photochemically driven UPO catalysis, the requirements up to 'speciality chemical' (TTN of 160 000) are satisfied for some cases of UPOcatalyzed reactions (Table 3 and Fig. 7). For wide expansion of UPO's scope to the production of "bulk chemicals," further development of new UPOs, highly active and stable proteins, and whole-cell platforms are necessary.

When combined with the given knowledge and advances of photocatalysis, the unique catalytic activities and structures of monooxygenases, peroxygenases, and dehydrogenases promise a future route for greener synthesis using solar energy. Current challenges of the limited number of redox enzymes, narrow substrate range, and low stability are expected to be overcome by genome mining of new enzymes, protein engineering, and the development of whole-cell biocatalysts. A sustainable photobioreactor with optimized light source, robust photocatalyst, and waste-free electron donor is needed for driving preparative chemistry of peroxygenases and dehydrogenases.

Author contributions

The manuscript was written through contributions of all authors. All authors have given approval to the final version of the manuscript.

Conflicts of interest

The authors have no conflicts to declare.

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References

- E. Roduner, W. Kaim, B. Sarkar, V. B. Urlacher, J. Pleiss, R. Gläser, W.-D. Einicke, G. A. Sprenger, U. Beifuß, E. Klemm, C. Liebner, H. Hieronymus, S.-F. Hsu, B. Plietker and S. Laschat, *ChemCatChem*, 2013, 5, 82–112.
- 2 M. P. van der Helm, B. Klemm and R. Eelkema, *Nat. Rev. Chem.*, 2019, **3**, 491–508.
- 3 X. Ye and C.-H. Tan, Chem. Sci., 2021, 12, 533-539.
- 4 J. Dong, E. Fernández-Fueyo, F. Hollmann, C. E. Paul, M. Pesic, S. Schmidt, Y. Wang, S. Younes and W. Zhang, *Angew. Chem., Int. Ed.*, 2018, 57, 9238–9261.
- 5 M. Hobisch, D. Holtmann, P. G. de Santos, M. Alcalde, F. Hollmann and S. Kara, *Biotechnol. Adv.*, 2021, **51**, 107615.
- 6 D. Holtmann and F. Hollmann, *ChemBioChem*, 2016, 17, 1391–1398.
- 7 S. H. Lee, D. S. Choi, S. K. Kuk and C. B. Park, *Angew. Chem., Int. Ed.*, 2018, **57**, 7958–7985.
- 8 L. Schmermund, V. Jurkas, F. F. Ozgen, G. D. Barone, H. C. Büchsenschütz, C. K. Winkler, S. Schmidt, R. Kourist and W. Kroutil, ACS Catal., 2019, 9, 4115–4144.
- 9 M. C. R. Rauch, M. M. E. Huijbers, M. Pabst, C. E. Paul, M. Pešić, I. W. C. E. Arends and F. Hollmann, *Biochim. Biophys. Acta, Proteins Proteomics*, 2020, 1868, 140303.
- 10 F. F. Özgen, M. E. Runda and S. Schmidt, *ChemBioChem*, 2021, 22, 790–806.
- 11 M. Bocola, F. Schulz, F. Leca, A. Vogel, M. W. Fraaije and M. T. Reetz, *Adv. Synth. Catal.*, 2005, 347, 979–986.
- 12 T. K. Le, J. H. Park, D. S. Choi, G. Y. Lee, W. S. Choi, K. J. Jeong, C. B. Park and C.-H. Yun, *Green Chem.*, 2019, 21, 515–525.
- 13 T. K. Le, J. Kim, N. A. Nguyen, T. H. H. Nguyen, E. G. Sun, S. M. Yee, H. S. Kang, S. J. Yeom, C. B. Park and C.-H. Yun, *ChemSusChem*, 2021, 14, 3054–3058.

- 14 E. M. J. Gillam, Z. Y. Guo and F. P. Guengerich, Arch. Biochem. Biophys., 1994, 312, 59–66.
- 15 M. Girhard, S. Schuster, M. Dietrich, P. Dürre and V. B. Urlacher, *Biochem. Biophys. Res. Commun.*, 2007, **362**, 114–119.
- 16 A. Dennig, M. Kuhn, S. Tassoti, A. Thiessenhusen, S. Gilch, T. Bulter, T. Haas, M. Hall and K. Faber, *Angew. Chem., Int. Ed.*, 2015, 54, 8819–8822.
- P. Molina-Espeja, E. Garcia-Ruiz, D. Gonzalez-Perez, R. Ullrich, M. Hofrichter and M. Alcalde, *Appl. Environ. Microbiol.*, 2014, 80, 3496–3507.
- 18 P. Molina-Espeja, S. Ma, D. M. Mate, R. Ludwig and M. Alcalde, *Enzyme Microb. Technol.*, 2015, 73–74, 29–33.
- 19 M. P. J. Deurzen, B. W. van Groen, F. Rantwijk and R. A. van Sheldon, *Biocatalysis*, 1994, **10**, 247–255.
- 20 D. Cannella, K. B. Möllers, N.-U. Frigaard, P. E. Jensen, M. J. Bjerrum, K. S. Johansen and C. Felby, *Nat. Commun.*, 2016, 7, 11134.
- 21 B. Bissaro, Z. Forsberg, Y. Ni, F. Hollmann, G. Vaaje-Kolstad and V. G. H. Eijsink, *Green Chem.*, 2016, **18**, 5357–5366.
- 22 B. Bissaro, E. Kommedal, Å. K. Røhr and V. G. H. Eijsink, *Nat. Commun.*, 2020, **11**, 890.
- 23 B. M. Blossom, D. A. Russo, R. K. Singh, B. van Oort, M. B. Keller, T. I. Simonsen, A. Perzon, L. F. Gamon, M. J. Davies, D. Cannella, R. Croce, P. E. Jensen, M. J. Bjerrum and C. Felby, ACS Sustainable Chem. Eng., 2020, 8, 9301–9310.
- 24 G. Son, J. Kim and C. B. Park, ACS Appl. Energy Mater., 2020, 3, 1215–1221.
- 25 A. J. Ganzhorn, D. W. Green, A. D. Hershey, R. M. Gould and
 B. V. Plapp, *J. Biol. Chem.*, 1987, 262, 3754–3761.
- 26 S. Gargiulo, I. W. C. E. Arends and F. Hollmann, *ChemCatChem*, 2010, 3, 338-342.
- 27 C. Ottone, D. Pugliese, M. Laurenti, S. Hernández,
 V. Cauda, P. Grez and L. Wilson, ACS Appl. Mater. Interfaces, 2021, 13, 10719–10727.
- 28 Q. Guo, L. Gakhar, K. Wickersham, K. Francis, A. Vardi-Kilshtain, D. T. Major, C. M. Cheatum and A. Kohen, *Biochemistry*, 2016, 55, 2760–2771.
- 29 F. Hollmann, A. Taglieber, F. Schulz and M. T. Reetz, *Angew. Chem., Int. Ed.*, 2007, **46**, 2903–2906.
- 30 A. Taglieber, F. Schulz, F. Hollmann, M. Rusek and M. T. Reetz, *ChemBioChem*, 2008, 9, 565–572.
- 31 J. H. Park, S. H. Lee, G. S. Cha, D. S. Choi, D. H. Nam, J. H. Lee, J.-K. Lee, C.-H. Yun, K. J. Jeong and C. B. Park, *Angew. Chem., Int. Ed.*, 2015, 54, 969–973.
- 32 F. F. Özgen, M. E. Runda, B. O. Burek, P. Wied, J. Z. Bloh,
 R. Kourist and S. Schmidt, *Angew. Chem., Int. Ed.*, 2020,
 59, 3982–3987.
- 33 F. Mascia, S. B. Pereira, C. C. Pacheco, P. Oliveira, J. Solarczek, A. Schallmey, R. Kourist, V. Alphand and P. Tamagnini, *Green Chem.*, 2022, 24, 6156–6167.
- 34 E. Erdem, L. Malihan-Yap, L. Assil-Companioni, H. Grimm,
 G. D. Barone, C. Serveau-Avesque, A. Amouric,
 K. Duquesne, V. de Berardinis, Y. Allahverdiyeva,
 V. Alphand and R. Kourist, *ACS Catal.*, 2022, 12, 66–72.

- 35 A. Hoschek, A. Schmid and B. Bühler, *ChemCatChem*, 2018, 10, 5366–5371.
- 36 N. H. Tran, D. Nguyen, S. Dwaraknath, S. Mahadevan, G. Chavez, A. Nguyen, T. Dao, S. Mullen, T. A. Nguyen and L. E. Cheruzel, *J. Am. Chem. Soc.*, 2013, **135**, 14484–14487.
- 37 Q. Lam, A. Cortez, T. T. Nguyen, M. Kato and L. Cheruzel, J. Inorg. Biochem., 2016, 158, 86–91.
- 38 M. Kato, D. Nguyen, M. Gonzalez, A. Cortez, S. E. Mullen and L. E. Cheruzel, *Biorog. Med. Chem.*, 2014, 22, 5687– 5691.
- 39 Q. Lam, M. Kato and L. Cheruzel, *Biochim. Biophys. Acta*, 2016, **1857**, 589–597.
- 40 S. H. Lee, Y. C. Kwon, D. M. Kim and C. B. Park, *Biotechnol. Bioeng.*, 2013, **110**, 383–390.
- 41 J. H. Lee, D. H. Nam, S. H. Lee, J. H. Park, C. B. Park and K. J. Jeong, *J. Ind. Eng. Chem.*, 2016, **33**, 28–32.
- 42 K. Jensen, P. E. Jensen and B. L. Møller, ACS Chem. Biol., 2011, 6, 533–539.
- 43 S. B. Mellor, A. Z. Nielsen, M. Burow, M. S. Motawia,
 D. Jakubauskas, B. L. Moller and P. E. Jensen, *ACS Chem. Biol.*, 2016, 11, 1862–1869.
- 44 A. Hoschek, B. Bühler and A. Schmid, *Angew. Chem., Int. Ed.*, 2017, **56**, 15146–15149.
- 45 Z. Zhao, D. Lan, X. Tan, F. Hollmann, U. T. Bornscheuer,
 B. Yang and Y. Wang, *ACS Catal.*, 2019, 9, 2916–2921.
- 46 F. Björkling, S. E. Godtfredsen and O. Kirk, *Chem. Commun.*, 1990, **19**, 1301–1303.
- 47 G. T. Höfler, A. But and F. Hollmann, *Org. Biomol. Chem.*, 2019, **17**, 9267–9274.
- 48 M.-C. Sigmund and G. J. Poelarends, *Nat. Catal.*, 2020, 3, 690–702.
- 49 O. Shoji and Y. Watanabe, J. Biol. Inorg Chem., 2014, 19, 529–539.
- 50 P. D. Shaw and L. P. Hager, *J. Biol. Chem.*, 1961, **236**, 1626–1630.
- 51 F. van Rantwijk and R. A. Sheldon, *Curr. Opin. Biotechnol.*, 2000, **11**, 554–564.
- 52 R. Ullrich, J. Nuske, K. Scheibner, J. Spantzel and M. Hofrichter, *Appl. Environ. Microbiol.*, 2004, 70, 4575– 4581.
- 53 F. Tonin, F. Tieves, S. Willot, A. van Troost, R. van Oosten, S. Breestraat, S. van Pelt, M. Alcalde and F. Hollmann, *Org. Process Res. Dev.*, 2021, 25, 1414–1418.
- 54 A. Beltrán-Nogal, I. Sánchez-Moreno, D. Méndez-Sánchez,
 P. G. de Santos, F. Hollmann and M. Alcalde, *Curr. Opin.* Struct. Biol., 2022, 73, 102342.
- 55 C. Aranda, J. Carro, A. González-Benjumea, E. D. Babot, A. Olmedo, D. Linde, A. T. Martínez and A. Gutiérrez, *Biotechnol. Adv.*, 2021, 51, 107703.
- 56 H. Joo, Z. Lin and F. H. Arnold, Nature, 1999, 399, 670-673.
- 57 A. W. Munro, K. J. McLean, J. L. Grant and T. M. Makris, *Biochem. Soc. Trans.*, 2018, **46**, 183–196.
- 58 O. Fujishiro, O. Shoji, S. Nagano, H. Sugimoto, Y. Shiro and
 Y. Watanabe, *J. Biol. Chem.*, 2011, 286, 29941–29950.

- 59 C. E. Paul, E. Churakova, E. Maurits, M. Girhard, V. B. Urlacher and F. Hollmann, *Bioorg. Med. Chem.*, 2014, 22, 5692–5696.
- 60 L. Hammerer, M. Friess, J. Cerne, M. Fuchs, G. Steinkellner,
 K. Gruber, K. Vanhessche, T. Plocek, C. K. Winkler and
 W. Kroutil, *ChemCatChem*, 2019, 22, 5642–5649.
- 61 J. Belcher, K. J. McLean, S. Matthews, L. S. Woodward, K. Fisher, S. E. J. Rigby, D. R. Nelson, D. Potts, M. T. Baynham, D. A. Parker, D. Leys and A. W. Munro, *J. Biol. Chem.*, 2014, 289, 6535–6550.
- 62 Y. Jiang, Z. Li, C. Wang, Y. J. Zhou, H. Xu and S. Li, *Biotechnol. Biofuels*, 2019, **12**, 79.
- 63 H. Onoda, O. Shoji, K. Suzuki, H. Sugimoto, Y. Shiro and Y. Watanabe, *Catal. Sci. Technol.*, 2018, **8**, 434–442.
- 64 M. A. Rude, T. S. Baron, S. Brubaker, M. Alibhai, S. B. Del Cardayre and A. Schirmer, *Appl. Environ. Microbiol.*, 2011, 77, 1718–1727.
- 65 N. Ma, Z. Chen, J. Chen, J. Chen, C. Wang, H. Zhou, L. Yao,
 O. Shoji, Y. Watanabe and Z. Cong, *Angew. Chem., Int. Ed.*, 2018, 57, 7628–7633.
- 66 Z. Forsberg, M. Sorlie, D. Petrovic, G. Courtade, F. L. Aachmann, G. Vaaje-Kolstad, B. Bissaro, A. K. Rohr and V. G. H. Eijsink, *Curr. Opin. Struct. Biol.*, 2019, 59, 54– 64.
- 67 P. Chylenski, B. Bissaro, M. Sorlie, A. K. Rohr, A. Varnai, S. J. Horn and V. G. H. Eijsink, *ACS Catal.*, 2019, 9, 4970– 4991.
- 68 B. Wang, Z. Wang, G. J. Davies, P. H. Walton and C. Rovira, *ACS Catal.*, 2020, **10**, 12760–12769.
- 69 L. Ciano, G. J. Davies, W. B. Tolman and P. H. Walton, *Nat. Catal.*, 2018, 1, 571–577.
- 70 B. L. Cantarel, P. M. Coutinho, C. Rancurel, T. Bernard, V. Lombard and B. Henrissat, *Nucleic Acids Res.*, 2009, 37, D233–D238.
- 71 B. Bissaro, Å. K. Røhr, G. Müller, P. Chylenski, M. Skaugen,
 Z. Forsberg, S. J. Horn, G. Vaaje-Kolstad and
 V. G. H. Eijsink, *Nat. Chem. Biol.*, 2017, 13, 1123–1128.
- 72 Y. Gaber, B. Rashad, R. Hussein, M. Abdelgawad, N. S. Ali,
 T. Dishisha and A. Várnaie, *Biotechnol. Adv.*, 2020, 43, 107583.
- 73 V. G. H. Eijsink, D. Petrovic, Z. Forsberg, S. Mekasha,
 Å. K. Røhr, A. Várnai, B. Bissaro and G. Vaaje-Kolstad, *Biotechnol. Biofuels*, 2019, 12, 58.
- 74 H. L. Wapshott-Stehli and A. M. Grunden, *Enzyme Microb. Technol.*, 2021, 145, 109744.
- 75 B. O. Burek, S. Bormann, F. Hollmann, J. Z. Bloh and D. Holtmann, *Green Chem.*, 2019, 21, 3232–3249.
- 76 S. J. Freakley, S. Kochius, J. van Marwijk, C. Fenner, R. J. Lewis, K. Baldenius, S. S. Marais, D. J. Opperman, S. T. L. Harrison, M. Alcalde, M. S. Smit and G. J. Hutchings, *Nat. Commun.*, 2019, **10**, 4178.
- 77 F. Hollmann and A. Schimid, *J. Inorg. Biochem.*, 2009, **103**, 313–315.
- 78 A. E. W. Horst, S. Bormann, J. Meyer, M. Steinhagen, R. Ludwig, A. Drews, M. Ansorge-Schumacher and D. Holtmann, *J. Mol. Catal. B: Enzym.*, 2016, **133**, S137–S142.

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- 79 T. Krieg, S. Huttmann, K.-M. Mangold, J. Schrader and D. Holtmann, *Green Chem.*, 2011, **13**, 2686–2689.
- 80 L. Getrey, T. Krieg, F. Hollmann, J. Schrader and D. Holtmann, *Green Chem.*, 2014, **16**, 1104–1108.
- 81 C. Kormann, D. W. D. W. Bahnemann and M. R. Hoffmann, *Environ. Sci. Technol.*, 1988, 22, 798–806.
- 82 W. R. Frisell, C. W. Chung and C. G. Mackenzie, J. Biol. Chem., 1959, 234, 1297–1302.
- 83 D. I. Perez, M. M. Grau, I. W. C. E. Arends and F. Hollmann, *Chem. Commun.*, 2009, 6848–6850.
- 84 E. Churakova, M. Kluge, R. Ullrich, I. Arends, M. Hofrichter and F. Hollmann, *Angew. Chem., Int. Ed.*, 2011, 50, 10716– 10719.
- 85 E. Fernández-Fueyo, Y. Ni, A. G. Baraibara, M. Alcalde, L. M. van Langen and F. Hollmann, *J. Mol. Catal. B: Enzym.*, 2016, 134, 347–352.
- 86 Y. Ni, E. Fernandez-Fueyo, A. G. Baraibar, R. Ullrich, M. Hofrichter, H. Yanase, M. Alcalde, W. J. H. van Berkel and F. Hollmann, *Angew. Chem., Int. Ed.*, 2016, 55, 798–801.
- 87 M. Pesic, S. J.-P. Willot, E. Fernández-Fueyo, F. Tieves, M. Alcalde and F. Hollmann, *Z. Naturforsch., C: J. Biosci.*, 2019, 74, 101–104.
- 88 M. M. C. H. van Schie, A. T. Kaczmarek, F. Tieves, P. G. de Santos, C. E. Paul, I. W. C. E. Arends, M. Alcalde, G. Schwarz and F. Hollmann, *ChemCatChem*, 2020, **12**, 3186–3189.
- 89 A. Al-Shameri, S. J.-P. Willot, C. E. Paul, F. Hollmann and L. Lauterbach, *Chem. Commun.*, 2020, **56**, 9667.
- 90 W. Zhang, B. O. Burek, E. Fernández-Fueyo, M. Alcalde, J. Z. Bloh and F. Hollmann, *Angew. Chem., Int. Ed.*, 2017, 56, 15451–15455.
- 91 W. Zhang, E. Fernandez-Fueyo, Y. Ni, M. van Schie, J. Gacs, R. Renirie, R. Wever, F. G. Mutti, D. Rother, M. Alcalde and F. Hollmann, *Nat. Catal.*, 2018, **1**, 55–62.
- 92 B. O. Burek, S. R. de Boer, F. Tieves, W. Zhang, M. van Schie,
 S. Bormann, M. Alcalde, D. Holtmann, F. Hollmann,
 D. W. Bahnemann and J. Z. Bloh, *ChemCatChem*, 2019, 11, 3093–3100.
- 93 M. M. C. H. van Schie, W. Zhang, F. Tieves, D. S. Choi, C. B. Park, B. O. Burek, J. Z. Bloh, I. W. C. E. Arends, C. E. Paul, M. Alcalde and F. Hollmann, *ACS Catal.*, 2019, 9, 7409–7417.
- 94 S. J. P. Willot, E. Fernandez-Fueyo, F. Tieves, M. Pesic, M. Alcalde, I. W. C. E. Arends, C. B. Park and F. Hollmann, ACS Catal., 2019, 9, 890–894.
- 95 M. Hobisch, M. M. C. H. van Schie, J. Kim, K. R. Andersen, M. Alcalde, R. Kourist, C. B. Park, F. Hollmann and S. Kara, *ChemCatChem*, 2020, **12**, 4009–4013.
- 96 H. E. Bonfield, K. Mercer, A. Diaz-Rodriguez, G. C. Cook,
 B. S. J. McKay, P. Slade, G. M. Taylor, W. X. Ooi,
 J. D. Williams, J. P. M. Roberts, J. A. Murphy,
 L. Schmermund, W. Kroutil, T. Mielke, J. Cartwright,
 G. Grogan and L. J. Edwards, *ChemPhotoChem*, 2020, 4, 45–51.
- 97 B. Yuan, D. Mahor, Q. Fei, R. Wever, M. Alcalde, W. Zhang and F. Hollmann, *ACS Catal.*, 2020, **10**, 8277–8284.
- 98 L. Schmermund, S. Reischauer, S. Bierbaumer, C. K. Winkler, A. Diaz-Rodriguez, L. J. Edwards, S. Kara,

T. Mielke, J. Cartwright, G. Grogan, B. Pieber and W. Kroutil, *Angew. Chem., Int. Ed.*, 2021, **60**, 6965–6969.

- 99 J. Kim, T. V. T. Nguyen, Y. H. Kim, F. Hollmann and C. B. Park, *Nat. Synth.*, 2022, 1, 217–226.
- 100 D. S. Choi, Y. Ni, E. Fernandez-Fueyo, M. Lee, F. Hollmann and C. B. Park, *ACS Catal.*, 2017, 7, 1563–1567.
- 101 M. Girhard, E. Kunigk, S. Tihovsky, V. V. Shumyantseva and
 V. B. Urlacher, *Biotechnol. Appl. Biochem.*, 2013, 60, 111–118.
- 102 I. Zachos, S. K. Gaßmeyer, D. Bauer, V. Sieber, F. Hollmann and R. Kourist, *Chem. Commun.*, 2015, **51**, 1918–1921.
- 103 K. Koninger, M. Grote, I. Zachos, F. Hollmann and R. Kourist, *J. Visualized Exp.*, 2016, **111**, e53439.
- 104 S. Bojarra, D. Reichert, M. Grote, A. G. Baraibar, A. Dennig,
 B. Nidetzky, C. Mugge and R. Kourist, *ChemCatChem*, 2018,
 10, 1192–1201.
- 105 W. J. Ong, L. L. Tan, Y. H. Ng, S. T. Yong and S. P. Chai, *Chem. Rev.*, 2016, **116**, 7159–7329.
- 106 W. Zhang, J. Gacs, I. W. C. E. Arends and F. Hollmann, *ChemCatChem*, 2017, **9**, 3821–3826.
- 107 S. Le Caër, Water, 2011, 3, 235–253.
- 108 W. Zhang, H. Liu, M. M. C. H. van Schie, P.-L. Hagedoorn, M. Alcalde, A. G. Denkova, K. Djanashvili and F. Hollmann, ACS Catal., 2020, 10, 14195–14200.
- 109 S. Bormann, A. G. Baraibar, Y. Ni, D. Holtmann and F. Hollmann, *Catal. Sci. Technol.*, 2015, **5**, 2038–2052.
- 110 P. Tufvesson, J. Lima-Ramos, M. Nordblad and J. M. Woodley, *Org. Process Res. Dev.*, 2011, **15**, 266–274.
- 111 Y. Ni, D. Holtmann and F. Hollmann, *ChemCatChem*, 2014, 6, 930–943.
- 112 T. Hilberath, A. van Troost, M. Alcalde and F. Hollmann, *Front. Catal.*, 2022, **2**, 882992.
- 113 M. M. C. H. van Schie, J.-D. Spöring, M. Bocola, P. D. de María and D. Rother, *Green Chem.*, 2021, 23, 3191–3206.
- 114 E. Churakova, I. W. C. E. Arends and F. Hollmann, *ChemCatChem*, 2013, 5, 565–568.
- 115 S. C. Perry, D. Pangotra, L. Vieira, L.-I. Csepei, V. Sieber, L. Wang, C. P. de León and F. C. Walsh, *Nat. Rev. Chem.*, 2019, 3, 442–458.
- 116 J. Kim, S. H. Lee, F. Tieves, C. E. Paul, F. Hollmann and C. B. Park, *Sci. Adv.*, 2019, **5**, eaax0501.
- 117 J. Kim and C. B. Park, *Curr. Opin. Chem. Biol.*, 2019, **49**, 122–129.
- 118 F. Hollmann, I. W. C. E. Arends and K. Buehler, *ChemCatChem*, 2010, **2**, 762–782.
- 119 I. Schröder, E. Steckhan and A. Liese, *J. Electroanal. Chem.*, 2003, **541**, 109–115.
- 120 S. Aksu, I. W. C. E. Arends and F. Hollmann, *Adv. Synth. Catal.*, 2009, **351**, 1211–1216.
- 121 G. Hilt, T. Jarbawi, W. R. Heineman and E. Steckhan, *Eur. J. Chem.*, 1997, **3**, 79–88.
- 122 J. B. Jones and K. E. Taylor, *J. Chem. Soc., Chem. Commun.*, 1973, 205–206.
- 123 M. Rauch, S. Schmidt, I. W. C. E. Arends, K. Oppelt, S. Kara and F. Hollmann, *Green Chem.*, 2017, **19**, 376–379.

- 124 P. Könst, S. Kara, S. Kochius, D. Holtmann, I. W. C. E. Arends, R. Ludwig and F. Hollmann, *ChemCatChem*, 2013, 5, 3027–3032.
- 125 J. Kim, S. H. Lee, F. Tieves, D. S. Choi, F. Hollmann, C. E. Paul and C. B. Park, *Angew. Chem., Int. Ed.*, 2018, 57, 13825–13828.
- 126 D. Wang, J. Kim and C. B. Park, *ACS Appl. Mater. Interfaces*, 2021, **13**, 58522–58531.
- 127 J. Kim, Y. W. Lee, E.-G. Choi, P. Boonmongkolras,
 B. W. Jeon, H. Lee, S. T. Kim, S. K. Kuk, Y. H. Kim,
 B. Shin and C. B. Park, *J. Mater. Chem. A*, 2020, 8, 8496–8502.
- 128 J. Kim, Y. Um, S. Han, T. Hilberath, Y. H. Kim, F. Hollmann and C. B. Park, *ACS Appl. Mater. Interfaces*, 2022, **14**, 11465– 11473.
- 129 S. K. Kuk, J. Jang, J. Kim, Y. Lee, Y. S. Kim, B. Koo, Y. W. Lee, J. W. Ko, B. Shin, J.-K. Lee and C. B. Park, *ChemSusChem*, 2020, 13, 2940–2944.
- 130 J. Kim, J. Jang, T. Hilberath, F. Hollmann and C. B. Park, *Nat. Synth.*, 2022, **1**, 776–786.
- 131 F. Schulz, F. Leca, F. Hollmann and M. T. Reetz, *Beilstein J. Org. Chem.*, 2005, 1, 10.
- 132 B. R. Riebel, P. R. Gibbs, W. B. Wellborn and A. S. Bommarius, *Adv. Synth. Catal.*, 2003, **345**, 707–712.
- 133 M. K. Julsing, S. Cornelissen, B. Buhler and A. Schmid, *Curr. Opin. Chem. Biol.*, 2008, **12**, 177–186.