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Review

Direct Biocatalytic Processes for CO₂ Capture as a Green Tool to Produce Value-Added Chemicals

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Abstract: Direct biocatalytic processes for CO₂ capture and transformation in value-added chemicals may be considered a useful tool for reducing the concentration of this greenhouse gas in the atmosphere. Among the other enzymes, carbonic anhydrase (CA) and formate dehydrogenase (FDH) are two key biocatalysts suitable for this challenge, facilitating the uptake of carbon dioxide from the atmosphere in complementary ways. Carbonic anhydrases accelerate CO₂ uptake by promoting its solubility in water in the form of hydrogen carbonate as the first step in converting the gas into a species widely used in carbon capture storage and its utilization processes (CCSU), particularly in carbonation and mineralization methods. On the other hand, formate dehydrogenases represent the biocatalytic machinery evolved by certain organisms to convert CO₂ into enriched, reduced, and easily transportable hydrogen species, such as formic acid, via enzymatic cascade systems that obtain energy from chemical species, electrochemical sources, or light. Formic acid is the basis for fixing C₁-carbon species to other, more reduced molecules. In this review, the state-of-the-art of both methods of CO₂ uptake is assessed, highlighting the biotechnological approaches that have been developed using both enzymes.

Keywords: carbonic anhydrase; formate dehydrogenase; carbon capture storage and its utilization; cofactor regeneration



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1. Reducing Carbon Dioxide from the Air: The Challenge

One of the main challenges faced by humanity in the 21st century is climate change. Over the last two centuries, the temperature of the Earth's crust has risen progressively and, since 1980, alarmingly, at a rate of 0.18 °C per decade. Indeed, last year's average temperature was 1.04 °C higher than the median temperature in the period prior to 1880 [1]. Temperature elevation drives an increase in extreme weather evidenced by a series of well-known events (draughts, floodings, torrential downpours, etc.), the melting of large extensions of frozen water, with the subsequent ascent of the sea, changes in ecosystems with undefined outcomes and, in this sense, uncertainty on how these changes will affect our way of living and welfare, with estimations that are clearly detrimental [2]. Moreover, the acidification of seas and oceans is also a problem, with coral reef weakening already having been detected, as well as the low level of oxygen present in marine life [3,4].

At the end of the nineteenth century, S.A. Arrhenius quantified the contribution of "carbonic acid" (nowadays, carbon dioxide) to the greenhouse effect and was the first in indicating that "The production of carbonic acid by the combustion of coal would therefore suffice to cover the loss of carbonic acid by weathering and by peat formation seven times over. Those are the two chief factors deciding the consumption of carbonic acid, and we thus recognize that the percentage of carbonic acid in the air must be increasing at a constant rate as long as the consumption of coal, petroleum, etc., is maintained at its present figure, and at a still more rapid rate if this consumption should continue to increase as it does now". He also concluded that

this would lead to an increase in the temperature of Earth's atmosphere [5,6]. Since then, a huge amount of evidence correlating both air CO₂ concentration and global warming has accumulated [2,7]. Moreover, there is a direct relationship between human activity, carbon dioxide concentration, and climate change, that is, the anthropogenic origin of global warming is well established. Atmospheric CO₂ concentration has increased from 280 ppm (year 1750) to 415 ppm (2021), this value being the highest concentration reached in the last three million years [8]. This carbon dioxide increase is essentially related to the emissions of this gas to the atmosphere as a consequence of the use of fossil fuels by humans [9]. Other gases, such as methane and nitrous oxide, also contribute to global warming, albeit to a minor extent (11 and 7%, respectively) [10]. The objective reached at the Paris Climate Agreement in 2015 to maintain an increase in overall temperature below 2.0 °C with respect to preindustrial levels has recently been revised in the sense that such increments should not exceed 1.5 °C [11].

In this scenario, any scientific strategy that allows for reducing the concentration of CO₂ in the atmosphere is an object of interest, although the most relevant solution is to avoid burning fossil energy sources that release CO₂, and to substitute them with other sustainable ones. Although more efficient and responsible use of fossil energy sources by society is also essential to contributing to decreasing CO₂ emissions, the responsibility of capturing the excess CO₂ already emitted is also inescapable.

Among other approaches, biocatalysts are useful tools for reducing the accumulation of atmospheric CO₂ through either carbon capture and storage (CCS) or carbon capture and its utilization (CCU, Figure 1A). Both approaches are often applied together, known as carbon capture storage and utilization (CCSU) [12]. These methodologies require the passing of CO₂ from gas to carbon solid and/or chemically reduced forms, a task that directly implies chemistry in all its fields.

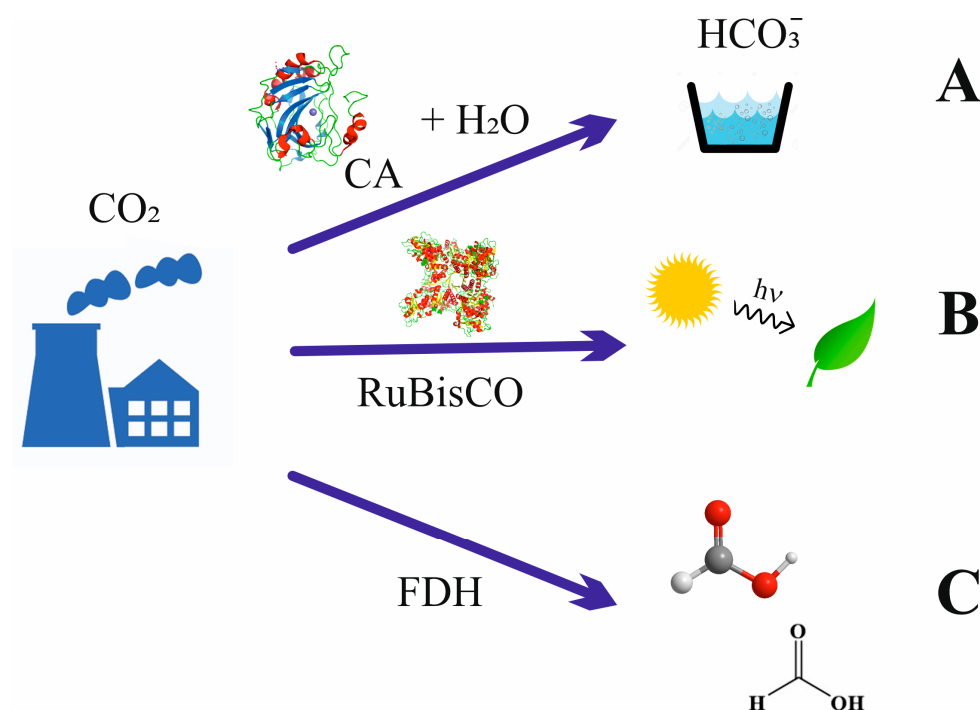


Figure 1. Three pathways to incorporating CO₂ into soluble carbon forms by using enzymes. (A) Formation of hydrogen carbonate using carbonic anhydrase; (B) photosynthesis performed using Ru-BisCO protein; (C) reduction to achieve C₁ in reduced form, formic acid, catalyzed using FDHs. Approaches A and C are the aims of the present work (see text for details).

Research in this field has exponentially increased in the last decade. Indeed, Figure 2A shows the number of articles published, directly or indirectly, that relate either to car-

bon storage or carbon utilization per year, while Figure 2B,C display the percentages of these articles classified by research area. As observed, research on carbon capture is especially intense in areas such as catalysts, synthesis, electrochemistry, and energy and fuels. Nowadays, there are two main approaches for CO₂ transformation, namely biological and chemical transformations [13]. In turn, biological CO₂ fixation can be photosynthetic or not photosynthetic, while the chemical uptake can be divided into hydrogenation, carboxylation, mineralization, chemical reduction, and photochemical reduction. Although hydrogenation is a well-established technology, nowadays, it is still mostly not green, as it is obtained from the cracking of fossil fuels. The sustainable synthesis and transport of hydrogen is indeed one of the main challenges related to energy sources. The main target of hydrogenation is the production of methanol (see Section 7.2), although other reduced molecules can also be obtained.

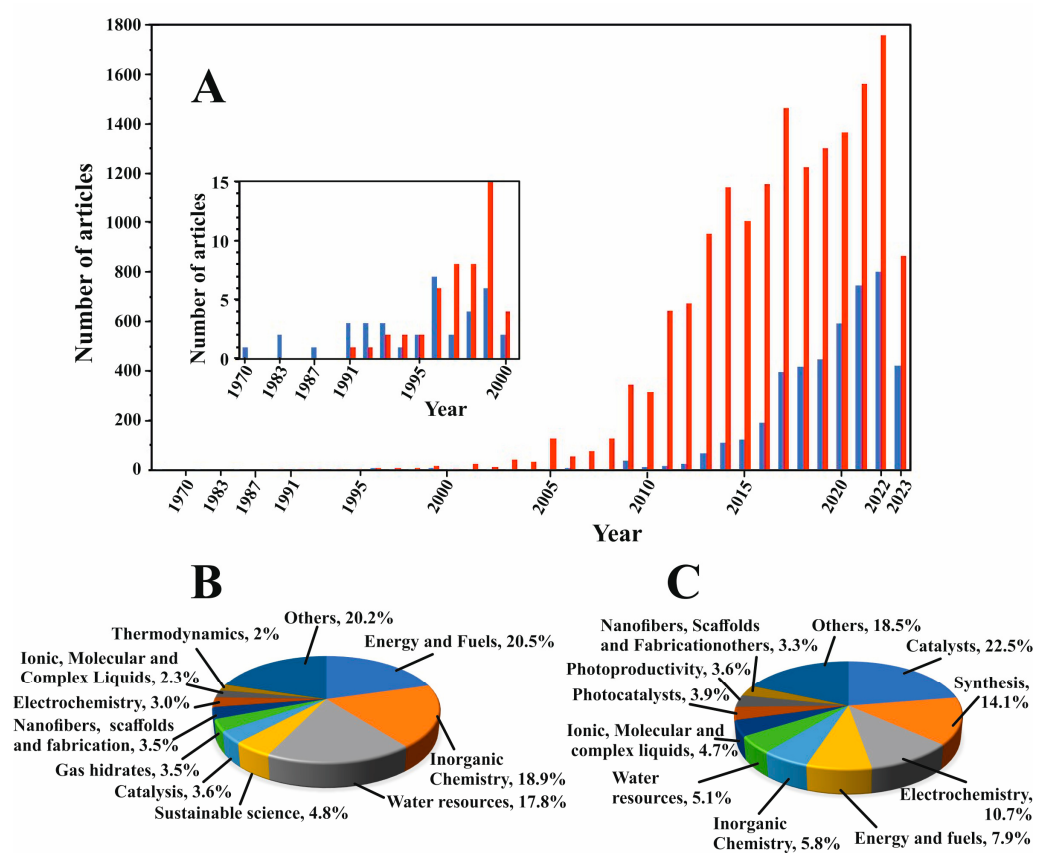
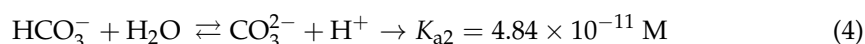
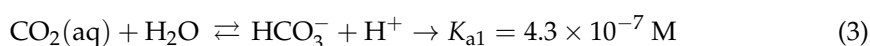
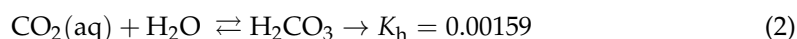
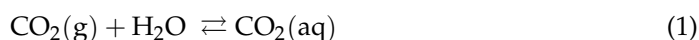


Figure 2. (A) Number of articles published per year related to carbon capture storage (red) and carbon capture utilization (blue); the insert is a vertical expansion, scale 0–15, by year, published in the last century (the year 2023 contains the number of articles at halfway through the year). (B,C) Percentage of articles related to CCS (B) and CCU (C) classified by research area. Source: compiled by the authors, based on the Web of Science database (<https://clarivate.com>), accessed on 2 July 2023.

Carboxylation is the process of directly converting CO₂ into organic value-added compounds. Organic carbonates and polymers are obtained through this process. An example is the production of non-isocyanate polyurethanes (NIPUs) from glycerol carbonates derivatives obtained through CO₂ cycloaddition to glycidol moieties [14]. In this case, the authors designed a sustainable chemoenzymatic protocol for the synthesis of glycerol carbonate acrylate (GCA) and glycerol carbonate methacrylate (GCMA) from glycidol and CO₂ by using ionic liquid (IL) technologies and enzymes, providing conversions of up to 100% under low-pressure values (1–10 bar). These methods are still in their first stages of development and hence a relatively novel field to work in.

Mineralization is described in Section 6.3 and is used in construction, for instance, for generating cement. This technique allows for the uptake of large quantities of CO₂ for obtaining sustainable materials, although it is a highly consuming energy at a high scale. Chemical, electrochemical and photochemical reduction combined with enzymes is described below (see Sections 7.2–7.4).

The genuine physical and chemical properties of CO₂ make it a molecule that cannot be easily captured or retained. Thus, CO₂ is a nonpolar molecule that resides in a gaseous state under P and T standard conditions because of the weak van der Waals interactions established between the molecules themselves. In addition, due to its null polarity, CO₂ has a low diffusion coefficient in polar solvents (1.26×10^{-5} cm²/s in water under standard conditions), while its solubility follows Henry's law (76.5 mM in water at 0 °C and partial pressure of 1 atm) [15]. More importantly, its kinetics of capture by water, while increasing with pH, is extremely low at neutral or acidic pH values, behaving as a Brönsted acid according to Equations (1) and (2) (Scheme 1). The K_h value (Equation (2)) indicates that only a minimal fraction of CO₂(aq) is present in an aqueous solution of carbonic acid (ca. 1/600 of the molecules at neutral pH values). Both hydrated carbon dioxide and carbonic acid behave as weak Brönsted acids because of the low deprotonation constants (see Equations (3) and (4)) [16].



Scheme 1. Acid–base equilibria for CO₂ water-soluble species as a function of the pH (Equations (1)–(4)) and reduction CO₂ equation to generate formic acid using the cofactor NADH (Equation (5)).

The equilibrium shown in Equation (3) is essential for the role of CO₂ as a buffer between blood and cells, where the interconversion between CO₂ and HCO₃[−] (hydro carbonate or bicarbonate) forms should be fast in living beings. Under neutral or weak basic conditions, the formation of bicarbonate anions is slow (the first order kinetic constant, k_1 , is of the order 10^{-2} s^{−1}), while it increases in basic media [17]. Consequently, living organisms must have a biocatalyst that allows for fast exchange between acid and basic species. Carbon dioxide gas can be fixed either in aqueous soluble (hydrogen carbonate) or much more insoluble (carbonate) forms using only alkalization or using other chemical processes that incorporate carbon into value-added compounds. In nature, carbonic anhydrase (CA) catalyzes the CO₂/hydrogen carbonate reaction (Equation (3)), enabling CO₂ capture within reasonable times [17,18]. Thus, CA is used as a tool for incorporating CO₂ as a hydrogen carbonate anion in soluble species in a multitude of chemical approaches [19,20]. In contrast, carbonates are usually highly insoluble. It follows that the use of alkaline reactants (Equation (4)) is the easiest and, consequently, main method for sequestering CO₂ by forming the corresponding salts or their derivatives.

Alternatively, CO₂ can be fixed to hydrogen-enriched energy forms via several natural mechanisms [21,22]. Although the most extended and productive mechanism in nature (green plants and algae) is photosynthesis, where ribulose-1,5-bisphosphate carboxylase-oxygenase (RuBisCo) converts CO₂ and water into C₃-carbohydrates by taking energy from light (Figure 1B) [22–24], here, the focus is on nonphotosynthetic enzymes. The conversion of CO₂ into formate anion (C₁ species, Equation (5), Scheme 1) is the simplest way in which nature captures CO₂ into reduced, highly energetic molecules, and formate dehydrogenases (FDH) are the main actors in this performance (Figure 1C) [25–29].

Formic acid, or its basic form formate, is a simple molecule for efficient hydrogen transport. Moreover, formate synthesis is the first step for obtaining other more complex and energetically enriched molecules [22,30]. Carbon dioxide and formic acid have similar formation energies (Figure 3A), with the most similar energetic states for C₁-carbon forms. This converts formic acid as the easiest starting point for obtaining other C₁ more reduced carbon species. Nevertheless, reaction 5, in the forward sense, is highly endergonic under physiological conditions. Indeed, the redox potential at pH 7 for CO₂ reduction to formate is −430 mV [31], while that of NADH is only −320 mV (Figure 3B). More importantly, for the reasons discussed below, this first step is kinetically, energetically, and economically expensive, and it becomes the bottleneck for obtaining value-added products in CO₂ regeneration. Several approaches for circumventing this issue are discussed below. However, other no less relevant problems arise when this reaction is performed for biotechnological purposes. FDHs from different organisms can solve these problems and incorporate formate anions into their biosynthetic routes. Thus, nature provides solutions to the previous difficulties. It is a researcher's task to adjust these solutions at laboratory and industrial levels for the benefit of society.

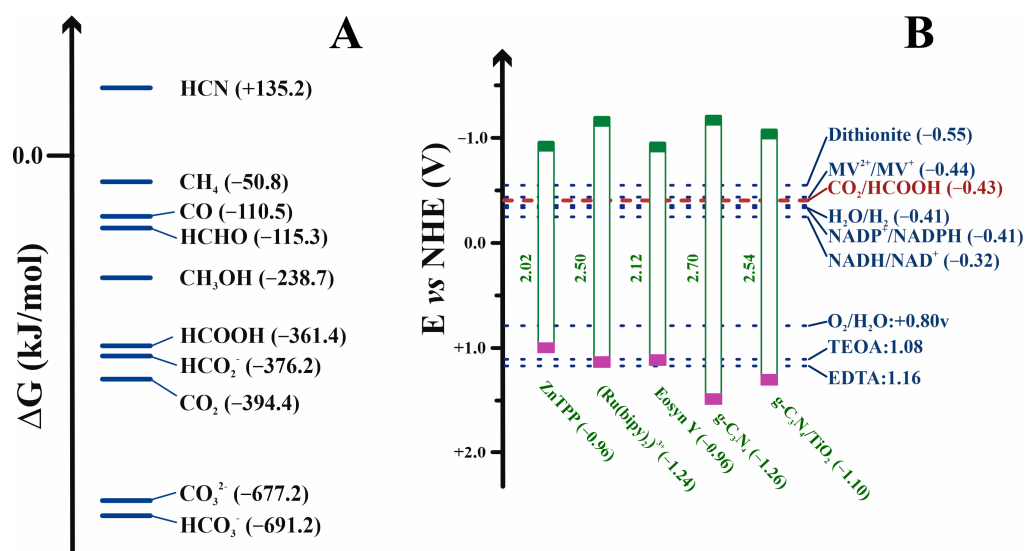


Figure 3. (A) Free formation energies of some C₁ compounds under standard conditions. (B) Redox potentials vs. the normal hydrogen electrode (NHE) at pH 7 of some relevant compounds, photosynthesizers, and sacrificial agents (see Section 7.4) commented in the text [32,33]. The redox potential values are shown between parentheses. The HOMO and LUMO energy orbitals for some dyes are indicated in green and purple, respectively.

In the following pages, the enzymes that participate in both processes of CO₂ capture using CA (CCUS processes) and FDH (formate synthesis) are described. How these natural systems circumvent the thermodynamics and kinetics problems derived from reactions 3 and 5 are also discussed, paying special attention to the state-of-the-art concerning the biotechnological applications of these enzymes for CCS and CCU technologies, as well as for reducing CO₂ to formic acid.

2. Carbonic Anhydrases: Efficient Devices for CO₂ Uptake

2.1. Classification and Structure of Carbonic Anhydrases

Carbonic anhydrases (EC 4.2.1.1) are found in all living kingdoms [18,34–36]. They catalyze the reaction of interconversion between CO₂ and HCO₃[−] (Equation (3)) in nature and are generally monomeric proteins with molecular weights roughly comprised between 30–50 kDa, depending on their class. CAs have been classified into: α -CAs (found in animal cells, algae, and eubacteria), β -CAs [37] (found in higher plants, microalgae, eubacteria, archaeobacteria, and fungi), γ -CAs [37] (algae), δ -CA [38–40] (found in the marine diatom

Thalassiosira weissflogii), ϵ -CAs [41] (found, for example, in the carboxysomal shell of *Halothiobacillus neapolitanus*), ζ -CA [42] (found in *Thalassiosira weissflogii*), η -CA [43] (found in *Plasmodium falciparum*), θ -CA [44], and ι -CA [45–47] (found in *Burkholderia territorii* and *Phaeodactylum tricornutum*, among other bacteria). CAs are zinc(II) enzymes, although some of them can contain other metal ions active in physiological roles: Cd(II) has been found in δ -CA, although its role is disputed [48,49]; Fe(II) is present in some γ -CAs growing under anaerobic conditions [50]; Co(II) has also been replaced in many α -CAs with excellent activities [51], although its original role has not been proven. Remarkably, a group of proteins called “COG4337” has been described as ι -CAs, and strikingly, those from the cyanobacterium *Anabaena* sp. PCC7120 and the chlorarachniophyte alga *Bigeloviella natans* display activity without any metal involved [52]. Table 1 lists the main features of some representative CAs.

Table 1. Features of representative carbonic anhydrases.

Type	Organism	MW (kDa)	Activity (WAU/mg)	Activity (k^{-1})	Metal Coordination	Ref.
α	Bovine	29.8	2540		Zn(II), 3 His, H ₂ O	[53]
	<i>Homo sapiens</i> (HCAI)	28.7	920	2.0×10^5		[54]
	<i>Homo sapiens</i> (HCAII)	29.1	8000	1.4×10^6		[54]
	<i>Persephonella marina</i>	26.9	1748			[55]
	<i>Thermosulfurimonas dismutans</i>	27.9	2032			[53]
	<i>Thermovibrio ammonificans</i>	25.3	1016			[53]
	<i>Bacillus halodurans</i>	37.0	3425			[56]
	<i>Sulphurhydrogenibium yellowstonense</i>	26.0	7254			[57]
β	<i>Bacillus subtilis</i>	37.0	714		Zn(II), His, 2 Cys	[58]
	<i>Acetobacterium woodii</i>	22.0	1814			[55]
	<i>Methanobacterium thermoautotrophicum</i>	19.9	580			[55]
	<i>Aspergillus fumigatus</i>	23.0	20			[59]
γ	<i>Geobacillus kaustophilus</i>	22.0	179		Zn(II) or Fe(II), 3 His, H ₂ O	[60]
	<i>Thermus thermophilus</i> HB8	24.3 ^a	0.9			[61]
	<i>Methanosarcina thermophila</i>	40.0	4872			[62]
	<i>Burkholderia pseudomallei</i>	28.2		5.3×10^5		[63]
	<i>Vibrio cholerae</i>	26.3		7.39×10^5		[63]
	<i>Porphyromonas gingivalis</i>	26.2		4.1×10^5		[63]
δ	<i>Thalassiosira weissflogii</i> , TWCA1	27.0		1.3×10^5	Zn(II), 3 His, H ₂ O	[64]
δ	<i>Emiliana huxleyi</i>	18.3		1.3×10^6 (b)	Zn(II), 3 His, H ₂ O	[65]
ζ	<i>Thalassiosira weissflogii</i> , CDCA1	69 ^a		1.5×10^6	Cd(II) or Zn(II), His, 2 Cys	[42,66]
η	<i>Plasmodium falciparum</i>	26.2		1.4×10^5	Zn(II), 2 His, Gln, H ₂ O	[43,67]
θ	<i>Thalassiosira pseudonana</i>	26.0	122		Cd(II) or Zn(II), His, 2 Cys	[38]
	<i>Phaeodactylum tricornutum</i>	31.1	30.9			[47]
ι	<i>Burkholderia territorii</i>	28.2		3.0×10^5	Cd(II) or Zn(II), His, 2 Cys	[46]
	<i>Anabaena</i> sp. PCC 7120 ^c	19.3		16.7		[52]
	<i>Bigeloviella natans</i> ^d	55.3		85.8		[52]

^a A trimer formed by three chains, R1, R2, and R3. ^b CA activity in the cytoplasm. ^c Activity of the GOG4337 recombinant protein all2909 (ref. [52]). ^d Activity of the GOG4337 recombinant protein Bn86287 (ref. [52]).

Mammalian CAs belong to the α -CA class. Depending on their location and primary sequences, several isoforms of human carbonic anhydrases (HCA) have been described,

with HCAII being the most studied and best-characterized CA. The HCAII tertiary structure (Figure 4A) consists of a unique domain containing ten β -strands that twist to form a β -sheet (eight of them organized in an antiparallel arrangement and the other two in parallel) [68,69]. Surrounding these β -sheets, up to eight other α -helices are located on the surface of the protein.

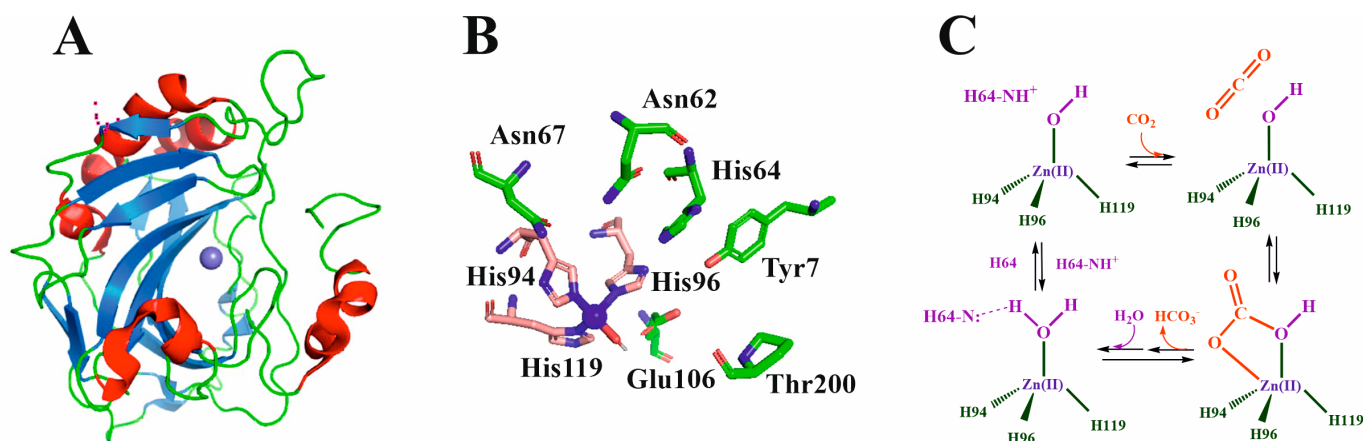


Figure 4. (A) Three-dimensional structure of human CA (5ZXW pdb code [68]). (B) Active center of the enzyme showing the most relevant amino acids (see text for details). (C) CA mechanism of action.

The structure of HCA is the most studied of all HCA isoforms, and its active site can be defined as a cone-shaped cleft, 15 Å deep, formed by a hydrophilic region (Tyr7, Asn62, His64, Asn67, Thr199, and Thr200) and a hydrophobic region (Val121, Val143, Leu198, Val207, and Trp209). Although the core of the active site in α -CAs is highly conserved, there is variability in the polarity and hydrophobicity of its periphery [70]. The catalytic Zn(II) ion is located in a deep-centered slot that is accessible to the solvent. The Zn(II) ion is coordinated to three histidine residues (Figure 4B). While His94 and 96 are coordinated through their N ϵ 2 imidazole nitrogen to the metal ion, His119 is bound through its N δ 1 imidazole nitrogen (HCAII numeration) [48]. A water molecule completes the tetrahedral coordination of Zn(II). A second coordination sphere is formed by amino acids that are not directly coordinated with the metal ion but are essential in the catalytic process. Through the formation of a network of hydrogen bonds, residues Tyr7, Asn62, His64, Asn67, Glu106, Thr199, and Thr200 stabilize the mediator species in such a way that the reaction can occur.

The HCA family includes a subclass of three noncatalytic isoforms (HCAs VIII, X, and XI) called CA-related proteins (CA-RPs), whose classification is based on their sequence. The noncatalytic behavior is due to the absence of one or more histidines that coordinate the Zn(II) ion of a catalytic HCA isoform. For instance, in HCA-RP VIII, the Zn-coordinating His94 (HCA II numbering) is replaced by an arginine (Arg116, according to HCA-RP VIII numbering). This residue avoids CO₂ hydration in the first step of CA catalysis. [71]. Although the biological functions of CA-RPs have not been defined, these isoforms are of high interest in different scientific research fields. Recently, the X-ray crystal structure of only one HCA-RP (HCA-RP VIII) was determined [72]. HCA-RP VIII is expressed in the cerebellum [73] and has been identified as a binding partner for the inositol 1,4,5 triphosphate (IP3) receptor type [74]. It should be mentioned that the stability and structures of other HCAs such as HCA III or extracellular HCAs (i.e., IV, VI, IX and XIV) have not been as extensively studied as HCAs I and II. For instance, the bovine CA III showed a similar unfolding profile to that of HCA II, providing a molten globule intermediate and an unfolded state at a C_m of 2.6 M guanidinium chloride (GuHCl) concentration [75]. On the contrary, the isoforms VIII, X, and XI showed two distinct transitions, and their sensitivity to guanidinium chloride chemical denaturalization was higher than that of HCA II (C_m 0.4 M for HCA-RP VIII and 0.9 M for HCA II) [71].

2.2. CA Mechanism of Action

Carbonic anhydrase accelerates reaction 3 by more than six orders of magnitude (Table 1) with respect to its rate without the biocatalyst [18,76] (from ca. $3.6 \times 10^{-2} \text{ M}^{-1} \text{ s}^{-1}$ to $1.0 \times 10^6 \text{ M}^{-1} \text{ s}^{-1}$ in the absence and presence of CA) [17], allowing for the reaction to take place under physiological conditions. Otherwise, CO_2 cannot be assimilated as bicarbonate by living organisms.

The kinetic parameters of CA can be determined by measuring hydratase activity (Equation (3), Scheme 1). The method basically consists of saturating a buffered water solution (typically Tris.HCl 0.020 M, pH 8.3) kept in an ice bath with CO_2 at a fixed high pH value, then adding CA, and measuring the time that the solution takes to reach a low pH value (generally ca. 6.3) because of the conversion of carbon dioxide to hydrogen carbonate anion that takes the capture of protons (Equation (3)) and hence lowers pH. The Wilbur–Anderson hydratase activity unit [77] is calculated as the ratio $((t_0 - t)/t_0)/(\text{mg enzyme})$, where t_0 and t are the measured times that an indicator present in the solution shifts its color for the control (without CA) and the sample (in the presence of CA), respectively. This method provides a qualitative measurement of CA activity, but it is not strictly transferable from one set of experiments to another [78]. This method depends on the degree of saturation of CO_2 (which in turn changes with temperature and the time at which CO_2 is bubbled), the nature of the buffer, its concentration, and its initial pH value. Together, these units are roughly—not strictly—comparable. Most classes of CAs also exhibit esterase activity (δ -CA lacks esterase activity), whose measurement is more direct and contrastable [79]. This is generally performed by the hydrolysis of *p*-nitrophenyl acetate, which releases free *p*-nitrophenol, with maximum absorption at 400 nm, which is easily measurable [80]. Accurate experimental requirements for the latter experiments (concentration of the reactants, enzyme, and pH) are easily reproducible and should depend only on contrastable and exchangeable conditions.

The mechanism of action of CA has been deciphered [81] and basically consists of two stages (Figure 4C). In the first phase, a CO_2 molecule, partially stabilized by interactions with groups of the enzyme active site, reaches the active center, and subsequently, the hydroxide group bound to the Zn(II) ion attacks the CO_2 carbon atom (nucleophilic attack). Then, hydrogen carbonate is formed, and a water molecule replaces it through Zn(II) coordination. In the second step, a proton is transferred from the Zn(II)-coordinated water molecule to the solvent. This is the rate-limiting step. His64 is the amino acid responsible for accepting this proton, which is finally transferred to the bulk solvent. Due to this mechanism, at neutral/soft acid pH levels, the rate of $\text{CO}_2/\text{HCO}_3^-$ conversion is enhanced by more than six orders of magnitude, making life possible. At pH values higher than 9.5, a direct hydroxide attack can form carbonate anions at rates comparable to those performed by CA at neutral pH values.

3. Formate Dehydrogenases: Natural Machines for Reducing CO_2

The specific reduction in carbon dioxide converts it into formate/formic acid (Equation (5), Scheme 1). This C_1 metabolism reaction occurs in hydrogenotrophic methanogens (*Euryarcheota*) and autotrophic acetogens (bacteria) and is carried out by the enzyme formate dehydrogenase (EC 1.17.1.9) [24,82–85]. Energy and high reduction power (i.e., a cofactor in the NAD(P)H form) are required to perform this process. In contrast, formate dehydrogenases also catalyze the backward reaction (5), oxidizing formate anions and obtaining energy from them. FDHs are divided into two main groups: nonmetal- or NAD-dependent FDHs, and metal-dependent FDHs. The structure, nature, and activity of these two FDH sets are completely different. More importantly, the ability of these two groups to reduce CO_2 is manifestly diverse, as discussed below.

3.1. Metal-Independent/NAD-Dependent FDHs

Metal-independent FDHs belong to the family of D-specific 2-oxoacyl dehydrogenases. They are present in bacteria, yeast, plants, and mammals, are globular proteins with ca.

350–400 amino acids, depending on the species, and usually form homodimers [25,26,86]. Each FDH monomer contains two domains, one destined to allocate the substrate (the catalytic domain) and another pocket that allows for the binding of the NADH cofactor. The function of metal-independent FDHs is generally associated with mechanisms for obtaining energy from the oxidation of formate anions in methanogenic pathways (backward reaction 5), that is, these FDHs are machines efficient in catalyzing backward reaction, although much less effective in performing CO₂ reduction [25,26,84,87]. The cofactor of all known metal-independent FDHs is NAD(P)⁺/NAD(P)H, and, consequently, they are also called NAD(P)-dependent FDHs. FDH from the methylotrophic yeast *Candida boidinii* (*Cb*) is the most studied and best characterized nondependent FDH, since it was the first FDH expressed in *Escherichia coli*, it is commercially available and relatively inexpensive [88,89]. As shown in Figure 5A, the enzyme (364 amino acids) consists of 15 α -helices and 13 β -strands [90]. There is a deep groove between both domains that allows both the substrate and the cofactor to be bound in this cavity with short contacts between them. Figure 5B displays the active site of the enzyme, including both the NAD⁺ binding site and the most significant amino acids concerning catalytic properties. NAD⁺ cofactor strongly binds the protein through its adenine, ribose, and phosphate moieties in such a way that it can only be removed by extensive washing with, for instance, 0.2 M sodium chloride. Formate binds closely to the nicotinamide group through the positively charged residue Arg258. The mechanism of formate oxidation (Figure 5C) consists of hydride transfer to the nicotinamide oxidized group and the release of the CO₂ formed. The intermediated anion is stabilized by an arginine (Arg258 *Cb*FDH numbering). The groove where the nicotinamide group and the formate anion are located is hydrophobic; hence, the hydride anion cannot interact with the solvent, so the reaction can take place. Both high formate and NAD⁺ strong binding with positive residues of the enzyme (Gln287, His311) reduce the ability to exchange the products of the reaction. In turn, this is one of the main factors that decrease the efficient recycling of the cofactor, making these FDHs, in general, not excellent biocatalysts for CO₂ reduction.

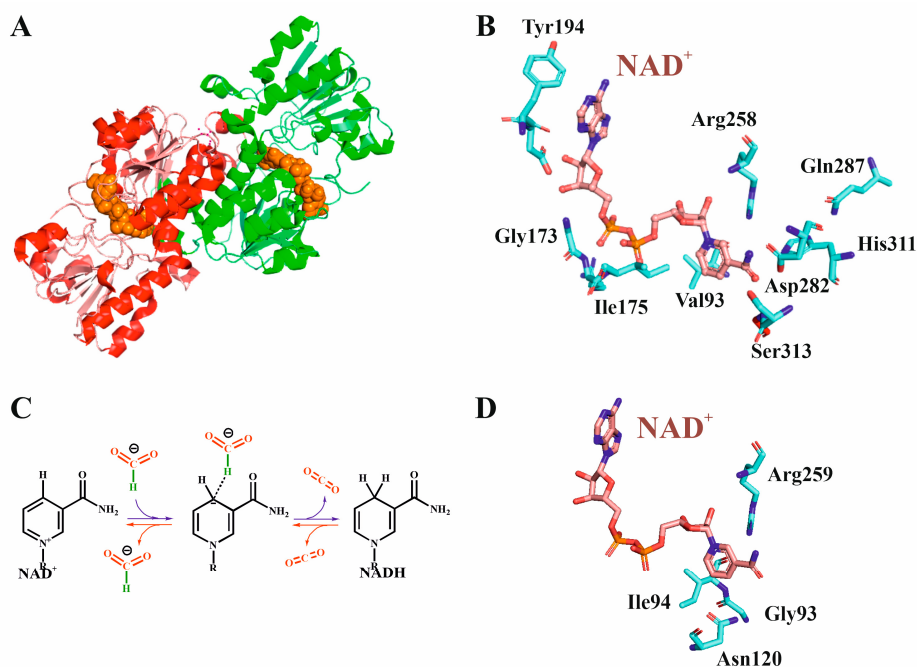


Figure 5. (A) Three-dimensional structure of *Cb*FDH with NAD⁺ and azide anion (6D4C pdb code [90]). The two dimers are colored in red and green, while the NAD⁺ cofactor is colored in

orange. (B) Active center of *Cb*FDH showing the arrangement of the cofactor (pink) as well as the most relevant amino acids closest the NAD^+ , as discussed in the text. (C) Mechanism of action of NAD^+ -dependent FDHs for formate oxidation (blue arrows) or formate reduction (inverse sense, red arrows); the NAD nicotinamide group in its oxidized and reduced state is displayed. (D) Active center of *Ct*FDH (6T8Z pdb code [91]) showing the four residues mutated in both works cited in the text [92,93], the numbering corresponding to *Ct*FDH.

With wild-type *Cb*FDH being not ideal for forward reaction 5 [94,95], specific mutants increase its capability to capture CO_2 . For instance, the double mutant V120S-N187D was shown to substantially increase CO_2 reduction [96]. Other FDHs have k_{cat} and k_{M} values that are higher than wild-type *Cb*FDHs and its mutants (Table 2). Choe et al. studied *Thiobacillus* sp. KNK65MA FDH (*Ts*FDH) and concluded that it presented 84-fold higher catalytic efficiency for CO_2 reduction than *Cb*FDH [67]. NAD-FDH mutants with higher efficiency have been successfully designed [89,96]. Binay and coworkers obtained and purified mutants of *Candida methylica* (*Cm*FDH) and another four mutants from *Chaetomium thermophilum* (*Ct*FDH) in amino acid positions close to the cofactor binding [92,93]. The highest activity was found for the Asn120Cys mutant in *Ct*FDH, for which the k_{cat} value increased 6.5-fold, the same increment observed for the k_{M} value, which indicated that the efficiency of CO_2 reduction was due to the lower affinity for the substrate, that is, for the ability to release formate. As observed in Figure 5D, Asn120 is located close to the nicotinamide NAD group; specifically, the amide nitrogen of Asn120 is as close as 3.8 Å from the nicotinamide ring. Its mutation by a smaller cysteine residue introduces more space in the active site and, consequently, higher conformational flexibility, which facilitates the release of formate anion [92]. His96 interacts with the hydrogen carbonate anion stabilizing the hydride in the transition state, as confirmed by molecular dynamics performed on the double mutant. Based on kinetic and molecular dynamic studies, the authors concluded that subtle structural changes around the Asn120 position allowed for the location of two molecules of hydrogen carbonate instead of one of formate, favoring the CO_2 forward reaction taking place. On the other hand, replacing key residues G93H/I94Y in *Cm*FDH, located in the catalytic pocket of FDH, increased the catalytic efficiency ($k_{\text{cat}}/k_{\text{M}}$) of the wild-type protein 5.4-fold for the reduction of HCO_3^- . Here, K_{M} values do not vary significantly, while $k_{\text{cat}}/k_{\text{M}}$ does. The authors suggested that there is a reorganization in the active site that enlarges the space and allows for the reactant (carbonate anion) to adopt a better orientation for catalysis, and so it becomes easier for the HCO_3^- to reach the nicotinamide ring and the reaction is produced in a faster way [93].

Table 2. Kinetic and thermodynamic parameters of most relevant FDHs, classified into their two groups. Molecular weights and pH conditions in which these conditions were determined are also indicated. For metal-dependent FDHs, the types and number of their subunits as well as the cofactors of their active centers are also indicated.

A. NAD-Dependent FDHs.												
Organism	MW (kDa) ^a	Formate Oxidation				CO ₂ reduction				ref		
		pH	k_{cat} (s ⁻¹) (U mg ⁻¹) ^b	K_M (mM)	k_{cat}/K_M (mM ⁻¹ s ⁻¹)	pH	k_{cat} (s ⁻¹) (U mg ⁻¹)	K_M (mM)	k_{cat}/K_M (mM ⁻¹ s ⁻¹)			
<i>Myceliophthora thermophila</i>	42	10.5	0.32	7.2	0.04	7.0	0.10	0.43	0.23	[97]		
<i>Ancylobacter aquaticus</i>	45	6.0	(21.6)			6.0	(23)		4.5	[98]		
<i>Candida boidinii</i>	41	7.0	1.081 (6.1)	8.55	0.13	5.5	0.015	2.6	0.006	[98]		
<i>Thiobacillus</i> sp. KNK65MA	45	6.5	1.769 (10.9)	16.24	0.11	5.5	0.32	0.95	0.34	[98]		
<i>Candida methylica</i>	42	8.0	1.31 (13.2)	7.01	0.19	8.0	0.008	0.078	0.01	[97]		
<i>Chaetomium thermophilum</i>	45	5.0	2.04(3.1)	3.30	0.62	5.0	0.023	3.29	0.069	[93,99]		
<i>Ceriporiopsis subvermisporea</i>	40	6.5	(1.3)			6.0	(0.8)			[98]		
<i>Moraxella</i> sp. C-1	45	5.5	(14.3)			5.5	(2.8)			[98]		
<i>Paracoccus</i> sp. 12-A	45	5.5	(12.2)			5.5	(6.5)			[98]		
B. Metal-Dependent FDHs												
Organism	MW ^a (kDa)	Subunits ^b	Cofactors	Formate Oxidation				CO ₂ reduction				
				pH	k_{cat} (s ⁻¹) (U mg ⁻¹) ^a	K_M (mM)	k_{cat}/K_M (mM ⁻¹ s ⁻¹)	pH	k_{cat} (s ⁻¹) (U mg ⁻¹) ^a	K_M (mM)	k_{cat}/K_M (mM ⁻¹ s ⁻¹)	
<i>Syntrophobacter fumaroxidans</i> FDH-1	175	($\alpha\beta\gamma$)2	W, SeCys 4 [Fe2S2] [Fe4S4]	7.0	(700)	0.04	-		282 (900)	-	-	[100]
<i>Syntrophobacter fumaroxidans</i> FDH-2	125	($\alpha\beta$)2	W, SeCys 2 [Fe2S2]	7.0	(2700)	0.01	-		282 (89)	-	-	[100]
<i>Desulfovibrio desulfuricans</i>	135	($\alpha\beta\gamma$)3	Mo, SeCys 2 MGDs 4 c-heme 2 [Fe4S4]	8.0	543	0.0571	9526	7.0	46.6	0.0157	2968	[101]
<i>Escherichia coli</i> FDH-H	79	$\alpha\beta$	Mo, SeCys 2 MGDs 1 [Fe4S4]	7.5	2800	26	107.7	7.5	1.0	8.3	0.12	[102,103]
<i>Desulfovibrio vulgaris</i> Hildenborough	97.4	($\alpha\beta\chi$)3	Mo, SeCys 4 [Fe4S4] 4 c-heme	7.6	1310 (77)	0.017	77.06		315(1.0)	0.42	750	[104,105]
<i>Acetobacterium woodii</i>	169	($\alpha\beta$)3	Mo SeCys [4Fe-4S])	7.0	(600)	1.0	-	7.0	372 (132)	3.8	97.9	[106]

Table 2. Cont.

B. Metal-Dependent FDHs												
Organism	MW ^a (kDa)	Subunits ^b	Cofactors	Formate Oxidation				CO ₂ reduction				ref
				pH	$k_{cat}(s^{-1})$ (U mg ⁻¹) ^a	$K_M(mM)$	k_{cat}/K_M (mM ⁻¹ s ⁻¹)	pH	$k_{cat}(s^{-1})$ (U mg ⁻¹) ^a	$K_M(mM)$	k_{cat}/K_M (mM ⁻¹ s ⁻¹)	
<i>Cupriavidus necator</i>	178	($\alpha\beta\gamma$) ₃	Mo, 4 [Fe ₄ S ₄] FMN 3 [Fe ₂ S ₂]	7.0	140	0.082	1707	7.0	11	2.7	4.07	[107]
<i>Rhodobacter capsulatus</i>	180	($\alpha\beta\gamma$) ₂	Mo, 4 Fe ₄ S ₄ , 1 Fe ₂ S ₂ , 2 MGD, FMN	5.0	36.5	281	0.13	7.7	1.48	-	-	[108,109]
<i>Pseudomonas oxalaticus</i>	315	-	Mo, 2 FMN	-	-	0.135	-	6.2	3.0	40	0.075	[110,111]
<i>Clostridium ljungdahlii</i>	80	-	W Cys 2 MGD [Fe ₄ S ₄]	9.0	14.77	1.40	10.55	7.0	0.73	7.27	0.17	[112,113]
<i>Clostridium autoethanogenum</i>	74		W 2 MGD [Fe ₄ S ₄]	9.0	1.04	4.51	0.231	7.0	4.00	23.15	0.17	[113]
<i>Clostridium coskatii</i>	62		W 2 MGD [Fe ₄ S ₄]	9.0	0.62	5.57	0.111	7.0	5.62	59.65	0.094	[113]
<i>Clostridium ragsdalei</i>	74		W	9.0	11.88	44.83	0.265	7.0	3.28	31.20	0.11	[113]
<i>Desulfovibrio gigas</i>	121	($\alpha\beta$) ₂	W, SCys 4 [Fe ₄ S ₄]	8.0	(34.1)	-	-	-	-	-	-	[114]
<i>Moorella thermoacetica</i>		($\alpha\beta$) ₂	W, SeCys 4 [Fe ₄ S ₄]	7.5	(1100)	-	-	-	-	-	-	[115]

Sub-Table (A): ^a Molecular weight of the monomer (most FDHs are associated with dimers). ^b Data between parenthesis refer to the activity in units (μmol min⁻¹) of formate/CO₂ oxidized/reduced per mg of FDH; Sub-Table (B): ^a Molecular weight of the monomer. ^b The number refers to the number of subunits, that is, the composition of dimers or trimers of each FDH. MGD: molybdopterin guanine dinucleotide. FMN: flavin mononucleotide.

3.2. Metal-Dependent FDHs

FDHs containing metals constitute the other large set of FDHs [28,84,116,117]. All metal-dependent FDHs catalyze the interconversion between formate and CO₂. The sense of the reaction (Equation (5), Scheme 1) depends on the external conditions. In general, in biological conditions, formate oxidation (backward reaction 5) is favored, and thus, some organisms obtain their energy from this exergonic reaction. However, metal-dependent FDHs can also catalyze CO₂ reduction, and most of them do so, although to a different extent. There are a small number of microorganisms (hydrogenotrophic methanogens, *Euryarcheota*, and autotrophic acetogens, bacteria) that use FDH not for generating energy (i.e., not for oxidizing formate), but for using C₁ carbon species as a primary source of their carbon metabolism [116,117]. All these microorganisms have metal-dependent FDHs.

This type of FDH is much more complex than NAD-FDHs. Indeed, they have more than 700 amino acids arranged in different domains that, in turn, contain several cofactors and/or metal centers such as ferredoxins, heme groups, flavin mononucleotides, etc., depending on the species [118]. For instance, FDH N from *E. coli* comprises three domains (Figure 6A): the α -domain, which contains the Mo cofactor (see below) and one [4Fe-4S] cluster; the β -domain with four [4Fe-4S] centers; and the γ -domain, with two *b*-hemes (Figure 6B) [119]. These enzymes receive the electrons from these metal clusters and not necessarily from NADH (although, in some cases, NADH can also be the cofactor), and thus, these FDHs are called nondependent NADHs. Excellent reviews describing the three-dimensional structures of these FDHs, their metal centers, their functions, and the biotechnological achievements of these enzymes have been published [28,84,116,117].

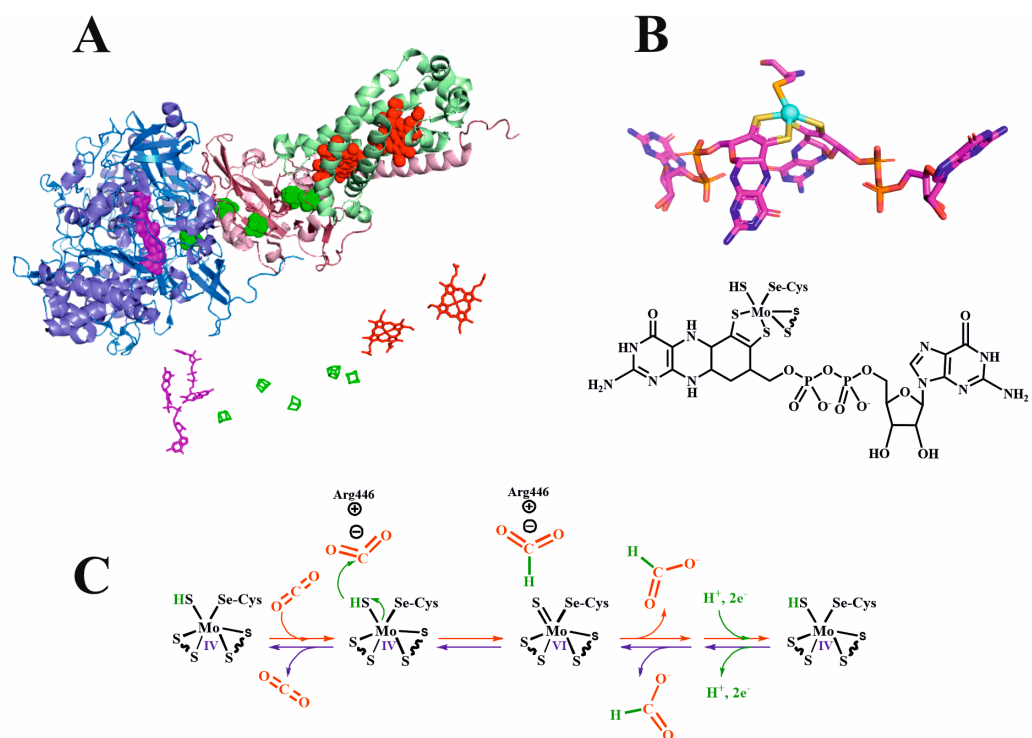


Figure 6. (A) **Up:** three-dimensional structure of FDH N from *E. coli* (1KQF pdb code [119]). The three polypeptide chains are shown in blue (chain A), pink (chain B), and green (chain C); the molybdopterin (in chain A), the five F₄S₄ clusters (in chains A and B), and the two heme (chain C) metal cofactors are displayed in pink, green, and red, respectively. (A) **Down:** a detailed picture of the cofactors at the same scale. (B) **Up:** graphic of the molybdenum cofactor in FDH N from *E. coli*. (B) **Down:** chemical formula of the Mo cofactor. (C) FDH mechanism of action for CO₂ reduction (red arrows) or formate oxidation (inverse sense, blue arrows) [27]; the direction of the reaction depends on the experimental conditions (see text for details).

Importantly, despite their heterogeneity, these FDHs share common features concerning the active center. They all contain a molybdenum or a tungsten metal ion bound to two dithiolene atoms provided by two pyranopterin guanidine dinucleotides, a sulfur or selenium donor atom, and a disulfide anion (Figure 6B) [104]. CysSe residue is present in both Mo- and W-FDHs; hence, it is not specific to a determined metal. On the other hand, according to kinetics parameters (k_{cat} and k_M) the presence of CysSe instead of the native amino acid cysteine, is not crucial for FDH activity.

Metal-dependent FDHs catalyze Equation (5) (Scheme 1) in both directions; however, unlike wild-type NAD-dependent FDHs, they are much more efficient in catalyzing forward Equation (5) than the latter, clearly evident in Table 2. This table presents the kinetic parameters, as well as the composition of the active site and cofactors, specifically for metal-dependent FDHs. Hence, for biotechnological applications, that is, for reducing CO_2 , metal-dependent FDHs are much more active and, hence, more attractive than non-metal FDHs.

What are the key factors that enable metal-dependent FDHs to reduce CO_2 efficiently? These issues have been extensively studied in the literature [26–29,84]. Here, some notes on the crucial aspects are commented upon. First, the existence of different redox centers acts as a corridor for efficient electron transfer toward the CO_2 molecule. Second, and importantly, there is presence of a sulfido group that accepts a hydride anion (see Figure 6C). It is well known that, in Mo/W enzymes, a sulfido group accepts a hydride. In FDHs, metal oxidation states change from IV to VI; when they are in a reduced state, ligands tend to be protonated, while tending to deprotonate in the Mo/W(VI) oxidation state. Thus, metal sulfido can act as a donor/acceptor hydride. Indeed, spectroscopic studies are consistent with the transfer of a hydride from a sulfur atom [27,84,116,118]. On the other hand, there is no evidence of the direct coordination either of CO_2 or formate directly to the metal center. Altogether, this facilitates the acceptance/donation of a hydride directly towards the CO_2 carbon, which has an electronic net deficiency, and so is prone to attack by anions. It is also remarkable that the tungsten or molybdenum metal ion are indistinguishable concerning the catalytic activities while the presence of selenium cysteine does not appear to be relevant in CO_2 reduction.

Several schemes have been proposed for the mechanism of action of these FDHs [120,121]. However, it is robustly supported that CO_2 , rather than HCO_3^- , is the substrate of FDHs for forward Equation (5) [104,122]. This reaction takes place by abstracting (or adding, reverse reaction) a hydride anion, without the intervention of any oxygen atom [123]. Concordantly, the mechanism of action should consider these two fundamental facts. Moura et al. proposed a mechanism for the forward reaction in which the reduced CysSH coordinated to the Mo ion attacks the carbonyl atom and hydride transfer takes place (Figure 6C) [27]. The formate anion is then stabilized by the positive charge of Arg446 (FDH N from *E. coli* numeration), and afterward, when the protein is again reduced by the other cofactors and by the addition of another hydride to the coordinated Cys, the formate is released. In contrast, the reverse reaction is also produced by the opposite hydride attack from the formate anion on the same Cys (in this case, oxidized).

4. Improving CA Performance: Enzyme Immobilization

For industrial and biotechnological applications to be profitable, enzymes must be as stable and reusable as possible. CA and FDH in solution, like all soluble proteins, behave as solutes with full mobility in the solvent. Although an aqueous medium is, in general, the most suitable for enzyme action, the stability and activity of enzymes in this medium usually decrease rapidly. In addition, the use of enzymes in solution is always constrained by strict pH and temperature conditions. More importantly, enzymes in aqueous solutions can only be applied in the cycle of a specific reaction; hence, their applicability at the industrial level is highly limited. In contrast, enzymes immobilized on solid or gel supports extraordinarily increase their stability, amplifying the range of action of the biocatalyst conditions and the possibility of using more drastic, usually more efficient,

reaction conditions (for instance, increasing temperature) [124–126]. This immobilization allows for their easy separation from reactants and products, and, consequently, enzymes in this form can be reused for posterior cycles [127,128]. This drastically reduces the cost of the whole process, regardless of the biotechnological industrial reaction. Both CA and FDH have been immobilized on different supports. Based on the immobilization method, the following approaches can be considered: physical adsorption, covalent binding, entrapment, encapsulation, and crosslinking.

Excellent reviews on CA and FDH immobilization have recently been published [129–139]. Here, some illustrative cases regarding the relevance of immobilization in enzyme stabilization are highlighted according to the immobilization method (Figure 7). Because all the examples provided in Section 6 are related to CO₂ reduction by immobilized FDH, here, the focus is on CA immobilization. Table 3 summarizes relevant studies on CA immobilization using these methods.

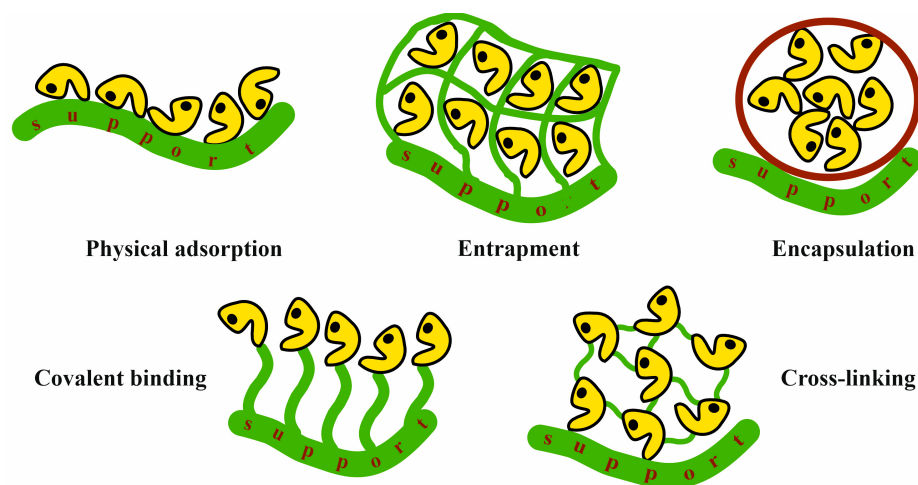


Figure 7. Different methods of enzyme immobilization.

Table 3. Immobilization of Carbonic Anhydrase using different methods.

Method	Support	Immobilization Conditions	Main Results	Ref
Physical adsorption	Mesoporous aluminosilicate	Phosphate buffer 100 mM, pH 7, 0.2 mL of enzyme (1 mg/mL) in 10 mg of support for 6 hr at 120 rpm.	CA loading up to 3 mg/mL optimal for CA activity. Carbonation activity was 55% (8 cycles of reuse)	[140]
	Carboxylic acid group-functionalized mesoporous silica (FMS)	0.6–2.0 mg FMS with functional groups incubated with 70–1000 IL of 2.0 mg/mL BCA II in water (pH 6.5) for 2 h at 21 °C under 1200 rpm shaking.	High protein loading density 0.5 mg of protein/mg. High stability and activity (95.6%) of the immobilized enzyme	[141]
	Silver nanoparticles amine-functionalized mesoporous SBA-15	10 mg dispersed separately in 2 mL free HCA in buffer (3 mg/mL HCA in 100 mM sodium phosphate, pH 6.4) followed by incubation at 25 °C with shaking for 4 h.	The activity of the materials was ~25-fold higher than the activity of the free HCA for converting CO ₂ to CaCO ₃ after 30 cycles.	[142]
	Mesoporous silica nanoparticles with polydopamine (PDA) and polyethyleneimine (PEI)	5 mg FDH/1.6 mg CA added to 2 mL of phosphate buffer (0.05 M, pH 6). PDA/PEI-mSiO ₂ (0.05 g) and 5 µL GA aqueous solution (20 wt%) were added (30 °C for 24 h).	Optimal FDH and CA concentration was 2.5 and 0.8 mg/mL respectively, (specific activity of 0.045 mM/h/mg). 10 reuse cycles with 86.7% of activity.	[143]
	Palmityl-substituted sepharose 4B	Denatured Bovine CA solution by incubation for 15–120 min (58–65 °C), Tris-sulfate buffer, pH 7.5 (0.0125–0.5 mg/mL).	Denaturation and renaturation processes of the enzyme in the matrix. After renaturation at 60 °C, 85% of immobilization was achieved (70% of the original activity after 1 h).	[144]
	Silica or titania particles	The clarified lysates expressed BCA-peptide (0.088 mg protein) were incubated with 30 mg silica or 20 mg titania for 10 min with shaking at room temperature in 25 mM sodium phosphate buffer (pH 6.5).	After 10 days, 90 ± 4% and 95 ± 3% of silica and titania particles' original activity remained. Immobilized BCA-peptide fusion protein shows ~95% of its residual activity with up to 5 cycles of reuse.	[145]
Entrapment or encapsulation	Porous polypropylene and a non-porous polydimethoxysilane hollow fiber membrane	Application of CA in situ to the shell side surface of each fiber.	CO ₂ absorption into K ₂ CO ₃ increased approximately three-fold when CA is adsorbed onto the Porous polypropylene or non-porous polydimethoxysilane PDMS membrane surface	[146]
	Magnetic nanoparticles (sol-gel ferria hydrosol)	200 mL of ferria hydrosol mixed with CAB solution (10 g/L, 0.05 M Tris, pH 7.4) and dried at 20 °C.	Large thermal stability of CA immobilized, active up to 95 °C	[147]
	Polyurethane foam (PUF)	1 mL of CA (1 g/L) or whole-cell solution in 20 mM Tris-sulfate buffer (pH 8.3) was added to a 50 mL containing 1 g of prepolymer. The swelling of PUF continued for 30 min, and then an additional 10 min was allowed for curing.	An immobilization efficiency of 3.4%, 16-fold higher than that for free enzymes. The reusability of the immobilized whole-cell catalyst shows no apparent decrease in activity after 9 reuses. The rate of CO ₂ capture was accelerated by 80%.	[148]
	Ni-based MOFs (Ni-BTC) nanorods,	100 mL cell lysate of His-HCA II (0.4 mg protein) and 900 mL Milli-Q water were incubated with Ni-based MOFs.	His-HCA II from cell lysate obtained an activity recovery of 99%. After storing for 10 days, the immobilized His-HCA II maintained 40% activity (free enzyme lost 91% activity). Immobilized His-HCA II retained 65% (8 cycles).	[149]
	Supported ionic-liquid membranes (SILMs)	0.25 mg CA/g IL.	SILMs resent permeability (PCO ₂ = 733.73 barrier) at high temperatures (up to 373 K) and a good transport selectivity towards CO ₂ against N ₂ .	[150]
	Cholinium-based ionic liquids	0.1 mg CA/g IL	CA samples promote an enhancement of 63% in the carbon dioxide transport rate.	[151]

Table 3. Cont.

Method	Support	Immobilization Conditions	Main Results	Ref
Covalent binding and cross-linking	Mesoporous SBA-15 surfaces covalently functionalized with amines	HCA immobilization was achieved by mixing 10 mg of amine-functionalized SBA-15 with a 0.1% glutaraldehyde (GA) solution (50 mM sodium phosphate, pH 8.0, 1 h). The product was treated with free HCA (3 mg/mL HCA, pH 7.0), incubated and shaking (1 h, 25 °C).	Immobilized HCA retained activities after long-term storage, exposure to high temperatures, and reuse (40 cycles). CO ₂ capture efficiency of immobilized HCA was 36 times higher than that of free HCA, 75% of the enzymatic activity was retained after 40 cycles.	[152]
	Chitosan	CA liquid was slowly added into the pH 5 chitosan solution with continuous stirring. Ratios of 1:0.05–1:2 chitosan:CA (g:mL)	Textile packing with covalently attached enzyme aggregates retained 100% of the initial 66.7% CO ₂ capture efficiency over 71 days and retained 85% of the initial capture efficiency after 1-year of ambient dry storage.	[153]
	Polyethyleneimine and polydopamine in MOF 808	PBS buffer (pH 8, 10 mM), PEI/PDA-MOF-808 or PDA-MOF-808 was disseminated. 200–400 µL of CA solution (1 mg/mL, Milli-Q) and 10 µL of aqueous solution of GA (0–25 wt%) were added and shaken for a while at 28 °C.	CaCO ₃ produced by CA@PEI/PDA-MOF-808 was 11.0-fold and 2.5-fold higher than free CA and PEI/PDA-MOF-808, respectively. After 8 consecutive rounds, the total production of CaCO ₃ by CA@PEI/PDA-MOF-808 was 92-fold higher than free CA.	[154]
	Alumino-siloxane hybrid aerogel beads	100 mg of Al/Si-NH ₂ beads were treated with 0.5% GA for 1 h. 4 mL of free BCA in the buffer (1 mg/mL 100 mM phosphate buffer) was added to GA-treated Al/Si-NH ₂ beads and stirred (3 h, 25 °C) for BCA immobilization.	Free BCA retained 70% of its maximum activity (immobilized BCA, 88%). Free BCA and BCA-Al/Si-NH ₂ retained 80% of the activities, after ten days. BCA-Al/Si-NH ₂ retained 89% of their enzyme activity up to 10 cycles	[155]
	Amine-functionalized by co-deposition of polydopamine (PDA) and polyethyleneimine (PEI)	Co-deposition of PDA and PEI and CA covalently anchored on the surface via GA. Surface was treated with a mixture of PDA (2 mg/mL) and PEI, pH 8.5. Amine-functionalized micro-reactor surface was contacted with a mixture of CA and GA (concentration 2.0% (v/v)).	A steady CO ₂ absorption rate for several hours and good reusability of the immobilized enzyme which maintains its original absorption performance after 10 cycles of operation.	[156]
	Polypropylene hollow fiber membranes using GA-activated chitosan	Aminated knitted hollow fiber membranes mats (204 cm ² fiber surface area) were incubated in 5% GA in 100 mM phosphate buffer, pH 8.5 for 1 h under constant rocking at room temperature.	Chitosan/CA coated fibers exhibited accelerated CO ₂ removal in scaled-down gas exchange devices in buffer and blood (115% enhancement vs. control, 37% enhancement vs. control, respectively).	[157]
	Magnetic Cross-Linked Enzyme Aggregates (CLEAs) to bovine carbonic anhydrase (BCA) and magnetic nanoparticle	- Adsorption: the NPs suspension (5 and 2 mg solids) added to 1 mL of 10 mM PBS (pH 7.4), 10 g/L BCA. - Precipitation: NP-enzyme suspension added dropwise to 9 mL containing the precipitating agent (pH 7.4), 1 h or 0.5 h under mixing. - Cross-linking: 25% vol GA added. The system kept under mixing for 3, 16 and 22 h. - CLEAs separated by MF.	BCA-CLEAs can increase the CO ₂ absorption rate concerning the one observed in the same reactor filled with only alkaline solvent. Biocatalyst reusability analysis showed that BCA-CLEA retained 95% of its initial activity after five CO ₂ absorption tests at 1000 mg BCA CLEA/L and as many liquid-solid separation steps by membrane filtration.	[158]
	Geopolymer micro-spheres (GMS) and covalent attachment by GA	GMS were introduced into a Tris-HCl buffer solution (0.05 M, pH 8.0) with CAs solution (1 mg/mL) in a 50 mL centrifuge tube. After shaking and reacting, the GA was added for the cross-linking between the GMS and CAs.	K_{cat}/K_m values were 61.50 and 12.36 M ⁻¹ s ⁻¹ for the immobilized and free CAs, respectively. At 60 °C, free CAs were inactivated (immobilized CAs kept 34.8% activity). Immobilized CAs maintain 68.73% of activity (8 cycles).	[159]
	Microbial transglutaminase (MTG) acts as a “cross-linking medium”	Iso-peptide bond between glutamine and the primary amine group of a lysine in artificial peptide tags. Equimolar amounts F-CA and M-FDH were added, and a metal constant temperature oscillator was used for the cross-linking reaction (reaction time 12 h, 25 °C, 400 rpm. The amount of MTG used was 1 U/mL.	The remaining CA activity was more than 93%, and the remaining FDH activity was more than 84%. The efficiency of the cross-linked enzyme is increased by 5.8 times compared with free enzymes. FDH thermal stability at different temperatures is improved the optimal found CA/FDH ratio was 1:2.	[160]

4.1. Physical Adsorption

Physical adsorption was the first method used to immobilize enzymes [161]. It consists of affixing the protein onto a solid matrix utilizing hydrophobic (van der Waals), electrostatic (ionic), or hydrogen bonding interactions [162]. Physical adsorption can involve partial conformational changes and/or denaturation of the enzyme; thus, special attention must be paid to avoid these events and confirm that the whole activity of the enzyme is retained after immobilization. The types of functional groups on the surface that produce the adhesion of the enzyme to the matrix is one of the crucial aspects of this immobilization method [124]. These groups can contain hydroxyl, carboxyl, amino, sulfhydryl, or imidazole groups, among others, and produce interactions with the rest of the amino acids of the protein. Weak interactions are optimal since strong ones could result in enzyme conformational changes or even denaturation. Indeed, weak interactions are nonspecific and reversible; therefore, proteins can be easily recovered. For instance, if the interactions are electrostatic, free protein can be released into the medium by simply increasing the ionic strength of the solution.

Pore structure can affect enzyme accessibility and, consequently, both the quantity and activity of the immobilized enzyme. Finally, the surface area is also a critical factor: the higher the surface area, the larger number of adsorption sites, increasing both the quantity of immobilized enzyme and the global activity of the carrier. Physical adsorption can increase the stability of the enzyme against changes in pH, temperature, or organic solvents; the enzymes can be easily separated from the reaction mixture, making reuse easy. Moreover, in some cases, adsorption can increase the activity of the enzyme due to stabilization of the active conformation of the enzyme. On the other hand, physical adsorption can sometimes result in loss of enzyme activity if the microenvironment is not adequate or in a decrease in activity due to diffusion limitations of the reactants towards the enzyme active center that can reduce the rate of substrate conversion. Finally, the cost of the enzyme immobilization process can be high when applied, in particular, on an industrial scale.

Mesoporous silica and aluminosilicates are excellent candidates for enzyme immobilization using adsorption methods [163]. Here, the size and structure of the pores are crucial. Mesoporous silica with larger pores can allow for higher enzyme accessibility to the adsorption sites, which typically increases the quantity of the immobilized enzyme and, consequently, its activity. Wanjari et al. immobilized CA in an ordered mesoporous synthesized aluminosilicate, obtaining acceptable kinetic values for the biocatalyst compared to the free enzyme, remaining stable for more than 25 days [140]. Yu et al. immobilized CA in silica functionalized with carboxylate groups, which provided a very high degree of enzyme uptake, and, although the enzyme slightly changes its conformation with respect to the free enzyme, their activities were almost equivalent (95.6% that of the free enzyme versus that of the immobilized enzyme) [141]. Vinoba and coworkers adsorbed bovine carbonic anhydrase (BCA) inside octa(aminophenyl)-silsesquioxane silica nanoparticles modified with silver or gold, which continued to be active after 20 recycling runs [142]. This adsorption approach has been developed extensively, a recent example being the fixation of CA together with FDH to produce formate in silica nanoparticles modified by polydopamine and polyethylamine [143]. Here, the production of formate was expedited up to 30-fold with respect to the free enzyme and activity was retained at 86.7% after 10 cycles.

Colloids are another type of support used to adsorb CA [138]. Crumblis et al. immobilized CA in gold sols, obtaining an enzyme with levels of activity comparable to that of the native one [164]. Curiously, denatured CA was also immobilized in modified Sepharose and subsequent enzyme renaturation using a cycle of heating and cooling, resulting in an active enzyme with elevated activity [144]. CA immobilization via electrostatic adsorption has been studied with nanoparticles using different charges [145]. Positively charged nanoparticles do not adsorb human CA II, while negatively charged ones do, showing kinetic activity that depends on the degree of hydration of both the enzyme and the particle surface. More recently, CA was fixed onto two different types of membrane via layer-by-layer assembly: the first with a porous membrane and the second without [146].

The carbonation rate of the porous membrane was three times higher than that of the enzyme alone. On the contrary, the nonporous membrane was less active (70–90%) than the native non-immobilized enzyme. The adhesive properties of the polysaccharide chitosan modified with different compounds have also been employed to immobilize CA [153,165]. Matrixes of chitosan with different coating methods and a given textile package have been shown to adsorb CA in a “drop-in-ready” method, with high efficiency for CO₂ scrubbing. The physical properties of these matrixes for CO₂ capture were maintained for more than 31 days, with high efficiency (>80%) at moderate temperatures.

4.2. Entrapment and Encapsulation

Immobilization by entrapment occurs when a polymer, gel, or metal organic framework (MOF) is generated in the presence of an enzyme [132,166]. In such cases, the protein can remain trapped within the hollows of the polymer. The nature of these interactions is not chemical in origin, but rather physical. The proteins have free movement at a local level, but the motion is highly restricted to the confined hollows, and most of the molecules are isolated and interact only with the matrix. Drozdov’s group immobilized CA into the pores of different sol–gel magnetite with singular magnetic properties using this method [147]. They studied the physical properties of the new material as well as the overall structure of the enzyme, mainly using infrared spectroscopy, concluding that the protein maintains its 3D arrangement in the generated nanoparticles. The immobilized enzyme was stable and catalytically active at 90 °C, which is the temperature at which the native free enzyme is completely denatured.

Encapsulation is similar to entrapment in the sense that molecules are also free in solution and their movements are restricted; however, molecules are captured in higher bags where they can interact with each other. Sol–gel matrices have also been employed for the encapsulation of CA with excellent results. Polyurethane foam has also been employed to entrap not only the enzyme itself but also *E. coli* cells expressing CA [148]. Indeed, whole-cell catalyst CO₂ hydration activity was measured by comparing both sole and whole-cell immobilized enzymes with respect to the free enzyme. The efficiency of hydratase activity (Equation (4)) was 16-fold higher for the whole-cell immobilized enzyme than for the free enzyme. Interestingly, the activity of the whole cell trapped in the PUF was approximately 100% for at least nine cycles. MOFs are structurally ordered materials formed from inorganic complexes bridged by organic ligands that are projected in three dimensions [167]. Hollows of defined sizes are arranged monotonously in MOFs. MOFs are employed in a multitude of applications, with the immobilization of proteins being one of the most promising [168,169]. CA has been encapsulated in different MOFs with different features, most of them being zeolites constituted by imidazoles, with acceptable or excellent results [149,154,170]. The enzyme encapsulation within MOFs generates enzyme diffusion through windows that have a smaller size than the cavity. Whether or not the term encapsulation can be properly applied to the immobilization of enzymes in MOFs depends on the ratio between the pore and the enzyme size. In any case, an MOF based on Ni(II) showed a high degree of reusability for CA, retaining more than 65% of its activity after eight cycles [149]. Zinc has also been used as a base for MOFs to immobilize CA. In this case, the Zn-OH groups of the hollow imitate the active site of the enzyme, which permits high CO₂ capture efficiency [171,172]. MOFs containing several lanthanides have also been employed. In this framework, the existence of a high level of electrostatic interactions substantially increases capacity for CO₂ uptake [173]. Here, taking advantage of the lanthanide contraction, the specific dimensions of the hollows can be modulated, with the Eu(III) derivative having the highest affinity towards carbon dioxide. In all these examples, infrared spectroscopy is one of the key techniques for characterizing the degree of CO₂ capture, as well as the distortions of the framework.

Ionic liquids (ILs) have also been employed to immobilize CA, although to a lesser extent [150,174–177]. While CO₂ is nonpolar, owing to the difference in electronegativity of the carbon and oxygen atoms, the charges of ILs can absorb CO₂ to a high degree; thus, this is a

field fertile for exploitation. Recently, CA was immobilized in poly(ionic liquids) (PILs) by mixing the monomer hydrophobic IL 1-vinyl-3-hexylimidazolium bis(trifluoromethylsulfon yl)imide with an ethylene glycol derivative that had previously been polymerized using crosslinking [133]. After generating the PIL, CA was entrapped within the hollows of the polymer. The yield of the resulting CA was highly dependent on the size of the porous material and the degree of humidity (the dry PIL was less efficient). The authors also tuned the degree of particle size using previous sonication and studied the kinetic parameters of CA-PIL. These values were comparable to that of the free enzyme, although the entrapped CA was stable for a month without detectable loss of activity, while the free enzyme decreased its activity by more than 30%. The CA-PIL was reused for five cycles with 60% activity.

4.3. Covalent Binding and Crosslinking

Covalent binding implies the formation of bonds between the groups of adequately functionalized supports and an enzyme. This is, by far, the most extended approach for immobilizing enzymes, particularly for CA [152,155,178,179]. Several protein functional groups can be used for this purpose. CA has been covalently bound to different supports by its amine groups by reaction with glutaraldehyde [156,180]. Generally, mesoporous supports containing hollows of controlled sizes are grafted with amine groups to obtain solid materials that are prone to covalently binding to enzymes using glutaraldehyde. This was carried out with the support SBA-15, in which three different amine compounds were inserted, followed by covalent immobilization of HCA [142,152,181]. The resulting material was morphologically characterized, and its activity, thermal stability, and reusability were also determined, obtaining better results than those of the free enzyme. Kimmel et al. immobilized CA on the surface of propylene fiber membranes. These membranes, commercially available, were coated with a siloxane layer and functionalized with amine groups. Posteriorly, CA was attached to these fibers via glutaraldehyde crosslinking under two conditions: with and without chitosan tethering. Then, the authors applied these fibers to CO₂ removal, finding enhancements of 115% and 37% versus the buffer and the blood controls, respectively [157]. Moreover, carboxylic groups activated by carbodiimide and *N*-hydroxysuccinimide agents were used to covalently immobilize CA in microtubes [182]. The resulting immobilized enzyme enhanced its ability to sequester CO₂ with respect to the free enzyme.

Finally, the crosslinking method is actually a special way of covalent binding. The proteins are bound to other large proteins to form high molecular weight complexes without any solid support or, properly, the enzymes themselves being a solid support. Typically, the protein is precipitated with the appropriate agent and then crosslinked, which can be performed with a purified protein or an extract of a still unpurified enzyme. This method confers high stability and a high degree of enzyme recovery. CA has been immobilized via crosslinking in numerous studies. Recently, Xu et al. encapsulated crosslinked CA in alginate beads and confirmed that this crosslinked CA enhanced the growth of microalgae cultures [183]. The crosslinked CA was stable during 10-cycle assays. Magnetic nanoparticles aggregated with CA were obtained by crosslinking the enzyme with glutaraldehyde, improving the yield of absorbing CO₂ up to 3.4-fold with respect to the free enzyme and retaining 95% activity after five cycles of reuse [158]. These magnetic nanoparticles are amply used because they allow for very simple separation and recovery of the biocatalyst by using an external magnetic field. They are considered excellent carriers and supporting matrices for enzyme immobilization, providing several advantages for the design of biocatalytic processes (i.e., large surface area, large surface-to-volume ratio, high mass transference, etc.). More recently, Chang et al. crosslinked CA and geopolymer microspheres with glutaraldehyde and performed a detailed study on the morphology, stability, and activity of the immobilized support [159]. After 60 days of storage at 25 °C, immobilized CA still presented 28.9% activity, whereas free enzyme activity was less than 10%. CA and FDH have also been crosslinked to reduce CO₂. Zhang et al. used microbial transglutaminase (MTG) as the crosslinking medium for CA and FDH labeled with peptide

tags and previously expressed in *E. coli* [160]. MTG catalyzes the formation of an isopeptide bond between the ϵ -amino group of lysine and a glutamine. The authors studied the activity and reusability of several crosslinked particles with different tags at CA/FDH ratios of 1:1, 1:2, and 1:3. Because CA is much more active than FDH, it is expected that the lower the CA/FDH ratio, the higher the formate yield obtained. However, the optimal found CA/FDH ratio was 1:2. The authors attributed the lower yields obtained for a 1:3 ratio to FDH steric hindrances in the crosslinked aggregates.

5. Carbon Capture Storage and Utilization: State-of-the-Art, Costs, and Perspectives

Methods for CO₂ capture are classified into precombustion, postcombustion, and oxy-combustion processes [12,184]. The precombustion approach is related to hydrogen gas production. This is obtained in numerous industrial processes such as electric power generation, ammonia or fertilizer synthesis, and petroleum refinement. The precombustion process refers to the conversion of the primary solid fuel (coal or biomass) by reforming it into a mixture of CO and H₂ gas (syngas). This gas reacts with the water stream at high temperatures and pressures to produce CO₂ and more H₂ (water–gas shift reaction). Finally, CO₂ is captured using several methods. Postcombustion CCS methodology denotes all the processes employed to capture CO₂ from exhaust gas (its major component being nitrogen) resulting from industrial chemical processes. CO₂ gas is emitted at relatively low temperature and pressure. Oxy-fuel combustion consists of the oxidation of fuel using pure oxygen instead of air, obtaining an almost pure CO₂ atmosphere without nitrogen gas. Most CCS methods have been developed for postcombustion gases and are referred to here, except where otherwise indicated. In many cases, the methodology can be the same for both post- and precombustion approaches, although with different designs depending on the P/T conditions of the exhaust gases. Physical adsorption and absorption [185] (geological storage [186] probably being the most relevant among absorption approaches) and cryogenic distillation [187] are the main methods used for CO₂ CCS.

Global CO₂ emissions from combustion processes grew by 0.9% in 2022, reaching a total of 36.8 Gt [188]. Energy used in industry, agriculture or land use, buildings, transport, direct industrial processes, waste, and others with 37.8%, 18.4%, 17.5%, 16.2%, 5.2%, 3.2%, and 1.7%, respectively, are the contributions to CO₂ emissions by the different sectors [189]. Only ca. 40 million Tm, i.e., 0.1%, was removed from the atmosphere using CCS methods in 2019. Thus, we are still very far from being efficient in eliminating the CO₂ expelled into the atmosphere.

The estimated present costs of CO₂ Tm removal vary nowadays from 40 to 80 USD depending on the method [190]; however, the net contribution to CO₂ elimination from the atmosphere is difficult to calculate, since net contributions in the whole process have to be taken into account. Hepburn and coworkers analyzed the perspectives, including cost, for different methods of CO₂ utilization [190]. They analyzed ten different methods of CO₂ utilization. For instance, chemical production, particularly the generation of urea, on one hand, and the production of polycarbonate polyols, on the other, are two fields in which CO₂ capture can be exploited. They estimated that CO₂ utilization in chemicals in 2050 could be around 0.3–0.6 Gt CO₂/yr with costs ranging from –80 to 320 USD per Tm of CO₂ (a negative value would indicate an additional economic benefit, while a positive value indicates that the cost of capturing and utilizing CO₂ would be higher than the value generated from it). Fuels, that is, CO₂-methanol plants, were also considered in their study, although they stated that many different scenarios can vary their prospects from 1 to 4.2 Gt/yr in 2050 and, in terms of cost, from 0 to 670 USD per Tm of CO₂.

6. Carbonic Anhydrase in Carbon Capture Storage

Although CA is not used in all previous technologies, its use is extended or has a good perspective in others, mainly chemical adsorption and mineralization, whose description is commented on below. Table 4 describes relevant studies performed with CA in CCS research.

Table 4. Relevant studies on CO₂ uptake processes performed using CA, and their main features.

Process	Idea of the Study	CO ₂ Uptake Conditions	Relevant Conclusions	CA Activities	Ref.
Chemical absorption	To examine the stability and activity of the enzyme under different pH values and amine solvents. Compare its performance to other common solvents.	CA tested for pH [7–11]. Temperature stability for up to 100 h of incubation. CA stability tested for 7 capture solvents (1 M or 3 M, 150 days, 40 °C).	CA stable at 60 °C, pH range [7–11]: residual activity at pH 5 or 12, ranging from 12 to 91%. The enzyme enhances reaction rates. NaCl, K ₂ CO ₃ , AMP, and MDEA show additive effects.	After 100 h (25 °C), CA activity was kept higher than 75% for 1M NaCl, AMP and MDEA and 125% in 1M K ₂ CO ₃	[191]
	To develop the CA-MOFs composite, with superior catalytic performance and high stability to promote CO ₂ absorption into a tertiary amine solution.	0.05 g L ^{−1} and 40 °C, P _{CO2} 15 kPa.	CA loading in ZIF-L-1 increased with the added enzyme amount. CA/ZIF-L-1 composite has higher catalytic activity and stability than the free CA. The immobilization of CA on ZIF-L-1 improves the CA conformational stability	CO ₂ absorption rate of CA/ZIF-L-1 in 1 M MEA and MDEA at 40 °C: 3.0×10^6 kmol s ^{−1} m ^{−2} . CA loading reached up to 87 mg g ^{−1} . The highest immobilized CA activity was 1.5 times that of the free CA	[192]
	Pilot-scale experiments with CA-enhanced MDEA for CC, bench-marking its mass transfer performance against the industrial standard 30 wt% MEA.	Experiments were done using 30 wt% MEA and MDEA solvents varying CA concentrations (0, 0.85, and 3.5 g/L) at different column L/G ratios.	Enzyme-enhanced MDEA solutions exhibit 80% of the mass transfer performance at 30 wt% MEA and have the potential to reduce the size and cost of absorbents in carbon capture.	CO ₂ capture efficiency higher than 90% for high L/G ratios. The mass transfer increased 20-fold for 25 and 50 wt% MDEA solutions by adding 0.2 g/L CA.	[193]
	Use of columns with membranes contractors with polyionic liquids (PIL), amines and CA for CO ₂ uptake.	PIL blend (F9:1(M10)) with immobilized CA in 30 wt% MDEA at low feed gas (15% CO ₂ in N ₂) pressure (1.3 bar)	Addition of the enzyme to the MDEA solution with PILs significantly improves CO ₂ uptake rate and reduced the equilibration time.	CA addition to MDEA solution improved the CO ₂ uptake rate by 1.7 times. CA and membrane improved the uptake rate by twice.	[194]
Carbonation	To present the effects of adding small quantities of immobilized CA on the absorption of CO ₂ into potassium carbonate.	CO ₂ absorption (partial pressure 90 kPa) in a 30% K ₂ CO ₃ solution. A wet-walled column (40–80 °C) with 38 g/L CA.	CA addition improves the CO ₂ absorption in K ₂ CO ₃ solvents. The rate of CA-catalyzed CO ₂ hydration increased with the CA concentration	Increasing CA concentration from 0.4 to 1.8 μM increases the absorption rate by 34% (40 °C)	[195]
	To explore the potential of using enzymes to catalyze the conversion of CO ₂ into bicarbonate and on the carbonation rate of brucite, Mg(OH) ₂ .	Gas flow (CO ₂ /N ₂ 10%/90%) at 10 psig. Experimental durations were 11, 7, and 3 days (low, medium, and high flow experiments).	CA accelerates the carbonation of brucite. Higher CO ₂ gas flow rates results in faster carbonation. Mineralogical compositions depend on the CO ₂ flow rate.	The carbonation rate of brucite using BCA was accelerated by up to 240% compared to controls.	[196]
	To propose a novel method based on microbially induced calcium precipitation to improve the cementitious properties of steel slag.	<i>Bacillus mucilaginosus</i> placed in a sealed container, the air pressure was reduced to −0.05 MPa. CO ₂ (99.99%) pumped to maintain the gas pressure (0.25 MPa, 30 °C, 32 h).	Ca-silicate promotes the growth of bacteria, enhances the production of polymers and improves bacterial adhesion. Ca-silicate-containing medium (CSCM) can be used as a microbial carrier.	Maximum bacterial growth rate in CSCM was 1.5 times higher than that in the control medium (CM). Bacterial activity in CSCM was 40% of that in CM (30 h). At a dosage of bacterial powders of 1 wt.%, carbonation reached the maximum.	[197]
	To develop and characterize a highly efficient and stable biocatalyst for CO ₂ sequestration using silica nanocomposite with auto-encapsulated CA. (CA)-based biocatalyst encapsulated in a biosilica with a peptide R5.	CaCO ₃ precipitation was carried out at 30 °C and monitored turbidimetrically at 600 nm. The final pH of the buffer was approximately 9.3	The encapsulated CA was not leached from the silica matrix. Encapsulation in silica effectively improved the thermostability and activity of the enzyme.	Encapsulation efficiency greater than 95%. Residual activity of the self-encapsulated CA was more than 50% (30 min, 80 °C). Activity of ngCA-R5@silica decreased only after 2.5 days at 60 °C. Encapsulated CA: 98% and 80% of its initial activity (1 day and 5 days of incubation, 50 °C). CaCO ₃ precipitation is reduced by 5.5-fold when ngCA-R5@silica (30 μg/mL) is present compared to the uncatalyzed reaction.	[198]

Table 4. Cont.

Process	Idea of the Study	CO ₂ Uptake Conditions	Relevant Conclusions	CA Activities	Ref.
Mineralization	Mineralization experiments were performed using <i>Curvibacter lanceolatus</i> strain HJ-1, including its secreted extracellular polymeric substances (EPS) and CA (CA).	Three types of mineralization experiments: with CA (duration 96 h), with EPS (96 h), and with bacteria (50 days).	Strain HJ-1, EPS, and CA promote carbonate precipitation. HJ-1 and EPS1 experiments contained calcite and aragonite. CA formed calcite only. HJ-1 and EPS are favorable for aragonite precipitation.	The mass of precipitate (inorganic plus organic substances) increases until ca. 55 mg. The maximum degree of calcification was approximately 6.6%. In the control groups without CA no precipitate was formed. In the absence of HCO ₃ [−] , the optimized calcification rate followed the order: HJ-1 (49.5%) > CA(6.6%) > EPS2(4.1%).	[199]
	Production of CA from the bacterium <i>Aeribacillus pallidus</i> TSHB1, a thermostable and alkaline-stable bacterium, highly effective in the formation of CaCO ₃ from aqueous CO ₂ .	Tris buffer (15 mL, pH 7.4) containing 0.9 g CaCl ₂ ·2H ₂ O, CA 0.05 mg (37 °C).	A 3.8-fold higher CA production by <i>A. pallidus</i> than that under unoptimized conditions. Enzyme thermostable that retains activity at alkaline pH: it is useful in carbon sequestration.	The partially purified enzyme produces precipitation of 42.5-mg CO ₃ ^{2−} mg ^{−1} protein. CO ₂ sequestration is efficient.	[200]
	Study of the biochemical properties, thermostability, and inhibition of CA from <i>Sulfurihydrogenibium azorense</i> , SazCA.	Hepes buffer 10 mM, NaBF ₄ 20 mM, pH 7.5. Phenol red (0.2 mM) as an indicator.	SazCA is highly thermostable and can survive incubation at 90–100 °C.	k_{cat} value $4.40 \times 10^6 \text{ s}^{-1}$, K_M value 12.5 mM, k_{cat}/K_M $3.5 \times 10^8 \text{ M}^{-1} \text{ s}^{-1}$, 5-fold faster than the second CA. SazCA is the second faster enzyme, after superoxide dismutase (SOD).	[201]
	To explore wollastonite (calcium silicate) carbonation for the removal of anthropogenic CO ₂ and evaluate the effectiveness of different natural (CA) and CA biomimetic catalysts (Zr-based MOFs) to enhance CO ₂ capture.	Water 170 mL, the catalyst (30 ppm), adjusted at pH 4 (25 °C). Wollastonite crystals (100 mg) added to the solution. CA is immobilized on the MOFs UiO-66 and MOF-808@Mg(OH) ₂ by an impregnation process.	CA accelerates carbonate precipitation but hinders carbonation of wollastonite. Thick carbonate coatings formed on the most reactive surfaces of wollastonite act as passivating layers leading to a reduction in the dissolution and carbonation rates.	The passivating effect explains why the conversion of wollastonite into calcite was so limited (up to 14 mol%). Zr-based MOFs accelerate the dissolution of wollastonite.	[202]

6.1. Chemical Absorption

Chemical absorption has been the most used CCS method for decades [203]. This procedure involves the scrubbing of exhaust gas at low pressures and temperatures with alkaline solutions typically containing amines and/or carbonates or hydroxide solutions [203]. Amines are weak bases that can capture protons from Brønsted acid CO₂. The reactions of primary/secondary or tertiary amines produce carbamates or bicarbonate anions, respectively, according to the reactions described below.

The amines typically used for these purposes are alkanolamines. An alcohol group increases water solubility and decreases vapor pressure compared to analogous amines. The main chemical solvent used as an absorber is monoethanolamine (MEA). The solutions typically consist of an aqueous solution of 20–30 wt% MEA. CO₂ is captured at low pressure (ca. 1 bar) and in a mixed gas containing other gases, such as N₂, SO_x, and NO_x. Secondary amine diethanolamine (DEA) and tertiary amine *N*-methyl diethanolamine (MDEA) are also amply used amines.

Several points should be considered when choosing a suitable solvent for CO₂ sequestration. First, the enthalpy of the reaction: the higher the enthalpy of the reaction, the higher the cost of solvent regeneration. This is a crucial point because it is estimated that 60–80% of the costs of these processes arise from solvent regeneration [204]. The enthalpy of (exothermic) reactions 6 and 7 increases from tertiary to secondary and primary amines; thus, primary amines improve both energetic and economic costs. Moreover, the power of corrosion also follows the same order (primary amines are the most corrosive). Another decisive issue is their ability to load CO₂. Tertiary amines have the highest capabilities in this regard [205]. According to these thermodynamic aspects, tertiary amines are the best ones to use. However, another crucial point is the kinetics of CO₂ sequestration; indeed, tertiary amines have low reaction rates and are kinetically much more inert than primary amines [206,207]. Owing to the slow kinetics of tertiary amines, primary amines (or secondary amines) are currently preferred. However, the high costs of cooperation and maintenance due to the ease of amine degradation and the formation of highly corrosive salts are drawbacks when operating with amines.

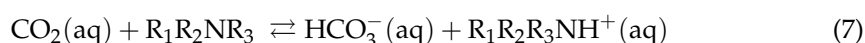
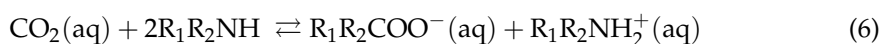
Consequently, for kinetic reasons, primary amines, specifically MEA, are by far the most widely used alkanolamines in the industry. Numerous plants have been developed using MEA solutions. These systems are typically coupled with industrial processes and CO₂ is captured with postcombustion gases. For instance, their use is extended to the iron and steel industries (responsible for approximately 31% of all industrial CO₂ emissions). These plants can recover 85–95% of the CO₂ in gas. As an example, a steel production plant recently established by Emirate Steel Industries has a yield plant using CCS based on amine absorption that captures 0.8 Mt CO₂/year [208].

In the last decade, CA has been revealed as a tool for accelerating CO₂ uptake in chemical absorption processes. Gundersen et al. studied CA stability and activity for a long time (150 days) as a function of pH, temperature, and the solvent, combining MEA and MDEA solutions, among others [191]. They concluded that CA was suitable for these purposes; the biocatalyst was stable and active between pH 7 and 11, with maximum activity at 40 °C. In addition, the enzyme preserved its activity between 12 and 91% of the original activity depending on the solvent employed. The absorption of CA was also accomplished in MOF ZIF-L-1 in the presence of MDEA [198]. In ZIF-L (a zeolitic imidazolate framework), the imidazolate groups enhance CA immobilization, and CO₂ uptake is hence greatly increased. The authors highlighted that this new MOF obtained excellent CO₂ absorption rates at 40 °C and a CO₂ partial pressure of 15 kPa, while the activity was maintained for six reuse cycles [192]. Additionally, a pilot-scale plant was set up with CA in solution in the presence of MDEA. The authors observed an enhancement in CO₂ capture in the presence of the enzyme and demonstrated the possibility of translating the laboratory results to higher scales [193]. However, because the free enzyme is damaged by amines, immobilization is necessary. Kim et al. also studied the

effect of CA on CO₂ absorption rates in the presence of MEA and MDEA, although they used a membrane contactor with hydrophobic and hydrophilic supports [194]. This system allows an expanded contact surface to enhance CO₂ absorption.

6.2. Chemical Carbonation

Chemical carbonation is probably the most efficient method for capturing CO₂. This is performed when CO₂ is bubbled through an alkaline solution, typically consisting of dissolved KOH or Ca(OH)₂, where potassium or calcium carbonates precipitate. The limiting step for capturing CO₂ in postcombustion processes is Equation (3). However, due to Equation (4), CO₂ capture is much faster in alkaline media since carbonates are formed, and so hydrogen carbonate concentration decreases, and Equation (3) is shifted towards the consumption of CO₂. Indeed, once bicarbonate anion (soluble) is formed, reactions such as Equations (6) and (7) (amine formation, Scheme 2) occur much faster in alkaline media. Even so, the limiting step, for kinetic reasons, continues to be the CO₂ gas uptake, as commented previously. Thus, the main challenge in applying alkaline solutions, either amines or carbonates, to the CCS approach is speeding up CO₂ conversion to bicarbonate [195]. Consequently, numerous studies on CA to increase the mass transfer of CO₂ capture have been proven not only at the laboratory level, and its feasibility has been demonstrated on an industrial scale [15,205]. Novozymes NS81239 CA (NCA) at 2 µM increased the absorption rate of CO₂ into potassium carbonate by ca. 30%, augmenting this uptake at temperatures in the range of 40–60 °C [195,196]. Power et al. demonstrated that bovine CA accelerated the carbonation rate of brucite Mg(OH)₂ from CO₂ gas by up to 240% [209]. In these studies, CA was supplied as a free enzyme; therefore, its regeneration was not studied. Biological tools have also been used to enhance carbonation. Jin et al. accelerated calcium carbonate precipitation by employing *Bacillus mucilaginosus* on steel slag powder [197], increasing the carbonation degree from 66.34 to 86.25%. Moreover, the mechanical properties and durability of the treated steel slag were enhanced. The CA immobilization, as described in the previous section, strongly improves the reusability of the enzyme as well as the chemical carbonation. For instance, Jo and coworkers proved the suitability of CA encapsulated in a biosilica matrix, obtaining good yields for carbonation compared to the free enzyme [198].



Scheme 2. Reactions of CO₂ with primary, secondary (Equation (6)), and tertiary (Equation (7)) amines in water solutions.

6.3. Mineralization

Biomineralization is a very slow and exothermic process by which carbonate minerals are formed from silicates and CO₂ under basic conditions [210]. The starting silicates usually contain divalent metals such as Ca(II) and Mg(II) or trivalent metals such as Fe(III). This event occurs in nature on a regular basis and is responsible for the formation of inorganic structures in living organisms such as exoskeletons in protozoa, algae or invertebrates, and shells or plant mineral structures. It is also responsible for the presence of large amounts of limestone on the Earth's surface [211,212]. When trying to emulate biomineralization, which takes place over very large timescales, the main drawback is speeding up the process. Artificial mineralization mimics nature, although in short periods. It involves the injection of CO₂ directly into geological formations to promote a carbonate-forming reaction with alkaline minerals [213]. This mineral sequestration would be a viable alternative for subsequent storage because the carbonate products formed would not require monitoring owing to their high stability and safety. On the other hand, in-ground or ex situ mineralization is based on the exposure of crushed rock material in a processing plant where CO₂ is introduced, facilitating the formation of carbonate minerals. Natural

minerals or alkaline solid waste can be used [213,214]. The use of natural silicates requires a large amount of material, which implies a very large operational size and an unfeasible economic mineral impact. Ex situ mineralization can also be carried out using alkaline wastes containing divalent metals such as ash originating from the coal or metallurgical industry, cement and concrete wastes, or iron and steel slag [215,216]. This method would reduce not only environmental CO₂ but also the accumulation of waste from industrial activities, although a major disadvantage is that its capacity is much smaller than that of CO₂ mineralization from silicates. At laboratory scale, this mineralization has been satisfactorily performed by directly extracting CO₂ from the air, and its direct extraction by passing the air through cooling towers using NaOH solutions has also been proposed for larger scales [217]. However, the same authors pointed out the elevated costs of this approach on a large scale.

The efficiency of the biomineralization process can also be accelerated by modifying certain parameters such as increasing the temperature, pressure, or retention time. The biomineralization process is also favored by the presence of purines, NaCl, or CA. The presence of CA accelerates the rate of hydration of CO₂ dissolved in water; therefore, possible modifications to CA to support high pH and temperature conditions without losing its advantageous functionality have been studied [218,219]. On the other hand, some varieties of carbonic anhydrase are inhibited in the presence of high concentrations of hydrogen carbonate, which becomes a problem for its use in industry. However, this can be circumvented, at least partially, by increasing the pH to values equal to or higher than 9.0, conditions under which some CAs are still stable and functional. Immobilization improves the stability of CA at high temperatures or alkaline conditions, as confirmed by Arias et al. when forming calcite in vitro by mineralization using CA immobilized in eggshell membranes [219]. Recombinant CAs have also been used to accelerate mineralization under extreme conditions. For instance, CA from the alkalistable *Aeribacillus pallidus* was genetically modified, achieving acceptable yields in the presence of pollutants such as NO_x and SO_x [200]. Similarly, CA from the thermophilic bacterium *Sulfurihydrogenibium azorense* was modified, and its half-life was found to be 8 days when the biomineralization process was carried out at 70 °C and 53 days at a reaction temperature of 50 °C [201]. Di Lorenzo et al. studied the effect of CA and a Zr-based MOF in the carbonation process of wollastonite (CaSiO₃) to produce calcite (CaCO₃) [202]. Although CA accelerated CO₂ uptake by the silicate, the total gas absorber quantity was lower than that of the MOF. Jin et al. also took advantage of CA to accelerate the carbonation of γ -dicalcium silicate, which is also present in steel slag. They used a powder containing alkali-resistant CA bacteria, increasing the yield by 19.0% [220].

7. Biotechnological Aspects of CO₂ Reduction

7.1. Hindrances to Biochemically Reducing CO₂

Forward Equation (5) presents several drawbacks that hinder its application outside the natural living environment, that is, employing it with biotechnological aims. First, owing to the low redox potential of the CO₂/HCO₃[−] pair (−430 mV), the reaction is highly endergonic under physiological conditions. Three main strategies have been developed to overcome this problem (Figure 8): coupling thermodynamically favorable reactions in the presence of an excess concentration of the reducing agent [136,221–224]; coupling the reaction to an electrochemical device that provides electrons from a battery anode (i.e., by supplying electric energy) [225–228]; and using photosensitive molecules able to absorb light and regenerate redox partners [31,229–233].

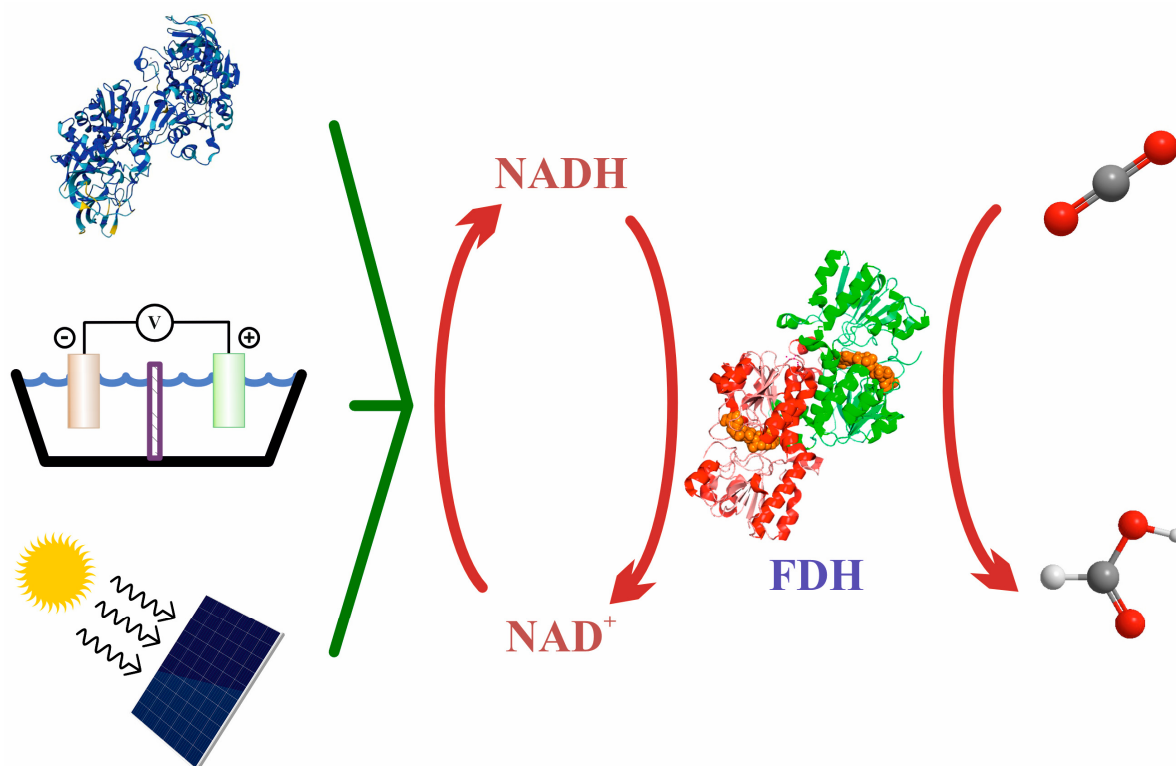


Figure 8. The three different (enzymatic, electrochemical, and photochemical) approaches to regenerating cofactor NADH and producing formate with FDH.

Regeneration of the cofactor NADH is crucial for obtaining formate with acceptable yields [234–237]. As previously mentioned, metal-dependent FDHs do not necessarily accept electrons from NADH (see Figure 6); nevertheless, whatever the cofactor is, either in nature or in biotechnological approaches, it has to be regenerated for adequate progress of the reaction. NADH, the primary source of electrons, is an expensive reactant, and thus economic aspects are also relevant in cofactor regeneration. The approaches to regenerating NADH are similar to those followed to facilitate the thermodynamics of the reaction. Two additional issues related to the oxidation of this cofactor also have to be kept in mind: NAD⁺ can strongly interact through its negative phosphate charges, with positive charges on the active site of the biocatalyst, inhibiting the biocatalyst, particularly in nonmetal-dependent FDHs [238]. Moreover, it is well known that NAD⁺ trends to form dimers (see below), which makes cofactor regeneration impossible [221,234]. Carbon dioxide uptake into the reaction medium is another crucial factor. As commented throughout this text, since CO₂ is a nonpolar gas, its solubility in water or similar solvents is low and, more importantly, the kinetics of solubilization and mass transfer are very slow [204]. Hence, the employment of systems (specific solvents or solutions) that can incorporate CO₂ into their structures is decisive. As discussed above (see Sections 6.1 and 6.2), amines have been used extensively [204].

Ionic liquids, solvents in which positive cations can interact with the oxygen atoms of a CO₂ molecule through their sole pair of electrons, are also excellent media to solve, in high quantities, carbon dioxide gas [214,239,240]. The selection of the appropriate conditions for the reaction is also crucial. Dealing with a gas, high pressures and low temperatures are ideal conditions for solubilizing CO₂, while with respect to pH, carbon dioxide solubilization is favored at high pH values (Equations (3) and (4)). However, optimal conditions are determined by FDH stability and activity, and temperatures must thus be moderated (lower than 50 °C), pressures cannot be high, and pH values must be between 6.0 and 7.5, the range at which FDH shows its highest yields. Finally, the use of

the adequate enzyme, that is, the FDH of the appropriate organism, is also a key factor in the success of the reaction.

Approaches to circumventing these problems and achieving significant formate yields as a starting point for obtaining value-added chemicals are discussed in the following sections.

7.2. Coupled reactions: Enzymatic Multicascades

Equation (5) is reversible although under physiological conditions is highly shifted towards the backward reaction. The first successful conversion of CO₂ from formate using FDH in a laboratory system was reported by Hopner and collaborators [100]. They used the metal-dependent FDH from *Pseudomonas oxalaticus* as biocatalyst and NADH concentrations in such a way that reaction 5, still being thermodynamically unfavorable, was not completely shifted towards the oxidation of formate. They devised a sealed system containing ¹⁴CO₂ and measured the radioactivity of the H¹⁴COOH formed. In this pioneering study, the kinetic parameters ($k_{\text{cat}}/K_{\text{M}}$) of the enzyme and its pH activity profile were determined. However, the cofactor was not regenerated and the rates between the forward and reverse reactions in their working conditions were 1:30, with a turnover number for CO₂ reduction as low as 3 s^{−1}, which makes these results insufficient.

The simplest and most straightforward way to increase formic acid generation is to eliminate the products from the reaction media, that is, regenerating the NADH cofactor using a reducing chemical agent present in the medium. This approach was developed in the 1980s with redox biocatalytic systems [241,242]. The reaction desired to take place is coupled with another “inverse” reaction, in which the reactant is added at high concentrations. Thus, the reaction is thermodynamically favored, and the cofactor is regenerated (Figure 9A). Chenault and Whitsides, pioneers in using this technique with formate dehydrogenase, employed CbFDH to regenerate NADH by coupling formate oxidation with the reduction in lactate to pyruvate using D-lactate dehydrogenase and obtained acceptable results, with nicotinamide residual activity of 55% after each run [242]. However, in their reaction, FDH was used for formate oxidation (backward Equation (5)), and the coupled reaction was used to regenerate NAD⁺. Since then, the reduction of CO₂ (forward Equation (5)) using FDHs in a conjugated oxidation–reduction reaction system for regenerating the NADH cofactor has been carried out with several enzymes. Yu et al. cloned the FDH gene from *Cupriavidus necator* in *E. coli* and coupled the reduction of CO₂ with the oxidation of D-glucose to D-δ-gluconolactone using glucose dehydrogenase (GcDH) to regenerate NADH [243] (Figure 9A). The expressed enzyme, an O₂-tolerant, Mo-dependent FDH, was able to effectively reduce formic acid comparable to that of the nonrecombinant protein. Glutamate dehydrogenase (GDH) is also used to regenerate NADH. GDH catalyzes the oxidation of glutamate to α-ketoglutarate and ammonia, thereby reducing NAD⁺ to NADH. GDH is highly stable over a wide range of pH values and at temperatures as high as 85 °C, as well as being widely commercially available and inexpensive [236,244].

Nonetheless, the most extensive coupled reaction approach is the well-known and widely employed enzymatic cascade reaction that drives from CO₂ to formic acid, catalyzed using FDH, from this species to formaldehyde, i.e., formaldehyde dehydrogenase (FaldDH), the biocatalyst, and, lately, to reduce this molecule to methanol, by the action of the alcohol dehydrogenase (ADH) (Figure 9B). The oxidases mentioned above are typically used for NADH regeneration.

El-Zahab et al. co-immobilized FDH, FaldDH, and ADH together with GDH into polystyrene particles to reduce CO₂ to methanol [245] (Figure 9B). The NADH cofactor was also immobilized, although separately. The results obtained with the immobilized enzymes were similar to those obtained for the free enzymes; however, importantly, the yield of the reaction was maintained at over 80% after 11 cycles. Ji et al. coupled the same reaction cascade [246], but in their study, the four redox enzymes were entrapped in hollow nanofibers together with CA to facilitate CO₂ absorption. The methanol yield was 36.17%, retaining ca. 80% of the productivity after 10 reuses, with an accumulative yield of more than of more than 900% for NADH regeneration. In another study using the same cascade,

Ren and collaborators encapsulated the same biocatalysts in MOF ZIF-8 and investigated the effect of polyethyleneimine (PEI) on anchoring the NADH cofactor and, hence, on the yield of the reaction [247]. Compared to the free system, the yield of this reactor system increased 4.6-fold and the activity after eight cycles was retained by 50%.

Ionic liquids are known to solubilize CO₂ [248,249]. Taking advantage of this, Pinelo's laboratory immobilized the four mentioned enzymes as well as the cofactor in a series of modified ILs composed of choline (CH) and amino acids (CHGlu, CHPro, CHGly, and CHHis) [250]. They generated a membrane reactor in which the products were removed in situ and the reaction was displaced towards the desired product. The yield of CHGlu increased up to fivefold compared to that of with the control aqueous system when NADH was regenerated. In another study, immobilization of the four enzymes was performed on superparamagnetic nanoparticles. Here, the yield was low (2.3% of methanol per NADH molecule); however, under CO₂ pressure (126 psi), the reaction yield increased 64-fold after 30 min of reaction [250].

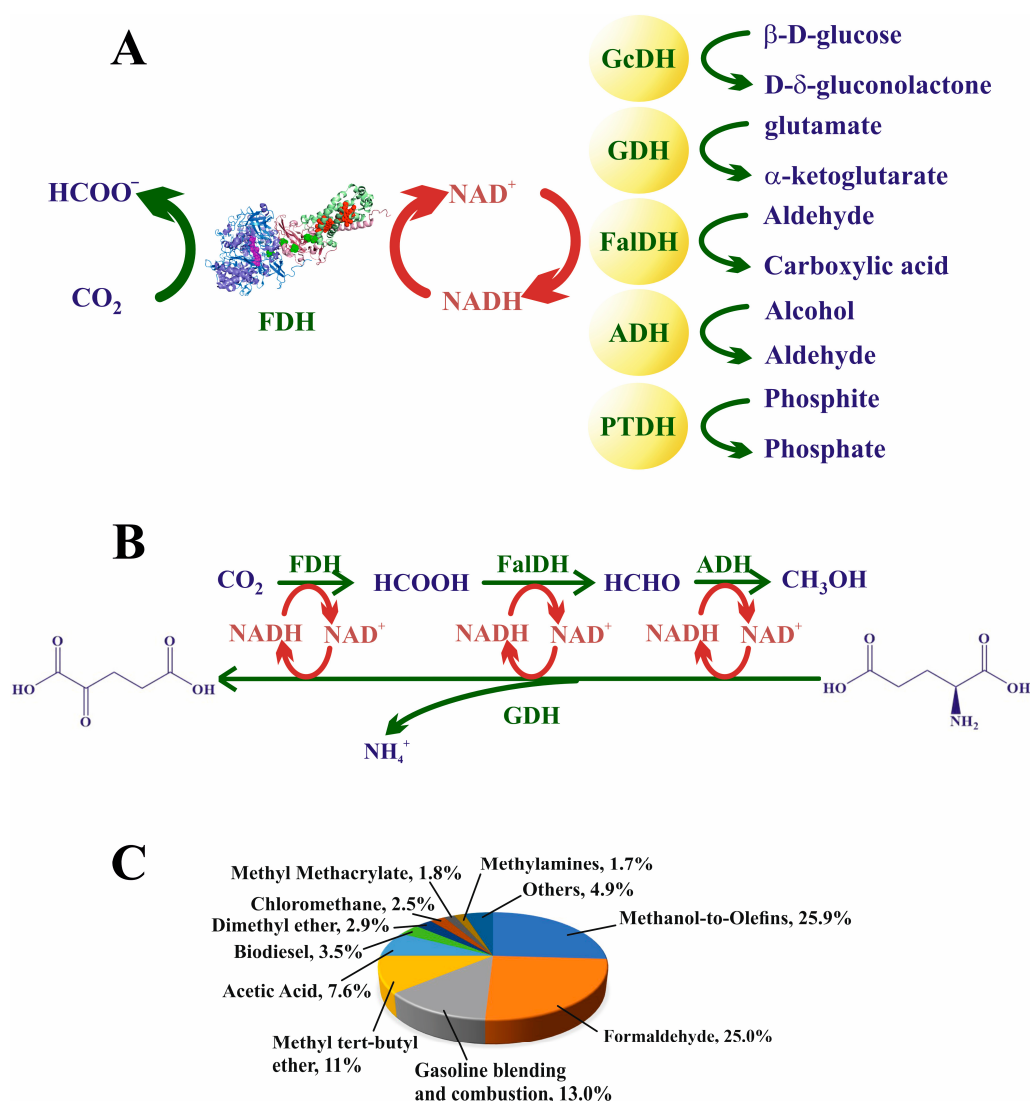


Figure 9. (A) Several enzymes whose oxidation reactions have been coupled to FDH CO₂ reduction for regenerating NADH. (B) A classical enzyme multicascade for the reduction of CO₂ to methanol; here, GDH is coupled for regenerating NADH. (C) Uses of methanol at the global level in the year 2020 shown in percentages according to their production and demand [251].

Phosphite dehydrogenase (PTDH, Figure 10A) has also been used for NADH regeneration in a multicascade approach to obtain methanol from CO₂. Cazelles et al. studied the

effect of regenerating NADH using three different coupled reactions, namely phosphite oxidation, catalyzed using phosphite reductase; glycerol oxidation to dihydroxyacetone, performed using glycerol dehydrogenase; and a natural photosystem (chloroplasts) extracted from spinach leaves that oxidize water to molecular oxygen [252]. They encapsulated the three enzymes in phospholipid–silica nanocapsules and obtained excellent activities with PTDH with respect to the free enzymes in solution (55 times higher activities) under 5 bar of CO₂ pressure for 3 h, although the other two systems were not so efficient. Singh et al. also employed PTDH for regenerating NADH [253]. They expressed recombinant proteins (FDH, FaldH from different bacteria, and ADH from yeast) in *E. coli* and performed assays with free enzymes in water solution and in the presence of many different cosolvents. IL 1-ethyl-3-methylimidazolium acetate (EMIM-Ac) was found to be the most effective in increasing methanol production. Indeed, the yield was enhanced more than twofold (from 3.28 mM of methanol to 7.86 mM, 6 h of reaction) in the presence of 1% EMIM-Ac because of the ability of EMIM cations to interact with CO₂, increasing solubility. Finally, lactate dehydrogenase (LDH) was also used to regenerate NADH in a multicascade reaction [254]. This enzyme was immobilized in a sol–gel matrix, and CO₂ reduction was acceptable after 1 h of reaction, as indicated by the authors.

Using CO₂ as a substrate for the generation of methanol is an attractive process because of its potential use as a fuel and the multitude of products obtained from it at the industrial level. Figure 9C shows the chemicals produced from methanol at a global level in 2020 [251]. Methanol is a precursor of numerous compounds such as olefines (25.9% of the demand for methanol, the year 2020), formaldehyde (25.0%), gasoline blending (13.1%), biodiesel, and so on. Therefore, it is of great commercial interest due to its potential application both in the energy industry as a fuel and in environmental CO₂ sequestration to mitigate high atmospheric CO₂ levels. The methanol production in the year 2023 was 98.9×10^6 Tm and it is projected that by the year 2027, more than 8×10^6 Tm will be obtained from e-methanol (produced from captured carbon dioxide and hydrogen produced from renewable electricity, ca 5×10^6 Tm) and biomethanol (produced from sustainable biomass, ca 3×10^6 Tm). For example, currently, 4000 Tm of methanol is produced annually using biocatalysts based on copper and zinc oxide in Iceland, recycling some 5500 tons of CO₂ annually. Other approaches to achieving the same effect have also been undertaken in Germany and China [222]. The use of enzymes is an advantage for the conversion of CO₂ to methanol because of the high selectivity of the catalyzed reaction. This reduction of CO₂ to methanol is considered a green chemical process and occurs at atmospheric temperature and pressure [253].

7.3. Electrochemical Regeneration of NADH Cofactor

NADH can be regenerated at the cathode of an electrolytic cell by applying adequate electric voltage (Figure 10A). This method can be extensively applied and permits the easy separation of products [225–228]. In principle, as the supported potential difference can be as high as desired within technical constraints, the cofactor regeneration under appropriate conditions could be high, although the complete energy cycle is probably not. If a cell is supported by renewable energy, this approach can also be considered green. In the electrolytic cell, the cofactor can be the primary acceptor of electrons from the electrode (direct electrochemical regeneration, Figure 10B) or, in contrast, other molecules can accept the electrons, the cofactor being reduced by these mediators (indirect mode Figure 10B).

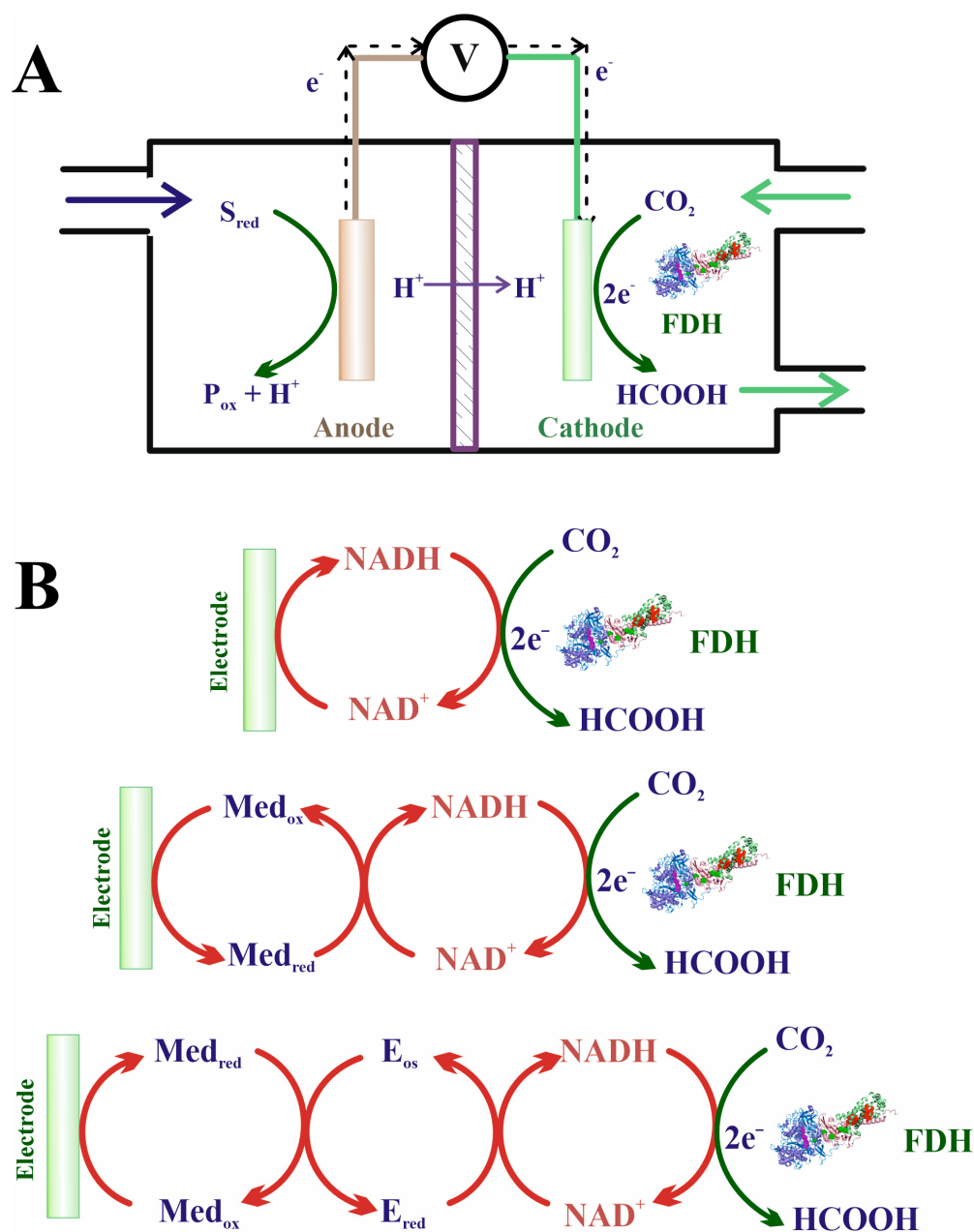


Figure 10. (A) Schematic view of an electrochemical device for obtaining formic acid from CO_2 . (B) Scheme of three possible mechanisms for transferring electrons from the cathode to the cofactor and finally to FDH: without any mediator species, with a mediator (typically a photosynthesizer molecule), and also with an enzyme intercalated between the mediator and the cofactor.

The direct mode has two intrinsic disadvantages that are very difficult to overcome: the formation of $(NAD)_2$ -inactive dimers and the necessity of using high overpotentials. The electrochemical reduction of NAD^+ molecule proceeds in two stages (Figure 11). In the first step, NAD^+ captures an electron and an NAD^* radical is formed. In the second step, an additional electron and a proton are accepted by the NAD^* radical. However, NAD^* can dimerize and, furthermore, due to the adsorption of NAD^+ onto the electrode, this dimerization process is favored over the uptake of the second electron. The need to use high overpotentials for direct $NADH$ reduction is another limitation of this method. Several studies modifying the electrode nature concluded that mass transfer between the cathode and the cofactor was a crucial step for favoring reduction versus dimer formation. These problems can be partially circumvented by selecting an appropriate electrode [255].

For instance, Ag or Pt electrodes coated on Cu foams were successfully employed for NADH regeneration [256]. The existence of a mediator on the electrode surface is another key point for avoiding dimer formation. Mediators such as (2,2-bipyridyl) (pentamethylcyclopentadienyl)rhodium, $[\text{Cp}^*\text{Rh}(\text{bpy})]$, and methylviologen (MV) have been shown to decrease the overpotential for NADH regeneration, facilitating electron transfer and NADH recovery [221].

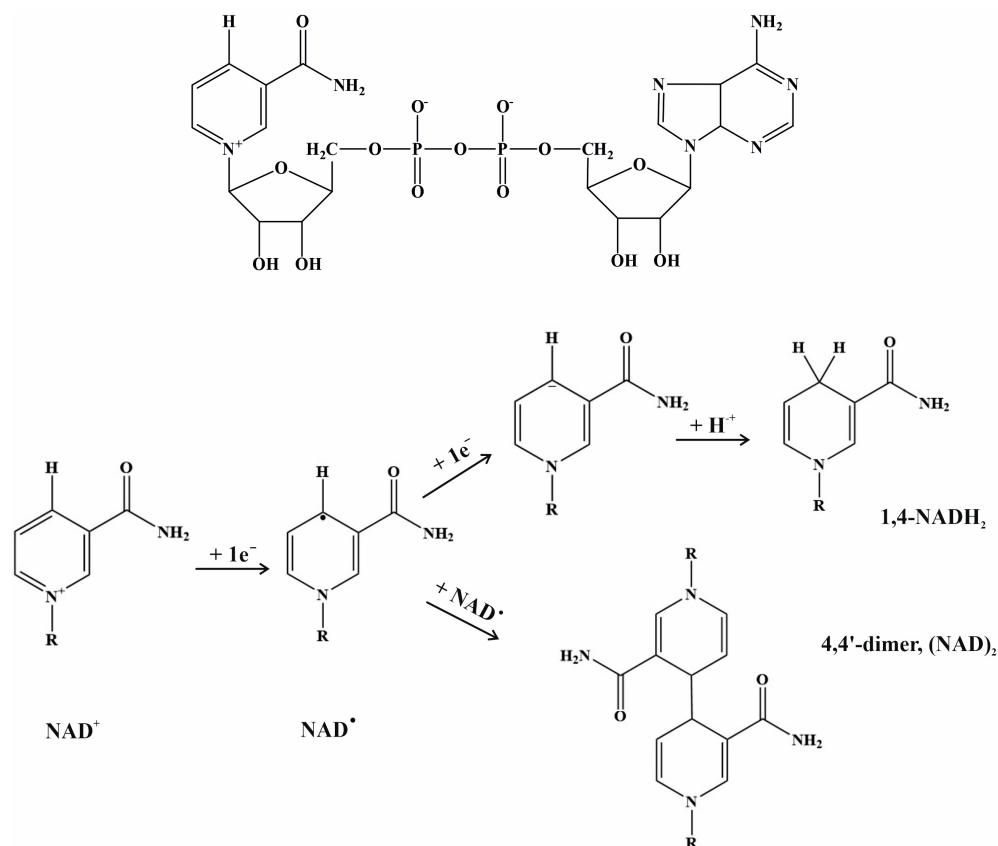


Figure 11. The two reduction steps of NAD⁺. The addition of one electron produces the radical NAD[•]. This radical can either be reduced by the addition of an additional electron and a proton (up), to produce 1,4-NADH (up) or combine with another radical to generate the 4,4'-dimer (NAD)₂.

In an elegant experiment, Song et al. generated a cysteine residue in FDH from *Thiobacillus* sp. KNK65MA and attached the mutated enzyme to copper nanoparticles (CuNPs) deposited on the electrode surface [257]. Polyethylene glycol (PEG) was then used to crosslink the FDH-NAD⁺ system so that the cofactor could swing from CuNP to the enzyme and vice versa. No mediator was used. This system produced 8.5 mM formate, several times that achieved with a free enzyme.

The enzyme cascade for methanol production from CO₂ (Figure 10B) has also been widely performed using an electrochemical approach. Addo et al. coupled the multienzyme cascade for producing methanol from CO₂ to a polyneutral red electrode to regenerate NADH [258]. Electroenzymatic reduction in CO₂ was also achieved with maximum Faradaic efficiency ($99 \pm 5\%$) by the immobilization of Mo-dependent FDH from *E. coli* at the surface of a carbon electrode [259]. Here, reduction was achieved through the mediator cobaltocene being covalently bound to the linear polymer poly(allylamine), which transferred electrons from the cathode to the enzyme. In another interesting study, CO₂ reduction was electrochemically achieved by using copper deposited in a glassy carbon electrode, the Rh(III) complex $[\text{Cp}^*\text{Rh}(\text{bpy})\text{Cl}]^+$ (Cp* = pentamethylcyclopentadienyl; bpy = bipyridine) as a mediator and CbFDH as a biocatalyst, with yields threefold higher than those of previous analogous works with copper foil electrodes [260]. Chen et al. also used

an Rh(III) complex as a mediator for the electrochemical regeneration of NADH [261]. MOF NU-1006 containing *Cb*FDH was deposited on the electrode surface of fluorine-doped tin oxide glass. NU-1006 is a mesoporous material with a channel size that can accommodate *Cb*FDH ($6 \times 4 \times 11$ nm). The electrode was obtained using a multilayer system in which the entrapped enzyme was the latter and was in contact with the solution containing CO_2 . Regeneration of the cofactor was optimum because of the modified electrode, with a formate generation rate of $79 \pm 3.4 \text{ mM h}^{-1}$. All these experiments have the advantage of easy regeneration of NADH, due to the possibility of creating a sufficient negative redox potential. However, its translation towards more complex systems, such as a multicascade system coupled to other oxidoreductases, no longer seems biotechnologically accessible.

Barin et al. immobilized *Cb*FDH on polystyrene nanofibers and NADH in a copper foam electrode [262]. Although the activity of this system was inferior to that of the free enzyme, the immobilized enzyme was stable for a long period (41% of the initial yield after 20 days) and had acceptable reusability after eight cycles (53% of the initial activity). The authors also observed an inhibitory effect of NADH at concentrations higher than 0.51 mM. Indeed, several studies have demonstrated the inhibition of FDH by the cofactor at concentrations higher than millimolar. On the other hand, Zhang et al. encapsulated FDH, FaldDH, and ADH in MOF ZIF-8 and used the Rh complex ($\text{Cp}^*\text{Rh}(2,2'\text{-bipyridyl-5,5'-dicarboxylic acid})\text{Cl}_2$) grafted on the cathode as a mediator to perform NADH regeneration [263]. The concentration of methanol obtained was fivefold (from 0.061 to 0.320 mM) with the encapsulated enzyme compared to the free enzyme. Moreover, when NADH was electrochemically regenerated, an increase of 0.742 mM was observed. This again demonstrated that electrochemical NADH regeneration is probably the best approach to achieving this goal.

7.4. Photochemical NADH Regeneration

Photochemical reactions are another suitable approach to reducing CO_2 [31,229–232]. The energy arises from light, which is an inexpensive, renewable, and clean source. This requires a supply of electrons and a photosensitizer mediator scheme (Figure 12). The electron donor, *D*, also called the sacrificial agent, is typically a stable solute present in large concentrations that can be easily oxidized using a photosensitizer in its excited state [223]. Amines such as triethanolamine (TEOA), triethylamine (TEA), and ethylenediaminetetraacetic acid (EDTA) are the most commonly employed, although many other species, such as 3-(*N*-morpholino) propanesulfonic acid, 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid, 2-(*N*-morpholino)ethanesulfonic acid mercaptoethanol, phosphite, propanol, and even molecular hydrogen gas have also been used [30]. The redox potentials of these systems must be low enough to be oxidized using the photosensitizer so that a pool of electrons is always present in the solution. This electron donor supplier must be present at high concentrations, typically not lower than 100 mM, for the system to be effective. A water molecule has also been proposed, imitating natural photosynthesis, as a sacrificial agent; however, the redox potential of $\text{H}_2/\text{H}_2\text{O}$ is too low to produce an efficient system.

The key to these reactions is photosensitizers, molecules that are excited by light [31,232]. These species act as electron mediators, capturing electrons from the sacrificial agent and transferring them to the cofactor (Figure 12). Then, the cofactor is regenerated, and CO_2 can be reduced using FDH. The mediator is a molecule that is stable in its oxidized state and can be photoactivated, passing from the ground to an excited state in which this molecule has a much higher redox potential, being able to oxidize the sacrificial agent and becoming reduced. In this reduced state, the mediator is highly reducing and immediately relinquishes the electrons to the cofactor. This transfer proceeds with a proton cession, such that the cofactor in its oxidation state (NAD^+ or others) is reduced (NADH or analogous). The photosensitizers must fulfill three requirements [30,264]: First, the band gap between the HOMO and LUMO must be low enough to accept an electron from visible light (see Figure 3). This is normally satisfied in extended π – π -conjugated systems or semiconductors. Second, in the excited state, its redox potential must be sufficiently high to oxidize the

sacrificial agent under solution conditions. Finally, the ground state redox potential must be sufficiently low to reduce the cofactor. Other chemical (stability), economical (inexpensive), and environmental (clean) requirements must also be satisfied.

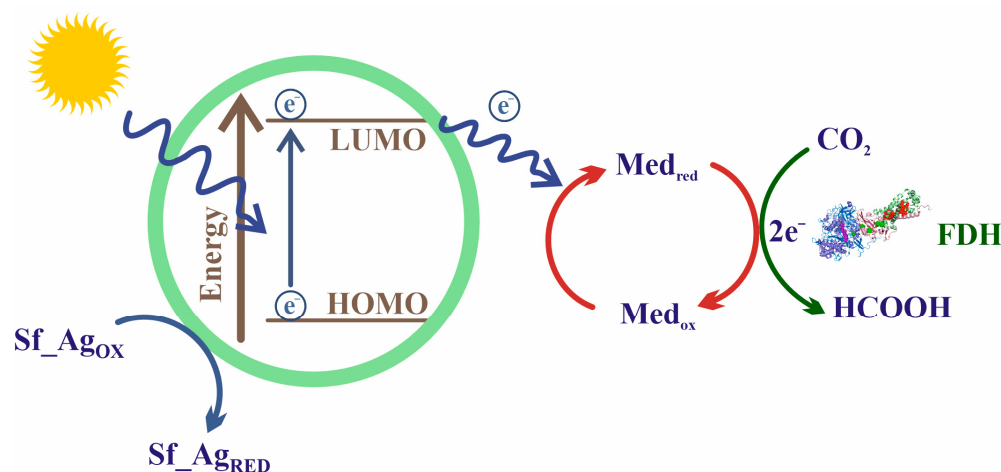


Figure 12. Scheme of the photochemical reduction of CO₂ using FDH.

Various photosensitizers have also been used. Rh(III) and Ru(III) bipyridyl complexes are well known for their photosensitivity; consequently, their use has been extended [264–269]. Guo et al. employed $\text{Cp}^*\text{Rh}(\text{bpy})(\text{H}_2\text{O})]^{2+}$ (Cp = cyclopentadienyl; bpy = 2,2'-bipyridyl) as a synthesizer in a system where FDH and FalDH were immobilized on polyethylene membranes doped with the widely used semiconductor TiO₂ [270]. A comparison between the results using water or EDTA as a sacrificial agent and as a function of pH was described, the latter being much more efficient for formaldehyde production. The optimal FDH:FalDH ratio was 1:0.3, reaching up to 6.5% formaldehyde production after 4 h of reaction. Photosensitizers composed of xanthene dyes have also been used. Kim and col. combined eosin Y with cobaloxime complexes for regenerating NADH using TEOA as the sacrificial electron donor [271]. The system exhibited an acceptable turnover number for formate generation (*ca.* 1.6) and an optimal NADH production (0.038 mM/h). In situ changes in eosin Y infrared and UV-visible spectroscopy properties eosin Y was also used to follow in situ formic acid generation using infrared and UV-visible absorption spectroscopies [272]. Interestingly, EDTA was used not only as an electron sacrificial agent but also as a source of CO₂, without any additional electron carriers.

Nanomaterials, which act as porphyrin-based photosensitizers, are another set of well-developed approaches for acting in these photochemical reactions. Ji and coworkers designed a biomimetic chlorosome by combining porphyrin, eosin Y, and $[\text{Cp}^*\text{RhCl}_2]_2$ to generate supramolecular assemblies [273]. Using TEOA as the ultimate electron donor agent, an enzyme cascade was coupled, obtaining 38 μM methanol from CO₂ after 2 h of reaction. MOFs have revolutionized many technochemical applications [157]. The MOFs used for CO₂ reduction are also basic pivots in this respect. MOFs contain molecules and holes with different chemical and physical features (hydrophilicity, hydrophobicity, acid-base properties, photochemical features, etc.) that make them ideal for catalysis in general, and photobiocatalysis in particular. An excellent recent example is Xing's work [229], where a porphyrin ligand was covalently bound to a Zr-based MOF and, posteriorly, the complex $\text{Cp}^*\text{Rh}(\text{bpydc})\text{Cl}_2$ (bpydc = 2,2'-bipyridine-5,5'-dicarboxylic acid) was incorporated into the organic frame, generating a system that can be activated by light owing to the porphyrin, the Rh(III) complex, and original aromatic ligands of the MOF. FDH was then immobilized via electrostatic entrapment. Using TEOA, up to 244 $\mu\text{g}/\text{mL}$ of formic acid was formed after 4 h of reaction. A similar approach employing an MOF based on pyrene skeleton linkers, NU-1006, has also been developed [261]. In this case, the Rh(III) complex attached to the MOF was used as an electron mediator, reducing NADH and being reduced by a pyrene photosensitizer. FDH was also trapped in the MOF and the reaction took place

with a yield of formic acid production of 0.144 ± 0.003 mM after 24 h, while the cofactor was regenerated at a rate of 28 mMh^{-1} . In another recent study, MOF Mil-125-NH₂ was functionalized with the Rh complex in such a way that it was covalently fixed to the secondary sphere of the MOF and FDH was subsequently immobilized [274]. The system obtained a formic acid yield of 9.5 mM in 24 h, whereas the NADH regeneration was 64%.

Graphitic carbon nitride (g-C₃N₄)-based materials are other fascinating agents with extremely interesting properties, such as photocatalysts [275]. These materials have also been used in biocatalysis and to generate formate from CO₂. Zeng et al. used g-C₃N₄ doped with the metal dichalcogenide WS₂, which provides g-C₃N₄ with specific semiconductor properties that can be used as photosensitizers [276]. Using [Cp*Rh(phen)H₂O]²⁺ (phen = 1,10-phenanthroline) as a mediator and TEOA as the sacrificial agent, a multi-cascade system for obtaining methanol from CO₂ was designed with acceptable results (methanol productivity $372.1 \mu\text{mol h}^{-1} \text{ gcat}^{-1}$). In a similar approach, Meng and col. designed nanospheres with thiophene incorporated into hollows in a double shell that acted as the photosynthesizer [277]. This nanomaterial was coupled to [Cp*Rh(bpy)H₂O]²⁺, which in turn was coupled to the cofactor to reduce CO₂. Optimal NADH yield regeneration was obtained (74%). Silver nanoclusters combined with TiO₂ and g-C₃N₄ have also been employed in efficient devices for formate production [278]. These nanoclusters are good light sensors, making it possible to obtain good yields in CO₂ uptake with the metal-dependent FDH from *Clostridium ljungdahlii*. Carbon nitride has also been used in microbial electrosynthesis. Here, the photoanode chamber was formed using an activated carbon fiber (ACF) supported by NiCoWO₄ in g-C₃N₄ [279]. The oxidation of water by light in this chamber provides the electrons for reducing CO₂ in the cathode chamber, formed by a g-C₃N₄/ACF (without NiCoWO₄), in which a culture of *E. coli* was adhered as a biofilm. FDH from *E. coli* produced the reduction of CO₂ due to the electrons arriving from the biocathode and the protons, also produced in the cathode by the photochemical decomposition of water, which crossed a cation exchange membrane. The witty system, first applied in CO₂ reduction, provided highly efficient formate synthesis (12.8 mM per day).

When dealing with photochemical reactions, a critical point is the reactive oxygen species (ROS) that can be generated by photoexcitation processes. To avoid this, systems mimicking chloroplasts have been developed, creating divided compartments where reactions occur separately. The photoactivation process was achieved by combining a Rh(III) complex conjugated onto g-C₃N₄ previously modified using thiophene, with triethanolamine used as the electron sacrificial agent [280]. The key to this approach arises from the encapsulation of FDH into a MOF (MAF-7), which protects the enzyme from photoactivation reactions. NADH shuttled electrons from the reduction compartment towards FDH, obtaining 16.75 mM of formic acid after 9 h of illumination. Immobilization of FDH together with CA in the TOF ZIF-8 was also carried out by Yu et al. in another system imitating photosynthesis and with g-C₃N₄ as the photosynthesizer [281]. CA accelerated the interconversion of CO₂/HCO₃[−], allowing for an effective mass transfer rate between the gas and liquid phases. The authors reported production of 243 μM formic acid with excellent system stability since the yield production was above 80% after 10 batches.

Graphene has also been employed as a photocatalyst in systems that imitate photosynthesis and produce formic acid from CO₂ [210]. Ji's group conceived and developed a nanofiber polyurethane system as a support where the cationic electrolyte, a polyaniline, was deposited onto graphene oxide, while the cascade of proteins FDH, FaldH, and ADH was entrapped in the hollows of the nanofiber to generate methanol [246,282,283]. Finally, polymers that mimic photosystems in CO₂ reduction have also been candidates for biotechnological applications. Kim et al. reported a photosystem based on polydiacetylenes and covalently attached (phen)Ru(bpy)₂ as the mediator [284]. The system exhibited acceptable regeneration values for NADH.

It should also be noted that semiartificial photocatalysis centers have been also designed to reduce CO₂. Sokol et al. generated a sophisticated electrocell in which the photoanode was the photosystem II from *Thermosynechococcus elongatus* embedded in a

redox polymer composed of the complex $[\text{Os}(\text{bipy})_2\text{Cl}]\text{Cl}$ bound to a polymer of a derivative of allylamine was, in turn, deposited on TiO_2 [285]. Here, light (680 nm) oxidized water to O_2 , generating electrons that were shuttled towards the cathode, where FDH from *Desulfovibrio vulgaris* had adhered to TiO_2 but, in this case, coated with a fluorine tin oxide. Excellent Faraday efficiencies ($>70\%$) and acceptable yields ($0.185 \mu\text{mol}/\text{cm}^2$) were obtained, although the system exhibited the progressive photodegradation of PSII.

Due to its high substrate and product selectivity as a biocatalyst, FDH has been used in recent years to obtain chemical products from reduced forms of CO_2 , such as formic acid. In addition to its own applicability, this acid is used as a springboard product to obtain a wide variety of derivatives. For example, CO_2 can be converted to oxalic acid through the coupling of two formate molecules [286]. This can occur via a metal–formate intermediate (typically sodium or potassium alkaline metals), obtained from the electro- or photocatalytic reduction of CO_2 coupled to FDH. After obtaining this metal–formate, two molecules are coupled to give oxalate, which, after acidification, gives rise to oxalic acid. Oxalic acid produced through this coupling route can be used in various industries, including pharmaceuticals, as a component of some antibiotics, or textiles and in other industries such as food (beer or wine production) or chemicals [286]. In addition, from the reduction of this oxalic acid, a wide variety of derivatives are obtained. The first oxalic acid reduction product is glyoxylic acid, which, after reduction, produces glycolic acid and also ethylene glycol. These chemicals are the starting points in the production of agrochemicals, flavors, cosmetics, and polymers [287].

7.5. CO_2 Reduction by Whole-Cell Bacteria

Finally, it should be pointed out that recent studies focused on CO_2 reduction encompass an even more open overview of what nature provides us in this area. Indeed, whole cells can also be used for fixing CO_2 , using H_2 as the reductant agent [288–290], electrochemical reduction [291], or even a combination of H_2 and photocatalysis [290]. Whole-cell biocatalysis is a promising method developed in the last few years that shows high efficacy and selectivity and can be used in soft conditions. Moreover, since there is no requirement for purifying enzymes, the most expensive step in biocatalytic processes, the whole-cell approach is a hopeful methodology that will probably have the highest efficiency/cost ratio. It is also relatively easy to adjust the experimental conditions to be developed if the suitable organism or correct molecular biology tools in them are adequately employed. Table 5 reports the most relevant results obtained in the CO_2 reduction to formate using the whole-cell approach. The culture of these microorganisms supplied by H_2 gas reduces CO_2 obtaining extraordinarily high concentrations of formate. Methylobacteria species are satisfactory in this aspect [290].

Table 5. CO₂ reduction to obtain formate using whole-cell approaches.

Reduction	Organism	Significant Assay Features/Conditions	Formate Production	Ref
Hydrogenation	Overexpressed genes of FDH from <i>E. coli</i> , <i>Clostridium carboxidivorans</i> , <i>Pyrococcus furiosus</i> and <i>Methanobacterium thermos-formicicum</i> in <i>E. coli</i> JM109(DE3)	H ₂ atmosphere. Wet cell pellet (0.5 g wet cells/mL) resuspended in 50 mM sodium phosphate buffer, pH 7.0, containing 0.25 M sodium bicarbonate as a source of CO ₂ . Incubation 37 °C.	The highest formic yield (FDH from <i>P. furiosus</i>) was more than 1 g L ⁻¹ h ⁻¹	[292]
Hydrogenation	<i>Escherichia coli</i>	CO ₂ and H ₂ placed under pressure (up to 10 bar). First, pressurized the system to rapidly convert 100% conversion of gaseous CO ₂ to formic acid. Next, NaOH addition to the <i>E. coli</i> cell suspension (pH > 8).	Formate concentration to 150 (4 bar) and 200 (6 bar) mmol L ⁻¹ . Increasing the pressure to 10 bar (122.88 mmol L ⁻¹ CO ₂ and 3.61 mmol L ⁻¹ H ₂): > 0.5 mol L ⁻¹ formate at 23 h of reaction.	[293]
Hydrogenation	<i>Acetobacterium woodii</i> and <i>Thermoanaerobacter kivui</i>	Cells grown with 28 mM glucose or 0.1 M pyruvate (50 mM imidazole, 20 mM KCl, 20 mM MgSO ₄ , 2 mM DTE, 4 µM resazurin, pH 7.0). 1 mg/mL in an anoxic medium. Shaking for 10 min (60 °C), with additional 300 mM KHCO ₃ . The experiment started by replacing the gas phase with H ₂ + CO ₂ (80:20%, 2 × 10 ⁵ Pa).	Optimal formate production rates of 234 mmol g ⁻¹ _{protein} h ⁻¹	[270]
Hydrogenation	H ₂ -dependent CO ₂ reductase from <i>Acetobacterium woodii</i> expressed in <i>E. coli</i> JM109	Whole-cell <i>E. coli</i> in presence of formate (10 mM) and methylviologen (10 mM) as electron acceptor. <i>E. coli</i> cells incubated with H ₂ + CO ₂ (80:20%, 1.1 × 10 ⁵ Pa).	6 mM formic acid in 60 h. Addition of 2.5 mM HCO ₃ ⁻ increased 4-fold the formate generation.	[294]
Hydrogenation	<i>Acetobacterium woodii</i> and <i>Thermoanaerobacter kivui</i>	50 mM imidazole, 20 mM MgSO ₄ , 20 mM KCl, 20 mM NaCl, 2 mM DTE, pH 7.0; or 50 mM K-phosphate, 20 mM KCl, 20 mM NaCl, 2 mM DTE, pH 7.0) was maintained at 30 and 60 °C for <i>A. woodii</i> and <i>T. kivui</i> , respectively. Gas flow rate was maintained at a value of 25 mL/min	60 mM and 80 formic acid generation after 5 h of reaction for <i>A. woodii</i> and <i>T. kivui</i> , respectively.	[295]
Electrochemical	<i>Methylobacterium extorquens</i> AM1,	System: 1 mM H ₂ SO ₄ with a platinum electrode placed in the anode; protons supplied to the cathode through a proton-exchange membrane. 1.9 g wet cells, 10 mM methyl viologen (MV) added to the cathode reactor as an electron mediator	Formate concentrations of up to 60 mM, 80 h (1.9 g wet cells, 10 mM MV, pH 7.0)	[290]

Table 5. Cont.

Reduction	Organism	Significant Assay Features/Conditions	Formate Production	Ref
Electrochemical	<i>Shewanella oneidensis</i> MR-1	The electrochemical cell with two compartments divided by a proton-exchange memberane. Copper plates and Ag/AgCl electrodes. Whole-cell <i>S. oneidensis</i> MR-1 (wet-cell, 0.5 g) and MV, 10 mM). RT, anaerobic conditions.	Formic acid at 0.59 mM h ⁻¹ for 24 h. Medium supplemented with fumarate and nitrate: 1.9 mM h ⁻¹ for 72 h. LB supplemented with 40 mM fumarate, 1mM nitrate and 20 mM DL-lactate: 136.84 mM formic acid at 72 h. (3.8 mM h ⁻¹ g ⁻¹ _{wet-cell})	[296]
Electrochemical	<i>E. coli</i>	NaHCO ₃ electrolyte saturated with N ₂ or CO ₂ media at three different poised potentials, i.e., 0.4, 0.8 and 1.0 V vs. Ag/AgCl. <i>E. coli</i> -immobilized FePc-CDC/ACF and ACF electrodes in the presence of the NR mediator (FePc: iron pfthalocyanine; CDC: carbide-derived carbon; ACF: activated carbon fiber; NR: Neutral Red).	Maximum formate production rate of ~30 mg/L-h under CO ₂ flow (120 mg/L-h) with NR mediator	[297]
Electrochemical	<i>Methylobacterium extorquens</i> AM1	Nafion 115 (proton permeable) membrane. Cathode: 0.6 g cell pellet (potassium phosphate 200 mM, pH 7.0) as catholyte. Electron mediator: ethyl viologen 10 mM. 0.6 g cell pellet suspended into catholyte (200 mM-potassium phosphate at pH 7) saturated by CO ₂ purging (30 min). Water splitting reaction in 100 mM-H ₂ SO ₄ .	Formate production: 6 mM h ⁻¹	[298]
Electrochemical	<i>Shewanella loihica</i> PV-4	Cathode: biohydrogel formed by <i>Shewanella loihica</i> PV-4 immobilized in graphene oxide.	High Faradaic efficiency (~99.5%) and 46-fold increase of formate titer without exogenous mediator (4.2 mM formate. 36 h)	[299]
Hydrogen/carbohydrate fermentation	<i>Saccharomyces cerevisiae</i>	Two phases: Glucose fermentation for generating CO ₂ CO ₂ Ru catalysis hydrogenation	26% of the CO ₂ was hydrogenated. Addition of His 150 mM: 128 mM in formic acid at 48 h.	[300]
Photocatalytic hydrogenation	<i>Shewanella oneidensis</i> MR-1	Anaerobic conditions: N ₂ -filled chamber and samples irradiated from outside the chamber by a KL5125 Cold 150W light source. Irradiation into MV, with triethanolamine (TEOA) as sacrificial agent in 50 mM HEPES, 50 mM NaCl, pH 7, 23 °C.	Formate (~1500 nmol) was produced when MR-1 was incubated with CO ₂ (~5000 nmol) after 48 h incubation	[301]

Leo et al. genetically modified *E. coli* cells to express the hydrogen-dependent CO₂ reductase from *Acetobacterium woodii* [294]. In this bacterial culture, increasing the cell density up to 30 mg/mL resulted in a formate production yield of 6 mM/h, whereas the addition of 2.5 mM HCO₃[−] increased the formate generation rate fourfold. In this way, the authors confirmed that *E. coli* CA also played an important role in providing the correct substrate to FDH. Müller's laboratory took advantage of the metabolic machinery of the acetogenic bacteria *Acetobacterium woodii* and *Thermoanaerobacter kivui* to convert H₂ and CO₂ from syngas into formic acid, obtaining production rates of 234 mmol g^{−1}_{protein} h^{−1} [288]. *E. coli* has been also used for formate generation with and without overexpression of exogenous FDH genes. Indeed, as commented previously (see Section 3.2), *E. coli* has two metal-dependent FDHs capable of reducing CO₂ to formate. This bacterium has been used in the whole-cell strategy with extraordinarily high formate generation. Indeed, more than 0.5 molL^{−1} of formate concentration was achieved when *E. coli* cells were grown under CO₂ and H₂ gas at 10 bar pressure for 23 h [293]. *E. coli* JM109(DE3) strain has also been used as a host bacterium where to overexpress FDHs from *Clostridium carboxidivorans*, *Pyrococcus furiosus* (Pf) and *Methanobacterium thermosformicum*, also obtaining excellent yields [292]. Indeed, the highest formate generation yield of this study, obtained with PfFDH, was more than 1 gL^{−1}h^{−1}.

Electrochemical reduction is another approach for obtaining high yields of formate within the whole-cell frame. An excellent and very recently released review describes in detail this methodology [291]. Table 5 highlights the remarkable features of the—up to now—few articles published in this area concerning formic acid generation. In these studies, yields are as good as for those using the H₂ reduction strategy. For instance, Le et al. achieved to obtain 3.8 mM h^{−1} g^{−1}_{wet-cell} of formic acid in the cathode electrode at 72 h when *Shewanella oneidensis* MR-1 was grown in LB media supplemented with nitrate 1 mM and DL-lactate 20 mM [296].

It is also remarkable that two other different approaches to applying whole-cell have been developed. Guntermann and col. devised a two-phase system where *Saccharomyces cerevisiae* D-glucose fermentation produces ethanol and CO₂ in the water phase. This was coupled to a hydrogen gas source that, in the presence of a Ru catalyst solved in the tetradecane phase, generated formic acid at concentrations of 128 mM after 48 h of reaction [300]. On the other hand, light-driven photocatalytic hydrogenation has been applied to a culture of *Shewanella oneidensis* MR-1 in anaerobic conditions [301]. Using methyl viologen as a photoactivated molecule and triethanolamine (TEOA) as the sacrificial agent, Rowe et al. obtained formic acid at concentrations higher than 1.5 mmol for 48 h of reaction.

Finally, it is worth remarking that the use of microorganisms that incorporate formic acid into their metabolic routes has also been another object of research in the last years. Formatotrophic organisms (capable of assimilating formate for use as a carbon source) offer a new approach to formate utilization, although they are difficult to cultivate, which partially restricts their applicability [302,303]. One strategy is to biotechnologically adapt microorganisms to assimilate formate by adapting their metabolism to the formatotrophic growth model through metabolic engineering tools. Both methanol and formate can be assimilated in the central metabolism through various metabolic pathways, and the bio-production of other compounds such as ethanol, acetone, isopropanol, or short- and medium-chain fatty acids and alcohols from these compounds is very promising [304,305]. To this end, the introduction of pathways, such as the Calvin cycle, the serine cycle, the acetyl-CoA reductive pathway or the glycine pathway in hosts, or the design of new pathways, is proposed [304,305]. The most commonly used hosts are *E. coli*, *S. cerevisiae* and *Cupriavidus necator*, as they are easy to modify with genetic engineering tools and are industrially applicable [306].

Synthetic pathways, such as the reductive glycine pathway (rGly, pathway for formate assimilation), can support higher biological yields than natural pathways and could therefore be implemented in a variety of microorganisms. The rGly pathway has been

introduced into the *E. coli* host by redesigning its central metabolism to be able to assimilate formate. In terms of strategy, the pathway to be integrated is divided into modules and introduced together into the host bacteria to express the enzymes necessary to form the complete metabolic pathway [304,305]. With this methodology, a strain capable of growing from formate with a doubling time of ~70 h and a growth yield of ~1.5 g cell dry weight (gCDW) per mole of formate was achieved [303]. In this way, products such as lactate or isobutanol, both pyruvate derivatives, were obtained. Cotton et al. used *C. necator* to modify it with rGly, so that it was able to grow under formate-rich conditions. In this respect, the production of methylketones, isoprenoids and terpenes, isobutanol, alkanes, and alkenes from CO₂ using *C. necator* seems particularly promising. With the same microorganism, the Calvin cycle was also used for the assimilation of formate or methanol, although with low energy efficiencies (20–35%) [307]. On the other hand, Collas et al. very recently devised an intelligent approach in which they engineered a crotonate biosynthetic pathway in *C. necator* [306]. This mechanism permits the conversion of formic acid in crotonate with extraordinary yields in a continuous process. Indeed, using this approach, they obtained 148 mg/L of product. This is a new starting point for the generation of new value-added chemicals.

Although whole-cell technology is still incipient, new opportunities that open a window to capture CO₂ under standard conditions, where nature develops, are enormous.

8. Conclusions and Perspectives

A panoramic overview of the state-of-the-art on carbon capture and its reduction in C₁ forms by employing the biocatalysts CA and FDH was presented. CO₂ uptake is one of the main challenges that science, in general, and chemistry, in particular, face today. Nature offers appropriate tools and clues for finding approaches to solve this huge problem posed to mankind. CCSU techniques are well developed in the laboratory and, to a lesser extent, at a large scale, although their application is still far from being appropriately adapted for obtaining high yields without great energetic costs. While CA immobilization is currently a reality and it stabilizes the enzyme allowing its reuse, the employment of recombinant CAs with higher resistance to temperature and pressure is a field that still remains to be fully developed. These molecular biology tools will help speed up CO₂/HCO₃[−] conversion at an industrial scale in the next few years.

Carbon dioxide reduction is also a reality at the laboratory scale, and research lines are well established, although they are still far from showing high yields. The formic acid molecule is a rich form for transporting hydrogen in an efficient way and is the primary step for producing many more reduced molecules in such a way that carbon is recycled and reused as an energy H₂ vector. Mass transfer from the gas to the solution is a limiting step in reaching the substrate to the biocatalyst. On the other hand, improvements in the stability and high performance of FDHs, as well as in their stability and reuse via immobilization, have been developed, with a significant explosion of this research in the last decade, despite still being a relatively virgin area. However, its application to larger scales has not been established.

Measures to mitigate climate change are urgently needed. CAs and FDHs are excellent devices for capturing CO₂ and transforming it into fuel, which is an interesting way to reduce two problems into one. According to the exponential progress existing in the research in this area nowadays, surprising approaches and solutions to these problems will probably be found not in the next decades but in years.

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References

1. National Center for Environmental Information. Annual 2021 Global Climate Report. Available online: <https://www.ncei.noaa.gov/access/monitoring/monthly-report/global/202113> (accessed on 24 February 2023).
2. Intergovernmental Panel on Climate Change (IPCC). Climate Change 2022, Mitigation of Climate Change. Available online: https://www.ipcc.ch/report/ar6/wg3/downloads/report/IPCC_AR6_WGIII_FullReport.pdf (accessed on 24 February 2023).
3. Fellowes, T.E.; Anggadi, F.; Byrne, M.; Vila-Concejo, A.; Bruce, E.; Baker, E. Stability of Coral Reef Islands and Associated Legal Maritime Zones in a Changing Ocean. *Environ. Res. Lett.* **2022**, *17*, 093003. [\[CrossRef\]](#)
4. Borges, F.O.; Sampaio, E.; Santos, C.P.; Rosa, R. Impacts of Low Oxygen on Marine Life: Neglected, but a Crucial Priority for Research. *Biol. Bull.* **2022**, *243*, 104–119. [\[CrossRef\]](#) [\[PubMed\]](#)
5. Arrhenius, S.; Holden, E.S. On the Influence of Carbonic Acid in the Air Upon the Temperature of the Earth. *Publ. Astron. Soc. Pac.* **1897**, *9*, 14–24. [\[CrossRef\]](#)
6. Arrhenius, S. The Celestial Bodies, In Particular the Earth, as Abodes of Living Beings. In *Worlds in the Making: The Evolution of the Universe*; Harper: New York, NY, USA, 1908; pp. 39–63.
7. Friedlingstein, P.; Andrew, R.M.; Rogelj, J.; Peters, G.P.; Canadell, J.G.; Knutti, R.; Luderer, G.; Raupach, M.R.; Schaeffer, M.; van Vuuren, D.P.; et al. Persistent Growth of CO₂ Emissions and Implications for Reaching Climate Targets. *Nat. Geosci.* **2014**, *7*, 709–715. [\[CrossRef\]](#)
8. Lindsey, R. Climate Change: Atmospheric Carbon Dioxide. Science and Information for a Climate-Smart Nation. Available online: <https://www.climate.gov/news-features/understanding-climate/climate-change-atmospheric-carbon-dioxide> (accessed on 24 February 2023).
9. IPCC. Climate Change 2007: Synthesis Report. Summary for Policymakers. Valencia (Spain) 2007. Available online: <https://www.ctc-n.org/resources/ipcc-fourth-assessment-report-climate-change-2007-synthesis-report-summary-policymakers> (accessed on 24 February 2023).
10. EPA. United States Environmental Protection Agency. Overview of Greenhouse Gases. Available online: <https://www.epa.gov/ghgemissions/overview-greenhouse-gases> (accessed on 24 February 2023).
11. Hoegh-Guldberg, O.; Jacob, D.; Taylor, M.; Bolanos, T.G.; Bindi, M.; Brown, S.; Camilloni, I.A.; Diedhiou, A.; Djalante, R.; Ebi, K.; et al. The Human Imperative of Stabilizing Global Climate Change at 1.5 Degrees C. *Science* **2019**, *365*, eaaw6974. [\[CrossRef\]](#)
12. Rajabloo, T.; Valee, J.; Marenne, Y.; Coppens, L.; De Ceuninck, W. Carbon Capture and Utilization for Industrial Applications. *Energy Rep.* **2023**, *9*, 111–116. [\[CrossRef\]](#)
13. Kamkeng, A.D.N.; Wang, M.; Hu, J.; Du, W.; Qian, F. Transformation technologies for CO₂ utilisation: Current status, challenges and future prospects. *Chem. Eng. J.* **2021**, *409*, 128138. [\[CrossRef\]](#)
14. Villa, R.; Porcar, R.; Nieto, S.; Donaire, A.; García-Verdugo, E.; Luis, S.V.; Lozano-Rodríguez, P. Sustainable chemo-enzymatic synthesis of glycerol carbonate (meth)acrylate from glycidol and carbon dioxide enabled by ionic liquid technologies. *Green Chem.* **2021**, *11*, 4191–4200. [\[CrossRef\]](#)
15. Knoche, W. Chemical Reactions of CO₂ in Water. In *Biophysics and Physiology of Carbon Dioxide, Proceedings in Life Sciences*; Bauer, C., Gros, G., Bartels, H., Eds.; Springer: Berlin/Heidelberg, Germany, 1980. [\[CrossRef\]](#)
16. Greenwood, N.N.; Earnshaw, A. *Chemistry of the Elements*, 2nd ed.; Butterworth-Heinemann, Elsevier: Oxford, UK, 1997. [\[CrossRef\]](#)
17. Talekar, S.; Jo, B.H.; Dordick, J.S.; Kim, J. Carbonic Anhydrase for CO₂ Capture, Conversion and Utilization. *Curr. Opin. Biotechnol.* **2022**, *74*, 230–240. [\[CrossRef\]](#)
18. Lindsag, S. Structure and Mechanism of Carbonic Anhydrase. *Pharmacol. Ther.* **1997**, *74*, 1–20. [\[CrossRef\]](#)
19. De Oliveira Maciel, A.; Christakopoulos, P.; Rova, U.; Antonopoulou, I. Carbonic Anhydrase to Boost CO₂ Sequestration: Improving Carbon Capture Utilization and Storage (CCUS). *Chemosphere* **2022**, *299*, 134419. [\[CrossRef\]](#)
20. Yoshimoto, M.; Walde, P. Immobilized Carbonic Anhydrase: Preparation, Characteristics and Biotechnological Applications. *World J. Microbiol. Biotechnol.* **2018**, *34*, 151. [\[CrossRef\]](#)
21. Appel, A.M.; Bercaw, J.E.; Bocarsly, A.B.; Dobbek, H.; DuBois, D.L.; Dupuis, M.; Ferry, J.G.; Fujita, E.; Hille, R.; Kenis, P.J.A.; et al. Frontiers, Opportunities, and Challenges in Biochemical and Chemical Catalysis of CO₂ Fixation. *Chem. Rev.* **2013**, *113*, 6621–6658. [\[CrossRef\]](#)
22. Bierbaumer, S.; Nattermann, M.; Schulz, L.; Zschoche, R.; Erb, T.J.; Winkler, C.K.; Tinzl, M.; Glueck, S.M. Enzymatic Conversion of CO₂: From Natural to Artificial Utilization. *Chem. Rev.* **2023**, *123*, 5702–5754. [\[CrossRef\]](#)
23. Badger, M.R.; Sharwood, R.E. Rubisco, the Imperfect Winner: It's All about the Base. *J. Exp. Bot.* **2023**, *74*, 562–580. [\[CrossRef\]](#)
24. Bauwe, H. Photorespiration—Rubisco's Repair Crew. *J. Plant. Physiol.* **2023**, *280*, 153899. [\[CrossRef\]](#)
25. Alpdagtas, S.; Binay, B. NAD⁺-Dependent Formate Dehydrogenase: A Review. *Biocatal. Biotransform.* **2021**, *39*, 260–268. [\[CrossRef\]](#)
26. Alekseeva, A.A.; Savin, S.S.; Tishkov, V.I. NAD⁺-Dependent Formate Dehydrogenase from Plants. *Acta Nat.* **2011**, *3*, 38–54. [\[CrossRef\]](#)

27. Maia, L.B.; Moura, I.; Moura, J.J.G. Molybdenum and Tungsten-Containing Formate Dehydrogenases: Aiming to Inspire a Catalyst for Carbon Dioxide Utilization. *Inorganica Chim. Acta* **2017**, *455*, 350–363. [\[CrossRef\]](#)
28. Jormakka, M.; Byrne, B.; Iwata, S. Formate Dehydrogenase—A Versatile Enzyme in Changing Environments. *Curr. Opin. Struct. Biol.* **2003**, *13*, 418–423. [\[CrossRef\]](#)
29. Yang, J.Y.; Kerr, T.A.; Wang, X.S.; Barlow, J.M. Reducing CO₂ to HCO₂⁻ at Mild Potentials: Lessons from Formate Dehydrogenase. *J. Am. Chem. Soc.* **2020**, *142*, 19438–19445. [\[CrossRef\]](#) [\[PubMed\]](#)
30. Wang, Y.; Li, M.; Zhao, Z.; Liu, W. Effect of Carbonic Anhydrase on Enzymatic Conversion of CO₂ to Formic Acid and Optimization of Reaction Conditions. *J. Mol. Catal. B Enzym.* **2015**, *116*, 89–94. [\[CrossRef\]](#)
31. Thauer, R.K.; Jungermann, K.; Decker, K. Energy-Conservation in Chemotropic Anaerobic Bacteria. *Bacteriol. Rev.* **1977**, *41*, 100–180. [\[CrossRef\]](#)
32. Zhang, Y.; Zhao, Y.; Li, R.; Liu, J. Bioinspired NADH Regeneration Based on Conjugated Photocatalytic Systems. *Solar RRL* **2021**, *5*, 2000339. [\[CrossRef\]](#)
33. Lee, S.H.; Choi, D.S.; Kuk, S.K.; Park, C.B. Photobiocatalysis: Activating Redox Enzymes by Direct or Indirect Transfer of Photoinduced Electrons. *Angew. Chem. Int. Ed.* **2018**, *57*, 7958–7985. [\[CrossRef\]](#)
34. Supuran, C.T.; Capasso, C. An Overview of the Bacterial Carbonic Anhydrases. *Metabolites* **2017**, *7*, 56. [\[CrossRef\]](#)
35. DiMario, R.J.; Machingura, M.C.; Waldrop, G.L.; Moroney, J.V. The Many Types of Carbonic Anhydrases in Photosynthetic Organisms. *Plant Sci.* **2018**, *268*, 11–17. [\[CrossRef\]](#)
36. Kupriyanova, E.; Pronina, N.; Los, D. Carbonic Anhydrase—A Universal Enzyme of the Carbon-Based Life. *Photosynthetica* **2017**, *55*, 3–19. [\[CrossRef\]](#)
37. Zimmerman, S.A.; Ferry, J.G. The Beta and Gamma Classes of Carbonic Anhydrase. *Curr. Pharm. Des.* **2008**, *14*, 716–721. [\[CrossRef\]](#)
38. McGinn, P.J.; Morel, F.M.M. Expression and Regulation of Carbonic Anhydrases in the Marine Diatom *Thalassiosira pseudonana* and in Natural Phytoplankton Assemblages from Great Bay, New Jersey. *Physiol. Plant.* **2008**, *133*, 78–91. [\[CrossRef\]](#)
39. Roberts, S.B.; Lane, T.W.; Morel, F.M.M. Carbonic Anhydrase in the Marine Diatom *Thalassiosira weissflogii* (Bacillariophyceae). *J. Phycol.* **1997**, *33*, 845–850. [\[CrossRef\]](#)
40. Beauchemin, M.; Morse, D. δ -Carbonic Anhydrases: Structure, Distribution, and Potential Roles. In *Carbonic Anhydrases as Biocatalysts: From Theory to Medical and Industrial Applications*; Université de Montréal: Montréal, QC, Canada, 2015; pp. 337–349. [\[CrossRef\]](#)
41. Dou, Z.; Heinhorst, S.; Williams, E.B.; Murin, C.D.; Shively, J.M.; Cannon, G.C. CO₂ Fixation Kinetics of *Halothiobacillus neapolitanus* Mutant Carboxysomes Lacking Carbonic Anhydrase Suggest the Shell Acts as a Diffusional Barrier for CO₂. *J. Biol. Chem.* **2008**, *283*, 10377–10384. [\[CrossRef\]](#)
42. Alterio, V.; Langella, E.; Buonanno, M.; Esposito, D.; Nocentini, A.; Berrino, E.; Bua, S.; Polentarutti, M.; Supuran, C.T.; Monti, S.M.; et al. Zeta-Carbonic Anhydrases Show CS₂ Hydrolase Activity: A New Metabolic Carbon Acquisition Pathway in Diatoms? *Comput. Struct. Biotechnol. J.* **2021**, *19*, 3427–3436. [\[CrossRef\]](#)
43. Del Prete, S.; Vullo, D.; Fisher, G.M.; Andrews, K.T.; Poulsen, S.-A.; Capasso, C.; Supuran, C.T. Discovery of a New Family of Carbonic Anhydrases in the Malaria Pathogen *Plasmodium falciparum*—The η -Carbonic Anhydrases. *Bioorg. Med. Chem. Lett.* **2014**, *24*, 4389–4396. [\[CrossRef\]](#)
44. Nawaly, H.; Tanaka, A.; Toyoshima, Y.; Tsuji, Y.; Matsuda, Y. Localization and Characterization Theta Carbonic Anhydrases in *Thalassiosira pseudonana*. *Photosynth. Res.* **2023**, *156*, 217–229. [\[CrossRef\]](#)
45. Nocentini, A.; Supuran, C.T.; Capasso, C. An Overview on the Recently Discovered Iota-Carbonic Anhydrases. *J. Enzym. Inhib. Med. Chem.* **2021**, *36*, 1988–1995. [\[CrossRef\]](#)
46. Del Prete, S.; Nocentini, A.; Supuran, C.T.; Capasso, C. Bacterial Iota-Carbonic Anhydrase: A New Active Class of Carbonic Anhydrase Identified in the Genome of the Gram-Negative Bacterium *Burkholderia Territorii*. *J. Enzym. Inhib. Med. Chem.* **2020**, *35*, 1060–1068. [\[CrossRef\]](#)
47. Kikutani, S.; Nakajima, K.; Nagasato, C.; Tsuji, Y.; Miyatake, A.; Matsuda, Y. Thylakoid Luminal Theta-Carbonic Anhydrase Critical for Growth and Photosynthesis in the Marine Diatom *Phaeodactylum Tricornutum*. *Proc. Natl. Acad. Sci. USA* **2016**, *113*, 9828–9833. [\[CrossRef\]](#)
48. Alterio, V.; Langella, E.; De Simone, G.; Monti, S.M. Cadmium-Containing Carbonic Anhydrase CDCA1 in Marine Diatom *Thalassiosira weissflogii*. *Mar. Drugs* **2015**, *13*, 1688–1697. [\[CrossRef\]](#)
49. Angeli, A.; Buonanno, M.; Donald, W.A.; Monti, S.M.; Supuran, C.T. The Zinc—but Not Cadmium—Containing ζ -Carbonic from the Diatom *Thalassiosira weissflogii* Is Potently Activated by Amines and Amino Acids. *Bioorg. Chem.* **2018**, *80*, 261–265. [\[CrossRef\]](#)
50. Nathan, V.K.; Ammini, P. Carbon Dioxide Sequestering Ability of Bacterial Carbonic Anhydrase in a Mangrove Soil Microcosm and Its Bio-Mineralization Properties. *Water Air Soil Pollut.* **2019**, *230*, 192. [\[CrossRef\]](#)
51. Banci, L.; Bertini, I.; Luchinat, C.; Donaire, A.; Martinez, M.-J. The Factors Governing the Coordination Number in the Anion Derivatives of Carbonic Anhydrase. *Comments Inorg. Chem.* **1990**, *9*, 245–261. [\[CrossRef\]](#)
52. Hirakawa, Y.; Senda, M.; Fukuda, K.; Yu, H.Y.; Ishida, M.; Taira, M.; Kinbara, K.; Senda, T. Characterization of a Novel Type of Carbonic Anhydrase That Acts without Metal Cofactors. *BMC Biol.* **2021**, *19*, 105. [\[CrossRef\]](#) [\[PubMed\]](#)
53. Jo, B.H.; Kim, I.G.; Seo, J.H.; Kang, D.G.; Cha, H.J. Engineered *Escherichia coli* with Periplasmic Carbonic Anhydrase as a Biocatalyst for CO₂ Sequestration. *Appl. Environ. Microbiol.* **2013**, *79*, 6697–6705. [\[CrossRef\]](#)

54. Ekinici, D.; Beydemir, S.; Alim, Z. Some Drugs Inhibit in vitro Hydratase and Esterase Activities of Human Carbonic Anhydrase-I and II. *Pharmacol. Rep.* **2007**, *59*, 580–587. [\[PubMed\]](#)
55. Steger, F.; Reich, J.; Fuchs, W.; Rittmann, S.K.-M.R.; Gübitz, G.M.; Ribitsch, D.; Bochmann, G. Comparison of Carbonic Anhydrases for CO₂ Sequestration. *Int. J. Mol. Sci.* **2022**, *23*, 957. [\[CrossRef\]](#)
56. Faridi, S.; Satyanarayana, T. Novel Alkalistable α -Carbonic Anhydrase from the Polyextremophilic Bacterium *Bacillus Halodurans*: Characteristics and Applicability in Flue Gas CO₂ Sequestration. *Environ. Sci. Pollut. Res.* **2016**, *23*, 15236–15249. [\[CrossRef\]](#)
57. Capasso, C.; De Luca, V.; Carginale, V.; Cannio, R.; Rossi, M. Biochemical Properties of a Novel and Highly Thermostable Bacterial Alpha-Carbonic Anhydrase from *Sulfurihydrogenibium yellowstonense* YO3AOP1. *J. Enzym. Inhib. Med. Chem.* **2012**, *27*, 892–897. [\[CrossRef\]](#)
58. Ramanan, R.; Kannan, K.; Vinayagamoorthy, N.; Ramkumar, K.M.; Sivanesan, S.D.; Chakrabarti, T. Purification and Characterization of a Novel Plant-Type Carbonic Anhydrase from *Bacillus subtilis*. *Biotechnol. Bioprocess Eng.* **2009**, *14*, 32–37. [\[CrossRef\]](#)
59. Kim, S.S.; Kim, N.J.; Hong, S.; Kim, S.S.; Sung, J.; Jin, M.S. The Structural Basis of the Low Catalytic Activities of the Two Minor Beta-Carbonic Anhydrases of the Filamentous Fungus *Aspergillus fumigatus*. *J. Struct. Biol.* **2019**, *208*, 61–68. [\[CrossRef\]](#)
60. Sridharan, U.; Ragunathan, P.; Kuramitsu, S.; Yokoyama, S.; Kumarevel, T.; Ponnuraj, K. Structural and Functional Characterization of a Putative Carbonic Anhydrase from *Geobacillus kaustophilus* Reveals Its Cambialistic Function. *Biochem. Biophys. Res. Commun.* **2021**, *547*, 96–101. [\[CrossRef\]](#)
61. Wang, W.; Zhang, Y.; Wang, L.; Jing, Q.; Wang, X.; Xi, X.; Zhao, X.; Wang, H. Molecular Structure of Thermostable and Zinc-Ion-Binding Gamma-Class Carbonic Anhydrases. *Biometals* **2019**, *32*, 317–328. [\[CrossRef\]](#)
62. Alber, B.E.; Ferry, J.G. A Carbonic Anhydrase from the Archaeon *Methanosarcina thermophila*. *Proc. Natl. Acad. Sci. USA* **1994**, *91*, 6909–6913. [\[CrossRef\]](#)
63. Del Prete, S.; Vullo, D.; Di Fonzo, P.; Osman, S.M.; AlOthman, Z.; Donald, W.A.; Supuran, C.T.; Capasso, C. Sulfonamide Inhibition Profile of the Gamma-Carbonic Anhydrase Identified in the Genome of the Pathogenic Bacterium *Burkholderia pseudomallei* the Etiological Agent Responsible of Melioidosis. *Bioorg. Med. Chem. Lett.* **2017**, *27*, 490–495. [\[CrossRef\]](#)
64. Del Prete, S.; Vullo, D.; De Luca, V.; Supuran, C.T.; Capasso, C. Biochemical Characterization of the Delta-Carbonic Anhydrase from the Marine Diatom *Thalassiosira weissflogii*, TweCA. *J. Enzym. Inhib. Med. Chem.* **2014**, *29*, 906–911. [\[CrossRef\]](#)
65. Zhang, H.; Blanco-Ameijeiras, S.; Hopkinson, B.M.; Bernasconi, S.M.; Mejia, L.M.; Liu, C.; Stoll, H. An Isotope Label Method for Empirical Detection of Carbonic Anhydrase in the Calcification Pathway of the Coccolithophore *Emiliania huxleyi*. *Geochim. Cosmochim. Acta* **2021**, *292*, 78–93. [\[CrossRef\]](#)
66. Xu, Y.; Feng, L.; Jeffrey, P.D.; Shi, Y.; Morel, F.M.M. Structure and Metal Exchange in the Cadmium Carbonic Anhydrase of Marine Diatoms. *Nature* **2008**, *452*, 56–61. [\[CrossRef\]](#)
67. Supuran, C.T.; Capasso, C. The η -Class Carbonic Anhydrases as Drug Targets for Antimalarial Agents. *Expert Opin. Ther. Targets* **2015**, *19*, 551–563. [\[CrossRef\]](#)
68. Kitahara, M.; Fudo, S.; Yoneda, T.; Nukaga, M.; Hoshino, T. Anisotropic Distribution of Ammonium Sulfate Ions in Protein Crystallization. *Cryst. Growth Des.* **2019**, *19*, 6004–6010. [\[CrossRef\]](#)
69. Woods, L.A.; Dolezal, O.; Ren, B.; Ryan, J.H.; Peat, T.S.; Poulsen, S.-A. Native State Mass Spectrometry, Surface Plasmon Resonance, and X-Ray Crystallography Correlate Strongly as a Fragment Screening Combination. *J. Med. Chem.* **2016**, *59*, 2192–2204. [\[CrossRef\]](#)
70. Avvaru, B.S.; Busby, S.A.; Chalmers, M.J.; Griffin, P.R.; Venkatakrishnan, B.; Agbandje-McKenna, M.; Silverman, D.N.; McKenna, R. Apo-Human Carbonic Anhydrase II Revisited: Implications of the Loss of a Metal in Protein Structure, Stability, and Solvent Network. *Biochemistry* **2009**, *48*, 7365–7372. [\[CrossRef\]](#) [\[PubMed\]](#)
71. Berghem, N.C.; Hallberg, M.; Wisén, S. Molecular Characterization of the Human Carbonic Anhydrase-Related Protein (HCA-RP VIII). *Biochim. Biophys. Acta* **1998**, *1384*, 294–298. [\[CrossRef\]](#) [\[PubMed\]](#)
72. Picaud, S.S.; Muniz, J.R.; Kramm, A.; Pilka, E.S.; Kochan, G.; Oppermann, U.; Yue, W.W. Crystal Structure of Human Carbonic Anhydrase-Related Protein VIII Reveals the Basis for Catalytic Silencing. *Proteins* **2009**, *76*, 507–511. [\[CrossRef\]](#) [\[PubMed\]](#)
73. Lakkis, M.M.; Berghem, N.C.; O'Shea, K.S.; Tashian, R.E. Expression of the Acatalytic Carbonic Anhydrase VIII Gene, Car8, during Mouse Embryonic Development. *Histochem. J.* **1997**, *29*, 135–141. [\[CrossRef\]](#) [\[PubMed\]](#)
74. Hirota, J.; Ando, H.; Hamada, K.; Mikoshiba, K. Carbonic Anhydrase-Related Protein is a Novel Binding Protein for Inositol 1,4,5-Trisphosphate Receptor Ttype 1. *Biochem. J.* **2003**, *372*, 435–441. [\[CrossRef\]](#)
75. Borén, K.; Andersson, P.; Larsson, M.; Carlsson, U. Characterization of a Molten Globule State of Bovine Carbonic Anhydrase III: Loss of Asymmetrical Environment of the Aromatic Residues Has a Profound Effect on Both the Near- and Far-UV CD Spectrum. *Biochim. Biophys. Acta* **1999**, *1430*, 111–118. [\[CrossRef\]](#)
76. Capasso, C.; Supuran, C.T. Inhibition of Bacterial Carbonic Anhydrases as a Novel Approach to Escape Drug Resistance. *Curr. Top. Med. Chem.* **2017**, *17*, 1237–1248. [\[CrossRef\]](#)
77. Wilbur, K.M.; Anderson, N.G. Electrometric and Colorimetric Determination of Carbonic Anhydrase. *J. Biol. Chem.* **1948**, *176*, 147–154. [\[CrossRef\]](#)
78. Fuchs, W.; Steger, F.; Reich, J.; Ribitsch, D.; Rittmann, S.K.-M.R.; Bochmann, G. A Simple and Straightforward Method for Activity Measurement of Carbonic Anhydrases. *Catalysis* **2021**, *11*, 819. [\[CrossRef\]](#)
79. Verpoorte, J.A.; Mehta, S.; Edsall, J.T. Esterase Activities of Human Carbonic Anhydrases B and C. *J. Biol. Chem.* **1967**, *242*, 4221–4229. [\[CrossRef\]](#)

80. Effendi, S.S.W.; Ng, I.-S. The Prospective and Potential of Carbonic Anhydrase for Carbon Dioxide Sequestration: A Critical Review. *Process Biochem.* **2019**, *87*, 55–65. [\[CrossRef\]](#)
81. Supuran, C.T.; De Simone, G. Carbonic Anhydrases: An Overview. In *Carbonic Anhydrases as Biocatalysts: From Theory to Medical and Industrial Applications*; Elsevier: Florence, Italy, 2015; pp. 3–13. [\[CrossRef\]](#)
82. Niks, D.; Hille, R. Reductive Activation of CO₂ by Formate Dehydrogenases. In *Enzymes of Energy Technology; Methods in Enzymology*; Armstrong, F., Ed.; Elsevier Acad. Press. Inc.: San Diego, CA, USA, 2018; Volume 613, pp. 277–295. [\[CrossRef\]](#)
83. Nielsen, C.F.; Lange, L.; Meyer, A.S. Classification and Enzyme Kinetics of Formate Dehydrogenases for Biomanufacturing via CO₂ Utilization. *Biotechnol. Adv.* **2019**, *37*, 107408. [\[CrossRef\]](#) [\[PubMed\]](#)
84. Maia, L.B.; Moura, I.; Moura, J.J.G. Molybdenum and Tungsten-Containing Enzymes: An Overview. In *Molibdenum and Tungsten Enzymes: Biochemistry*; Hille, R., Schulzke, C., Kirk, M.L., Eds.; RSC Metallobiology Series; Royal Society of Chemistry: Cambridge, UK, 2017; Volume 5, pp. 1–80.
85. Lemaire, O.N.; Jespersen, M.; Wagner, T. CO₂-Fixation Strategies in Energy Extremophiles: What Can We Learn From Acetogens? *Front. Microbiol.* **2020**, *11*, 486. [\[CrossRef\]](#) [\[PubMed\]](#)
86. Moon, M.; Park, G.W.; Lee, J.; Lee, J.-S.; Min, K. Recent Progress in Formate Dehydrogenase (FDH) as a Non-Photosynthetic CO₂ Utilizing Enzyme: A Short Review. *J. CO₂ Util.* **2020**, *42*, 101353. [\[CrossRef\]](#)
87. Alpdağtaş, S.; Turunen, O.; Valjakka, J.; Binay, B. The Challenges of Using NAD⁺-Dependent Formate Dehydrogenases for CO₂ Conversion. *Crit. Rev. Biotechnol.* **2022**, *42*, 953–972. [\[CrossRef\]](#)
88. Sakai, Y.; Murdanoto, A.P.; Konishi, T.; Iwamatsu, A.; Kato, N. Regulation of the Formate Dehydrogenase Gene, FDH1, in the Methylophilic Yeast *Candida boidinii* and Growth Characteristics of an FDH1-Disrupted Strain on Methanol, Methylamine, and Choline. *J. Bacteriol.* **1997**, *179*, 4480–4485. [\[CrossRef\]](#)
89. Tishkov, V.I.; Popov, V.O. Protein Engineering of Formate Dehydrogenase. *Biomol. Eng.* **2006**, *23*, 89–110. [\[CrossRef\]](#)
90. Pagano, P.; Guo, Q.; Ranasinghe, C.; Schroeder, E.; Robben, K.; Hase, F.; Ye, H.; Wickersham, K.; Aspuru-Guzik, A.; Major, D.T.; et al. Oscillatory Active-Site Motions Correlate with Kinetic Isotope Effects in Formate Dehydrogenase. *ACS Catal.* **2019**, *9*, 11199–11206. [\[CrossRef\]](#)
91. Yilmazer, B.; Isupov, M.N.; De Rose, S.A.; Bulut, H.; Benninghoff, J.C.; Binay, B.; Littlechild, J.A. Structural insights into the NAD⁺-dependent Formate Dehydrogenase Mechanism Revealed from the NADH Complex and the Formate NAD⁺-Ternary Complex of the *Chaetomium thermophilum* Enzyme. *J. Struct. Biol.* **2020**, *212*, 107657. [\[CrossRef\]](#)
92. Pala, U.; Yilmazer, B.; Corbacioglu, M.; Ruuponen, J.; Valjakka, J.; Turunen, O.; Binay, B. Functional Effects of Active Site Mutations in NAD⁺-Dependent Formate Dehydrogenases on Transformation of Hydrogen Carbonate to Formate. *Protein Eng. Des. Sel.* **2018**, *31*, 327–335. [\[CrossRef\]](#)
93. Cakar, M.M.; Ruuponen, J.; Mangas-Sanchez, J.; Birmingham, W.R.; Yildirim, D.; Turunen, O.; Turner, N.J.; Valjakka, J.; Binay, B. Engineered Formate Dehydrogenase from *Chaetomium Thermophilum*, a Promising Enzymatic Solution for Biotechnical CO₂ Fixation. *Biotechnol. Lett.* **2020**, *42*, 2251–2262. [\[CrossRef\]](#)
94. Guo, Q.; Gakhar, L.; Wickersham, K.; Francis, K.; Vardi-Kilshtain, A.; Major, D.T.; Cheatum, C.M.; Kohen, A. Structural and Kinetic Studies of Formate Dehydrogenase from *Candida boidinii*. *Biochemistry* **2016**, *55*, 2760–2771. [\[CrossRef\]](#)
95. Sato, R.; Amao, Y. Studies on the Catalytic Mechanism of Formate Dehydrogenase from *Candida boidinii* Using Isotope-Labelled Substrate and Co-Enzyme. *Catal. Today* **2023**, *411–412*, 113796. [\[CrossRef\]](#)
96. Jiang, W.; Lin, P.; Yang, R.; Fang, B. Identification of Catalysis, Substrate, and Coenzyme Binding Sites and Improvement Catalytic Efficiency of Formate Dehydrogenase from *Candida boidinii*. *Appl. Microbiol. Biotechnol.* **2016**, *100*, 8425–8437. [\[CrossRef\]](#)
97. Altas, N.; Aslan, A.S.; Karatas, E.; Chronopoulou, E.; Labrou, N.E.; Binay, B. Heterologous Production of Extreme Alkaline Thermostable NAD⁺-Dependent Formate Dehydrogenase with Wide-Range pH Activity from *Myceliophthora thermophila*. *Process Biochem.* **2017**, *61*, 110–118. [\[CrossRef\]](#)
98. Choe, H.; Joo, J.C.; Cho, D.H.; Kim, M.H.; Lee, S.H.; Jung, K.D.; Kim, Y.H. Efficient CO₂-Reducing Activity of NAD-Dependent Formate Dehydrogenase from *Thiobacillus* sp. KNK65MA for Formate Production from CO₂ Gas. *PLoS ONE* **2014**, *9*, e103111. [\[CrossRef\]](#)
99. Aslan, A.S.; Valjakka, J.; Ruuponen, J.; Yildirim, D.; Turner, N.J.; Turunen, O.; Binay, B. *Chaetomium thermophilum* Formate Dehydrogenase Has High Activity in the Reduction of Hydrogen Carbonate (HCO₃[−]) to Formate. *Protein Eng. Des. Sel.* **2017**, *30*, 47–55. [\[CrossRef\]](#)
100. de Bok, F.A.M.; Hagedoorn, P.L.; Silva, P.J.; Hagen, W.R.; Schiltz, E.; Fritsche, K.; Stams, A.J.M. Two W-Containing Formate Dehydrogenases (CO₂-Reductases) Involved in Syntrophic Propionate Oxidation by *Syntrophobacter fumaroxidans*. *Eur. J. Biochem.* **2003**, *270*, 2476–2485. [\[CrossRef\]](#)
101. Maia, L.B.; Fonseca, L.; Moura, I.; Moura, J.J.G. Reduction of Carbon Dioxide by a Molybdenum-Containing Formate Dehydrogenase: A Kinetic and Mechanistic Study. *J. Am. Chem. Soc.* **2016**, *138*, 8834–8846. [\[CrossRef\]](#)
102. Bassegoda, A.; Madden, C.; Wakerley, D.W.; Reisner, E.; Hirst, J. Reversible Interconversion of CO₂ and Formate by a Molybdenum-Containing Formate Dehydrogenase. *J. Am. Chem. Soc.* **2014**, *136*, 15473–15476. [\[CrossRef\]](#)
103. Axley, M.J.; Grahame, D.A.; Stadtman, T.C. *Escherichia coli* Formate-Hydrogen Lyase. Purification and Properties of the Selenium-Dependent Formate Dehydrogenase Component. *J. Biol. Chem.* **1990**, *265*, 18213–18218. [\[CrossRef\]](#) [\[PubMed\]](#)

104. Oliveira, A.R.; Mota, C.; Mourato, C.; Domingos, R.M.; Santos, M.F.A.; Gesto, D.; Guigliarelli, B.; Santos-Silva, T.; Romao, M.J.; Cardoso Pereira, I.A. Toward the Mechanistic Understanding of Enzymatic CO₂ Reduction. *ACS Catal.* **2020**, *10*, 3844–3856. [\[CrossRef\]](#)
105. da Silva, S.M.; Pimentel, C.; Valente, F.M.A.; Rodrigues-Pousada, C.; Pereira, I.A.C. Tungsten and Molybdenum Regulation of Formate Dehydrogenase Expression in *Desulfovibrio Vulgaris* Hildenborough. *J. Bacteriol.* **2011**, *193*, 2909–2916. [\[CrossRef\]](#) [\[PubMed\]](#)
106. Schuchmann, K.; Mueller, V. Direct and Reversible Hydrogenation of CO₂ to Formate by a Bacterial Carbon Dioxide Reductase. *Science* **2013**, *342*, 1382–1385. [\[CrossRef\]](#)
107. Yu, X.; Niks, D.; Mulchandani, A.; Hille, R. Efficient Reduction of CO₂ by the Molybdenum-Containing Formate Dehydrogenase from *Cupriavidus necator* (*Ralstonia eutropha*). *J. Biol. Chem.* **2017**, *292*, 16872–16879. [\[CrossRef\]](#)
108. Radon, C.; Mittelstadt, G.; Duffus, B.R.; Burger, J.; Hartmann, T.; Mielke, T.; Teutloff, C.; Leimkuhler, S.; Wendler, P. Cryo-EM Structures Reveal Intricate Fe-S Cluster Arrangement and Charging in *Rhodobacter capsulatus* Formate Dehydrogenase. *Nat. Commun.* **2020**, *11*, 1912. [\[CrossRef\]](#)
109. Hartmann, T.; Leimkuhler, S. The Oxygen-Tolerant and NAD⁺-Dependent Formate Dehydrogenase from *Rhodobacter capsulatus* Is Able to Catalyze the Reduction of CO₂ to Formate. *FEBS J.* **2013**, *280*, 6083–6096. [\[CrossRef\]](#)
110. Ruschig, U.; Muller, U.; Willnow, P.; Hopner, T. CO₂ Recution to Formate by NADH Catalyzed by Formate Dehydrogenase from *Pseudomonas oxalaticus*. *Eur. J. Biochem.* **1976**, *70*, 325–330. [\[CrossRef\]](#)
111. Müller, U.; Willnow, P.; Ruschig, U.; Höpner, T. Formate Dehydrogenase from *Pseudomonas oxalaticus*. *Eur. J. Biochem.* **1978**, *83*, 485–498. [\[CrossRef\]](#)
112. Cakar, M.M.; Mangas-Sanchez, J.; Birmingham, W.R.; Turner, N.J.; Binay, B. Discovery of a New Metal and NAD⁺-Dependent Formate Dehydrogenase from *Clostridium ljungdahlii*. *Prep. Biochem. Biotechnol.* **2018**, *48*, 327–334. [\[CrossRef\]](#)
113. Min, K.; Moon, M.; Park, G.W.; Lee, J.-P.; Kim, S.J.; Lee, J.-S. Newly Explored Formate Dehydrogenases from *Clostridium* Species Catalyze Carbon Dioxide to Formate. *Bioresour. Technol.* **2022**, *348*, 126832. [\[CrossRef\]](#)
114. Almendra, M.J.; Brondino, C.D.; Gavel, O.; Pereira, A.S.; Tavares, P.; Bursakov, S.; Duarte, R.; Caldeira, J.; Moura, J.J.G.; Moura, I. Purification and Characterization of a Tungsten-Containing Formate Dehydrogenase from *Desulfovibrio gigas*. *Biochemistry* **1999**, *38*, 16366–16372. [\[CrossRef\]](#)
115. Yamamoto, I.; Saiki, T.; Liu, S.M.; Ljungdahl, L.G. Purification and Properties of NADP-Dependent Formate Dehydrogenase from *Clostridium thermoaceticum*, a Tungsten Selenium Iron Protein. *J. Biol. Chem.* **1983**, *258*, 1826–1832. [\[CrossRef\]](#)
116. Cordas, C.M.; Moura, J.J.G. Molybdenum and Tungsten Enzymes Redox Properties—A Brief Overview. *Coord. Chem. Rev.* **2019**, *394*, 53–64. [\[CrossRef\]](#)
117. Hartmann, T.; Schwanhold, N.; Leimkuhler, S. Assembly and Catalysis of Molybdenum or Tungsten-Containing Formate Dehydrogenases from Bacteria. *Biochim. Biophys. Acta Proteins Proteom.* **2015**, *1854*, 1090–1100. [\[CrossRef\]](#)
118. Kirk, M.L.; Hille, R. Spectroscopic Studies of Mononuclear Molybdenum Enzyme Centers. *Molecules* **2022**, *27*, 4802. [\[CrossRef\]](#)
119. Jormakka, M.; Tornroth, S.; Byrne, B.; Iwata, S. Molecular Basis of Proton Motive Force Generation: Structure of Formate Dehydrogenase-N. *Science* **2002**, *295*, 1863–1868. [\[CrossRef\]](#)
120. Niks, D.; Hille, R. Molybdenum- and Tungsten-Containing Formate Dehydrogenases and Formylmethanofuran Dehydrogenases: Structure, Mechanism, and Cofactor Insertion. *Protein Sci.* **2019**, *28*, 111–122. [\[CrossRef\]](#)
121. Robinson, W.E.; Bassegoda, A.; Reisner, E.; Hirst, J. Oxidation-State-Dependent Binding Properties of the Active Site in a Mo-Containing Formate Dehydrogenase. *J. Am. Chem. Soc.* **2017**, *139*, 9927–9936. [\[CrossRef\]](#)
122. Meneghello, M.; Oliveira, A.R.; Jacq-Bailly, A.; Pereira, I.A.C.; Leger, C.; Fourmond, V. Formate Dehydrogenases Reduce CO₂ Rather than HCO₃[−]: An Electrochemical Demonstration. *Angew. Chem. Int. Ed.* **2021**, *60*, 9964–9967. [\[CrossRef\]](#)
123. Tiberti, M.; Papaleo, E.; Russo, N.; De Gioia, L.; Zampella, G. Evidence for the Formation of a Mo-H Intermediate in the Catalytic Cycle of Formate Dehydrogenase. *Inorg. Chem.* **2012**, *51*, 8331–8339. [\[CrossRef\]](#)
124. Khan, M.R. Immobilized Enzymes: A Comprehensive Review. *Bull. Natl. Res. Cent.* **2021**, *45*, 207. [\[CrossRef\]](#)
125. Bie, J.; Sepodes, B.; Fernandes, P.C.B.; Ribeiro, M.H.L. Enzyme Immobilization and Co-Immobilization: Main Framework, Advances and Some Applications. *Processes* **2022**, *10*, 494. [\[CrossRef\]](#)
126. Porcar, R.; Lavandera, I.; Lozano, P.; Altava, B.; Luis, S.V.; Gotor-Fernandez, V.; Garcia-Verdugo, E. Supported Ionic Liquid-like Phases as Efficient Solid Ionic Solvents for the Immobilisation of Alcohol Dehydrogenases towards the Development of Stereoselective Bioreductions. *Green Chem.* **2021**, *23*, 5609–5617. [\[CrossRef\]](#)
127. Sheldon, R.A.; Basso, A.; Brady, D. New Frontiers in Enzyme Immobilisation: Robust Biocatalysts for a Circular Bio-Based Economy. *Chem. Soc. Rev.* **2021**, *50*, 5850–5862. [\[CrossRef\]](#) [\[PubMed\]](#)
128. Lozano, P.; Garcia-Verdugo, E.; Bernal, J.M.; Izquierdo, D.F.; Isabel Burguete, M.; Sanchez-Gomez, G.; Luis, S.V. Immobilised Lipase on Structured Supports Containing Covalently Attached Ionic Liquids for the Continuous Synthesis of Biodiesel in ScCO₂. *ChemSusChem* **2012**, *5*, 790–798. [\[CrossRef\]](#)
129. Ren, S.; Jiang, S.; Yan, X.; Chen, R.; Cui, H. Challenges and Opportunities: Porous Supports in Carbonic Anhydrase Immobilization. *J. CO₂ Util.* **2020**, *42*, 101305. [\[CrossRef\]](#)
130. Molina-Fernandez, C.; Luis, P. Immobilization of Carbonic Anhydrase for CO₂ Capture and Its Industrial Implementation: A Review. *J. CO₂ Util.* **2021**, *47*, 101475. [\[CrossRef\]](#)

131. Rasouli, H.; Nguyen, K.; Iliuta, M.C. Recent Advancements in Carbonic Anhydrase Immobilization and Its Implementation in CO₂ Capture Technologies: A Review. *Sep. Purif. Technol.* **2022**, *296*, 121299. [\[CrossRef\]](#)
132. Yuan, Y.; Wang, F.; Li, H.; Su, S.; Gao, H.; Han, X.; Ren, S. Potential Application of the Immobilization of Carbonic Anhydrase Based on Metal Organic Framework Supports. *Process Biochem.* **2022**, *122*, 214–223. [\[CrossRef\]](#)
133. Molina-Fernandez, C.; Peters, A.; Debecker, D.P.; Luis, P. Immobilization of Carbonic Anhydrase in a Hydrophobic Poly(Ionic Liquid): A New Functional Solid for CO₂ Capture. *Biochem. Eng. J.* **2022**, *187*, 108639. [\[CrossRef\]](#)
134. Rouf, S.; Greish, Y.E.; Al-Zuhair, S. Immobilization of Formate Dehydrogenase in Metal Organic Frameworks for Enhanced Conversion of Carbon Dioxide to Formate. *Chemosphere* **2021**, *267*, 128921. [\[CrossRef\]](#)
135. Singh, R.K.R.; Tiwari, M.K.; Singh, R.K.R.; Lee, J.-K. From Protein Engineering to Immobilization: Promising Strategies for the Upgrade of Industrial Enzymes. *Molecules* **2013**, *18*, 1232–1277. [\[CrossRef\]](#)
136. Di Spiridione, C.; Aresta, M.; Dibenedetto, A. Improving the Enzymatic Cascade of Reactions for the Reduction of CO₂ to CH₃OH in Water: From Enzymes Immobilization Strategies to Cofactor Regeneration and Cofactor Suppression. *Molecules* **2022**, *27*, 4913. [\[CrossRef\]](#)
137. Zhu, X.; Du, C.; Gao, B.; He, B. Strategies to Improve the Mass Transfer in the CO₂ Capture Process Using Immobilized Carbonic Anhydrase. *J. Environ. Manag.* **2023**, *332*, 117370. [\[CrossRef\]](#)
138. Ren, S.; Chen, R.; Wu, Z.; Su, S.; Hou, J.; Yuan, Y. Enzymatic Characteristics of Immobilized Carbonic Anhydrase and Its Applications in CO₂ Conversion. *Colloid. Surface B* **2021**, *204*, 111779. [\[CrossRef\]](#)
139. Shen, J.; Salmon, S. Biocatalytic Membranes for Carbon Capture and Utilization. *Membranes* **2023**, *13*, 367. [\[CrossRef\]](#) [\[PubMed\]](#)
140. Wanjari, S.; Prabhu, C.; Satyanarayana, T.; Vinu, A.; Rayalu, S. Immobilization of Carbonic Anhydrase on Mesoporous Aluminosilicate for Carbonation Reaction. *Micropor. Mesopor. Mat.* **2012**, *160*, 151–158. [\[CrossRef\]](#)
141. Yu, Y.; Chen, B.; Qi, W.; Li, X.; Shin, Y.; Lei, C.; Liu, J. Enzymatic Conversion of CO₂ to Bicarbonate in Functionalized Mesoporous Silica. *Micropor. Mesopor. Mat.* **2012**, *153*, 166–170. [\[CrossRef\]](#)
142. Vinoba, M.; Bhagiyalakshmi, M.; Jeong, S.K.; Yoon, Y.I.; Nam, S.C. Carbonic Anhydrase Conjugated to Nanosilver Immobilized onto Mesoporous SBA-15 for Sequestration of CO₂. *J. Mol. Catal. B Enzym.* **2012**, *75*, 60–67. [\[CrossRef\]](#)
143. Mao, M.; Zhai, T.; Meng, L.; Meng, Z.; Liu, W. Controllable Preparation of Mesoporous Silica and Its Application in Enzyme-Catalyzed CO₂ Reduction. *Chem. Eng. J.* **2022**, *437*, 135479. [\[CrossRef\]](#)
144. Azari, F.N.-G.M. Reversible Denaturation of Carbonic Anhydrase Provides a Method for Its Adsorptive Immobilization. *Biotechnol. Bioeng.* **1999**, *62*, 193–199. [\[CrossRef\]](#)
145. Kim, J.K.; Abdelhamid, M.A.A.; Pack, S.P. Direct Immobilization and Recovery of Recombinant Proteins from Cell Lysates by Using EctP1-Peptide as a Short Fusion Tag for Silica and Titania Supports. *Int. J. Biol. Macromol.* **2019**, *135*, 969–977. [\[CrossRef\]](#)
146. Yong, J.K.J.; Stevens, G.W.; Caruso, F.; Kentish, S.E. In Situ Layer-by-Layer Assembled Carbonic Anhydrase-Coated Hollow Fiber Membrane Contactor for Rapid CO₂ Absorption. *J. Memb. Sci.* **2016**, *514*, 556–565. [\[CrossRef\]](#)
147. Ivanovski, V.; Shapovalova, O.E.; Drozdov, A.S. Structural Rearrangements of Carbonic Anhydrase Entrapped in Sol-Gel Magnetite Determined by ATR-FTIR Spectroscopy. *Int. J. Mol. Sci.* **2022**, *23*, 5975. [\[CrossRef\]](#)
148. Moon, H.; Kim, S.; Jo, B.H.; Cha, H.J. Immobilization of Genetically Engineered Whole-Cell Biocatalysts with Periplasmic Carbonic Anhydrase in Polyurethane Foam for Enzymatic CO₂ Capture and Utilization. *J. CO₂ Util.* **2020**, *39*, 101172. [\[CrossRef\]](#)
149. Jiao, M.; He, J.; Sun, S.; Vriesekoop, F.; Yuan, Q.; Liu, Y.; Liang, H. Fast Immobilization of Human Carbonic Anhydrase II on Ni-Based Metal-Organic Framework Nanorods with High Catalytic Performance. *Catalysts* **2020**, *10*, 401. [\[CrossRef\]](#)
150. Abdelrahim, M.Y.; Martins, C.F.; Neves, L.A.; Capasso, C.; Supuran, C.T.; Coelho, I.M.; Crespo, J.G.; Barboiu, M. Supported Ionic Liquid Membranes Immobilized with Carbonic Anhydrases for CO₂ Transport at High Temperatures. *J. Memb. Sci.* **2017**, *528*, 225–230. [\[CrossRef\]](#)
151. Martins, C.F.; Neves, L.A.M.; Estevão, L.A.; Rosatella, A.; Alves, V.D.; Afonso, C.A.M.; Crespo, J.G.; Coelho, I.M. Effect of Water Activity on Carbon Dioxide Transport in Cholinium-Based Ionic Liquids with Carbonic Anhydrase. *Sep. Purif. Technol.* **2016**, *168*, 74–82. [\[CrossRef\]](#)
152. Vinoba, M.; Bhagiyalakshmi, M.; Jeong, S.K.; Yoon, Y.I.; Nam, S.C. Capture and Sequestration of CO₂ by Human Carbonic Anhydrase Covalently Immobilized onto Amine-Functionalized SBA-15. *J. Phys. Chem. C* **2011**, *115*, 20209–20216. [\[CrossRef\]](#)
153. Shen, J.; Yuan, Y.; Salmon, S. Durable and Versatile Immobilized Carbonic Anhydrase on Textile Structured Packing for CO₂ Capture. *Catalysts* **2022**, *12*, 1108. [\[CrossRef\]](#)
154. Xv, J.; Zhang, Z.; Pang, S.; Jia, J.; Geng, Z.; Wang, R.; Li, P.; Bilal, M.; Cui, J.; Jia, S. Accelerated CO₂ Capture Using Immobilized Carbonic Anhydrase on Polyethyleneimine/Dopamine Co-Deposited MOFs. *Biochem. Eng. J.* **2022**, *189*, 108719. [\[CrossRef\]](#)
155. Shamna, I.; Jeong, S.K.; Margandan, B. Covalent Immobilization of Carbonic Anhydrase on Amine Functionalized Alumino-Siloxane Aerogel Beads for Biomimetic Sequestration of CO₂. *J. Ind. Eng. Chem.* **2021**, *100*, 288–295. [\[CrossRef\]](#)
156. Iliuta, I.; Rasouli, H.; Iliuta, M.C. Intensified CO₂ Capture in Wall-Coated Microreactors with Immobilized Carbonic Anhydrase: Experimental and Modeling. *Sep. Purif. Technol.* **2023**, *307*, 122590. [\[CrossRef\]](#)
157. Kimmel, J.D.; Arazawa, D.T.; Ye, S.-H.; Shankaraman, V.; Wagner, W.R.; Federspiel, W.J. Carbonic Anhydrase Immobilized on Hollow Fiber Membranes Using Glutaraldehyde Activated Chitosan for Artificial Lung Applications. *J. Mater. Sci. Mater. Med.* **2013**, *24*, 2611–2621. [\[CrossRef\]](#)

158. Peirce, S.; Russo, M.E.; Istatico, R.; Lafuente, R.F.; Salatino, P.; Marzocchella, A. Structure and Activity of Magnetic Cross-Linked Enzyme Aggregates of Bovine Carbonic Anhydrase as Promoters of Enzymatic CO₂ Capture. *Biochem. Eng. J.* **2017**, *127*, 188–195. [\[CrossRef\]](#)
159. Chang, S.; He, Y.; Li, Y.; Cui, X. Study on the Immobilization of Carbonic Anhydrases on Geopolymer Microspheres for CO₂ Capture. *J. Clean. Prod.* **2021**, *316*, 128163. [\[CrossRef\]](#)
160. Zhang, X.; Shao, W.; Chen, B.; Wang, M. Cross-Linking of Carbonic Anhydrase and Formate Dehydrogenase Based on Amino Acid Specific Recognition: Conversion of Carbon Dioxide to Formic Acid. *Enzym. Microb. Technol.* **2021**, *146*, 109763. [\[CrossRef\]](#)
161. Nelson, J.M.; Griffin, E.G. Adsorption of Invertase. *J. Am. Chem. Soc.* **1916**, *38*, 1109–1115. [\[CrossRef\]](#)
162. Jesionowski, T.; Zdzarta, J.; Krajewska, B. Enzyme immobilization by adsorption: A review. *Adsorption* **2014**, *20*, 801–821. [\[CrossRef\]](#)
163. Nabavi Zadeh, P.S.; Åkerman, B. Immobilization of Enzymes in Mesoporous Silica Particles: Protein Concentration and Rotational Mobility in the Pores. *J. Phys. Chem. B* **2017**, *121*, 2575–2583. [\[CrossRef\]](#) [\[PubMed\]](#)
164. Crumbliss, A.L.; Perine, S.C.; Stonehuerner, J.; Tubergen, K.R.; Zhao, J.; Henkens, R.W.; O'Daly, J.P. Colloidal Gold as a Biocompatible Immobilization Matrix Suitable for the Fabrication of Enzyme Electrodes by Electrodeposition. *Biotechnol. Bioeng.* **1992**, *40*, 483–490. [\[CrossRef\]](#)
165. Suvannasara, P.; Juntapram, K.; Praphairaksit, N.; Siralermukul, K.; Muangsin, N. Mucoadhesive 4-Carboxybenzenesulfonamide-Chitosan with Antibacterial Properties. *Carbohydr. Polym.* **2013**, *94*, 244–252. [\[CrossRef\]](#) [\[PubMed\]](#)
166. Li, J.-J.; Yin, L.; Wang, Z.-F.; Jing, Y.-C.; Jiang, Z.-L.; Ding, Y.; Wang, H.-S. Enzyme-Immobilized Metal-Organic Frameworks: From Preparation to Application. *Chem. Asian J.* **2022**, *17*, e202200751. [\[CrossRef\]](#)
167. Zhou, H.-C.; Long, J.R.; Yaghi, O.M. Introduction to Metal–Organic Frameworks. *Chem. Rev.* **2012**, *112*, 673–674. [\[CrossRef\]](#)
168. Zhang, W.; Taheri-Ledari, R.; Saeidirad, M.; Qazi, F.S.; Kashtiaray, A.; Ganjali, F.; Tian, Y.; Maleki, A. Regulation of Porosity in MOFs: A Review on Tunable Scaffolds and Related Effects and Advances in Different Applications. *J. Environ. Chem. Eng.* **2022**, *10*, 108836. [\[CrossRef\]](#)
169. Silva, A.R.M.; Alexandre, J.Y.N.H.; Souza, J.E.S.; Lima Neto, J.G.; de Sousa Junior, P.G.; Rocha, M.V.P.; dos Santos, J.C.S. The Chemistry and Applications of Metal–Organic Frameworks (MOFs) as Industrial Enzyme Immobilization Systems. *Molecules* **2022**, *27*, 4529. [\[CrossRef\]](#)
170. Liu, Q.; Bai, X.; Pham, H.; Hu, J.; Dinu, C.Z. Active Nanointerfaces Based on Enzyme Carbonic Anhydrase and Metal–Organic Framework for Carbon Dioxide Reduction. *Nanomaterials* **2021**, *11*, 1008. [\[CrossRef\]](#)
171. Huang, Y.; Zhang, S.; Chen, H.; Zhao, L.; Zhang, Z.; Cheng, P.; Chen, Y. A Zinc Coordination Complex Mimicking Carbonic Anhydrase for CO₂ Hydrolysis and Sequestration. *Inorg. Chem.* **2019**, *58*, 9916–9921. [\[CrossRef\]](#)
172. Bien, C.E.; Chen, K.K.; Chien, S.-C.; Reiner, B.R.; Lin, L.-C.; Wade, C.R.; Ho, W.S.W. Bioinspired Metal–Organic Framework for Trace CO₂ Capture. *J. Am. Chem. Soc.* **2018**, *140*, 12662–12666. [\[CrossRef\]](#)
173. Han, L.; Pham, T.; Zhuo, M.; Forrest, K.A.; Suepaul, S.; Space, B.; Zaworotko, M.J.; Shi, W.; Chen, Y.; Cheng, P.; et al. Molecular Sieving and Direct Visualization of CO₂ in Binding Pockets of an Ultramicroporous Lanthanide Metal–Organic Framework Platform. *ACS Appl. Mater. Interfaces* **2019**, *11*, 23192–23197. [\[CrossRef\]](#) [\[PubMed\]](#)
174. Wu, Z.; Nan, Y.; Zhao, Y.; Wang, X.; Huang, S.; Shi, J. Immobilization of Carbonic Anhydrase for Facilitated CO₂ Capture and Separation. *Chin. J. Chem. Eng.* **2020**, *28*, 2817–2831. [\[CrossRef\]](#)
175. Neves, L.A.; Afonso, C.; Coelho, I.M.; Crespo, J.G. Integrated CO₂ Capture and Enzymatic Bioconversion in Supported Ionic Liquid Membranes. *Sep. Purif. Technol.* **2012**, *97*, 34–41. [\[CrossRef\]](#)
176. Bednár, A.; Nemestóthy, N.; Bakonyi, P.; Fülöp, L.; Zhen, G.; Lu, X.; Kobayashi, T.; Kumar, G.; Xu, K.; Bélafi-Bakó, K. Enzymatically-Boosted Ionic Liquid Gas Separation Membranes Using Carbonic Anhydrase of Biomass Origin. *Chem. Eng. J.* **2016**, *303*, 621–626. [\[CrossRef\]](#)
177. de Castro, A.M.; Neves, L.A.; Corvo, M.C.; Cabrita, E.J.; Crespo, J.G. Effect of Carbonic Anhydrase on CO₂ Absorption Promoted by Choline Hydroxide Using Supported Liquid Membranes. *Sep. Purif. Technol.* **2022**, *280*, 119921. [\[CrossRef\]](#)
178. Migliardini, F.; De Luca, V.; Carginale, V.; Rossi, M.; Corbo, P.; Supuran, C.T.; Capasso, C. Biomimetic CO₂ Capture Using a Highly Thermostable Bacterial α -Carbonic Anhydrase Immobilized on a Polyurethane Foam. *J. Enzym. Inhib. Med. Chem.* **2014**, *29*, 146–150. [\[CrossRef\]](#)
179. Zaidi, S.; Srivastava, N.; Khare, S.K. Microbial Carbonic Anhydrase Mediated Carbon Capture, Sequestration & Utilization: A Sustainable Approach to Delivering Bio-Renewables. *Bioresour. Technol.* **2022**, *365*, 128174. [\[CrossRef\]](#)
180. Rasouli, H.; Iliuta, I.; Bougie, F.; Garnier, A.; Iliuta, M.C. Enhanced CO₂ Capture in Packed-Bed Column Bioreactors with Immobilized Carbonic Anhydrase. *Chem. Eng. J.* **2022**, *432*, 134029. [\[CrossRef\]](#)
181. Vinoba, M.; Bhagiyalakshmi, M.; Jeong, S.K.; Yoon, Y.I.; Nam, S.C. Immobilization of Carbonic Anhydrase on Spherical SBA-15 for Hydration and Sequestration of CO₂. *Colloids Surf. B Biointerfaces* **2012**, *90*, 91–96. [\[CrossRef\]](#)
182. Uygun, M.; Singh, V.V.; Kaufmann, K.; Uygun, D.A.; De Oliveira, S.D.S.; Wang, J. Micromotor-Based Biomimetic Carbon Dioxide Sequestration: Towards Mobile Microscrubbers. *Angew. Chem.-Int. Ed.* **2015**, *54*, 12900–12904. [\[CrossRef\]](#)
183. Xu, X.; Kentish, S.E.; Martin, G.J.O. Direct Air Capture of CO₂ by Microalgae with Buoyant Beads Encapsulating Carbonic Anhydrase. *ACS Sustain. Chem. Eng.* **2021**, *9*, 9698–9706. [\[CrossRef\]](#)
184. Sifat, N.S.; Haseli, Y. A Critical Review of CO₂ Capture Technologies and Prospects for Clean Power Generation. *Energies* **2019**, *12*, 4143. [\[CrossRef\]](#)

185. Fu, L.; Ren, Z.; Si, W.; Ma, Q.; Huang, W.; Liao, K.; Huang, Z.; Wang, Y.; Li, J.; Xu, P. Research Progress on CO₂ Capture and Utilization Technology. *J. CO₂ Util.* **2022**, *66*, 102260. [CrossRef]
186. Fagorite, I.V.; Chijioke, C.F.; Opara, I.A.; Onyekuru, S.O.; Oguzie, E.E. Environmental and Safety Issues Associated with Geological Carbon Storage: A Review. *Euro-Mediterr. J. Environ. Integr.* **2022**, *7*, 445–461. [CrossRef]
187. Farrukh, S.; Wu, D.; Al-Dadah, R.; Gao, W.; Wang, Z. A Review of Integrated Cryogenic Energy Assisted Power Generation Systems and Desalination Technologies. *Appl. Therm. Eng.* **2023**, *221*, 119836. [CrossRef]
188. International Energy Agency. CO₂ Emissions in 2022. Available online: <https://iea.blob.core.windows.net/assets/3c8fa115-35c4-4474-b237-1b00424c8844/CO2Emissionsin2022.pdf> (accessed on 30 June 2023).
189. International Energy Agency. Global Energy-Related CO₂ Emissions by Sector. Available online: <https://www.iea.org/data-and-statistics/charts/global-energy-related-co2-emissions-by-sector> (accessed on 30 June 2023).
190. Hepburn, C.; Adlen, E.; Beddington, J.; Carter, E.A.; Fuss, S.; Mac Dowell, N.; Minx, J.C.; Smith, P.; Williams, C.K. The Technological and Economic Prospects for CO₂ Utilization and Removal. *Nature* **2019**, *575*, 87–97. [CrossRef]
191. Gundersen, M.T.; Von Solms, N.; Woodley, J.M. Enzymatically Assisted CO₂ Removal from Flue-Gas. In *Energy Procedia*; Department of Chemical and Biochemical Engineering, Technical University of Denmark: Lyngby, Denmark, 2014; Volume 63, pp. 624–632. [CrossRef]
192. Zhang, S.; Du, M.; Shao, P.; Wang, L.; Ye, J.; Chen, J.; Chen, J. Carbonic Anhydrase Enzyme-MOFs Composite with a Superior Catalytic Performance to Promote CO₂ Absorption into Tertiary Amine Solution. *Environ. Sci. Technol.* **2018**, *52*, 12708–12716. [CrossRef]
193. Gladis, A.; Lomholdt, N.F.; Fosbøl, P.L.; Woodley, J.M.; von Solms, N. Pilot Scale Absorption Experiments with Carbonic Anhydrase-Enhanced MDEA-Benchmarking with 30 wt% MEA. *Int. J. Greenh. Gas Control.* **2019**, *82*, 69–85. [CrossRef]
194. Kim, T.-J.; Lang, A.; Chikukwa, A.; Sheridan, E.; Dahl, P.I.; Leimbrink, M.; Skiborowski, M.; Roubroeks, J. Enzyme Carbonic Anhydrase Accelerated CO₂ Absorption in Membrane Contactor. In *Energy Procedia*; SINTEF Materials and Chemistry: Trondheim, Norway, 2017; Volume 114, pp. 17–24. [CrossRef]
195. Thee, H.; Smith, K.H.; Da Silva, G.; Kentish, S.E.; Stevens, G.W. Carbonic Anhydrase Promoted Absorption of CO₂ into Potassium Carbonate Solutions. *Greenh. Gases Sci. Technol.* **2015**, *5*, 108–114. [CrossRef]
196. Hu, G.; Nicholas, N.J.; Smith, K.H.; Mumford, K.A.; Kentish, S.E.; Stevens, G.W. Carbon Dioxide Absorption into Promoted Potassium Carbonate Solutions: A Review. *Int. J. Greenh. Gas Control* **2016**, *53*, 28–40. [CrossRef]
197. Jin, P.; Zhang, S.; Liu, Y.; Zhang, W.; Wang, R. Application of *Bacillus mucilaginosus* in the Carbonation of Steel Slag. *Appl. Microbiol. Biotechnol.* **2021**, *105*, 8663–8674. [CrossRef] [PubMed]
198. Jo, B.H.; Seo, J.H.; Yang, Y.J.; Baek, K.; Choi, Y.S.; Pack, S.P.; Oh, S.H.; Cha, H.J. Bioinspired Silica Nanocomposite with Autoencapsulated Carbonic Anhydrase as a Robust Biocatalyst for CO₂ Sequestration. *ACS Catal.* **2014**, *4*, 4332–4340. [CrossRef]
199. Yang, G.; Li, L.; Li, F.; Zhang, C.; Lyu, J. Mechanism of Carbonate Mineralization Induced by Microbes: Taking *Curvibacter lanceolatus* Strain HJ-1 as an Example. *Micron* **2021**, *140*, 102980. [CrossRef]
200. Bose, H.; Satyanarayana, T. Suitability of the Alkalistable Carbonic Anhydrase from a Polyextremophilic Bacterium *Aeribacillus pallidus* TSHB1 in Biomimetic Carbon Sequestration. *Bioprocess Biosyst. Eng.* **2016**, *39*, 1515–1525. [CrossRef]
201. De Luca, V.; Vullo, D.; Scozzafava, A.; Carginale, V.; Rossi, M.; Supuran, C.T.; Capasso, C. An Alpha-Carbonic Anhydrase from the Thermophilic Bacterium *Sulphurihydrogenibium azorense* Is the Fastest Enzyme Known for the CO₂ Hydration Reaction. *Bioorg. Med. Chem.* **2013**, *21*, 1465–1469. [CrossRef]
202. Di Lorenzo, F.; Ruiz-Agudo, C.; Ibañez-Velasco, A.; Gil-San Millán, R.; Navarro, J.A.R.; Ruiz-Agudo, E.; Rodríguez-Navarro, C. The Carbonation of Wollastonite: A Model Reaction to Test Natural and Biomimetic Catalysts for Enhanced CO₂ Sequestration. *Minerals* **2018**, *8*, 209. [CrossRef]
203. Wang, M.; Lawal, A.; Stephenson, P.; Sidders, J.; Ramshaw, C. Post-Combustion CO₂ Capture with Chemical Absorption: A State-of-the-Art Review. *Chem. Eng. Res. Des.* **2011**, *89*, 1609–1624. [CrossRef]
204. Heldebrandt, D.J.; Koeck, P.K.; Glezakou, V.-A.; Rousseau, R.; Malhotra, D.; Cantu, D.C. Water-Lean Solvents for Post-Combustion CO₂ Capture: Fundamentals, Uncertainties, Opportunities, and Outlook. *Chem. Rev.* **2017**, *117*, 9594–9624. [CrossRef]
205. Sharif, M.; Zhang, T.; Wu, X.; Yu, Y.; Zhang, Z. Evaluation of CO₂ Absorption Performance by Molecular Dynamic Simulation for Mixed Secondary and Tertiary Amines. *Int. J. Greenh. Gas. Control* **2020**, *97*, 103059. [CrossRef]
206. Bernhardsen, I.M.; Krokvik, I.R.T.; Jens, K.-J.; Knuutila, H.K. Performance of MAPA Promoted Tertiary Amine Systems for CO₂ Absorption: Influence of Alkyl Chain Length and Hydroxyl Groups. *Energy Procedia* **2017**, *114*, 1682–1688. [CrossRef]
207. Liu, B.; Cui, Z.; Tian, W. The Kinetics Investigation of CO₂ Absorption into TEA and DEEA Amine Solutions Containing Carbonic Anhydrase. *Processes* **2021**, *9*, 2140. [CrossRef]
208. Bui, M.; Adjiman, C.S.; Bardow, A.; Anthony, E.J.; Boston, A.; Brown, S.; Fennell, P.S.; Fuss, S.; Galindo, A.; Hackett, L.A.; et al. Carbon Capture and Storage (CCS): The Way Forward. *Energy Environ. Sci.* **2018**, *11*, 1062–1176. [CrossRef]
209. Power, I.M.; Harrison, A.L.; Dipple, G.M. Accelerating Mineral Carbonation Using Carbonic Anhydrase. *Environ. Sci. Technol.* **2016**, *50*, 2610–2618. [CrossRef] [PubMed]
210. Ehrlich, H.; Bailey, E.; Wysokowski, M.; Jesionowski, T. Forced Biomineralization: A Review. *Biomimetics* **2021**, *6*, 46. [CrossRef]
211. Bose, H.; Satyanarayana, T. Microbial Carbonic Anhydrases in Biomimetic Carbon Sequestration for Mitigating Global Warming: Prospects and Perspectives. *Front. Microbiol.* **2017**, *8*, 1615. [CrossRef]

212. Wang, F.; Dreisinger, D.B.; Jarvis, M.; Hitchins, T. The Technology of CO₂ Sequestration by Mineral Carbonation: Current Status and Future Prospects. *Can. Metall. Q.* **2018**, *57*, 46–58. [\[CrossRef\]](#)
213. Hills, C.D.; Tripathi, N.; Carey, P.J. Mineralization Technology for Carbon Capture, Utilization, and Storage. *Front. Energy Res.* **2020**, *8*, 142. [\[CrossRef\]](#)
214. Bhatia, S.K.; Bhatia, R.K.; Jeon, J.-M.; Kumar, G.; Yang, Y.-H. Carbon Dioxide Capture and Bioenergy Production Using Biological System—A Review. *Renew. Sust. Energy Rev.* **2019**, *110*, 143–158. [\[CrossRef\]](#)
215. Zajac, M.; Krol, M.; Bullerjahn, F.; Deja, J. Effect of Temperature on Carbon Dioxide Mineralisation in Recycled Cement Paste. *Adv. Cem. Res.* **2023**, *35*, 1–12. [\[CrossRef\]](#)
216. Yin, B.; Xu, H.; Fan, F.; Qi, D.; Hua, X.; Xu, T.; Liu, C.; Hou, D. Superhydrophobic Coatings Based on Bionic Mineralization for Improving the Durability of Marine Concrete. *Constr. Build. Mater.* **2023**, *362*, 129705. [\[CrossRef\]](#)
217. Heldebrant, D.J.; Kothandaraman, J.; Mac Dowell, N.; Brickett, L. Next Steps for Solvent-Based CO₂ Capture; Integration of Capture, Conversion, and Mineralisation. *Chem. Sci.* **2022**, *13*, 6445–6456. [\[CrossRef\]](#)
218. Rodriguez-Navarro, C.; Cizer, O.; Kudlacz, K.; Ibanez-Velasco, A.; Ruiz-Agudo, C.; Elert, K.; Burgos-Cara, A.; Ruiz-Agudo, E. The Multiple Roles of Carbonic Anhydrase in Calcium Carbonate Mineralization. *CrystEngComm* **2019**, *21*, 7407–7423. [\[CrossRef\]](#)
219. Fernández, M.S.; Montt, B.; Ortiz, L.; Neira-Carrillo, A.; Arias, J.L. Effect of Carbonic Anhydrase Immobilized on Eggshell Membranes on Calcium Carbonate Crystallization In Vitro. In *Biomining*; Endo, K., Kogure, T., Nagasawa, H., Eds.; Springer: Singapore, 2018; pp. 31–37.
220. Jin, P.; Wang, R.; Zhang, S.; Chen, Y. Effect of Carbonic Anhydrase Bacteria on the Carbonation Process of γ -C2S. *Adv. Cem. Res.* **2021**, *34*, 15–27. [\[CrossRef\]](#)
221. Sharma, V.K.; Hutchison, J.M.; Allgeier, A.M. Redox Biocatalysis: Quantitative Comparisons of Nicotinamide Cofactor Regeneration Methods. *ChemSusChem* **2022**, *15*, e2022008. [\[CrossRef\]](#) [\[PubMed\]](#)
222. Navarro-Jaen, S.; Virginie, M.; Bonin, J.; Robert, M.; Wojcieszak, R.; Khodakov, A.Y. Highlights and Challenges in the Selective Reduction of Carbon Dioxide to Methanol. *Nat. Rev. Chem.* **2021**, *5*, 564–579. [\[CrossRef\]](#)
223. Corrado, M.L.; Knaus, T.; Schwaneberg, U.; Mutti, F.G. High-Yield Synthesis of Enantiopure 1,2-Amino Alcohols from L-Phenylalanine via Linear and Divergent Enzymatic Cascades. *Org. Process Res. Dev.* **2022**, *26*, 2085–2095. [\[CrossRef\]](#)
224. Singh, P.; Srivastava, R. Utilization of Bio-Inspired Catalyst for CO₂ Reduction into Green Fuels: Recent Advancement and Future Perspectives. *J. CO₂ Util.* **2021**, *53*, 101748. [\[CrossRef\]](#)
225. Armstrong, F.A.; Cheng, B.; Herold, R.A.; Megarity, C.F.; Siritanaratkul, B. From Protein Film Electrochemistry to Nanoconfined Enzyme Cascades and the Electrochemical Leaf. *Chem. Rev.* **2023**, *123*, 5421–5458. [\[CrossRef\]](#)
226. Meneghello, M.; Leger, C.; Fourmond, V. Electrochemical Studies of CO₂-Reducing Metalloenzymes. *Chem. Eur. J.* **2021**, *27*, 17542–17553. [\[CrossRef\]](#)
227. Rasheed, T.; Shafi, S.; Anwar, M.T.; Rizwan, K.; Ahmad, T.; Bilal, M. Revisiting Photo and Electro-Catalytic Modalities for Sustainable Conversion of CO₂. *Appl. Catal. A-Gen.* **2021**, *623*, 118248. [\[CrossRef\]](#)
228. Cadoux, C.; Milton, R.D. Recent Enzymatic Electrochemistry for Reductive Reactions. *ChemElectroChem* **2020**, *7*, 1974–1986. [\[CrossRef\]](#)
229. Xing, X.; Liu, Y.; Lin, R.-D.; Zhang, Y.; Wu, Z.-L.; Yu, X.-Q.; Li, K.; Wang, N. Development of an Integrated System for Highly Selective Photoenzymatic Synthesis of Formic Acid from CO₂. *ChemSusChem* **2023**, *16*, e2022019. [\[CrossRef\]](#)
230. Wang, Q.; Pan, Z. Advances and Challenges in Developing Cocatalysts for Photocatalytic Conversion of Carbon Dioxide to Fuels. *Nano Res.* **2022**, *15*, 10090–10109. [\[CrossRef\]](#)
231. Fang, X.; Kalathil, S.; Reisner, E. Semi-Biological Approaches to Solar-to-Chemical Conversion. *Chem. Soc. Rev.* **2020**, *49*, 4926–4952. [\[CrossRef\]](#)
232. Ozgen, F.F.; Runda, M.E.; Schmidt, S. Photo-Biocatalytic Cascades: Combining Chemical and Enzymatic Transformations Fueled by Light. *ChemBioChem* **2021**, *22*, 790–806. [\[CrossRef\]](#)
233. Chen, H.; Huang, Y.; Sha, C.; Moradian, J.M.; Yong, Y.-C.; Fang, Z. Enzymatic Carbon Dioxide to Formate: Mechanisms, Challenges and Opportunities. *Renew. Sust. Energy Rev.* **2023**, *178*, 113271. [\[CrossRef\]](#)
234. Immanuel, S.; Sivasubramanian, R.; Gul, R.; Dar, M.A. Recent Progress and Perspectives on Electrochemical Regeneration of Reduced Nicotinamide Adenine Dinucleotide (NADH). *Chem. Asian J.* **2020**, *15*, 4256–4270. [\[CrossRef\]](#)
235. Lee, Y.S.; Gerulskis, R.; Minteer, S.D. Advances in Electrochemical Cofactor Regeneration: Enzymatic and Non-Enzymatic Approaches. *Curr. Opin. Biotechnol.* **2022**, *73*, 14–21. [\[CrossRef\]](#)
236. Zhang, Z.; Zhang, X.; Ji, X. Developing and Regenerating Cofactors for Sustainable Enzymatic CO₂ Conversion. *Processes* **2022**, *10*, 230. [\[CrossRef\]](#)
237. Bachosz, K.; Zdzarta, J.; Bilal, M.; Meyer, A.S.; Jesionowski, T. Enzymatic Cofactor Regeneration Systems: A New Perspective on Efficiency Assessment. *Sci. Total Environ.* **2023**, *868*, 161630. [\[CrossRef\]](#)
238. Findrik, Z.; Vasic-Racki, D. Overview on Reactions with Multi-Enzyme Systems. *Chem. Biochem. Eng. Q.* **2009**, *23*, 545–553.
239. Neves, L.A.; Afonso, C.A.M.; Coelho, L.M.; Crespo, J.G. CO₂ Capture by Enzymatic Bioconversion in a Membrane Contactor with Task Specific Ionic Liquids. In *Procedia Engineering*; Universidade Nova de Lisboa: Lisbon, Portugal, 2012; Volume 44, pp. 557–558. [\[CrossRef\]](#)
240. Qadir, M.I.; Dupont, J. Thermo- and Photocatalytic Activation of CO₂ in Ionic Liquids. *Angew. Chem. Int. Ed.* **2023**, e202301497. [\[CrossRef\]](#)

241. Chenault, H.K.; Whitesides, G.M. Regeneration of Nicotinamide Cofactors for Use in Organic Synthesis. *Appl. Biochem. Biotechnol.* **1987**, *14*, 147–197. [CrossRef] [PubMed]
242. Chenault, H.K.; Simon, E.S.; Whitesides, G.M. Cofactor Regeneration for Enzyme-Catalysed Synthesis. *Biotechnol. Genet. Eng. Rev.* **1988**, *6*, 221–270. [CrossRef]
243. Yu, X.; Niks, D.; Ge, X.; Liu, H.; Hille, R.; Mulchandani, A. Synthesis of Formate from CO₂ Gas Catalyzed by an O₂-Tolerant NAD-Dependent Formate Dehydrogenase and Glucose Dehydrogenase. *Biochemistry* **2019**, *58*, 1861–1868. [CrossRef]
244. Liao, Q.; Liu, W.; Meng, Z. Strategies for Overcoming the Limitations of Enzymatic Carbon Dioxide Reduction. *Biotechnol. Adv.* **2022**, *60*, 108024. [CrossRef]
245. El-Zahab, B.; Donnelly, D.; Wang, P. Particle-Tethered NADH for Production of Methanol from CO₂ Catalyzed by Coimmobilized Enzymes. *Biotechnol. Bioeng.* **2008**, *99*, 508–514. [CrossRef]
246. Ji, X.; Su, Z.; Wang, P.; Ma, G.; Zhang, S. Tethering of Nicotinamide Adenine Dinucleotide inside Hollow Nanofibers for High-Yield Synthesis of Methanol from Carbon Dioxide Catalyzed by Coencapsulated Multienzymes. *ACS Nano* **2015**, *9*, 4600–4610. [CrossRef]
247. Ren, S.; Wang, Z.; Bilal, M.; Feng, Y.; Jiang, Y.; Jia, S.; Cui, J. Co-Immobilization Multienzyme Nanoreactor with Co-Factor Regeneration for Conversion of CO₂. *Int. J. Biol. Macromol.* **2020**, *155*, 110–118. [CrossRef]
248. Shukia, S.K.; Khokarale, S.G.; Bui, T.Q.; Mikkola, J.-P.T. Ionic Liquids: Potential Materials for Carbon Dioxide Capture and Utilization. *Front. Mater.* **2019**, *6*, 42. [CrossRef]
249. Zeng, S.; Zhang, X.X.; Bai, L.; Zhang, X.X.; Wang, H.; Wang, J.; Bao, D.; Li, M.; Liu, X.; Zhang, S. Ionic-Liquid-Based CO₂ Capture Systems: Structure, Interaction and Process. *Chem. Rev.* **2017**, *117*, 9625–9673. [CrossRef]
250. Zhang, Z.; Muschiol, J.; Huang, Y.; Sigurdardottir, S.B.; von Solms, N.; Daugaard, A.E.; Wei, J.; Luo, J.; Xu, B.-H.; Zhang, S.; et al. Efficient Ionic Liquid-Based Platform for Multi-Enzymatic Conversion of Carbon Dioxide to Methanol. *Green Chem.* **2018**, *20*, 4339–4348. [CrossRef]
251. The Methanol Institute. Available online: <https://www.methanol.org/applications/> (accessed on 7 July 2023).
252. Cazelles, R.; Drone, J.; Fajula, F.; Ersen, O.; Moldovan, S.; Galarneau, A. Reduction of CO₂ to Methanol by a Polyenzymatic System Encapsulated in Phospholipids-Silica Nanocapsules. *New J. Chem.* **2013**, *37*, 3721–3730. [CrossRef]
253. Singh, R.K.R.; Singh, R.K.R.; Sivakumar, D.; Kondaveeti, S.; Kim, T.; Li, J.; Sung, B.H.; Cho, B.-K.; Kim, D.R.; Kim, S.C.; et al. Insights into Cell-Free Conversion of CO₂ to Chemicals by a Multienzyme Cascade Reaction. *ACS Catal.* **2018**, *8*, 11085–11093. [CrossRef]
254. Dave, B.C.; Rao, M.S.; Burt, M.C. Converting Carbon Dioxide to Methanol Comprises Serial Reduction of the Carbon Dioxide to Methanol By Dehydrogenase Enzymes in the Presence of Reduced Nicotinamide Adenine Dinucleotide. US20070042479(A1), 18 August 2005.
255. Alvarez-Malmagro, J.; Oliveira, A.R.; Gutierrez-Sanchez, C.; Villajos, B.; Pereira, I.A.C.; Velez, M.; Pita, M.; De Lacey, A.L. Bioelectrocatalytic Activity of W-Formate Dehydrogenase Covalently Immobilized on Functionalized Gold and Graphite Electrodes. *ACS Appl. Mater. Interfaces* **2021**, *13*, 11891–11900. [CrossRef]
256. Barin, R.; Rashid-Nadimi, S.; Biria, D.; Asadollahi, M.A. Direct Electrochemical Regeneration of 1,4-NADH at the Copper Foam and Bimetallic Copper Foam. *Electrochim. Acta* **2017**, *247*, 1095–1102. [CrossRef]
257. Song, H.; Ma, C.; Liu, P.; You, C.; Lin, J.; Zhu, Z. A Hybrid CO₂ Electroreduction System Mediated by Enzyme-Cofactor Conjugates Coupled with Cu Nanoparticle-Catalyzed Cofactor Regeneration. *J. CO₂ Util.* **2019**, *34*, 568–575. [CrossRef]
258. Addo, P.K.; Arechederra, R.L.; Waheed, A.; Shoemaker, J.D.; Sly, W.S.; Minter, S.D. Methanol Production via Bioelectrocatalytic Reduction of Carbon Dioxide: Role of Carbonic Anhydrase in Improving Electrode Performance. *Electrochem. Solid-State Lett.* **2011**, *14*, E9–E13. [CrossRef]
259. Yuan, M.; Sahin, S.; Cai, R.; Abdellaoui, S.; Hickey, D.P.; Minter, S.D.; Milton, R.D. Creating a Low-Potential Redox Polymer for Efficient Electroenzymatic CO₂ Reduction. *Angew. Chem. Int. Ed.* **2018**, *57*, 6582–6586. [CrossRef]
260. Kim, S.-H.; Chung, G.-Y.; Kim, S.-H.; Vinothkumar, G.; Yoon, S.-H.; Jung, K.-D. Electrochemical NADH Regeneration and Electroenzymatic CO₂ Reduction on Cu Nanorods/Glassy Carbon Electrode Prepared by Cyclic Deposition. *Electrochim. Acta* **2016**, *210*, 837–845. [CrossRef]
261. Chen, Y.; Li, P.; Zhou, J.; Buru, C.T.; Dordevic, L.; Li, P.; Zhang, X.; Cetin, M.M.; Stoddart, J.F.; Stupp, S.I.; et al. Integration of Enzymes and Photosensitizers in a Hierarchical Mesoporous Metal–Organic Framework for Light-Driven CO₂ Reduction. *J. Am. Chem. Soc.* **2020**, *142*, 1768–1773. [CrossRef]
262. Barin, R.; Biria, D.; Rashid-Nadimi, S.; Asadollahi, M.A. Enzymatic CO₂ Reduction to Formate by Formate Dehydrogenase from *Candida Boidinii* Coupling with Direct Electrochemical Regeneration of NADH. *J. CO₂ Util.* **2018**, *28*, 117–125. [CrossRef]
263. Zhang, Z.; Li, J.; Ji, M.; Liu, Y.; Wang, N.; Zhang, X.; Zhang, S.; Ji, X. Encapsulation of Multiple Enzymes in a Metal–Organic Framework with Enhanced Electro-Enzymatic Reduction of CO₂ to Methanol. *Green Chem.* **2021**, *23*, 2362–2371. [CrossRef]
264. Zhang, T.R.; Lu, S.Y. Sacrificial Agents for Photocatalytic Hydrogen Production: Effects, Cost, and Development. *Chem. Catalysis* **2022**, *2*, 1502–1505. [CrossRef]
265. Hernandez-Ibanez, N.; Gomis-Berenguer, A.; Montiel, V.; Ania, C.O.; Iniesta, J. Fabrication of a Biocathode for Formic Acid Production upon the Immobilization of Formate Dehydrogenase from *Candida boidinii* on a Nanoporous Carbon. *Chemosphere* **2022**, *291*, 133117. [CrossRef]

266. Cheng, Y.; Shi, J.; Wu, Y.; Wang, X.; Sun, Y.; Cai, Z.; Chen, Y.; Jiang, Z. Intensifying Electron Utilization by Surface-Anchored Rh Complex for Enhanced Nicotinamide Cofactor Regeneration and Photoenzymatic CO₂ Reduction. *Research* **2021**, *2021*, 8175709. [\[CrossRef\]](#)
267. Ottone, C.; Pugliese, D.; Laurenti, M.; Hernandez, S.; Cauda, V.; Grez, P.; Wilson, L. ZnO Materials as Effective Anodes for the Photoelectrochemical Regeneration of Enzymatically Active NAD⁺. *ACS Appl. Mater. Interfaces* **2021**, *13*, 10719–10727. [\[CrossRef\]](#)
268. Tensi, L.; Macchioni, A. Extremely Fast NADH-Regeneration Using Phosphonic Acid as Hydride Source and Iridium-Pyridine-2-Sulfonamide Catalysts. *ACS Catal.* **2020**, *10*, 7945–7949. [\[CrossRef\]](#)
269. Liao, Q.; Guo, M.; Mao, M.; Gao, R.; Meng, Z.; Fan, X.; Liu, W. Construction and Optimization of a Photo-Enzyme Coupled System for Sustainable CO₂ Conversion to Methanol. *Process Biochem.* **2023**, *129*, 44–55. [\[CrossRef\]](#)
270. Guo, M.; Gu, F.; Meng, L.; Liao, Q.; Meng, Z.; Liu, W. Synthesis of Formaldehyde from CO₂ Catalyzed by the Coupled Photo-Enzyme System. *Sep. Purif. Technol.* **2022**, *286*, 120480. [\[CrossRef\]](#)
271. Kim, J.A.; Kim, S.; Lee, J.; Baeg, J.-O.; Kim, J.A. Photochemical Production of NADH Using Cobaloxime Catalysts and Visible-Light Energy. *Inorg. Chem.* **2012**, *51*, 8057–8063. [\[CrossRef\]](#) [\[PubMed\]](#)
272. Laun, K.; Duffus, B.R.; Kumar, H.; Oudsen, J.-P.H.; Karafoulidi-Retsou, C.; Waffo, A.T.; Hildebrandt, P.; Ly, K.H.; Leimkuehler, S.; Katz, S.; et al. A Minimal Light-Driven System to Study the Enzymatic CO₂ Reduction of Formate Dehydrogenase. *ChemCatChem* **2022**, *14*, e202201067. [\[CrossRef\]](#)
273. Ji, X.; Wang, J.; Kang, Y.; Mei, L.; Su, Z.; Wang, S.; Ma, G.; Shi, J.; Zhang, S. Enhanced Solar Energy Harvest and Electron Transfer through Intra- and Intermolecular Dual Channels in Chlorosome-Mimicking Supramolecular Self-Assemblies. *ACS Catal.* **2018**, *8*, 10732–10745. [\[CrossRef\]](#)
274. Lin, G.; Zhang, Y.; Hua, Y.; Zhang, C.; Jia, C.; Ju, D.; Yu, C.; Li, P.; Liu, J. Bioinspired Metalation of the Metal-Organic Framework MIL-125-NH₂ for Photocatalytic NADH Regeneration and Gas-Liquid-Solid Three-Phase Enzymatic CO₂ Reduction. *Angew. Chem. Int. Ed.* **2022**, *61*, e202206283. [\[CrossRef\]](#) [\[PubMed\]](#)
275. Ong, W.-J.; Tan, L.-L.; Ng, Y.H.; Yong, S.-T.; Chai, S.-P. Graphitic Carbon Nitride (g-C₃N₄)-Based Photocatalysts for Artificial Photosynthesis and Environmental Remediation: Are We a Step Closer To Achieving Sustainability? *Chem. Rev.* **2016**, *116*, 7159–7329. [\[CrossRef\]](#)
276. Zeng, P.; Ji, X.; Su, Z.; Zhang, S. WS₂/g-C₃N₄ Composite as an Efficient Heterojunction Photocatalyst for Biocatalyzed Artificial Photosynthesis. *RSC Adv.* **2018**, *8*, 20557–20567. [\[CrossRef\]](#)
277. Meng, J.; Tian, Y.; Li, C.; Lin, X.; Wang, Z.; Sun, L.; Zhou, Y.; Li, J.; Yang, N.; Zong, Y.; et al. A Thiophene-Modified Doubleshell Hollow g-C₃N₄ Nanosphere Boosts NADH Regeneration via Synergistic Enhancement of Charge Excitation and Separation. *Catal. Sci. Technol.* **2019**, *9*, 1911–1921. [\[CrossRef\]](#)
278. Gao, Y.; Li, W.; Sun, X.; Zhao, Y.; Ji, H.; Luo, H.; Wu, G.; Wan, L.; Zhang, L. Boosting the Performance of Formate Dehydrogenase by Silver Nanoclusters for Photoreduction of CO₂ to Formate. *ACS Sustain. Chem. Eng.* **2022**, *10*, 14888–14896. [\[CrossRef\]](#)
279. Gupta, P.; Verma, N. Conversion of CO₂ to Formate Using Activated Carbon Fiber-Supported g-C₃N₄-NiCoWO₄ Photoanode in a Microbial Electrosynthesis System. *Chem. Eng. J.* **2022**, *446*, 137029. [\[CrossRef\]](#)
280. Tian, Y.; Zhou, Y.; Zong, Y.; Li, J.; Yang, N.; Zhang, M.; Guo, Z.; Song, H. Construction of Functionally Compartmental Inorganic Photocatalyst-Enzyme System via Imitating Chloroplast for Efficient Photoreduction of CO₂ to Formic Acid. *ACS Appl. Mater. Interfaces* **2020**, *12*, 34795–34805. [\[CrossRef\]](#)
281. Yu, S.; Lv, P.; Xue, P.; Wang, K.; Yang, Q.; Zhou, J.; Wang, M.; Wang, L.; Chen, B.; Tan, T. Light-Driven Enzymatic Nanosystem for Highly Selective Production of Formic Acid from CO₂. *Chem. Eng. J.* **2021**, *420*, 127649. [\[CrossRef\]](#)
282. Ji, X.; Kang, Y.; Su, Z.; Wang, P.; Ma, G.; Zhang, S. Graphene Oxide and Polyelectrolyte Composed One-Way Expressway for Guiding Electron Transfer of Integrated Artificial Photosynthesis. *ACS Sustain. Chem. Eng.* **2018**, *6*, 3060–3069. [\[CrossRef\]](#)
283. Ji, X.; Su, Z.; Wang, P.; Ma, G.; Zhang, S. Integration of Artificial Photosynthesis System for Enhanced Electronic Energy-Transfer Efficacy: A Case Study for Solar-Energy Driven Bioconversion of Carbon Dioxide to Methanol. *Small* **2016**, *12*, 4753–4762. [\[CrossRef\]](#) [\[PubMed\]](#)
284. Kim, S.; Lee, S.; Anjong, T.F.; Jang, H.Y.; Kim, J.-Y.; Lee, C.; Park, S.; Lee, H.J.; Yoon, J.; Kim, J. Artificial Photocatalytic System Using Polydiacetylene-(NH-Phen)Ru(Bpy)₂ for Cofactor Regeneration and CO₂ Reduction. *J. Phys. Chem. C* **2016**, *120*, 28407–28414. [\[CrossRef\]](#)
285. Sokol, K.P.; Robinson, W.E.; Oliveira, A.R.; Warnan, J.; Nowaczyk, M.M.; Ruff, A.; Pereira, I.A.C.; Reisner, E. Photoreduction of CO₂ with a Formate Dehydrogenase Driven by Photosystem II Using a Semi-Artificial Z-Scheme Architecture. *J. Am. Chem. Soc.* **2018**, *140*, 16418–16422. [\[CrossRef\]](#) [\[PubMed\]](#)
286. Schuler, E.; Ermolich, P.A.; Shiju, N.R.; Gruter, G.-J. Monomers from CO₂: Superbases as Catalysts for Formate-to-Oxalate Coupling. *ChemSusChem* **2021**, *14*, 1517–1523. [\[CrossRef\]](#)
287. Murcia-Valderrama, M.A.; van Putten, R.-J.; Gruter, G.-J.M. The Potential of Oxalic—And Glycolic Acid Based Polyesters (Review). Towards CO₂ as a Feedstock (Carbon Capture and Utilization—CCU). *Eur. Polym. J.* **2019**, *119*, 445–468. [\[CrossRef\]](#)
288. Schwarz, F.M.; Mueller, V. Whole-Cell Biocatalysis for Hydrogen Storage and Syngas Conversion to Formate Using a Thermophilic Acetogen. *Biotechnol. Biofuels* **2020**, *13*, 32. [\[CrossRef\]](#)
289. Luxuan, L.; Zhijun, X.; Huang, X. Whole-Cell-Based Photosynthetic Biohybrid Systems for Energy and Environmental Applications. *ChemPlusChem* **2021**, *86*, 1021–1036.

290. Hwang, H.; Yeon, Y.; Lee, S.; Choe, H.; Cho, D.; Park, S.; Kim, Y. Electro-Biocatalytic Production of Formate from Carbon Dioxide using an Oxygen-Stable Whole-Cell Biocatalyst. *Bioresour. Technol.* **2015**, *185*, 35–39. [\[CrossRef\]](#)
291. Le, T.Q.A. Recent Applications and Strategies to Enhance Performance of Electrochemical Reduction of CO₂ Gas into Value-Added Chemicals Catalyzed by Whole-Cell Biocatalysts. *Processes* **2023**, *11*, 766. [\[CrossRef\]](#)
292. Alissandratos, A.; Kim, H.-K.; Easton, C.J. Formate Production through Carbon Dioxide Hydrogenation with Recombinant Whole Cell Biocatalysts. *Bioresour. Technol.* **2014**, *164*, 7–11. [\[CrossRef\]](#)
293. Roger, M.; Brown, F.; Gabrielli, W.; Sargent, F. Efficient Hydrogen-Dependent Carbon Dioxide Reduction by *Escherichia coli*. *Curr. Biol.* **2018**, *28*, 140–145.e2. [\[CrossRef\]](#)
294. Leo, F.; Schwarz, F.M.; Schuchmann, K.; Mueller, V. Capture of Carbon Dioxide and Hydrogen by Engineered *Escherichia coli*: Hydrogen-Dependent CO₂ Reduction to Formate. *Appl. Microbiol. Biotechnol.* **2021**, *105*, 5861–5872. [\[CrossRef\]](#)
295. Schwarz, F.M.; Oswald, F.; Müller, V. Acetogenic Conversion of H₂ and CO₂ into Formic Acid and Vice Versa in a Fed-Batch-Operated Stirred-Tank Bioreactor. *ACS Sustain. Chem. Eng.* **2021**, *9*, 6810–6820. [\[CrossRef\]](#)
296. Le, Q.A.T.; Kim, H.G.; Kim, Y.H. Electrochemical Synthesis of Formic Acid from CO₂ Catalyzed by *Shewanella oneidensis* MR-1 Whole-Cell Biocatalyst. *Enzym. Microb. Technol.* **2018**, *116*, 1–5. [\[CrossRef\]](#)
297. Singh, S.; Noori, M.T.; Verma, N. Efficient Bio-Electroreduction of CO₂ to Formate on a Iron Phthalocyanine-Dispersed CDC in Microbial Electrolysis System. *Electrochim. Acta* **2020**, *338*, 135887. [\[CrossRef\]](#)
298. Phan, U.T.; Jeon, B.W.; Kim, Y.H. Microbial Engineering of *Methylobacterium extorquens* AM1 to Enhance CO₂ Conversion into Formate. *Enzym. Microb. Technol.* **2023**, *168*, 110264. [\[CrossRef\]](#)
299. Chen, H.; Li, J.; Fan, Q.; Zheng, T.; Zhang, Y.; Yong, Y.-C.; Fang, Z. A feasible strategy for microbial electrocatalytic CO₂ reduction via whole-cell-packed and exogenous-mediator-free rGO/*Shewanella* biohydrogel. *Chem. Eng. J.* **2023**, *460*, 141863. [\[CrossRef\]](#)
300. Guntermann, N.; Mengers, H.G.; Franciò, G.; Blank, L.M.; Leitner, W. Bio-energy Conversion with Carbon Capture and Utilization (BECCU): Integrated Biomass Fermentation and Chemo-Catalytic CO₂ Hydrogenation for Bioethanol and Formic Acid Coproduction. *Green Chem.* **2021**, *23*, 9860–9864. [\[CrossRef\]](#)
301. Rowe, S.F.; Le Gall, G.; Ainsworth, E.V.; Davies, J.A.; Lockwood, C.W.J.; Shi, L.; Elliston, A.; Roberts, I.N.; Waldron, K.W.; Richardson, D.J.; et al. Light-Driven H₂ Evolution and C=C or C=O Bond Hydrogenation by *Shewanella oneidensis*: A Versatile Strategy for Photocatalysis by Nonphotosynthetic Microorganisms. *ACS Catal.* **2017**, *7*, 7558–7566. [\[CrossRef\]](#)
302. Tuyishime, P.; Sinumvayo, J.P. Novel Outlook in Engineering Synthetic Methylobacteria and Formatotrophs: A Course for Advancing C₁-Based Chemicals Production. *World J. Microbiol. Biotechnol.* **2020**, *36*, 118. [\[CrossRef\]](#) [\[PubMed\]](#)
303. Yishai, O.; Lindner, S.N.; Gonzalez de la Cruz, J.; Tenenboim, H.; Bar-Even, A. The formate bio-economy. *Curr. Opin. Chem. Biol.* **2016**, *35*, 1–9. [\[CrossRef\]](#) [\[PubMed\]](#)
304. Shengyuan Guo, S.; Asset, T.; Atanassov, P. Catalytic Hybrid Electrocatalytic/Biocatalytic Cascades for Carbon Dioxide Reduction and Valorization. *ACS Catalysis* **2021**, *11*, 5172–5188. [\[CrossRef\]](#)
305. Liang, B.; Zhao, Y.; Yang, J. Recent Advances in Developing Artificial Autotrophic Microorganism for Reinforcing CO₂ Fixation. *Front. Microbiol.* **2020**, *11*, 592631. [\[CrossRef\]](#)
306. Collas, F.; Dronsella, B.B.; Kubis, A.; Schann, K.; Arto, N.; Claassens, N.J.; Kensy, F.; Orsi, E. Engineering the Biological Conversion of Formate into Crotonate in *Cupriavidus necator*. *Metab. Eng.* **2023**, *in press*. [\[CrossRef\]](#)
307. Cotton, C.A.; Claassens, N.J.; Benito-Vaquerizo, S.; Bar-Even, A. Renewable Methanol and Formate as Microbial Feedstocks. *Curr. Opin. Biotechnol.* **2020**, *62*, 168–180. [\[CrossRef\]](#)

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