

Cite this: *Chem. Sci.*, 2020, **11**, 2587

All publication charges for this article have been paid for by the Royal Society of Chemistry

Received 12th November 2019

Accepted 12th February 2020

DOI: 10.1039/c9sc05746c

rsc.li/chemical-science

# The Hitchhiker's guide to biocatalysis: recent advances in the use of enzymes in organic synthesis

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Enzymes are excellent catalysts that are increasingly being used in industry and academia. This perspective is primarily aimed at synthetic organic chemists with limited experience using enzymes and provides a general and practical guide to enzymes and their synthetic potential, with particular focus on recent applications.

## 1. Introduction

Faced with having to perform hundreds of reactions in what is essentially one pot (the prokaryotic cell), life has had to provide catalysts that are highly efficient and very selective. There is an enzyme suited to every natural compound. These enzymes can also catalyse non-natural synthetic reactions, and enzymes are even evolving *in vivo* to specifically handle non-natural industrial chemicals.<sup>1</sup> Moreover, biocatalysts can all function in the same readily available green solvent, water, a capability that the chemical industry needs to attain if it is to be sustainable.<sup>2</sup>

In this article we look at how the organic chemist can obtain these nifty catalysts, what they do and how they work, and how they can be adapted to industrial use, integrated into chemical processes, and modified to meet specific process requirements.

## 2. Sources and commercial availability of enzymes in the metagenome era

Although the chemist James Sumner had already isolated the enzyme urease in 1926 and proved its protein structure, it would take some time before this knowledge was translated into new chemical processes. However, by the beginning of the millennium quite a few companies were using biocatalysis<sup>3</sup> and Straathof *et al.* could already list 134 industrial biocatalytic

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processes or biotransformations.<sup>4</sup> Biocatalysis research accelerated from about 2000 onwards, largely as a result of the spectacular advances in (meta)genome sequencing and protein engineering<sup>5</sup> and by 2019 Woodley observed that several hundred processes are now in operation,<sup>6</sup> especially in the pharmaceutical industry.<sup>7</sup>

The extent of their impact on chemicals manufacture is immense. The largest use of isolated enzymes in a single process is the liquefaction and saccharification of starch to produce glucose. Another process uses glucose isomerase to produce 10<sup>7</sup> tons per annum of high fructose corn syrup from glucose.<sup>8</sup> The enzymatic conversion of waste cellulosic biomass to provide sugars for bioethanol will no doubt eventually surpass this in scale. In the chemical industry, nitrile hydratases are used to catalyse the conversion of acrylonitrile to the industrial monomer, acrylamide, on a multi hundred thousand ton per annum scale (see Section 5). In the pharmaceutical industry, penicillin G amidase is used in the industrial synthesis of thousands of tons of a variety of beta lactam antibiotics, such as amoxicillin.<sup>9</sup> Enzymes are also used on a large scale in the more traditional applications: food and beverage processing, detergents, pulp and paper manufacture, and in the textiles and leather industries.

Enzymes can be sourced from plants, *e.g.* carrots<sup>10</sup> and soy beans,<sup>11</sup> fungi<sup>12,13</sup> (such as baker's yeast, filamentous fungi and mushrooms), bacteria<sup>14</sup> and archaea.<sup>15</sup> The early use of readily available enzymes from animal organs, *e.g.* pig liver, obtained from abattoirs, has been superseded by advances in biotechnology, which enabled the cheaper production of enzymes through over-expression using recombinant DNA technology in microbial expression hosts.<sup>16</sup> Future developments are likely to involve the production of enzymes in plant hosts that can be grown, harvested and easily processed to give ultra-cheap enzymes.<sup>17</sup>

The globally traded enzyme market is worth about \$7 bn (this excludes production for in-house use). The majority of commercially traded enzymes are produced and sold by Novozymes, followed by DuPont (incorporating Danisco and Genencor). Other large scale producers include DSM, Amano and BASF (incorporating Verenum), Advanced Enzyme Technologies Limited and AB Enzymes. Other companies such as Thermo Fischer Scientific,

Hoffmann La Roche, Procter and Gamble, BBI Enzymes, Puratos, Novus International and Chr. Hansen are also producers of significant quantities of commercial enzymes. These companies tend to produce enzyme formulations for large scale applications, such as food and beverage production, household care (*e.g.* detergents) and bioenergy or feed, but increasingly also for pharmaceutical production and biotechnology. Carbohydrases, proteases, and to a lesser extent, lipases are the major enzyme classes sold and used on a large scale. The advantage of using these enzymes is that they are readily available, consistent from batch to batch, and are produced in sufficient quantities to support commercial biocatalytic processes. The downside is the limited range of enzymes available. Commercial enzyme producers often kindly provide enzyme samples to researchers in quantities that are sufficient for student projects. However, industry limits the available information on the enzymes for reasons of intellectual property and hence the species origin and protein sequence is becoming increasingly difficult to identify. This can be a problem if the product is discontinued, or if the enzyme is modified during formulation improvement. However, if an enzyme is already being used at scale in a chemical process the company can usually accommodate continued production.

Smaller companies, such as Codexis (USA), Biocatalysts (Wales, UK) and BRAIN (Germany), are focused on enzyme development. Interestingly, Johnson Matthey, the speciality chemicals company that produces catalysts, now has a wide range of biocatalysts among their products.

The smallest companies tend to be university spin-outs and include Enzymicals AG (Germany) and Prozomix (UK). They have the advantage of offering wide novel enzyme selections, catering to previously unexplored enzyme classes. They are also often willing to develop enzyme reactions according to client requirements. Enzymes may be purchased as single enzymes (often as kits containing co-factors, test substrates and other required reagents) and as enzyme selections (usually 10–50 enzymes of the same activity type). Research Chemical companies, such as Sigma-Aldrich, also sell enzymes, sourced from companies of all sizes.

Caution should be taken as many enzyme preparations are only partially purified from the cell, and may therefore contain multiple enzyme types and exhibit more than one activity.<sup>18</sup> Moreover, some enzymes have the capacity for more than one enzyme activity (*e.g.* fatty acid synthase).

New enzymes with different substrate profiles or even catalytic mechanisms can be obtained in various ways. Microbial collections are often cross-referenced for the catalytic activities each microbial isolate has shown. Increasingly the genomes of all species are being sequenced and annotated for the genes and the proteins (including enzymes) which they encode and the data deposited in publicly accessible data bases that can be “mined” for biocatalysts. Examples of such databases are the National Centre for Biological Information (NCBI)<sup>19</sup> and Uniprot.<sup>20</sup> If the gene sequence is known it can be ordered on-line and the plasmid ready for cloning in a suitable host will be delivered to your doorstep in a couple of weeks.

If an ideal enzyme cannot be found, new enzymes can be obtained directly from the environment using metagenomics,



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whereby DNA is extracted from an environmental sample of *e.g.* soil or water and fragments are filed into a “library” in individual host bacteria (*e.g.* *E. coli*). The library can be screened against target molecules, often using colorimetric assays in large arrays or automated chromatography systems. This enables the discovery of novel sequences and, with a suitable chemical screen, novel activities.<sup>21–23</sup> For example, the recent surge of interest in transamination motivated the discovery of novel, robust transaminases derived from the metagenome of the microbial population of the human mouth,<sup>24</sup> and domestic drain.<sup>25</sup> If a suitable enzyme has not been found it is now usual to modify the enzyme at the genetic level using advanced directed evolution techniques.<sup>26</sup> These use *in vitro* mutation techniques, often through recombination of variants, to create permutations of the enzyme’s amino acid sequence. Although the theoretical number of potential variants is astronomical, using enzyme structural data to plan mutations allows this to be reduced to a more tractable number.<sup>27,28</sup>

### 3. Enzyme structure, functions, mechanisms and scope in organic synthesis

Enzymes consist of linear polypeptide chains (proteins) of about 300 alpha L-amino acids (hence a mass of about 33 kDa).<sup>29</sup> This varies considerably as 4-oxalocrotonate tautomerase is only 62 amino acids long<sup>30</sup> while fatty acid synthase is 2461 amino acids long. They can also form associations of multiple proteins, and will therefore often be larger in practice. Proteins are typically comprised of the 20 proteinogenic amino acids that are encoded for on the DNA, but there is an additional pair, selenocysteine and pyrrolysine, that are formed during translation and may be located in the enzyme’s active site.

Typically only a handful of these amino acids perform catalytic functions in any one protein, while the other 99% are often structural or regulatory. Some amino acids, such as histidine, are frequent contributors to the catalytic chemistry of enzymes, while others, such as glycine, do not contribute.<sup>31,32</sup> Structural amino acids align the catalytic amino acids in order to enable the catalytic mechanism. Their physical proximity and spatial arrangement also limits the range of acceptable substrates, thereby creating catalytic specificity.

The catalytic active sites provide an exclusive reaction environment and the active site is therefore usually buried at the end of a channel lined with bulky and aromatic amino acids that controls access of substrates and solvents. Consequently the amino acid composition in the channel varies with the enzyme type (*e.g.* more hydrophilic for ligases but hydrophobic for oxidoreductases).<sup>33</sup> Mechanical components, *e.g.* lipid activated lids to the active site of lipases, also help define substrate type.<sup>34</sup> These structures endow enzymes with exquisite chemo-, regio- and most importantly stereo-selectivity.<sup>35</sup> Engineering of these channels is a developing area of research.<sup>36</sup>

The non-catalytic amino acids also provide the structural backbone of the enzyme in that the hydrophobic regions fold inwards, away from the hydrophilic aqueous environment, to

provide the core; while the hydrophilic regions form the surface and enhance enzyme solubility in the aqueous environment.

Enzyme catalysis can also include non-protein elements, referred to as cofactors, which greatly expand the catalytic repertoire of that enzyme class. There are 27 small molecule organic enzyme cofactors<sup>37</sup> and a number of commonly used metals<sup>38</sup> including Mg, V, Mn, Fe, Ni, Co, Cu, Mo, Zn, Se and W,<sup>39</sup> many of which are tightly bound to the enzyme. Exploration of the repertoire of biocatalysts is far from reaching its limits and is not constrained by existing biochemistry. Further expansion can be realised through cofactor engineering,<sup>40–42</sup> whereby existing metal or organic cofactors can be replaced by artificial analogs. Another possibility is to reprogram the amino acids that constitute the enzyme backbone with non-canonical amino acids (ncAAs).<sup>43</sup>

In short, largely as a result of *in silico* metagenome mining and directed evolution, biocatalysis has evolved into a cost-effective and broadly applicable technology that has been successfully integrated into mainstream organic synthesis.<sup>7,44</sup> Indeed, the rapidly expanding biocatalysis toolbox<sup>45</sup> has created a situation where it is eminently feasible to apply a biocatalytic retrosynthetic approach to identify enzymatic routes by ‘deconstructing’ target molecules.<sup>46–49</sup>

### 4. Advantages and limitations of enzymes: improving performance

Organic chemists who are unfamiliar with enzymes are often discouraged by real or perceived limitations with regard to their application in organic synthesis. Indeed, it is still widely believed that enzymatic processes are slow and occur at very low substrate concentrations in water, resulting in long reaction times and poor catalyst and volumetric productivities (kg product per kg enzyme and kg product per L per h), respectively. This is certainly not observed with enzymes catalysing their natural substrates under physiological conditions. It is, however, often true of wild-type enzymes transforming molecules which are structurally very different to their natural counterparts under the demanding conditions of industrial processes with elevated temperatures and in the presence of organic solvents.

Nonetheless, advances in protein engineering and directed evolution techniques over the last two decades have dramatically changed this situation. Enzymes can be evolved, in a relatively short time and at relatively limited expense, to perform well with non-natural substrates under non-natural conditions. *i.e.* at substrate concentrations of  $>100 \text{ g L}^{-1}$ , volumetric productivities of  $>10 \text{ g L}^{-1}$  and catalyst productivities of  $>100 \text{ kg product per kg enzyme}.$ <sup>50</sup>

Another perceived drawback is that enzymes have very limited stability under the conditions encountered in organic synthesis. It is true that many, particularly wild-type enzymes, may be deactivated by unfolding (denaturing) at elevated temperatures, in the presence of organic solvents or under high shear caused by mixing or gas bubble surfaces. However, a number of extracellular enzyme classes, such as proteases,



amylases, lipases, laccases and cellulases, are generally more robust and have been widely applied in detergents and industrial processes.

A good source of more stable enzymes is from organisms that live in extreme environments (extremophiles), such as thermophiles that thrive around geological vents.<sup>51</sup> For example, enzymes were discovered that function at temperatures up to 120 °C, at extremes of pH between 0 and 12 and at pressures of 1000 bar, as well as in pure solvents and in slurries. Some have already been harnessed for industrial use, such as the glucoamylases and amylases which perform admirably in the hydrolysis of starch at temperatures above 100 °C. Some enzymes, such as lipases, are stable in highly hydrophobic solvents.<sup>52</sup> Alternatively, enzymes can be evolved to increase their stability under various conditions of temperature, pH, in the presence of organic solvents, *etc.*

Yet another practical issue with enzymes is that it is challenging to recover them from aqueous media and they are, therefore, used on a single-use throw away basis. This is neither conducive to a waste-free circular economy nor a cost-effective process. One possible solution to the problem is to immobilise the enzyme by transforming it into a heterogeneous solid catalyst which can be readily recovered from an aqueous medium and reused.<sup>53–59</sup> This substantially reduces the enzyme costs per kg product and the environmental footprint, thereby driving competitiveness and sustainability. Immobilisation also results in a decrease in the flexibility of the enzyme molecule, affording an increase in operational stability by suppressing its propensity to unfold at elevated temperatures or in the presence of organic solvents. In short, immobilisation can enable applications that would not be economically viable with the free enzyme and it can facilitate their use in cost-effective continuous flow processing.<sup>60,61</sup>

Immobilisation typically involves binding the enzyme to a prefabricated carrier (support) by physical adsorption, ionic interactions or the formation of covalent bonding *via* the amino acids in the hydrophilic surface shell (usually lysine). Alternatively, the enzyme can be encapsulated in a polymeric matrix formed in the presence of the enzyme. However, the use of a support inevitably leads to 'dilution of activity', owing to the introduction of a large portion of non-catalytic ballast, giving rise to lower catalyst productivities.

Immobilisation by cross linking of enzyme molecules to each other, in contrast, affords carrier-free immobilised enzymes with high productivities and avoids the extra costs of a carrier. For example, cross-linked enzyme aggregates (CLEAs) are formed by precipitation of the enzyme from aqueous buffer, without perturbation of their tertiary structure, and cross-linking with a bifunctional reagent, such as glutaraldehyde.<sup>62</sup> Immobilised enzymes are more robust and can be recycled using filtration or centrifugation. Magnetic recovery is also possible if a magnetisable feature is introduced into the support or the cross-linking procedure.<sup>63</sup> Whole cell biocatalysts have the advantage that enzymes are immobilised within the cell wall and the cells can be recovered by centrifugation. Enzymes can also be made more robust through protein engineering techniques,<sup>64</sup> such as directed evolution, sometimes with dramatic

improvements. For example, carbonic anhydrase was evolved to tolerate temperatures up to 107 °C at pH > 10, with a  $4 \times 10^6$  improvement in stability over the wild-type enzyme.<sup>65</sup>

## 5. Stoichiometric and chemocatalytic vs. biocatalytic processes in organic synthesis

In order to be green and sustainable, organic syntheses should generate minimum waste and avoid the use of toxic and/or hazardous materials.<sup>66</sup> Major causes of waste in organic syntheses are the use of stoichiometric inorganic or organic reagents and solvents in multi-step syntheses. Hence, in order to minimise waste it is necessary to use step-<sup>67</sup> and atom-economic,<sup>68</sup> catalytic methods, preferably under solvent-free conditions. Where use of a solvent cannot be avoided it should involve limited amounts of green, environmentally acceptable solvents.<sup>69</sup>

Enzymatic processes are green and sustainable<sup>50</sup> conforming to 10 of the 12 principles of green chemistry.<sup>70</sup> The catalyst is produced from renewable biomass and is biocompatible, biodegradable and essentially non-hazardous and non-toxic. It provides an attractive alternative to the use of scarce precious metals as catalysts and avoids the costs of removing traces of noble metals from end products. Enzymatic reactions generally avoid the need for functional-group activation, protection, and deprotection steps, affording routes that are more step-economic and generate less waste than conventional organic syntheses. Moreover, they are conducted in standard reactors at close to ambient temperatures and pressures in water, rendering them both environmentally attractive and cost-effective. Furthermore, enzymatic processes are generally highly selective: distinguishing between regio- and stereoisomers in addition to various functional groups (chemo-selective). In particular enzymatic processes have become the method of choice in the synthesis of enantiomerically pure pharma intermediates based on the magnificent, close to absolute enantioselectivities that are obtainable with highly engineered enzymes.<sup>71–74</sup>

### 5.1 Hydrolytic processes

Traditionally, enzymatic processes were used industrially in the hydrolysis and formation of, *inter alia*, ester, amide and glycoside bonds, especially where very high selectivities were required (chemo-, regio- or stereo-selectivity). Typical examples include the processing of fats and oils and the synthesis of beta-lactam antibiotics.

Enzymatic processes have the advantage of producing substantially less waste compared with the use of mineral acids or metal compounds as catalysts. A prime example, involving the production of a commodity chemical, is the Mitsubishi process for the hydrolysis of acrylonitrile to acrylamide catalysed by a nitrile hydratase (NHase) from the bacterium *Rhodococcus rhodochrous* J1 (Scheme 1a).

The process, originally developed in the 1980s,<sup>75</sup> currently accounts for the global production of several hundred thousand



## a. Mitsubishi process for acrylamide



## b. Lonza process for nicotinamide (vitamin B3)



## c. Dupont process for 5-cyanovaleramide



Scheme 1 Nitrile hydratase catalysed hydrations of nitriles.

tons per annum of acrylamide. The latter is mainly applied in the production of polyacrylamide, which requires very pure monomer as starting material. Hence, an important advantage of the NHase process, compared to the copper-catalysed hydration that it replaced, is its close to perfect chemoselectivity (>99.99%). The NHase is used as whole (dead) cells because of the limited stability of the active enzyme which is a tetramer that readily dissociates outside the cell.

NHase-catalysed hydration of the corresponding nitrile is also used in the Lonza process for the industrial production of nicotinamide (vitamin B3)<sup>76</sup> where the enzymatic process (Scheme 1b) is more selective, with respect to competing formation of the corresponding acid, than chemocatalytic alternatives. Similarly, 5-cyanovaleramide, a herbicide intermediate is produced by DuPont *via* regioselective NHase catalysed hydration of adiponitrile (Scheme 1c),<sup>77</sup> generating fewer byproducts and less waste than processes using metal catalysts such as RANEY®Cu or MnO<sub>2</sub>.

Other well-known commercial applications of hydrolases include the use of the protease thermolysin for synthesis of the artificial sweetener aspartame,<sup>78</sup> penicillin G amidase/acylase (E.C. 3.5.1.11) for synthesis of semi-synthetic β-lactam antibiotics,<sup>79</sup> and amylase for the production of glucose *via* hydrolysis of starch.<sup>80</sup>

Pharmaceuticals tend to be lipophilic and therefore easy to handle in organic solvents<sup>81</sup> and lipases are enzymes that are particularly tolerant of hydrophobic environments.<sup>82</sup> Therefore, it is not surprising that pharmaceutical syntheses using biocatalysis have been dominated by lipase catalysed reactions.<sup>83–85</sup> Lipases perform hydrolysis and esterification reactions, generating alcohols, carboxylic acids and esters, often with high stereo-selectivity when required. Moreover, they are catalytically versatile and can catalyse reactions such as aminolysis (the conversion of esters to amides, using amines as the nucleophile) and direct amidation of carboxylic acids.<sup>86</sup>

An interesting recent development in reverse hydrolysis reactions, such as the formation of carboxylic esters, is the use

of the acyltransferase from *Mycobacterium smegmatis* (MsAcT) to catalyse the synthesis of flavour esters by transesterification in water.<sup>87</sup> MsAcT is characterised by a hydrophobic tunnel leading to the active site, where access of water is disfavoured. Hence, transesterifications are favoured over hydrolysis, even in water.<sup>88,89</sup>

In the case of chiral substrates, enzymes can be used to synthesise the pure enantiomers. In hydrolytic processes, such as ester and amide hydrolyses this generally involves a kinetic resolution that affords a maximum yield of 50%. The maximum yield of 50% of the unwanted enantiomer (the distomer) constitutes waste that has to be disposed of or recycled back to the racemate *via* racemisation. In some cases the racemisation step can be performed *in situ*, affording a so-called dynamic kinetic resolution.<sup>90,91</sup>

The application of lipases for enzymatic kinetic resolution (EKR) of chiral intermediates in commercial syntheses includes pregabalin<sup>92</sup> (Scheme 2a) and (*R*)-1-phenylethylamine<sup>93</sup> (Scheme 2b) both of which involved the use of commercially available lipases. Hence, hydrolases such as lipases can now be considered mainstream organic synthetic tools, and their popularity continues unabated.<sup>94–96</sup> Research is being directed increasingly to adapting the enzymes for industrial applications.<sup>53,97</sup>

Up until fairly recently it was generally thought that kinetic resolutions would be dominated by biocatalytic processes and asymmetric syntheses would be largely the domain of precious metal complexes with chiral ligands as catalysts in reactions such as asymmetric hydrogenations and hydroformylations. This situation has changed in the last two decades, resulting in

## a. Chemoenzymatic manufacturing process for pregabalin by Pfizer



## b. Lipase-catalysed resolution by enantioselective amide formation by BASF



Scheme 2 Enzymatic kinetic resolutions using lipases.



a decrease in the use of lipases for kinetic resolutions<sup>98</sup> and the broadening application of enzymatic redox processes in industrial organic synthesis as reflected in patent applications by industry.<sup>99–102</sup> Indeed, in one study of 547 biocatalysis based patents filed between 2000 and 2015, only 10% involved hydrolases, while 68% were based on oxidoreductases.<sup>103</sup> Hence, in the next section we will focus on recent developments in oxidoreductases.

## 5.2 Redox processes

**5.2.1 Functional group reductions.** Many pharmaceutical intermediates are chiral secondary alcohols or chiral primary amines that can be synthesised by asymmetric reduction or reductive amination of the corresponding ketones (Scheme 3).

In the 1990s chiral alcohols were typically produced either by stoichiometric reduction of ketones with chiral boron reagents or asymmetric hydrogenation catalysed by homogeneous chiral complexes of ruthenium and other precious metals.<sup>104</sup> Ketoreductases (KREDs), also known as alcohol dehydrogenases (ADHs) catalyse oxidative dehydrogenation of (secondary) alcohols and the reverse reaction: the enantioselective reduction of ketones. The reducing equivalents are provided by stoichiometric amounts of nicotinamide cofactors, NADH or NADPH and for industrial viability these need to be recycled *in situ*. This is achieved by adding a large excess of an alcohol co-substrate, such as isopropanol, or a second enzyme in combination with an inexpensive co-substrate (Scheme 4). The most used combinations are formate/formate dehydrogenase (FDH) and glucose/glucose dehydrogenase (GDH).

Thanks to key advances in protein engineering by directed evolution, together with gene sequencing and DNA synthesis, KREDs have been successfully applied to the synthesis of a variety of enantiomerically pure secondary alcohols as key intermediates for a variety of chiral drugs,<sup>105,106</sup> including atorvastatin,<sup>107</sup> montelukast,<sup>108</sup> ezetimibe, sulopenem and licarbazepine (Scheme 5).

A widely used method for producing amines involves chemocatalytic reductive amination of ketones with a mixture of hydrogen and ammonia. This suggests that what was missing from the enzyme toolbox was an asymmetric reductive amination of prochiral ketones with ammonia catalysed by NAD(P)H-dependent amine dehydrogenases (AmDHs). Such enzymes are rare *in vivo* where amine catabolism involves aerobic oxidation catalysed by amine oxidases. Amino acid dehydrogenases, in contrast, are well-known and form an obvious starting point for the development of AmDHs by directed evolution.

Bommarius and co-workers<sup>109</sup> used leucine dehydrogenase (LeuDH) as a protein scaffold for developing an enzymatic reductive amination of structurally related methyl isobutyl



Scheme 4 KRED catalysed reduction of prochiral ketones.



Scheme 5 Examples of enantiomerically pure secondary alcohols produced with engineered KREDs.

ketone. Similarly, phenylalanine dehydrogenase (PheDH) was used as a starting point for engineering a highly active and highly selective (>99.8% ee) AmDH for both aliphatic and benzylic ketones<sup>110</sup> in combination with glucose/GDH for cofactor regeneration (Scheme 6). The same group subsequently used domain shuffling of the two parent AmDHs to produce a novel chimeric AmDH, with altered substrate scope. Further practical improvements were realised using a combination of ammonium formate and formate dehydrogenase (FDH) as both a source of ammonia and cofactor regeneration in a biphasic organic solvent/water system.<sup>111</sup> Mutti and co-workers<sup>112</sup> similarly used three different AmDHs, in combination with formate and FDH, for the reductive amination of a wide variety of prochiral ketones, including aromatic, benzylic, aliphatic and ketones with bulky substituents. They further suggested that practical utility could be improved by co-immobilisation of the two enzymes or co-expression in a single microorganism. Likewise, Li and co-workers<sup>113</sup> developed an AmDH for the reductive amination of phenylacetone and 4-phenyl-2-butanone by protein engineering of PheDH (Scheme 6).

Another fairly recent development is the emergence of imine reductases (IREDs)<sup>114–119</sup> for the enantioselective reduction of cyclic imines derived from secondary amines and the asymmetric reductive amination of ketones with primary amines (Scheme 7).<sup>120–123</sup>

The concept of chiral amine synthesis *via* enzymatic reductive amination was taken to a new level of sophistication by



Scheme 3 Reduction and reductive amination of ketones.





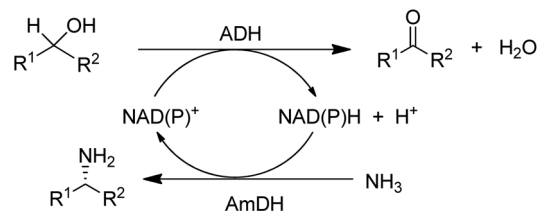
Scheme 6 Enzymatic primary amine synthesis with amine dehydrogenases.



Scheme 7 Imine reductases and amine dehydrogenase for reductive aminations with amines.

applying the concept of hydrogen borrowing (itself having been borrowed from the chemocatalytic literature).<sup>124</sup> For example, the combination of an ADH and an AmDH catalysed a redox-neutral conversion of a racemic alcohol to a single enantiomer of the corresponding amine (Scheme 8).<sup>125</sup> Ironically, the success of this method depends on the ADH being aselective<sup>126</sup> as it needs to catalyse the oxidation of both enantiomers of the alcohol, which is not a simple task as most ADHs are highly enantioselective. The overall efficiency of the process could be improved by co-immobilisation of the ADH and AmDH.<sup>127</sup> Overall the reaction constitutes a nucleophilic substitution of a hydroxyl group in a secondary alcohol, in this case by an amino group, with an equivalent of water as the sole co-product. Such a reaction is one of the nine reactions in a recently published list of 10 key research areas for a greening of the pharmaceutical industry.<sup>128</sup>

**5.2.2 Functional group oxidations.** The selective oxidation of primary and secondary alcohols to the corresponding



Overall:



Scheme 8 Enantioselective conversion of prochiral alcohols to chiral amines via hydrogen borrowing.

aldehydes and ketones is a key reaction in organic synthesis. Such reactions were traditionally carried out with stoichiometric amounts of inorganic oxidants, such as chromium VI reagents, but efficient chemocatalytic<sup>129–134</sup> and biocatalytic methods<sup>135</sup> have gained in importance as a direct result of the greening of the chemical industry.

The most commonly used enzymes for the oxidation of alcohols are alcohol oxidases (AOx) that use dioxygen as the oxidant and form H<sub>2</sub>O<sub>2</sub> as the co-product. In practice, catalase is added in order to rapidly remove the H<sub>2</sub>O<sub>2</sub> to avoid degradation of the enzyme. They are involved in carbohydrate oxidations where they tend to be very substrate specific. Examples include galactose oxidase (GOase) and glucose oxidase (GOX) which are copper and flavin-dependent, respectively.

In the last decade many new flavin- and Cu-dependent AOxs, with altered substrate specificities and stabilities have been developed with the aid of genome mining and directed evolution. Examples include GOase variants able to catalyse the oxidation of secondary alcohols,<sup>136</sup> amino alcohols<sup>137</sup> and 5-hydroxymethyl-furfural<sup>138</sup> and a choline oxidase which was evolved to become a broad spectrum primary alcohol oxidase (Scheme 9).

A variant of a flavin-dependent oxidase catalysed the oxidation of 5-hydroxymethylfurfural to 2,5-furandicarboxylic acid (FDCA), involving two alcohol oxidation steps and one aldehyde oxidation (Scheme 10).<sup>139</sup> Since many aldehydes are present in water mainly as the hydrate (gem-diol), it is not so surprising



Scheme 9 An evolved choline oxidase as a broad spectrum primary alcohol oxidase.





Scheme 10 HMFA oxidation with flavin dependent AOx.

that an alcohol oxidase can also catalyse the aerobic oxidation of aldehydes.

The above-mentioned variant of GOase also catalysed the ammoxidation of primary alcohols<sup>140</sup> to nitriles (Scheme 11).

Alternatively, a combination of the Cu-dependent oxidase, laccase, with stable nitroxyl radicals, such as TEMPO, catalyses the aerobic oxidation of primary alcohols to the corresponding aldehydes.<sup>141</sup> It was also effective in the oxidation of the secondary alcohol isosorbide to the corresponding diketone (Scheme 12).<sup>142</sup>

Alcohol dehydrogenases (ADHs) catalyse the oxidation of alcohols and require regeneration of the nicotinamide cofactor using an excess of a co-substrate.<sup>143–145</sup> Alternatively, NAD(P)H oxidase (NOx) can be employed to catalyse reoxidation of the cofactor by dioxygen.<sup>146,147</sup>

NAD-dependent aldehyde dehydrogenase (ALDH) catalyses the aerobic oxidation of aldehydes to the corresponding carboxylic acids and three recombinant ALDHs have been used, in combination with NOx for cofactor recycle, for the highly selective aerobic oxidation of a wide variety of aldehydes.<sup>148</sup> Combination of an ALDH with an ene-reductase (ERED) afforded a hydrogen-borrowing strategy for the conversion of  $\alpha,\beta$ -unsaturated aldehydes to the corresponding  $\alpha$ -substituted carboxylic acids in high chemo- and stereoselectivities (Scheme 13).<sup>149</sup>

Baeyer-Villiger Monooxygenases (BVMOs) are synthetically interesting flavin-dependent enzymes that catalyse the aerobic



Scheme 11 AOx catalysed ammoxidation of an alcohol.



Scheme 12 Laccase/TEMPO catalysed oxidations of alcohols.



Scheme 13 Combination of an ALDH with an ERED.

oxidation of ketones and cyclic ketones to esters,<sup>150,151</sup> and lactones, respectively. They can be combined with ADHs, for example as a fused ADH–BVMO,<sup>152,153</sup> to afford hydrogen-borrowing cascades for conversion of an alcohol to an ester or a smart synthesis of caprolactone from 1,6-hexanediol.<sup>154</sup> Similarly, an elegant redox neutral process<sup>155</sup> for highly



Scheme 14 Enantioselective sulfoxidation using an ADH/BVMO cascade.





Scheme 15 Desymmetrisation of a prochiral amine with an MAO.

enantioselective sulfoxidations with dioxygen was designed by combining an evolved cyclohexanone monooxygenase (CHMO) with a KRED (Scheme 14). It was used in the production of, for example, the single enantiomer drug esomeprazole (*S*-omeprazole).<sup>156</sup>

Flavin-dependent monoamine oxidases (MAOs) catalyse the aerobic oxidation of amines to the corresponding imines.<sup>157</sup> Turner and coworkers<sup>158</sup> used directed evolution to develop an MAO for the deracemisation of structurally diverse primary, secondary, and tertiary amines. The process is a kinetic resolution but can afford high yields of one enantiomer by combining it with repeated aselektive chemical reduction of the imine. MAO variants were also used for the enantioselective desymmetrisation of prochiral amines, *e.g.* in the synthesis of the hepatitis C drug, Boceprevir<sup>159</sup> (Scheme 15).

### 5.3 C–O, C–N and C–C bond forming reactions

The reactions discussed in the preceding two sections involved functional group transformations *via* hydrolytic or redox processes. In this section we shall discuss the formation of C–O and C–N containing functional groups by enzymatic conversion of hydrocarbons and the formation of C–C bonds.

**5.3.1 C–O bond formation.** Biocatalytic oxidations with O<sub>2</sub> or H<sub>2</sub>O<sub>2</sub> fall into two categories: dehydrogenations and

Scheme 17 Olefin *cis*-dihydroxylation catalysed by a dioxygenase.

oxyfunctionalisations.<sup>160</sup> The former mainly involve oxidation of functional groups, *e.g.* alcohols and amines (see preceding section). Oxyfunctionalisation, in contrast, is concerned with insertion of oxygen into hydrocarbon C–H or C–C bonds to form new functional groups. It involves reductive activation of dioxygen by *e.g.* cytochrome P450 monooxygenases, or the use of hydrogen peroxide with peroxygenases (E.C. 1.11.2.), to form activated electrophilic oxygen species such as high-valent iron oxo complexes (Scheme 16).<sup>161</sup>

Mono- and dioxygenases catalyse the formation of new C–O bonds by the oxidation of hydrocarbons by dioxygen. Cyt-P450-dependent monooxygenases, for example, are multicomponent biocatalysts in which reducing equivalents are supplied by NAD(P)H indirectly through a flavin or ferredoxin mediator (see Scheme 16). They catalyse a broad range of regio- and stereoselective hydroxylations of alkanes, alkenes and aromatic hydrocarbons employing dioxygen as the terminal oxidant, accounting for about 95% of all enzyme catalysed oxidations.<sup>162</sup>

Catalytically self-sufficient P450s containing the P450 and reductase domains in the same polypeptide chain, such as P450<sub>BM3</sub> from *Bacillus megaterium*,<sup>163</sup> are of particular interest in this context and a panel of new thermostable, self-sufficient P450s with broad substrate scope from thermophilic organisms has been reported.<sup>164</sup> However, the best result – 1.6 g L<sup>-1</sup> product concentration after 10 h in the hydroxylation of diclofenac – is still 1–2 orders of magnitude lower than what is needed for industrial viability but can be seen as a starting point for further engineering by directed evolution.<sup>165</sup>

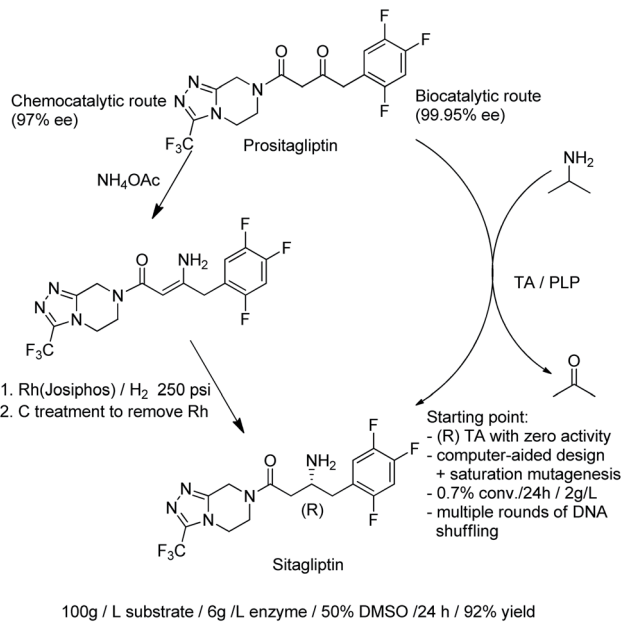
In addition to catalysing oxygen insertion into C–H bonds of alkanes and alkylaromatics and allylic C–H bonds in olefins, Cyt P450s<sup>166</sup> and peroxygenases<sup>167</sup> catalyse asymmetric epoxidations and an engineered non-haem iron dependent Rieske dioxygenase (RO) catalysed *cis*-dihydroxylations of olefins (Scheme 17).<sup>168</sup> The introduction of a single point mutation in the dioxygenase improved conversions (>99%) and selectivities (>95%). The method constitutes a greener, more sustainable alternative to the use of toxic osmium catalysts in the Sharpless *cis*-dihydroxylation.<sup>169</sup> Despite the synthetic potential of these transformations their reduction to commercial practice remains a significant challenge<sup>170</sup> as they are generally characterised by low product concentrations and space time yields. Although replacing dioxygen and oxygenases by hydrogen peroxide with peroxygenases<sup>171</sup> circumvents the need for cofactors it is also fraught with difficulties, such as low stabilities of the enzymes towards the hydrogen peroxide oxidant. The methods seem to be *en route* but have not yet reached their destination.

**5.3.2 C–N bond formations.** The direct conversion of alcohols to primary amines using an AmDH/ADH combination in a hydrogen borrowing strategy (see earlier) is an alternative to



Scheme 16 Monooxygenase and peroxygenase pathways.





Scheme 18 Synthesis of sitagliptin.

the use of  $\omega$ -transaminases (TAMs). The latter catalyse the transamination of prochiral ketones using an excess of a second amine,<sup>172</sup> usually isopropylamine, analogous to the use of isopropanol in KRED catalysed reduction of ketones. TAMs were the first enzymes to have their enantioselectivities optimised using directed evolution<sup>173</sup> and in the last decade they have emerged as excellent catalysts for the synthesis of enantiopure primary amines.<sup>35,174,175</sup> A commercially important example is the synthesis of a key intermediate for sitagliptin, the active ingredient of the antidiabetic agent, Januvia®. A highly engineered, (*R*)-selective  $\omega$ -TAM variant was developed, using a combination of protein engineering and computer-aided design, to enable conversion of the ketone precursor, prositagliptin to sitagliptin. The process replaced a rhodium-catalysed asymmetric hydrogenation (Scheme 18).<sup>176</sup>

Protein engineering was also used to develop highly active, (*S*)-selective TAMs for the asymmetric synthesis of bulky chiral amines in up to >99.9% ee.<sup>177</sup> A disadvantage of TAMs is that several equivalents of the sacrificial amine are generally required to drive the unfavourable equilibrium. Interestingly, robust TAMs, requiring less equivalents of isopropylamine, were recently developed from a domestic drain metagenome (see Section 2).

**5.3.3 C–C bond formations.** A wide variety of lyases, including hydroxynitrile lyases (HNLs; EC 4.1.2.46 and 4.1.2.47), a variety of aldolases, transketolases and carboli-gases, are able to catalyse the enantioselective formation of C–C bonds.<sup>178</sup> The main competition is from organocatalysis, in particular catalysis by the amino acid L-proline and derivatives thereof, which have also been shown to catalyse a wide variety of C–C bond forming reactions.<sup>179,180</sup> HNLs are robust enzymes that perform well in organic solvents or aqueous-organic biphasic systems. They catalyse the formation of  $\alpha$ -

## a. Enantioselective hydrocyanation



## b. DERA catalysed aldol reaction



Scheme 19 Enzymatic C–C bond forming reactions.

hydroxynitriles (cyanohydrins) by enantioselective addition of hydrogen cyanide to aldehydes and ketones.<sup>181</sup> Both *R*- and *S*-selective HNLs are widely distributed in nature and have been successfully applied in the industrial synthesis of a variety of products,<sup>182</sup> e.g. the *R*-cyanohydrin of 2-chlorobenzaldehyde, a pharmaceutical intermediate, produced by enantioselective hydrocyanation catalysed by an *R*-HNL CLEA (Scheme 19a).<sup>183</sup>

Among the many types of aldolases, 2-deoxy-D-ribose-5-phosphate aldolase (DERA) is of particular interest. *In vivo* it catalyses the synthesis of 2-deoxy-D-ribose-5-phosphate from acetaldehyde and glyceraldehyde-3-phosphate but *in vitro* it also catalyses the aldol reaction of acetaldehyde with numerous other simple aldehydes.<sup>184</sup> Its synthetic potential is limited by its low tolerance to industrially relevant aldehyde concentrations but this hurdle has been overcome by protein engineering. DSM, for example, developed a process for the synthesis of an advanced atorvastatin intermediate by aldol reaction of two equivalents of acetaldehyde with chloroacetaldehyde (Scheme 19b) that operates at 100 g L<sup>-1</sup> substrate concentration.<sup>185</sup>

**5.3.4 The multifaceted nature of biocatalysis.** As discussed in the preceding sections, industrially viable biocatalytic

## a. Hydrolase route



## b. Oxidoreductase route



## c. Lyase route



Scheme 20 Competing biocatalytic routes to atorvastatin intermediates.



methodologies are currently available for a wide variety of synthetic transformations. This enables the synthesis of target molecules *via* different approaches. A good example of this is the synthesis of atorvastatin in which hydrolase-, oxidoreductase- and lyase-based routes have all been developed (Scheme 20). In both the hydrolase and the oxidoreductase routes the C<sub>5</sub> intermediate (I) is formed and subsequently converted to the advanced C<sub>7</sub> intermediate (II). In contrast, the lyase-based route affords (II) directly.

## 6. Integration: biocatalytic and chemoenzymatic cascade processes

The 8<sup>th</sup> principle of Green Chemistry is reduction in the number of steps through reduced derivatisation. The high specificity of enzymes obviates the need for protection and deprotection steps.<sup>62</sup> Examples of the successful integration of biocatalysis steps into a traditional synthetic process are many, including diltiazem,<sup>186</sup> sitagliptin,<sup>173</sup> pregabalin,<sup>89</sup> atorvastatin<sup>104,187</sup> and other statins,<sup>188</sup> to name but a few.

In traditional multistep organic syntheses a simple step-by-step approach is used whereby intermediates are isolated and often purified before proceeding to the next step. The result is: low volumetric productivities, protracted recycle loops and considerable waste generation. Hence, integration of several catalytic steps into step economic, one-pot procedures, with no isolation of intermediates, will automatically lead to less waste formation. This telescoping of multistep syntheses has various practical advantages: fewer unit operations, smaller reactor volume, shorter cycle times, higher productivities and less waste.

This is nothing special in nature where cellular metabolism requires a vast array of enzymes working together in multi-step reactions in a single “pot” to make a multitude of products with high specificity and negligible accumulation of intermediates.

With a rapidly expanding toolbox of biocatalytic conversions, which proceed under approximately the same conditions – in water at roughly ambient temperature and pressure – the construction of enzymatic cascade reactions has become increasingly popular.<sup>44,189–194</sup> It provides tremendous advantages to chemists as the effort and yield losses associated with work-up after each step are eliminated, increasing the overall yield. Moreover, it can be used to drive equilibria towards product formation. For example, in a 3 step reaction involving enzymatic hydrocyanation of a ketone followed by two sequential enzymatic hydrolytic steps the equilibrium of the reversible hydrocyanation step was shifted towards product by its removal in the hydrolysis step.<sup>195</sup>

The outstanding benefits of biocatalytic cascades are illustrated in a synthesis of islatravir (an HIV reverse transcriptase translocation inhibitor), recently developed in a collaboration between Codexis and Merck (Scheme 21). The three-step biocatalytic cascade involves nine enzymes in a single aqueous solution without the need for isolation of intermediates. Four of the enzymes were ancillary, including sucrose phosphorylase, which catalysed a reaction



Scheme 21 Biocatalytic cascade synthetic route to islatravir.

incorporated into the cascade to shift the entire equilibrium forward. This route halved the total number of reaction steps required, greatly improved atom economy, and permitted an excellent overall islatravir yield of 51% from simple building blocks.<sup>196</sup>

The majority of biocatalytic cascade processes involve redox reactions, where reactions are coupled to integrate the use and regeneration of nicotinamide cofactors (NADH and NADPH) thereby making the overall process redox neutral. We have already discussed striking examples of such hydrogen borrowing in the preceding section.

Catalytic cascade processes can also involve the integration of biocatalytic and chemocatalytic steps to afford chemoenzymatic cascade processes.<sup>197–199</sup> Chemoenzymatic approaches have also proved useful in the synthesis of natural products.<sup>200–203</sup> A recent important publication by Lipshutz and coworkers has shown that the formation of micelles in the reaction medium can facilitate the combination of chemo- and bio-catalysis in a single reactor as the substrates, products and catalysts partition into the different compartments.<sup>204</sup> For an extensive review of the recent literature on biocatalytic cascade reactions see Kroutil and coworkers.<sup>205</sup>

## 7. New to nature reactions

Continued expansion of the biocatalysis toolbox for synthetic chemists is being made possible by the discovery of new enzymes with new functions,<sup>7,206</sup> as well as by repurposing of existing enzymes to catalyse reactions not observed in nature. These so-called “new-to-nature” reactions can arise naturally<sup>207,208</sup> but have been driven, in particular, by directed evolution.<sup>26,209</sup> This approach relies on enzyme catalytic promiscuity, the ability of an enzyme to catalyse more than one distinct type of chemical reaction within the same active site.<sup>210</sup> This alternative chemistry that is available to the enzyme is often present naturally at a very low level, but can be selected for and enhanced in the process of directed evolution.<sup>211</sup> An area also receiving considerable attention is the use of artificial enzymes for conducting new-to-nature reactions.<sup>212,213</sup>



Scheme 22 Stereoselective cyclopropanation reaction of P450<sub>BM3</sub>.

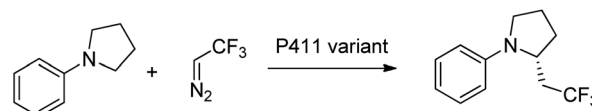
### 7.1 Haem metalloenzyme-mediated reactions

Cytochrome P450 monooxygenases catalyse a variety of oxidations of hydrocarbons with dioxygen whereby an oxygen atom is inserted into a C–H bond or a C–C double bond of an olefin. The key step in the latter reactions involves oxygen transfer from a high-valent oxoiron porphyrin complex to the double bond of the olefin, affording an epoxide. By analogy with this oxene transfer, it was shown that P450<sub>BM3</sub>, under anaerobic conditions, could catalyse carbene<sup>214</sup> and nitrene transfer reactions, to afford cyclopropanes and aziridines, respectively. For example, cyclopropane derivatives were formed by reaction of styrene with ethyl diazoacetate.<sup>215</sup> Depending on the mutation(s) introduced, either the *cis* or *trans* product could be obtained in high enantioselectivity (Scheme 22). In a similar manner, modified myoglobin, another haem protein, was also found to be an effective cyclopropanation catalyst.<sup>216</sup> The synthesis of a number of drug precursors containing the cyclopropane moiety was demonstrated at preparative scale using these engineered haem proteins.<sup>217</sup> In a variation on this theme, Hartwig and co-workers used an artificial iridium porphyrin-containing P450 enzyme optimised through directed evolution for alkene cyclopropanation.<sup>218</sup>

Similarly, enantioselective aziridination of styrene derivatives by reaction with tosyl azide was achieved using a P411 variant of cytochrome P450<sub>BM3</sub>.<sup>219</sup> Arnold and co-workers also exploited the nitrene-transfer capacity of a P450 enzyme variant



Scheme 24 Olefination of aromatic aldehydes.



Scheme 25 C–H fluoroalkylation of amines.

to catalyse intramolecular benzylic C–H amination,<sup>220</sup> a reaction first reported with wild type Cyt P450 by Breslow and co-workers in 1985.<sup>221</sup> Arnold *et al.* followed up this work with a demonstration of regioselective C–H amination either at the benzylic position or the homo-benzylic position (Scheme 23a).<sup>222</sup>

Similarly, engineered Cyt P450s catalysed intermolecular benzylic C–H amination *via* the insertion of nitrenoid species, derived from tosyl azide, into unactivated C–H bonds<sup>223</sup> to afford amines (Scheme 23b). This reaction has been extended to allow introduction of the NHTs group to position C-2 of *N*-methylindole.<sup>224</sup>

The utility of modified myoglobins was also expanded to include catalysis of olefination reactions of aromatic aldehydes using diazo acetate reagents (Scheme 24).<sup>225</sup>

Enzymatic carbene transfer reactions mediated by CytP450 mutants were also recently demonstrated with heterocyclic substrates such as indoles and pyrroles<sup>226</sup> and in the enantioselective C–H fluoroalkylation at the  $\alpha$ -C–H bond of amines (Scheme 25).<sup>227</sup>

Notwithstanding the elegance of such new-to-nature carbene and nitrene insertion reactions, it is worth noting that the reactions shown here are not possible in nature because of the absence of diazo esters and tosyl azide. However, it is interesting to note that Ohnishi *et al.* have recently proposed an *in vivo* nitrene transfer mechanism, involving the formation of an iron nitrenoid intermediate by cytochrome P450 BeZ catalysed elimination of acetic acid from an *O*-acetyl hydroxylamine functionality, to explain an intramolecular amination reaction.<sup>228</sup> The authors suggested the name P450 nitrene transferase for BeZ.

Other examples of non-natural reactions catalysed by engineered haem proteins can be found in recent reviews.<sup>26,229</sup>

### 7.2 Reactions catalysed by promiscuous hydrolases

A number of hydrolases, such as lipases, have been demonstrated to catalyse a wide range of reactions that do not occur *in vivo*.<sup>230,231</sup>

**7.2.1 Addition reactions.** Enzymatic Michael addition reactions, catalysed by a variety of hydrolases, have been widely reported with N and S nucleophiles.<sup>232</sup> Use of carbon nucleophiles in this reaction, resulting in C–C bond formation, was

#### a. Intramolecular C–H amination



#### b. Intermolecular C–H amination



Scheme 23 C–H aminations catalysed by Cyt P450s.





Scheme 26 Addition of thiols to vinyl esters.



Scheme 27 Lipase-catalysed Mannich reaction.

first reported by Berglund and co-workers using a Ser 105 Ala mutant of CalB.<sup>233</sup> They went on to show that this mutant enzyme could catalyse the Michael addition reaction between acetylacetone and methyl acrylate in aqueous medium, albeit in very low yield.<sup>234</sup> The mutant, lacking the nucleophilic serine residue in the active site, was unable to catalyse the natural hydrolysis reaction, but instead gave the Michael addition product, methyl 4-acetyl-5-oxohexanoate.

The lipase-catalysed addition of thiols to vinyl esters was achieved by Lin *et al.* in organic solvent, and depending on the choice of solvent either Markovnikov or anti-Markovnikov addition was observed (Scheme 26).<sup>235</sup>

**7.2.2 Aldol reactions.** The aldol reaction is catalysed *in vivo* by aldolases, but it is also catalysed by promiscuous lipases.<sup>236</sup> However, the lipase-catalysed reactions are not carried out with a high degree of enantioselectivity, with the best result to date being obtained by Yu and co-workers, who obtained the aldol product in 44% ee but low yield.<sup>237</sup> An extension of these hydrolase-catalysed aldol reactions to include the non-natural versions such as the Henry reaction has been achieved by He and co-workers in good yield using transglutaminase (TGase) from *Streptococcus thermophilus*.<sup>238</sup>

**7.2.3 Mannich reaction.** Yu and co-workers also reported the first lipase-catalysed Mannich reaction and they found that the best product yields were obtained using water (40–50%) as a co-solvent.<sup>239</sup> The reaction of various aromatic aldehydes with aniline and acetone (which played the dual role of reactant and solvent) proceeded in good yield when using the lipase from *Mucor miehei* (Scheme 27).

Further examples of non-natural conversions catalysed by promiscuous hydrolases can be found in a number of reviews.<sup>240–242</sup>

### 7.3 Other reactions

An artificial “metathase” was prepared by Ward and co-workers by the combination of streptavidin with a biotinylated Grubbs–Hoveyda metathesis catalyst as the cofactor, and ring-closing

metathesis was demonstrated *in cellulo*.<sup>243</sup> Siegel *et al.* designed artificial enzymes able to catalyse an intermolecular Diels–Alder reaction stereoselectively.<sup>244</sup> Photoexcitation has been demonstrated by Hyster *et al.* as a successful approach to accessing new non-natural oxidoreductase activity.<sup>245,246</sup> It is evident that the range of biocatalytic reactions useful to organic chemists is increasing by leaps and bounds.

## 8. Conclusions and future prospects

The phenomenal advances in new enzyme discovery, through the fruitful combination of metagenomics and bioinformatics for collection and analysis of gene sequences and directed evolution for enzyme improvement have completely revolutionised biocatalysis in the last two decades. Biocatalysis has been promoted to a method of choice in mainstream organic synthesis. Methodology development continues to play a pivotal role in the directed evolution of chemo-, regio- and stereoselective enzymes<sup>247</sup> and, since directed evolution produces huge amounts of data, it is ideally suited to benefit from machine learning.<sup>248–250</sup> Similarly, computer-aided retrosynthetic planning of biocatalytic routes can be used effectively to design new enzymatic processes.<sup>251</sup>

Another important technological development is the use of flow chemistry and microfluidics.<sup>252–255</sup> This will facilitate the design of biocatalytic cascade reactions as these can easily be performed in series in a flow reactor. Moreover, the protective and low shear environment of micro-reactors will mean that less effort will be required for increasing the robustness of the enzymes.

Other emerging technologies will help to drive the future of biocatalysis. Synthetic biology, for example, has advanced to the point where defined microorganisms can be composed from scratch. This permits the production of better defined biocatalysts with reduced contamination by other enzymes.<sup>256</sup> It has also allowed the redundancy in the gene code to be rationalised to provide dedicated codes for non-canonical amino acids,<sup>257</sup> the potential contribution of which to new biocatalysis was mentioned above.

Good primers on the subject of biocatalysis include books by Grogan<sup>258</sup> and Faber,<sup>259</sup> while a more comprehensive contemporary review has been compiled by Faber, Fessner and Turner<sup>260</sup> and Whittall, Sutton and Kroutil.<sup>261</sup> The excellent paper by Lin *et al.*<sup>262</sup> provides a convenient correlation of how enzyme catalysed reactions correspond with named organic chemical reactions that organic chemists are familiar with.

In conclusion, enzymes, long used in diverse large scale industries, are increasingly being applied in all areas of synthetic organic chemistry. If you are a synthetic chemist who is new to the application of enzymes in your research, *Don't Panic*, once you become familiar with biocatalysts you will find them indispensable.

## Conflicts of interest

The authors have no conflicts of interest.



## Acknowledgements

We would like to acknowledge Douglas Adams, the late acclaimed author, for inspiration, 40 years after the publication of the Hitchhiker's Guide to the Galaxy.

## References

- 1 L. Esquirol, T. S. Peat, M. Wilding, J.-W. Liu, N. G. French, C. J. Hartley, H. Onagi, T. Nebl, C. J. Easton, J. Newman and C. Scott, *J. Biol. Chem.*, 2018, **293**, 7880–7891.
- 2 M. A. F. Delgove, A. B. Laurent, J. M. Woodley, K. V. Bernaerts and Y. van der Meer, *ChemSusChem*, 2019, **12**, 1349–1360.
- 3 A. Schmid, J. S. Dordick, B. Hauer, A. Kiener, M. Wubbolts and B. Witholt, *Nature*, 2001, **409**, 258–268.
- 4 A. J. J. Straathof, S. Panke and A. Schmid, *Curr. Opin. Biotechnol.*, 2002, **13**, 548–556.
- 5 R. A. Sheldon and D. Brady, *Chem. Commun.*, 2018, **54**, 6088–6104.
- 6 J. M. Woodley, *Appl. Microbiol. Biotechnol.*, 2019, **103**, 4733–4739.
- 7 P. N. Devine, R. M. Howard, R. Kumar, M. P. Thompson, M. D. Truppo and N. J. Turner, *Nat. Rev. Chem.*, 2018, **2**, 409–421.
- 8 S. H. Bhosale, M. B. Rao and V. V. Deshpandi, *Microbiol. Rev.*, 1996, **60**, 280–300.
- 9 R. A. Sheldon and F. van Rantwijk, *Aust. J. Chem.*, 2004, **57**, 281.
- 10 A. T. Otori, F. G. Lobo, A. Carolina, G. do Amaral and C. de Souza de Oliveira, *J. Mol. Catal. B: Enzym.*, 2016, **127**, 93–97.
- 11 C. Sanfilippo, A. Paterna, D. M. Biondi and A. Patti, *Bioorg. Chem.*, 2019, **93**, 103325.
- 12 F. Garzón-Posse, L. Becerra-Figueroa, J. Hernández-Arias and D. Gamba-Sánchez, *Molecules*, 2018, **23**, 1241.
- 13 S. G. Burton, *Catal. Today*, 1994, **22**, 459–487.
- 14 C. C. C. R de Carvalho, *Microb. Biotechnol.*, 2017, **10**, 250–263.
- 15 J. A. Littlechild, *Archaea*, 2015, 147671.
- 16 L. Liu, H. Yang, H. D. Shin, R. R. Chen, J. Li, G. Du and J. Chen, *Bioengineered*, 2013, **4**, 212–223.
- 17 D. Tusé, T. Tu and K. A. McDonald, *BioMed Res. Int.*, 2014, 256135.
- 18 M. L. Bode, F. van Rantwijk and R. A. Sheldon, *Biotechnol. Bioeng.*, 2003, **84**, 710–713.
- 19 <https://blast.ncbi.nlm.nih.gov/Blast.cgi>last, accessed, 5 November 2019.
- 20 <https://www.uniprot.org/last>, accessed, 5 November 2019.
- 21 M. Tuffin, D. Anderson, C. Heath and D. A. Cowan, *Biotechnol. J.*, 2009, **4**, 1671–1683.
- 22 F. Berini, C. Casciello, G. L. Marcone and F. Marinelli, *FEMS Microbiol. Lett.*, 2017, **364**, fnx211.
- 23 K. Rashamuse, V. Magomani, T. Ronneburg and D. Brady, *Appl. Microbiol. Biotechnol.*, 2009, **83**, 491–501.
- 24 D. Baud, J. W. E. Jeffries, T. S. Moody, J. M. Ward and H. C. Hailes, *Green Chem.*, 2017, **19**, 1134–1143.
- 25 L. Leipold, D. Dobrijevic, J. W. E. Jeffries, M. Bawn, T. S. Moody, J. M. Ward and H. C. Hailes, *Green Chem.*, 2019, **21**, 75–86.
- 26 F. H. Arnold, *Angew. Chem., Int. Ed.*, 2018, **57**, 4143–4148.
- 27 U. T. Bornscheuer, B. Hauer, K. E. Jaeger and U. Schwaneberg, *Angew. Chem., Int. Ed.*, 2019, **58**, 36–40.
- 28 C. Zeymer and D. Hilvert, *Annu. Rev. Biochem.*, 2018, **87**, 131–157.
- 29 A. Tiessen, P. Pérez-Rodríguez and L. J. Delgado-Arredondo, *BMC Res. Notes*, 2012, **5**, 85.
- 30 L. H. Chen, G. I. Kenyon, F. Curtin, S. Harayama, M. E. G. Bembenek, C. P. Hajipour and C. P. Whitman, *J. Biol. Chem.*, 1992, **267**, 17716–17721.
- 31 A. J. M. Ribeiro, G. I. Holliday, N. Furnham, J. D. Tyzack, K. Ferris and J. M. Thornton, *Nucleic Acids Res.*, 2018, **46**(D1), D618–D623.
- 32 G. J. Bartlett, C. T. Porter, N. Borkakoti and J. M. Thornton, *J. Mol. Biol.*, 2002, **24**, 105–121.
- 33 L. Pravda, K. Berka, R. Svobodová Vařeková, D. Sehnal, P. Banáš, R. A. Laskowski, J. Koča and M. Otyepka, *BMC Bioinf.*, 2014, **15**, 379.
- 34 C. Cheng, T. Jiang, Y. Wu, L. Cui, S. Qin and B. He, *Bioengineered*, 2018, **119**, 1211–1217.
- 35 B. M. Nestl, S. C. Hammer, B. A. Nebel and B. Hauer, *Angew. Chem., Int. Ed.*, 2014, **53**, 3070–3095.
- 36 S. Kaushik, Z. Prokop, J. Damborsky and R. Chaloupkova, *FEBS J.*, 2017, **284**, 134–148.
- 37 J. D. Fischer, G. L. Holliday and J. M. Thornton, *Bioinformatics*, 2010, **26**, 2496–2497.
- 38 C. Andreini, I. Bertini, G. Cavallaro, G. L. Holliday and J. M. Thornton, *Bioinformatics*, 2009, **25**, 2088–2089.
- 39 C. Andreini, I. Bertini, G. Cavallaro, G. L. Holliday and J. M. Thornton, *J. Biol. Inorg. Chem.*, 2008, **13**, 1205–1218.
- 40 C. E. Paul and F. Hollmann, *Appl. Microbiol. Biotechnol.*, 2016, **100**, 4773–4778.
- 41 C. K. Prier and F. H. Arnold, *J. Am. Chem. Soc.*, 2015, **137**, 13992–14006.
- 42 C. Martinoli, H. M. Dudek, R. Orru, D. E. Edmondson, M. W. Fraaije and A. Mattevi, *ACS Catal.*, 2013, **3**, 3058–3062.
- 43 F. Agostini, J.-S. Völler, B. Koksche, C. G. Acevedo-Rocha, V. Kubyshekin and N. Budisa, *Angew. Chem., Int. Ed.*, 2017, **56**, 9680–9709.
- 44 A. M. Foley and A. R. Maguire, *Eur. J. Org. Chem.*, 2019, 3713–3734.
- 45 R. A. Sheldon and D. Brady, *ChemSusChem*, 2019, **17**, 2–25.
- 46 N. J. Turner and E. O'Reilly, *Nat. Chem. Biol.*, 2013, **9**, 285–288.
- 47 N. J. Turner and L. Humphreys, *Biocatalysis in organic synthesis: the retrosynthetic approach*, Royal Society of Chemistry, Cambridge, 2018.
- 48 A. P. Green and N. J. Turner, *Perspectives in Science*, 2016, **9**, 42–48.
- 49 R. O. M. A. de Souza, L. S. M. Miranda and U. T. Bornscheuer, *Chem.-Eur. J.*, 2017, **23**, 12040–12063.
- 50 R. A. Sheldon and J. M. Woodley, *Chem. Rev.*, 2018, **118**, 801–838.



- 51 A. Krüger, C. Schäfers, C. Schröder and G. Antranikian, *New Biotechnol.*, 2018, **40**, 144–153.
- 52 D. Brady, A. Rousseau, L. H. Steenkamp, J. Chaplin, S. Reddy, R. Mitra, B. Mboniswa, C. J. Parkinson, T. Moutlana, S. F. Marais and N. S. Gardener, *J. Mol. Catal. B: Enzym.*, 2012, **75**, 1–10.
- 53 A. Basso and S. Serban, *Mol. Catal.*, 2019, **479**, 110607.
- 54 R. DiCosimo, J. McAuliffe, A. J. Poulouse and G. Bohlmann, *Chem. Soc. Rev.*, 2013, **42**, 6437–6474.
- 55 U. Hanefeld, L. Gardossi and E. Magner, *Chem. Soc. Rev.*, 2009, **38**, 453–468.
- 56 R. A. Sheldon and S. van Pelt, *Chem. Soc. Rev.*, 2013, **42**, 6223–6235.
- 57 C. Garcia-Galan, A. Berenguer-Murcia, R. Fernandez-Lafuente and R. C. Rodrigues, *Adv. Synth. Catal.*, 2011, 2885–2904.
- 58 S. Prasad and I. Roy, *Recent Pat. Biotechnol.*, 2018, **12**, 33–56.
- 59 D. Brady and J. Jordaan, *Biotechnol. Lett.*, 2009, **31**, 1639–1650.
- 60 M. P. Thompson, I. Peñafiel, S. C. Cosgrove and N. J. Turner, *Org. Process Res. Dev.*, 2019, **23**, 9–18.
- 61 L. Tamborini, P. Fernandes, F. Paradisi and F. Molinari, *Trends Biotechnol.*, 2018, **36**, 74–88.
- 62 R. A. Sheldon, in *Biocatalysis: An Industrial Perspective*, ed. G. de Gonzalo and P. Dominguez de Maria, Royal Society of Chemistry, 2018, ch. 14.
- 63 R. A. Sheldon, *Catalysts*, 2019, **9**, 261.
- 64 M. J. Liszka, M. E. Clark, E. Schneider and D. S. Clark, *Annu. Rev. Chem. Biomol. Eng.*, 2012, **3**, 77–102.
- 65 O. Alvizo, L. J. Nguyen, C. K. Savile, J. A. Bresson, S. L. Lakhapatri, E. O. Solis, R. J. Fox, J. M. Broering, M. R. Benoit, S. A. Zimmerman, S. J. Novick, J. Liang and J. J. Lalonde, *Proc. Natl. Acad. Sci.*, 2014, **111**, 16436–16441.
- 66 R. A. Sheldon, *Green Chem.*, 2017, **19**, 18–43.
- 67 P. A. Wender, S. T. Handy and D. L. Wright, *Chem. Ind.*, 1997, 765–769.
- 68 B. M. Trost, *Science*, 1991, **254**, 1471–1477.
- 69 R. A. Sheldon, *Curr. Opin. Green Sust. Chem.*, 2019, **18**, 13–19.
- 70 P. Anastas and J. C. Warner, *Green Chemistry: Theory and Practice*, Oxford University Press, Oxford, UK, 1998.
- 71 J. Lalonde, *Curr. Opin. Biotechnol.*, 2016, **42**, 152–158.
- 72 M. D. Truppo, *ACS Med. Chem. Lett.*, 2017, **8**, 476–480.
- 73 K. Rosenthal and S. Lütz, *Curr. Opin. Green Sust. Chem.*, 2018, **11**, 58–64.
- 74 J. P. Adams, M. J. B. Brown, A. Diaz-Rodriguez, R. C. Lloyd and G. -D. Roiban, *Adv. Synth. Catal.*, 2019, **361**, 2421–2432.
- 75 T. Nagasawa, H. Shimizu and H. Yamada, *Appl. Microbiol. Biotechnol.*, 1993, **40**, 189–195.
- 76 B. Li, J. Su and J. Tao, *Org. Process Res. Dev.*, 2011, **15**, 291–293.
- 77 E. C. Hann, A. Eisenberg, S. K. Fager, N. E. Perkins, F. G. Gallagher, S. M. Cooper, J. E. Gavagan, B. Stieglitz, S. M. Hennessey and R. DiCosimo, *Bioorg. Med. Chem.*, 1999, **10**, 2239–2245.
- 78 Y. Isowa, M. Ohmori, T. Ichikawa, K. Mori, Y. Nonaka, K. Kihara, K. Oyama, H. Satoh and S. Nishimura, *Tetrahedron Lett.*, 1979, **20**, 2611–2612.
- 79 A. Bruggink, E. C. Roos and E. de Vroom, *Org. Process Res. Dev.*, 1998, **2**, 128–133.
- 80 C. Bertoldo and G. Antranikian, *Curr. Opin. Chem. Biol.*, 2002, **6**, 151–160.
- 81 C. A. Lipinski, *J. Pharmacol. Toxicol. Methods*, 2000, **44**, 235–249.
- 82 A. Salihu and M. Z. Alam, *Process Biochem.*, 2015, **50**, 86–96.
- 83 J. P. Adams, M. J. B. Brown, A. Diaz-Rodriguez, R. C. Lloyd and G.-D. Roiban, *Adv. Synth. Catal.*, 2019, **361**, 2421–2432.
- 84 R. N. Patel, *Bioorg. Med. Chem.*, 2018, **26**, 1252–1274.
- 85 A. C. Carvalho, T. de Fonseca, M. C. de Mattos, M. de Oliveira, T. L. de Lemos, F. Molinari, D. Romano and I. Serra, *Int. J. Mol. Sci.*, 2015, **16**, 29682–29716.
- 86 M. R. Petchey and G. Grogan, *Adv. Synth. Catal.*, 2019, **361**, 3895–3914.
- 87 I. C. Perdomo, S. Gianolio, A. Pinto, D. Romano, M. L. Contente, F. Paradisi and F. Molinari, *J. Agric. Food Chem.*, 2019, **67**, 6517–6522.
- 88 N. de Leeuw, G. Torrelo, C. Bisterfeld, V. Resch, L. Mestrom, E. Straulino, L. van der Weel and U. Hanefeld, *Adv. Synth. Catal.*, 2018, **360**, 242–249.
- 89 L. Mestrom, J. G. R. Claessen and U. Hanefeld, *ChemCatChem*, 2019, **11**, 2004–2010.
- 90 O. Verho and J.-E. Bäckvall, *J. Am. Chem. Soc.*, 2015, **137**, 3996–4009.
- 91 Y. Kim, J. Park and M.-J. Kim, *ChemCatChem*, 2011, **3**, 271–277.
- 92 C. A. Martinez, S. Hu, Y. Dumond, J. Tao, P. Kelleher and L. Tully, *Org. Process Res. Dev.*, 2008, **12**, 392–398.
- 93 M. Breuer, K. Ditrich, T. Habicher, B. Hauer, M. Keßeler, R. Stürmer and T. Zelinski, *Angew. Chem., Int. Ed.*, 2004, **43**, 788–824.
- 94 T. Itoh, *Chem. Rev.*, 2017, **117**, 10567–10607.
- 95 N. Kishi and H. Kojima, *ChemistrySelect*, 2019, **4**, 9570–9572.
- 96 W. P. Juma, V. Chhiba, D. Brady and M. L. Bode, *Tetrahedron: Asymmetry*, 2017, **28**, 1169–1174.
- 97 S. Dasetty, M. A. Blenner and S. Sarupria, *Mater. Res. Express*, 2017, **4**, 114008.
- 98 K. de Godoy Daiha, R. Angeli, S. D. de Oliveira and R. V. Almeida, *PLoS One*, 2015, **10**, e0131624.
- 99 D. L. Hughes, *Org. Process Res. Dev.*, 2016, **20**, 700–706.
- 100 D. L. Hughes, *Org. Process Res. Dev.*, 2018, **22**, 1063–1080.
- 101 P. Dominguez de María, G. de Gonzalo and A. R. Alcántara, *Catalysts*, 2019, **9**, 802.
- 102 G. W. Huisman and S. J. Collier, *Curr. Opin. Chem. Biol.*, 2013, **17**, 284–292.
- 103 R. Buller, K. Hechta, M. A. Mirata and H.-P. Meyer, in *Biocatalysis: An Industrial Perspective*, ed. G. de Gonzalo and P. Domínguez de María, Royal Society of Chemistry, 2017, pp. 3–43, ISBN 978-1-78262-619-0.
- 104 M. Yoshimura, S. Tanaka and M. Kitamura, *Tetrahedron Lett.*, 2014, **55**, 3635–3640.
- 105 R. Zhang, Y. Xu and R. Xiao, *Biotechnol. Adv.*, 2015, **33**, 1671–1684.
- 106 G. W. Huisman, J. Liang and A. Krebber, *Curr. Opin. Chem. Biol.*, 2010, **14**, 122.



- 107 S. K. Ma, J. Gruber, C. Davis, L. Newman, D. Gray, A. Wang, J. Grate, G. W. Huisman and R. A. Sheldon, *Green Chem.*, 2010, **12**, 81–86.
- 108 J. Liang, J. Lalonde, B. Borup, V. Mitchell, E. Mundorff, N. Trinh, D. A. Kochrekar, R. N. Cherat and G. G. Pai, *Org. Process Res. Dev.*, 2010, **14**, 193–198.
- 109 M. J. Abrahamson, E. Vazquez-Figueroa, N. B. Woodall, J. C. Moore and A. S. Bommarius, *Angew. Chem., Int. Ed.*, 2012, **51**, 3969–3972.
- 110 M. J. Abrahamson, J. W. Wong and A. S. Bommarius, *Adv. Synth. Catal.*, 2013, **355**, 1780–1786.
- 111 S. K. Au, B. R. Bommarius and A. S. Bommarius, *ACS Catal.*, 2014, **4**, 4021–4026.
- 112 T. Knaus, W. Böhmer and F. G. Mutti, *Green Chem.*, 2017, **19**, 453–463.
- 113 L. J. Ye, H. H. Toh, Y. Yang, J. P. Adams, R. Snajdrova and Z. Li, *ACS Catal.*, 2015, **5**, 1119.
- 114 P. N. Scheller, S. Fademrecht, S. Hofelzer, J. Pleis, F. Leipold, N. J. Turner, B. M. Nestl and B. Hauer, *ChemBioChem*, 2014, **15**, 2201–2204.
- 115 Z. Maugeri and D. Rother, *Adv. Synth. Catal.*, 2016, **358**, 2745–2750.
- 116 J. H. Schrittwieser, S. Velikogne and W. Kroutil, *Adv. Synth. Catal.*, 2015, **357**, 1655–1685.
- 117 J. Mangas-Sanchez, S. P. France, S. L. Montgomery, G. A. Aleku, H. Man, M. Sharma, J. I. Ramsden, G. Grogan and N. J. Turner, *Curr. Opin. Chem. Biol.*, 2017, **37**, 19–25.
- 118 G. Grogan and N. J. Turner, *Chem.–Eur. J.*, 2016, **22**, 1900–1907.
- 119 S. P. France, R. M. Howard, J. Steflík, J. M. Weise, J. Mangas-Sanchez, S. L. Montgomery, R. Crook, R. Kumar and N. J. Turner, *ChemCatChem*, 2018, **10**, 510–514.
- 120 P. N. Scheller, M. Lenz, S. C. Hammer, B. Hauer and B. M. Nestl, *ChemCatChem*, 2015, **7**, 3239.
- 121 M. Lenz, J. Meisner, L. Quertinmont, S. Lutz, J. Kastner and B. M. Nestl, *ChemBioChem*, 2017, **18**, 253–256.
- 122 D. Wetzl, M. Gand, A. Ross, H. Meller, P. Matzel, S. P. Hanlon, M. Meller, B. Wirz, M. Hçhne and H. Iding, *ChemCatChem*, 2016, **8**, 2023–2026.
- 123 P. Matzel, M. Gand and M. Hçhne, *Green Chem.*, 2017, **19**, 385–389.
- 124 M. H. S. A. Hamid, P. A. Slatford and J. M. J. Williams, *Adv. Synth. Catal.*, 2007, **349**, 1555–1575.
- 125 F. G. Mutti, T. Knaus, N. S. Scrutton, M. Breuer and N. J. Turner, *Science*, 2015, **349**, 1525–1529.
- 126 M. P. Thompson and N. J. Turner, *ChemCatChem*, 2017, **23**, 3833–3836.
- 127 W. Böhmer, T. Kraus and F. Mutti, *ChemCatChem*, 2018, **10**, 731–735.
- 128 M. C. Bryan, P. J. Dunn, D. Entwistle, F. Gallou, S. G. Koenig, J. D. Hayler, M. R. Hickey, S. Hughes, M. E. Kopach, G. Moine, P. Richardson, F. Roschangar, A. Steven and F. J. Weiberthm, *Green Chem.*, 2018, **20**, 5082–5103.
- 129 M. N. Kopylovich, A. P. C. Ribeiro, E. C. B. A. Alegria, N. M. R. Martins, L. M. D. R. S. Martins and A. J. L. Pombeiro, *Adv. Organomet. Chem.*, 2015, **63**, 91–174.
- 130 R. A. Sheldon, *Catal. Today*, 2015, **247**, 4–13.
- 131 G. J. ten Brink, I. W. C. E. Arends and R. A. Sheldon, *Science*, 2000, **287**, 1636–1639.
- 132 R. A. Sheldon, I. W. C. E. Arends, G. J. ten Brink and A. Dijkman, *Acc. Chem. Res.*, 2002, **35**, 774–781.
- 133 J. E. Steves and S. S. Stahl, in *Liquid Phase Aerobic Oxidation Catalysis*, ed. S. S. Stahl and P. J. Alsters, 2016, pp. 85–96.
- 134 J. E. Steves, Y. Preger, J. R. Martinelli, C. J. Welch, T. W. Root, J. M. Hawkins and S. S. Stahl, *Org. Process Res. Dev.*, 2015, **19**, 1548–1553.
- 135 J. Liu, S. Wu and Z. Li, *Curr. Opin. Chem. Biol.*, 2018, **43**, 77–86.
- 136 F. Escaleters and N. J. Turner, *ChemBioChem*, 2008, **9**, 857–860.
- 137 S. Herter, S. M. McKenna, A. R. Frazer, S. Lehmkuhler, A. J. Carnell and N. J. Turner, *ChemCatChem*, 2015, **7**, 2313–2317.
- 138 S. M. McKenna, S. Leimkuhler, S. Herter, N. J. Turner and A. Carnell, *Green Chem.*, 2015, **17**, 3271–3275.
- 139 W. P. Dijkman, C. Binda, M. W. Fraaije and A. Mattevi, *ACS Catal.*, 2015, **5**, 1833–1839.
- 140 J. Vilim, T. Kraus and F. Mutti, *Angew. Chem., Int. Ed.*, 2018, **57**, 14240–14244.
- 141 A. Dijkman, I. W. C. E. Arends and R. A. Sheldon, *Org. Biomol. Chem.*, 2003, **1**, 3232–3237.
- 142 J. Gross, K. Tauber, M. Fuchs, N. G. Schmidt, A. Rajagopalan, K. Faber, W. M. F. Fabian, J. Pfeiffer, T. Haas and W. Kroutil, *Green Chem.*, 2014, **16**, 2117–2121.
- 143 A. Weckbecker, H. Gröger and W. Hummel, *Adv. Biochem. Eng./Biotechnol.*, 2010, **120**, 195–242.
- 144 J. K. B. Cahn, C. A. Werlang, A. Baumschlager, S. Brinkman-Chen, S. I. Mayo and F. Arnold, *ACS Synth. Biol.*, 2017, **6**, 326–333.
- 145 W. Kroutil, H. Mang, K. Edegger and K. Faber, *Curr. Opin. Chem. Biol.*, 2004, **8**, 120–126.
- 146 W. Kroutil, H. Mang, K. Edegger and K. Faber, *Adv. Synth. Catal.*, 2004, **346**, 125–142.
- 147 J. D. Zhang, Z. M. Cui, X. J. Fan, H. L. Wu and H. H. Chang, *Bioprocess Biosyst. Eng.*, 2016, **39**, 603–611.
- 148 T. Knaus, V. Tseliou, L. D. Humphreys, N. S. Scrutton and F. G. Mutti, *Green Chem.*, 2018, **20**, 3931–3943.
- 149 T. Knaus, F. G. Mutti, L. D. Humphreys, N. J. Turner and N. S. Scrutton, *Org. Biomol. Chem.*, 2015, **13**, 223–233.
- 150 K. Bulke, M. Kadow, H. Mallin, S. Sass and U. T. Bornscheuer, *Org. Biomol. Chem.*, 2012, **10**, 6249–6265.
- 151 M. Bucko, P. Gemeiner, A. Scenkmayerova, T. Kajovic, F. Rudroff and M. D. Mihovilovic, *Appl. Microbiol. Biotechnol.*, 2016, **100**, 6585–6599.
- 152 H. Leisch, K. Morley and P. C. K. Lau, *Chem. Rev.*, 2011, **111**, 4165.
- 153 E. Y. Jeon, A. H. Beek, U. T. Bornscheuer and J. B. Park, *Appl. Microbiol. Biotechnol.*, 2015, **99**, 6267–6275.
- 154 A. Bornadel, R. Hatti-Kaul, F. Hollmann and S. Kara, *ChemCatChem*, 2015, **7**, 2442–2445.
- 155 R. Wilson, *Chim Oggi*, 2015, **33**, 50–52; see also G. de Gonzalo and A. Franconetti, *Enzyme Microb. Technol.*, 2018, **113**, 24–28.



- 156 Y. Zang, Y. Q. Wu, N. Xu, Q. Zhao, H. L. Yu and J. H. Xu, *ACS Sustainable Chem. Eng.*, 2019, **7**, 7218–7226.
- 157 G. Grogan, *Curr. Opin. Chem. Biol.*, 2018, **43**, 15–52.
- 158 D. Ghislieri, A. P. Green, M. Pontini, S. C. Willies, I. Rowles, A. Frank, G. Grogan and N. J. Turner, *J. Am. Chem. Soc.*, 2013, **135**, 10863–10869.
- 159 B. Mijts, S. Muley, J. Liang, L. M. Newman, X. Zhang, J. Lalonde, M. D. Clay, J. Zhu, J. Gruber, J. Colbeck, J. D. Munger Jr, J. Mavinahalli and R. Sheldon, *US Pat.*, 8574876 B2, 2013.
- 160 J. Dong, E. Fernandez-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.
- 161 S. Bormann, A. G. Baraibar, Y. Ni, D. Holtmann and F. Hollmann, *Catal. Sci. Technol.*, 2015, **5**, 2038–2052.
- 162 F. P. Guengerich, *ACS Catal.*, 2018, **8**, 10964–10976.
- 163 A. W. Munro, D. G. Leys, K. J. McLean, K. R. Marshall, T. W. B. Ost, S. Daff, C. S. Miles, S. K. Chapman, D. A. Lysek, C. C. Moser, C. C. Page and P. L. Dutton, *Trends Biochem. Sci.*, 2002, **27**, 250–257.
- 164 M. Tavanti, J. L. Porter, S. Sabatini, N. J. Turner and S. L. Flitsch, *ChemCatChem*, 2018, **10**, 1042–1051.
- 165 J. N. Kolev, J. M. Zaengle, R. Ravikumar and R. Fasan, *ChemBioChem*, 2014, **15**, 1001–1010.
- 166 T. Kubo, P. Meinhold, M. W. Peters and F. H. Arnold, *Chem.–Eur. J.*, 2006, **12**, 1216–1220.
- 167 S. Peter, M. Kinne, R. Ulrich, G. Kayser and M. Hofrichter, *Enzyme Microb. Technol.*, 2013, **52**, 370–376.
- 168 C. Gally, B. Nestl and B. Hauer, *Angew. Chem., Int. Ed.*, 2015, **54**, 12952–12956.
- 169 H. C. Kolb, M. S. Van Nieuwenhze and K. B. Sharpless, *Chem. Rev.*, 1994, **94**, 2483–2547.
- 170 D. Holtmann and F. Hollmann, *ChemBioChem*, 2016, **17**, 1391–1398.
- 171 Y. Wang, D. Lan, R. Durrani and F. Hollmann, *Curr. Opin. Chem. Biol.*, 2017, **37**, 1–9.
- 172 J. S. Shin and B. G. Kim, *Biotechnol. Bioeng.*, 1999, **65**, 206–211.
- 173 G. W. Matcham and A. R. S. Bowen, *Chem. Today*, 1996, **14**, 20–24.
- 174 S. A. Kelly, S. Pohle, S. Wharry, S. Mix, C. C. R. Allen, T. S. Moody and B. F. Gilmore, *Chem. Rev.*, 2018, **118**, 349–367.
- 175 J. E. Farnberger and W. Kroutil, *Eur. J. Org. Chem.*, 2015, 6965–6982.
- 176 C. K. Saville, J. M. Maney, E. C. Mundorff, J. C. Moore, S. Tam, W. R. Jarvis, J. C. Colbeck, A. Krebber, F. J. Fleitz, J. Brands, P. N. Devine, G. W. Huisman and G. J. Hughes, *Science*, 2010, **329**, 305–309.
- 177 I. V. Pavlidis, M. S. Weiß, M. Genz, P. Spurr, S. P. Hanlon, B. Wirz, H. Iding and U. Bornscheuer, *Nat. Chem.*, 2016, **8**, 1076–1082.
- 178 K. Fesko and M. Gruber-Khadjawi, *ChemCatChem*, 2013, **5**, 1248–1272.
- 179 S. K. Panday, *Tetrahedron: Asymmetry*, 2011, **22**, 1817–1847.
- 180 S. Bhowmick, A. Mondal, A. Ghosh and K. C. Bhowmick, *Tetrahedron: Asymmetry*, 2015, **26**, 1215–1244.
- 181 P. Bracco, H. Busch, J. von Langermann and U. Hanefeld, *Org. Biomol. Chem.*, 2016, **14**, 6375–6389.
- 182 E. Lanfranchia, K. Steiner, A. Glieder, I. Hajnal, R. A. Sheldon, S. van Pelt and M. Winkler, *Recent Pat. Biotechnol.*, 2013, **7**, 197–206.
- 183 L. M. van Langen, R. P. Selassa, F. van Rantwijk and R. A. Sheldon, *Org. Lett.*, 2005, **7**, 327–329.
- 184 M. Haridas, E. M. M. Abdelraheem and U. Hanefeld, *Appl. Microbiol. Biotechnol.*, 2018, **102**, 9959–9971.
- 185 S. Jennewein, M. Schurmann, M. Wolberg, I. Hilker, R. Luiten, M. Wubbolts and D. Mink, *Biotechnol. J.*, 2006, **1**, 537–548.
- 186 R. A. Sheldon, *Chimia*, 1996, **50**, 418EP0343714A1, 1989.
- 187 R. J. Fox, S. C. Davis, E. Mundorff, L. Newman, V. Gavrilovic, S. Ma, L. Chung, C. Ching, S. Tam, S. Muley, J. Grate, J. Gruber, J. Whitman, R. A. Sheldon and G. W. Huisman, *Nat. Biotechnol.*, 2007, **25**, 338–344.
- 188 P. Hoyos, V. Pace and A. R. Alcantara, *Catalysts*, 2019, **9**(260), 1–32.
- 189 R. A. Sheldon, in *Multi-Step Enzyme Catalysis: Biotransformations and Chemoenzymatic Synthesis*, ed. E. Garcia Junceda, Wiley-VCH, Weinheim, 2008, pp. 109–135.
- 190 J. Muschiol, C. Peters, N. Oberleitner, M. D. Mihovilovic, U. T. Bornscheuer and F. Rudroff, *Chem. Commun.*, 2015, **51**, 5798–5811.
- 191 R. Xue and J. M. Woodley, *Bioresour. Technol.*, 2012, **115**, 183–195.
- 192 V. Köhler and N. J. Turner, *Chem. Commun.*, 2015, **13**, 223–233.
- 193 R. C. Simon, N. Richter, E. Busto and W. Kroutil, *ACS Catal.*, 2014, **4**, 129–143.
- 194 G. E. R. Gordon, M. L. Bode, D. F. Visser, M. J. Lepuru, J. G. Zeevaart, N. Ragubeer, M. Ratsaka, D. R. Walwyn and D. Brady, *Org. Process Res. Dev.*, 2011, **15**, 258–265.
- 195 D. Brady, A. Beeton, C. Kgaje, J. Zeevaart, F. van Rantwijk and R. A. Sheldon, *Appl. Microbiol. Biotechnol.*, 2004, **64**, 76–85.
- 196 M. A. Huffman, A. Fryszkowska, O. Alvizo, M. Borra-Garske, K. R. Campos, K. A. Canada, P. N. Devine, D. Duan, J. H. Forstater, S. T. Grosser, H. M. Halsey, G. J. Hughes, J. Jo, L. A. Joyce, J. N. Kolev, J. Liang, K. M. Maloney, B. F. Mann, N. M. Marshall, M. McLaughlin, J. C. Moore, G. S. Murphy, C. C. Nawrat, J. Nazor, S. Novick, N. R. Patel, A. Rodriguez-Granillo, S. A. Robaire, E. C. Sherer, M. D. Truppo, A. M. Whittaker, D. Verma, L. Xiao, Y. Xu and H. Yan, *Science*, 2019, 1255–1259.
- 197 F. Rudroff, M. D. Mihovilovic, H. Gröger, R. Snajdrova, H. Iding and U. T. Bornscheuer, *Nat. Catal.*, 2018, **1**, 12–22.
- 198 A. V. Ruales-Salcedo, J. C. Higueta, J. Fontalvo and J. M. Woodley, *Z. Naturforsch.*, 2019, **74**, 77–84.
- 199 E. Garcia-Junceda, I. Lavandera, D. Rother and J. H. Schrittwieser, *J. Mol. Catal. B: Enzym.*, 2015, **114**, 1–6.
- 200 J. B. Pyser, S. A. Baker Dockrey, A. Rodríguez Benítez, L. A. Joyce, R. A. Wiscons, J. L. Smith and A. R. H. Narayan, *J. Am. Chem. Soc.*, 2019, **141**, 18551–18559.



- 201 T. J. Doyon, J. C. Perkins, S. A. Baker Dockrey, E. O. Romero, K. C. Skinner, P. M. Zimmerman and A. R. H. Narayan, *J. Am. Chem. Soc.*, 2019, **141**, 20269–20277.
- 202 J. Li, X. Zhang and H. Renata, *Angew. Chem., Int. Ed.*, 2019, **58**, 11657–11660.
- 203 C. R. Zwick III and H. Renata, *J. Org. Chem.*, 2018, **83**, 7407–7415.
- 204 M. Cortes-Clerget, N. Akporji, J. Zhou, F. Gao, P. Guo, M. Parmentier, F. Gallou, J.-Y. Berthon and B. H. Lipshutz, *Nat. Commun.*, 2019, **10**, 2169.
- 205 J. H. Schrittwieser, S. Velikogne, M. Hall and W. Kroutil, *Chem. Rev.*, 2018, **118**, 270–348.
- 206 U. T. Bornscheuer, *Philos. Trans. R. Soc., A*, 2017, **376**, 20170063.
- 207 H. Renata, Z. J. Wang and F. H. Arnold, *Angew. Chem., Int. Ed.*, 2015, **54**, 3351–3367.
- 208 S. Noor, M. C. Taylor, R. J. Russell, L. S. Jermini, C. J. Jackson, J. G. Oakshott and C. Scott, *PLoS One*, 2012, **7**, e39822.
- 209 S. C. Hammer, A. M. Knight and F. H. Arnold, *Curr. Opin. Green Sus. Chem.*, 2017, **7**, 23–30.
- 210 U. T. Bornscheuer and R. J. Kazlauskas, *Angew. Chem., Int. Ed.*, 2004, **43**, 6032–6040.
- 211 F. H. Arnold, *Q. Rev. Biophys.*, 2015, **48**, 404–410.
- 212 H. M. Key, P. Dydio, D. S. Clark and J. F. Hartwig, *Nature*, 2016, **534**, 534–537.
- 213 P. Srivastava, H. Yang, K. Ellis-Guardiola and J. C. Lewis, *Nat. Commun.*, 2015, **6**, 7789.
- 214 P. S. Coelho, E. M. Brustad, A. Kannan and F. H. Arnold, *Science*, 2013, **339**, 307–310.
- 215 T. K. Hyster and T. R. Ward, *Angew. Chem., Int. Ed.*, 2016, **55**, 7344–7357.
- 216 M. Bordeaux, V. Tyagi and R. Fasan, *Angew. Chem., Int. Ed.*, 2015, **127**, 1764–1768.
- 217 O. F. Brandenburg, R. Fasan and F. H. Arnold, *Curr. Opin. Biotechnol.*, 2017, **47**, 102–111.
- 218 H. M. Key, P. Dydio, Z. Liu, J. Y.-E. Rha, A. Nazarenko, V. Seyedkazemi, D. S. Clark and J. F. Hartwig, *ACS Cent. Sci.*, 2017, **3**, 302–308.
- 219 C. C. Farwell, R. K. Zhang, J. A. McIntosh, T. K. Hyster and F. H. Arnold, *ACS Cent. Sci.*, 2015, **1**, 89–93.
- 220 J. A. McIntosh, P. S. Coelho, C. C. Farwell, Z. J. Wang, J. C. Lewis, T. R. Brown and F. H. Arnold, *Angew. Chem., Int. Ed.*, 2013, **52**, 9309–9312.
- 221 E. W. Svastits, J. H. Dawson, R. Breslow and S. Gellman, *J. Am. Chem. Soc.*, 1985, **107**, 6427–6428.
- 222 T. K. Hyster, C. C. Farwell, A. R. Buller, J. A. McIntosh and F. H. Arnold, *J. Am. Chem. Soc.*, 2014, **136**, 15505–15508.
- 223 C. K. Prier, R. K. Zhang, A. R. Buller, S. Brinkman-Chen and F. H. Arnold, *Nat. Chem.*, 2017, **9**, 629–634.
- 224 O. F. Brandenburg, D. C. Miller, U. Markel, A. O. Chaib and F. H. Arnold, *ACS Catal.*, 2019, **9**, 8271–8275.
- 225 V. Tyagi and R. Fasan, *Angew. Chem., Int. Ed.*, 2016, **55**, 2512–2516.
- 226 O. F. Brandenburg, K. Chen and F. H. Arnold, *J. Am. Chem. Soc.*, 2019, **141**, 8989–8995.
- 227 J. Zhang, X. Huang, R. K. Zhang and F. H. Arnold, *J. Am. Chem. Soc.*, 2019, **141**, 9798–9802.
- 228 H. Tsutsumi, Y. Katsuyama, M. Izumikawa, M. Takagi, M. Fujie, N. Satoh, K. Shin-ya and Y. Ohnishi, *J. Am. Chem. Soc.*, 2018, **140**, 6631–6639.
- 229 J. G. Gober and E. M. Brustad, *Curr. Opin. Chem. Biol.*, 2016, **35**, 124–132.
- 230 B. P. Dwivedee, S. Soni, M. Sharma, J. Bhaumik, J. K. Laha and U. C. Banerjee, *ChemistrySelect*, 2018, **3**, 2441–2466.
- 231 D. Koszelewski and R. Ostaszewski, *ChemCatChem*, 2019, **11**, 2554–2558.
- 232 Q. Wu, B.-K. Liu and X.-F. Lin, *Curr. Org. Chem.*, 2010, **14**, 1966–1988.
- 233 M. Svedendahl, K. Hult and P. Berglund, *J. Am. Chem. Soc.*, 2005, **127**, 17988–17989.
- 234 M. Svedendahl, B. Jovanović, L. Fransson and P. Berglund, *ChemCatChem*, 2009, **1**, 252–258.
- 235 F.-W. Lou, B.-K. Liu, Q. Wu, D.-S. Lv and X.-F. Lin, *Adv. Synth. Catal.*, 2008, **350**, 1959–1962.
- 236 R. Kazlauskas, *Curr. Opin. Chem. Biol.*, 2005, **9**, 195–201.
- 237 C. Li, X.-W. Feng, N. Wang, Y.-J. Zhou and X.-Q. Yu, *Green Chem.*, 2008, **10**, 616–618.
- 238 R. C. Tang, Z. Guan, Y.-H. He and W. Zhu, *J. Mol. Catal. B: Enzym.*, 2010, **63**, 62–67.
- 239 K. Li, T. He, C. Li, X.-W. Feng, N. Wang and X.-Q. Yu, *Green Chem.*, 2009, **11**, 777–779.
- 240 E. Busto, V. Gotor-Fernández and V. Gotor, *Chem. Soc. Rev.*, 2010, **39**, 4504–4523.
- 241 M. López-Iglesias and V. Gotor-Fernández, *Chem. Rec.*, 2015, **15**, 743–759.
- 242 X. Ge, Y. Lai and X. Chen, *Chin. J. Org. Chem.*, 2013, **33**, 1686–1696.
- 243 M. Jeschek, R. Reuter, T. Heinisch, C. Trindler, J. Klehr, S. Panke and T. R. Ward, *Nature*, 2016, **537**, 661–665.
- 244 J. B. Siegel, A. Zanghellini, H. M. Lovick, G. Kiss, A. R. Lambert, J. L. St. Clair, J. L. Gallaher, D. Hilvert, M. H. Gelb, B. L. Stoddard, K. N. Houk, F. E. Michael and D. Baker, *Science*, 2010, **329**, 309–313.
- 245 T. K. Hyster, *Synlett*, 2019, **30**, A–G.
- 246 K. F. Biegasiewicz, S. J. Cooper, X. Gao, D. G. Oblinsky, J. H. Kim, S. E. Garfinkle, L. A. Joyce, B. A. Sandoval, G. D. Scholes and T. K. Hyster, *Science*, 2019, **364**, 1166–1169.
- 247 G. Qu, A. Li, C. G. Acevedo-Rocha, Z. Sun and M. T. Reetz, *Angew. Chem., Int. Ed.*, 2020, DOI: 10.1002/anie.201901491.
- 248 G. Li, Y. Dong and M. T. Reetz, *Adv. Synth. Catal.*, 2019, **361**, 2377–2386.
- 249 Y. Saito, M. Oikawa, H. Nakazawa, T. Niide, T. Kameda, K. Tsuda and M. Umetsu, *ACS Synth. Biol.*, 2018, **7**, 2014–2022.
- 250 F. Cadet, N. Fontaine, G. Li, J. Sanchis, M. N. F. Chong, R. Pandjaitan, I. Vetrivel, B. Offmann and M. T. Reetz, *Sci. Rep.*, 2018, **8**, 16757.
- 251 M. H. S. Segler, M. Preuss and M. P. Waller, *Nature*, 2018, **555**, 604–610.
- 252 X. W. Diefenbach, I. Farasat, E. D. Guetschow, C. J. Welch, R. T. Kennedy, S. Sun and J. C. Moore, *ACS Omega*, 2018, **3**, 1498–1508.
- 253 P. Žnidaršič-Plazl, *Biotechnol. J.*, 2019, **14**, 1800580.



- 254 T. Peschke, P. Bitterwolf, K. S. Rabe and C. M. Niemeyer, *Chem. Eng. Technol.*, 2019, **42**(10), 2009–2017.
- 255 C. W. Coley, D. A. Thomas, J. A. M. Lummiss, J. N. Jaworski, C. P. Breen, V. Schultz, T. Hart, J. S. Fishman, L. Rogers, H. Gao, R. W. Hicklin, P. P. Plehiers, J. Byington, J. S. Piotti, W. H. Green, A. J. Hart, T. F. Jamiso and K. F. Jensen, *Science*, 2019, **365**(6453), eaax1566.
- 256 A. Currin, N. Swainston, P. J. Dayace and D. B. Kell, *Chem. Soc. Rev.*, 2015, **44**, 1172–1239.
- 257 J. Fredens, K. Wang, D. de la Torre, L. F. H. Funke, W. E. Robertson, Y. Christova, T. Chia, W. H. Schmied, D. L. Dunkelmann, V. Beránek, C. Uttamapinant, A. Gonzalez Llamazares, T. S. Elliott and J. W. Chin, *Nature*, 2019, **569**, 514–518.
- 258 G. Grogan, *Practical Biotransformations: A Beginner's Guide*, Wiley-Blackwell, 2009, ISBN: 978-1-405-17125-0.
- 259 K. Faber, *Biotransformations in Organic Chemistry*, 2011, Springer, ISBN 978-3-319-61590-5.
- 260 *Biocatalysis in Organic Synthesis. Science of Synthesis*, ed. K. Faber, W.-D. Fessner and N. J. Turner, Thieme, 2015, vol. 1–3, ISBN 978-3132028715.
- 261 *Practical Methods for Biocatalysis and Biotransformations*, ed. J. Whittall, P. W. Sutton and W. Kroutil, Wiley, 2009, 2012 and 2016, vol. 1–3, ISBN: 9780470519271, 9781119991397 and ISBN: 9781118605257.
- 262 C.-I. Lin, R. M. McCarty and H.-W. Liu, *Angew. Chem., Int. Ed.*, 2017, **56**, 3446–3489.

