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Ionic Liquids as Bifunctional Cosolvents Enhanced CO₂ Conversion Catalysed by NADH-Dependent Formate Dehydrogenase

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Abstract: Efficient CO₂ conversion by formate dehydrogenase is limited by the low CO₂ concentrations that can be reached in traditional buffers. The use of ionic liquids was proposed as a manner to increase CO₂ concentration in the reaction system. It has been found, however, that the required cofactor (NADH) heavily degraded during the enzymatic reaction and that acidity was the main reason. Acidity, indeed, resulted in reduction of the conversion of CO₂ into formic acid and contributed to overestimate the amount of formic acid produced when the progression of the reaction was followed by a decrease in NADH absorbance (method N). Stability of NADH and the mechanism of NADH degradation was investigated by UV, NMR and by DFT calculations. It was found that by selecting neutral–basic ionic liquids and by adjusting the concentration of the ionic liquid in the buffer, the concentration of NADH can be maintained in the reaction system with little loss. Conversion of CO₂ to methanol in BmimBF₄ (67.1%) was more than twice as compared with the conversion attained by the enzymatic reaction in phosphate buffer (24.3%).

Keywords: ionic liquids; formate dehydrogenase; NADH degradation; CO₂ conversion

1. Introduction

Carbon dioxide (CO₂) emissions from combustion of fossil fuels and its greenhouse effect on climate change (i.e., global warming) are considered a current threat [1]. To minimize environmental problems and produce clean energy, efficient utilization of CO₂ and carbon regeneration has been the focus of a tremendous amount of research [2–5]. Recently, a strategy by which CO₂ may be converted enzymatically into valuable chemicals and fuels such as formate, formaldehyde and methanol has inspired many researchers [6,7]. Obert and Dave were the first to report a cascade reaction involving the three enzymes, formate dehydrogenase (FDH), formaldehyde dehydrogenase (FaldDH) and alcohol dehydrogenase (ADH), where the product of the first reaction serves as a substrate of downstream reaction. Reduced nicotinamide adenine dinucleotide (NADH) was used as the terminal electron donor for the enzymatic reaction [8]. However, enzymatic hydrogenation of CO₂ to formic acid (CH₂O₂), formaldehyde (CH₂O) and methanol (CH₃OH) is hampered by the low concentration of CO₂ that is available for the enzyme (formic acid dehydrogenase) in the reaction mixture. Such a low
concentration may account for the common poor conversion to formic acid and hence low conversions to formaldehyde and methanol [9].

CO$_2$ is highly soluble in ionic liquids (ILs) and solubility can be enhanced by adjusting the anion and substituents on the cation, for instance, by fluorinating the anion or cation components [10]. These components usually interact via electrostatic forces, van der Waals forces, hydrogen bonds and other physical effects, which can explain the high solubilities of CO$_2$ in ILs [11–13]. For instance, at just 50 bar of CO$_2$ pressure, CO$_2$ solubility was on the order of 50% mole fraction in [Bmim][PF$_6$] [14]. ILs have also been used in various enzymatic reactions involving enzymes such as lipases, celluloses, and alcohol dehydrogenases [15–17]. Zhao reported that ILs were successfully applied to lipase-catalysed enantioselective esterification reactions for enhancing the stability of lipase [18]. Zhu reported that cellulase showed a higher stability than cellobiase in aqueous 1-ethyl-3-methylimidazolium acetate media for cellulose hydrolysis [19].

It has therefore been promising to perform enzymatic conversion of CO$_2$ in an IL environment in order to obtain better conversion efficiency. For instance, Amado employed ILs as an alternative solvent for enzymatic conversion of CO$_2$ to methanol [20]. However, the conversion of CO$_2$ did not increase significantly compared to the result obtained with the conventional buffer. In addition, Nicole and Udo reported that the changes in concentration of ILs could cause FDH inactivation to different degrees [21]. Bahareh and Khosro reported that imidazolium based ILs had a severe inhibition effect on alcohol dehydrogenase [22]. Interestingly, however, the activity of enzyme was partially recovered with diluted ILs of less than 150 mM.

Former research has focused on the activity of dehydrogenase in ILs, but investigations on the interaction between ILs and coenzyme (NADH), which could play a key role in the conversion of CO$_2$, have been neglected. NADH is used not only as hydrogen donor in the reduction of CO$_2$ but is also employed for following the progression of the reaction and thus to quantify formate [23–25]. NADH has been reported to be unstable in dilute acid by Andersen and co-workers [26]. Incubation of N-substituted dihydropyridines in acid results in the addition of water across the 2–3 double bond to yield a 3-hydroxy-tetrahydropyridine. Alivisatos and co-workers have reported that the reaction of NADH in concentrated phosphate solutions first forms 3-hydroxy-tetrahydroxypyridine-tinamide which then subsequently rearranges to $\alpha$-O$_2$.3B-cyclo-tetrahydronicotinamide adenine dinucleotide [27]. Furthermore, the same mechanism of NADH degradation in acid has been demonstrated by Norman and co-workers [28]. Degradation of NADH in the enzymatic reaction not only reduced the conversion efficiency but also led to false results of high CO$_2$ conversion detected by method N (NADH UV absorption). Even though ILs were originally included in order to increase concentration of CO$_2$, degradation of NADH could counteract the positive effect and decrease the efficiency of the reaction. There is, therefore, a need to investigate the activity of NADH in ILs to understand the relationship between NADH and ILs. This will contribute to effectively conducting the enzymatic reaction and determining choice of ILs to be applied in the reaction. Additionally, UV-vis spectrophotometer is used for determination of the NADH concentration and yield of formate is calculated based on the amount of NADH consumed, which is defined as method N in this research. In order to find a precise method for detecting directly formate, a new detection method is thereby needed for determination of product (formate) in ILs in situations where NADH is degraded. The objective of present work was therefore to find a suitable method to detect formate in the presence of ILs, and to investigate the activity of NADH in ILs to improve the efficiency of the enzymatic reaction and the potential for application of ILs for CO$_2$ capture.

2. Results and Discussion

2.1. Comparison between CO$_2$ Conversions with and without IL

Determining CO$_2$ conversion by monitoring decrease of UV-absorbance of NADH during reaction (due to conversion of NADH into NAD, method N) overestimated dramatically the actual conversion of
CO₂ provided by measuring formic acid concentrations (method C) (Table 1). The overestimation was much higher when the reaction was performed in buffer (~60 times) as compared to the reaction carried out in a 20/80 (v/v) BmimBF₄/buffer mixture (~8 times). The values of CO₂ conversion provided by the C method are within a typical range for such an enzymatic reaction under thermodynamic equilibrium, which generally has a low conversion rate (0.002 mM·min⁻¹) at the conditions at which the reaction was performed (FDH = 1.3 μg/mL, Tris-buffer, 37 °C) [9,29]. Liu and co-workers reported a higher conversion rate for the same reaction, however the final yield of formate was at the same level (less than 1%) [30]. The higher values of CO₂ conversion by the method N suggested a possible degradation of NADH during reaction, which was further investigated.

<table>
<thead>
<tr>
<th>Entry</th>
<th>IL (V_IL/V_Buffer)</th>
<th>NADH Conv. (%) (SD)</th>
<th>Yield of Formate (%) (SD)</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>Phosphate buffer (Blank)</td>
<td>64.6 (±3.6)</td>
<td>1.1 (±0.5)</td>
</tr>
<tr>
<td>2</td>
<td>BmimBF₄ (20%)</td>
<td>25.4 (±3.2)</td>
<td>2.9 (±0.4)</td>
</tr>
</tbody>
</table>

* Reaction conditions: FDH (3 μL, 0.075 U/mL), NADH (2 μmol, 1 mM), CO₂ (1 bar), buffer (100 mM phosphate, pH = 7); 37 °C, 3 h. Standard deviation (SD).

2.2. Stability of NADH in ILs

In order to evaluate the effect of temperature on NADH degradation, aqueous mixtures of NADH with BmimBF₄ (5.0 equiv.) were monitored over a temperature range of 10–80 °C (Figure 1A). NADH degradation curves were obtained by in-situ UV–vis spectroscopy. A remarkable reduction of NADH was observed at temperatures higher than 40 °C, whereas such reduction was alleviated at lower temperature. At 37 °C, the kinetics approximated to a first order, with a degradation energy Ea of 59.5 kJ mol⁻¹ (Figure S5) and a degradation rate constant of 6.2 × 10⁻³ (min⁻¹). Degradation decreased exponentially with time, with 90% loss of NADH after 4 h (Figure 1B), suggesting that the decrease of the NADH concentration in the presence of FDH was not only due to CO₂ conversion, but also due to spontaneous degradation under the reaction conditions. Degradation of NADH does not only contribute to overestimate the final CO₂ conversion, but also has a direct impact in the actual conversion of CO₂ into formic acid, as it also plays a role as a substrate in the CO₂ reduction. In order to find likely structure-performance relationships and verify whether NADH degradation occurs in a wide range of ionic liquids, degradation was evaluated in five imidazolium-based ILs, including BmimBF₄ (Table 2). Results showed that the stability of NADH was not only dependent on temperature, but also highly dependent on the pH value of the reaction mixture. Under neutral–basic conditions, almost no degradation of NADH was detected (entry 1). In contrast, the concentration of NADH diminished markedly at lower pH values, even under weakly acidic conditions (entry 2). Notably, under strongly acidic conditions, the absorption at 340 nm assigned to NADH disappeared coincidently with the appearance of a blue-shifted peak at 332 nm (entries 3–5 and Figure S8). Finally, when DBULat (known to be a strong basic IL) was employed rather than imidazolium-based ILs (entry 6), only traces of degradation of NADH were observed (non-statistically significant).
2.3 Mechanism Discussion for NADH Degradation

2.3.1 Effect of Acidity on the Degradation of NADH

For better identification of the role of acidity on the stability of NADH during enzymatic conversion, the pH of the phosphate buffer was adjusted within the range 1.0–7.0 by the addition of proper amounts of phosphoric acid. In the absence of FDH, NADH degraded steadily with decreasing pH from 4.5 to 7.0 (Figure S6). At pH 4.5 (Figure 2), complete degradation of NADH was observed after incubation at 37 °C for 3 h. With further reduction of pH to 2.5, a new band appeared at 332 nm, and the intensity of this band gradually increased within the pH range from 2.5 to 4.5 (Figure 2). Such variation of NADH degradation with increasing acidity is consistent with our observations for the aqueous IL system in Table 2. However, the newly formed species referenced at 332 nm was not stable under extreme acid conditions, as indicated by decrease in absorbance at this wavelength over the pH range from 1.0 to 2.0.

<table>
<thead>
<tr>
<th>Entry</th>
<th>IL</th>
<th>Residue_{NADH} (%) (SD)</th>
<th>pH (SD)</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>BmimDCA</td>
<td>100.0 (±0.1)</td>
<td>7.8 (±0.20)</td>
</tr>
<tr>
<td>2</td>
<td>EmimOAc</td>
<td>12.8 (±3.1)</td>
<td>5.3 (±0.12)</td>
</tr>
<tr>
<td>3</td>
<td>EmimBF₄</td>
<td>0 (±0.18)</td>
<td>2.8 (±0.18)</td>
</tr>
<tr>
<td>4</td>
<td>BmimBF₄</td>
<td>0 (±0.18)</td>
<td>2.6 (±0.18)</td>
</tr>
<tr>
<td>5</td>
<td>BmimDMP</td>
<td>0 (±0.16)</td>
<td>2.3 (±0.16)</td>
</tr>
<tr>
<td>6</td>
<td>DBULat</td>
<td>99.7 (±0.6)</td>
<td>9.3 (±0.14)</td>
</tr>
</tbody>
</table>

*General conditions: V_{IL}/V_{H₂O}: 20%; NADH (1 mM); incubation time (3 h); temperature (37 °C). Residue_{NADH} (%) determined by absorption at 340 nm through UV-vis spectrum. Standard deviations (SD).
2.3.2. Proposed Mechanism of NADH Degradation

Next, a buffer solution (2 mL) of BmimBF$_4$ ($V_{IL}/V_{buffer} = 20\%$) and NADH (2 μmol) was prepared separately for in-situ UV-vis detection at $37 \ ^\circ C$. Initially, the presence of two typical peaks assignable to NADH at 340 and 260 nm, respectively, were observed. Upon incubation at $37 \ ^\circ C$ for 3 h, an apparent decrease of absorption intensity at 340 nm and a simultaneous increase of absorption at 260 nm were observed (Figure S7). Such variation was once referred to conversion of NADH to NAD$^+_d$ [31]. However, subsequent $^1H$ NMR studies indicated that a new compound besides NAD$^+_d$ was formed. Typical signals of the 1.4-dihyropyridine unit in NADH (δ: ppm), such as C(1)H$_2$ (2.80, 2.89), C(3)H (6.00) and C(4)H (6.95), were fully diminished (Figure 3, Figure S10, and Table S2). On the other hand, representative C-H signals of pyridinium in NAD$^+_d$ arising from hydride abstraction of NADH were not formed.

![Figure 3. $^1H$ NMR (600 MHz, D$_2$O, 298 K) (δ: ppm) study. (A) BmimBF$_4$; (B) NADH; (C), NAD$^+_d$; (D) a mixture of NADH and BmimBF$_4$ (molar ratio = 1:1); * signals for imidazolium C(sp$^2$)-H.]

DFT calculations were performed to better unravel the mechanisms of NADH degradation in acid conditions [32]. On the basis of previous literature and from our results, α-O$^2$-3B-cyclotetrahydronicotinamide adenine dinucleotide was most likely to be the final product and two potential reaction pathways for such conversion were evaluated (Figure 4) [28]. In pathway 1 (denoted by black curve in the Figure 4), the decomposition of NADH started with protonation of N-contained 6-membered ring at the β-position, with leading to b. Subsequent configuration isomerization from b to d$_1$ occurred through ring opening of O-contained 5-memered ring. Specifically, nucleophilic addition of H$_2$O to the iminium cation in b at the α-position and C-O bond breaking of the O-contained 5-membered ring in the presence of acid affords c. Next, dehydrogenation of c at the α-position of N-contained 6-membered ring readily proceeded to generate d$_1$ denoted as an α configuration, wherein the N-contained heterocycle and hydroxyl group sit at the same side of O-contained 5-memered ring (the β configuration in b represents two substituents sit at the opposite position). Optimization of the position of H$_2$O in the structure of d$_1$ is found favourable in energy. A ternary hydrogen-bonding interaction system -OH/H$_2$O/C=O was found in the relatively stable structure d. Finally, the intra-molecular nucleophilic attack of OH to iminium cation of d affords the product e, coincidently with one equivalent of proton being released. In comparison to pathway 1, configuration isomerization occurred directly in pathway 2 (denoted by a red curve in the Figure 4) through an O-contained ring opening intermediate f. As a result, d$_2$ with α configuration was formed, which is slightly different from d$_1$ in pathway 1 imposed by the interaction mode towards H$_2$O. Subsequently, d$_2$ follows the same route as d$_1$ to eventually provide e. All structures of transitional state for both pathways are also optimized (Table S1). It suggests that the reaction leading to d is more likely following pathway 1 since its energy barrier step (b→c) is 3.1 kJ/mol lower as compared with that (f→d$_2$) in pathway 2.
2.4. Enzymatic Reaction in BmimBF₄ and BmimDCA

2.4.1. Effect of NADH Degradation in BmimBF₄ and BmimDCA

pH was found to be the most crucial factor leading to NADH degradation. However, variation of pH value cannot be avoided given the intrinsic nature of the reaction. For instance, both the absorption of CO₂ into the aqueous solution and the formation of formic acid will result in a more acidic environment [33]. Interestingly, the results indicated that with CO₂, NADH degradation is significantly prevented in the presence of contained-BmimBF₄ buffer (Table 1, method N). To clarify and rationally understand this phenomenon, the influence of CO₂ absorption with and without BmimBF₄ on NADH degradation and pH variation were evaluated and compared. Experiments of NADH degradation were conducted in an aqueous solution with a range of volume ratio of BmimBF₄ to buffer in the presence or absence of CO₂ (Figure 5A). The degradation concentration-course plot (in black) in the case of without CO₂ offers a downward parabola. It demonstrated that the maximum degradation of NADH in a combination of BmimBF₄ and buffer is 22% when a volume ratio of 40% (V_IL/V_buffer) is reached. The in-situ pH evaluation of the solution (Figure 5B) indicated this degradation follows a function of pH value variation, with maximum degradation appearing at the lowest pH value. The pH value variation was attributed to the hydrolysis of BmimBF₄, which may occur with a different extent under these conditions [34]. On the other hand, a significant degradation of NADH was detected in the buffer as soon as CO₂ (1 bar) was introduced (Figure 5A, plot in red). Interestingly, such a degradation was gradually relieved with loading BmimBF₄ from 0 to 20% (V_IL/V_buffer). The stabilization has been attributed to the formation of ternary aggregates (CO₂/IL/H₂O), wherein the amphipathic BmimBF₄ could self-assemble at the interface between the nonpolar CO₂ and the aqueous phase [35]. In this manner, formation of carbonic acid was therefore avoided. Specifically, both hydrolysis of BmimBF₄ and the formation of carbonic acid will be inhibited when a higher ratio of V_IL/V_buffer was launched, with less NADH degradation.

In contrast, NADH is much more stable in the presence of a weakly basic BmimDCA (Figure 6A, plot in black), with less than 10% degradation of NADH occurring at maximum. The pH value of the solution progressively increased when the ratio of BmimDCA was increased (Figure 6B). A similar stabilization of NADH with IL functioning as the surface-active molecule was also observed when CO₂ was introduced (Figure 6A, plot in red).
proteins change dramatically upon increasing concentrations of ILs. Indeed, in pure ILs, enzymes can hardly be dissolved in homogenous phase without denaturation. In this enzymatic reaction, increasing concentration of IL to absorb CO₂ was detected in the buffer as soon as CO₂ was introduced (Figure 5A, plot in red). Interestingly, the results indicated that with CO₂ in contained-BmimBF₄ buffer for 3 h at 37 °C; NADH (1.4 mg, 2 μmol); CO₂ (1 bar); Buffer (2 mL).

To clarify, it is well known that water with small amounts of salts is considered to be the best media for proteins. However, high concentration of ILs (salts) can cause conformational changes of peptide chains that can result in enzyme denaturation [36]. Crystallisation and aggregation behaviour of proteins change dramatically upon increasing concentrations of ILs. Indeed, in pure ILs, enzymes can hardly be dissolved in homogenous phase without denaturation. In this enzymatic reaction, increasing concentration of IL to absorb CO₂ is expected to influence negatively the activity of the enzyme. This can explain that CO₂ conversion achieved the highest value when only a 20% of BmimBF₄ was used.

Figure 5. Effect of CO₂ on the degradation of NADH in BmimBF₄. (A) Degradation of NADH with CO₂ and without CO₂; (B) pH of solution at different concentrations of BmimBF₄; General conditions: NADH in contained-BmimBF₄ buffer for 3 h at 37 °C; NADH (1.4 mg, 2 μmol); CO₂ (1 bar); Buffer (2 mL).

Figure 6. Effect of CO₂ on degradation of NADH in BmimDCA. (A) Degradation of NADH with CO₂ and without CO₂; (B) pH of solution at different concentrations of BmimDCA; General conditions: NADH in contained-BmimDCA buffer for 3 h at 37 °C; NADH (1.4 mg, 2 μmol); CO₂ (1 bar); Buffer (2 mL).

2.4.2. Enzymatic Reaction in BmimBF₄ and BmimDCA

Enzymatic conversion of CO₂ to formate in phosphate buffer systems (IL/buffer) consisting of two alternative ILs at different ratios was subsequently conducted (Table 3). As indicated by experiments of NADH degradation (Figure 5A), it will lead to not only the formation of a ternary aggregates (CO₂/IL/H₂O) but also hydrolysis of BmimBF₄ when the concentration of BmimBF₄ was increased within a range (V_IL/V_buffer < 50%), which actually has an opposite effect towards the degradation of NADH. Therefore, taking these premises into consideration, to find a proper concentration of BmimBF₄ is required to attain an optimized conversion with least NADH being degraded (entry 3). In contrast to the acidic BmimBF₄, NADH was more stable in the presence of basic BmimDCA. And the conversion of NADH monotonously decreased when the concentration of BmimDCA was increased (entries 6–9), but the yield of formate increased with the concentration of BmimDCA from 10% to 40%, and dropped to the lowest at 60% BmimDCA.

<table>
<thead>
<tr>
<th>Entry</th>
<th>BmimBF₄ (%)</th>
<th>BmimDCA (%)</th>
<th>V_IL/V_buffer</th>
<th>pH</th>
<th>Yield of Formate (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>10</td>
<td>90</td>
<td>0.1</td>
<td>6.7 (±0.08)</td>
<td>34.0 (±3.9)</td>
</tr>
<tr>
<td>2</td>
<td>10</td>
<td>90</td>
<td>0.2</td>
<td>6.4 (±0.14)</td>
<td>34.0 (±3.9)</td>
</tr>
<tr>
<td>3</td>
<td>10</td>
<td>90</td>
<td>0.4</td>
<td>6.2 (±0.18)</td>
<td>34.0 (±3.9)</td>
</tr>
</tbody>
</table>

It is well known that water with small amounts of salts is considered to be the best media for proteins.
the phosphate buffer. Indeed, the yield of methanol when the reaction was conducted in 20% BmimBF₄ was two times higher than in the phosphate buffer. Therefore, production of formaldehyde is prevented, as the second sequential reaction requires a threshold concentration of formic acid to be activated [9]. Indeed, formic acid oxidation was 30 times faster than CO₂ reduction catalyzed by FDH [29]. For the second enzyme, FaldDH, the reaction (formic acid → formaldehyde) was also found to be less efficient than the reverse reaction (formaldehyde → formic acid). However, for the third enzyme, ADH, the forward reaction (formaldehyde → methanol) is much more favorable than the reverse reaction (methanol → formaldehyde). Therefore, to make this multi-enzymatic reaction efficient, formic acid and formaldehyde have to be consumed by the next sequential reaction, in order to shift the equilibrium of the cascade reaction and eventually achieving high conversion of CO₂ to methanol (Table 4, entry 1).

Another strategy to further increase conversion of CO₂ to methanol is to enhance CO₂ solubility, as low CO₂ solubility in water would also result in low formic acid concentration. As a consequence, production of formaldehyde is prevented, as the second sequential reaction requires a threshold concentration of formic acid to be activated [24]. Both factors (unfavourable equilibrium rates and need of a minimum concentration threshold) make the first reaction of the sequence (the one studied here) to play a decisive role on conversion of CO₂ to methanol. Addition of BmimBF₄ helps enhance the conversion in the first reaction by both stabilizing NADH and also increasing CO₂ solubility. Indeed, the yield of methanol when the reaction was conducted in 20% BmimBF₄ was two times higher than in the phosphate buffer. Other authors followed other strategies to increase conversion. For example, Ober and Dave immobilized the three enzymes in silica sol-gel matrixes, confining and reducing the volume of the

### Table 3. Enzymatic Conversion of CO₂ to Formate

<table>
<thead>
<tr>
<th>Entry</th>
<th>Solvent (v/v)</th>
<th>NADH Conv. (%) (SD)</th>
<th>Yield of Formate (%) (SD)</th>
<th>pH (SD)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Method N</td>
<td>Method C</td>
<td>Before (±)</td>
<td>After (±)</td>
</tr>
<tr>
<td>1</td>
<td>Phosphate buffer</td>
<td>64.6 (±3.6)</td>
<td>1.1 (±0.5)</td>
<td>7.0 (±0.04)</td>
</tr>
<tr>
<td>2</td>
<td>10% BmimBF₄</td>
<td>46.8 (±2.3)</td>
<td>0.2 (±0.3)</td>
<td>6.7 (±0.08)</td>
</tr>
<tr>
<td>3</td>
<td>20% BmimBF₄</td>
<td>25.4 (±3.2)</td>
<td>2.9 (±0.4)</td>
<td>6.5 (±0.06)</td>
</tr>
<tr>
<td>4</td>
<td>40% BmimBF₄</td>
<td>36.3 (±2.0)</td>
<td>2.3 (±0.4)</td>
<td>6.0 (±0.05)</td>
</tr>
<tr>
<td>5</td>
<td>60% BmimBF₄</td>
<td>24.6 (±3.0)</td>
<td>0.6 (±0.3)</td>
<td>5.9 (±0.05)</td>
</tr>
<tr>
<td>6</td>
<td>10% BmimDCA</td>
<td>34.0 (±3.9)</td>
<td>0.4 (±0.2)</td>
<td>7.0 (±0.04)</td>
</tr>
<tr>
<td>7</td>
<td>20% BmimDCA</td>
<td>16.2 (±1.3)</td>
<td>1.5 (±0.5)</td>
<td>7.2 (±0.06)</td>
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<tr>
<td>8</td>
<td>40% BmimDCA</td>
<td>4.9 (±2.8)</td>
<td>1.8 (±0.5)</td>
<td>7.5 (±0.08)</td>
</tr>
<tr>
<td>9</td>
<td>60% BmimDCA</td>
<td>1.6 (±1.4)</td>
<td>0.2 (±0.2)</td>
<td>7.8 (±0.09)</td>
</tr>
</tbody>
</table>

*a Reaction conditions: FDH (3 μL, 0.075 U/mL), NADH (2 μmol, 1 mM), CO₂ (1 bar), buffer (100 mM (Na₂HPO₄·NaH₂PO₄ = 39:61 (mol/mol)), 2 mM), 37 °C, 3 h. **Enzymatic reaction was repeated three times and standard deviations (SD) were calculated that presented in the Table 3. pH value of reaction mixture was added before and after saturating CO₂.

### Table 4. Enzymatic Conversion of CO₂ to Methanol

<table>
<thead>
<tr>
<th>Entry</th>
<th>Solvent (v/v)</th>
<th>Yield of Methanol (%) (SD)</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>Phosphate buffer</td>
<td>24.3 (±1.2)</td>
</tr>
<tr>
<td>2</td>
<td>BmimBF₄ (10%)</td>
<td>20.2 (±1.9)</td>
</tr>
<tr>
<td>3</td>
<td>BmimBF₄ (20%)</td>
<td>67.1 (±2.4)</td>
</tr>
<tr>
<td>4</td>
<td>BmimBF₄ (40%)</td>
<td>46.3 (±2.1)</td>
</tr>
<tr>
<td>5</td>
<td>BmimBF₄ (60%)</td>
<td>15.2 (±2.6)</td>
</tr>
</tbody>
</table>

*b Reaction conditions: FDH (3 μL, 0.075 U/mL), FaldDH (0.2 mg, 0.1 U/mL), ADH (0.2 mg, 30 U/mL), NADH (2 μmol, 1 mM), CO₂ (1 bar), buffer (100 mM phosphate, pH = 7); 37 °C, 3 h. Standard deviation (SD).
enzymes, in such a manner that the local concentration of reactants was enhanced [8]. Jiang followed a similar strategy, and was able to increase the yield of methanol up to 71.6% [24]. These studies also confirmed that breaking the unfavourable equilibrium by switching it to the right is a good manner to enhance the overall conversion.

3. Materials and Methods

3.1. Materials

Formate dehydrogenase (EC 1.2.1.2, homo-dimer, 76 kDa) from Candida boidinii (FDH), formaldehyde dehydrogenase (EC 1.2.1.46, homo-dimer, 150 kDa) from Pseudomonas sp. (FaldDH), alcohol dehydrogenase (EC 1.1.1.1, homo-tetramer, 141 kDa) from Saccharomyces cerevisiae (ADH), β-Nicotinamide adenine dinucleotide reduced form (NADH, >97 wt %), and β-Nicotinamide adenine dinucleotide hydrate (NAD⁺, >98 wt %) were purchased from Sigma Aldrich. Formate, sodium phosphates, citric acid, acetamide, isopropanol, sodium acetate, and acetic anhydride and were supplied by Sinopharm chemical agent (Shanghai, China). CO₂ gas (>99.99%) in a cylinder was supplied by Beijing Beiwen Gas Factory. 1-butyl-3-methylimidazolium tetrafluoroborate (BmimBF₄), 1-ethyl-3-methylimidazolium tetrafluoroborate (EmimBF₄), 1-ethyl-3-methylimidazolium acetate (EmimOAc), 1-butyl-3-methylimidazolium dicyanamide (BmimDCA), and 1-butyl-3-methylimidazolium dimethylphosphate (BmimDMP) (purity > 99.0 wt %) were purchased from Lanzhou Institute of Chemical Physics (Lanzhou, China). 1,8-Diazabicyclo[5.4.0]undec-7-ene lactate (DBULat) (purity > 98.0 wt %) were purchased from Linzhou Keneng material technology corporation. Ultra-pure water was obtained using a water purification system (Milli-Q Direct 8).

3.2. General Procedure for Enzymatic Conversion of CO₂ to Formate

The enzymatic reaction was conducted in a stainless steel reactor (25 mL) equipped with two valves and a pressure gauge. In the reactor, a buffer solution of FDH (3 µL) and NADH (1 mM) was bubbled with CO₂ for 5 min to remove the residual air, after which the pressure was adjusted to 1 bar. The reactor was positioned in water bath at 37 °C for 3 h. Samples were then taken from the reactor for analysis. Buffer (100 mM (Na₂HPO₄:NaH₂PO₄ = 39:61 (mol/mol)), pH = 7, 2 mL); Enzymatic reaction was repeated three times; standard deviations (SD) were calculated and presented in the Table 1.

3.3. Kinetic Degradation of NADH in BmimBF₄

Solution of NADH and BmimBF₄ was prepared in deionized water (2 mL) at concentration of 1 mM (1.4 mg, 2 µmol) and 5 mM (2 µL, 5.0 equiv.), respectively. The reaction was performed over a temperature range of 10–80 °C during which sample was measured every ten degree centigrade and kept for ten minutes. Additionally, another experiment was carried out at 37 °C for 9 h and the preparation of sample was same as above during which samples were measured every ten minutes (Figure 1B). Temperature (37 °C) in such experiment was referring to enzymatic reaction and the variation of NADH concentration with time was monitored by method N.

3.4. Stability of NADH in Aqueous ILs

In the reactor, a solution (2 mL) of 20 vol % ILs and 80 vol % deionized water was prepared to dissolve 2 µmol NADH (1 mM). Then, the reactor was positioned in water bath at 37 °C for 3 h. The sample was taken from the reactor and detected at 340 nm by an in-situ UV–vis spectroscopy (Table 2).

3.5. Effect of Acidity on the Degradation of NADH

Phosphate buffers in the pH range of 1.0–7.0 were separately prepared, namely, pH 7, 6.5, 5.0, 4.5, 4.0, 3.5, 3.0, 2.5, 2.0, 1.5, and 1.0, with addition of varied amounts of purity phosphoric acid in
phosphate solution (pH = 7). NADH (1 mM) in these buffers was separately incubated at 37 °C for 3 h. Residue NADH is measured at 340 nm by an in-situ UV–vis spectroscopy (Figures 2 and S6).

3.6. NMR for NADH Degradation in BmimBF$_4$

Equimolar of NADH and BmimBF$_4$ was mixed in deuter oxide. The mixture was heated at 50 °C for 3 h and then the product was purified through extraction of methylene dichloride. After removing methylene dichloride in a vacuum oven at 50 °C for 3 h, solid powder was obtained and detected by NMR.

3.7. Effect of NADH Degradation in BmimBF$_4$ and BmimDCA

Solutions (2 mL) of BmimBF$_4$ with phosphate buffer in different concentrations were prepared, namely, 0%, 5%, 10%, 15%, 20%, 30%, 40%, and 80% (V$_{IL}$/V$_{buffer}$). In the reactor, NADH (1 mM) in the prepared solution was bubbled with CO$_2$ for 5 min to remove the residual air, after which the pressure was adjusted to 1 bar. The reactor was positioned in water bath at 37 °C for 3 h. Without CO$_2$, it is no need to press CO$_2$ in the reactor. In the same way, solutions (2 mL) of BmimDCA with phosphate buffer in different concentrations were prepared, namely, 0%, 10%, 20%, 30%, 40%, and 60% (V$_{IL}$/V$_{buffer}$). In the reactor, NADH (1 mM) in the solution was bubbled with CO$_2$ for 5 min to remove the residual air, after which the pressure was adjusted to 1 bar. The reactor was positioned in water bath at 37 °C for 3 h. Furthermore, pH value of each solution was measured by pH meter (298.5 K, Mettler Toledo FE20, Mettler Toledo, Zurich, Zurich State, Switzerland) before starting reaction.

3.8. Enzymatic Reaction in BmimBF$_4$ and BmimDCA

Solutions (2 mL) of BmimBF$_4$ with phosphate buffer in different concentrations were prepared, namely, 0%, 10%, 40%, and 60% (V$_{IL}$/V$_{buffer}$). Solutions (2 mL) of BmimBF$_4$ with phosphate buffer were prepared in the same way. Procedure of enzymatic reaction is as same as described above (General procedure).

3.9. Analytical Method

Two methods, NADH (named as N) and colorimetric detection (named as C), were evaluated to determine the yield of formate product. The details of these methods are as follows:

3.10. Method N for Determination Concentration of NADH

Method N is an indirect method where the yield of formate is determined through quantifying reduction of NADH. It can be easily monitored by UV–vis spectroscopy at 340 nm referenced to NADH [23]. The assumption is that formation of formate during the enzyme-catalysed process would consume an equal-molar amount of NADH. The concentration of NADH was quantified by absorbance measurements, which was monitored at 340 nm by an in-situ UV–vis spectroscopy. After the enzymatic conversion to formate was completed, 0.5 mL of sample was taken and diluted with 1.5 mL of water. After dilution of NADH solution, absorbance at 340 nm is below 2 that is more precise. Then 2 mL of the solution was pipetted into a 10 mm quartz cell. The photometric measurement was monitored at 340 nm using a UV–vis spectrophotometer (Shimadzu UV, 2550 spectrophotometer Shimadzu Corporation, Tokyo, Tokyo metropolitan, Japan). The standard calibration curve is presented in Figure S1.

3.11. Method C for Determination Concentration of Formate

Method C was originally used for detecting formate in products from fermentation [37]. The assumption is that a complex compound, recorded at 515 nm using UV–vis spectroscopy, would be formed by the reaction of formate with a mixture of citric acid, isopropanol, acetic anhydride, sodium acetate in an appropriate ratio [38]. The yield of formate is then directly determined. First, 3.5 mL of 100% acetic anhydride, 50 μL of 30% (v/v) sodium acetate, and 1 mL of 2-propanol solution
containing 0.5% (w/v) citric acid and 10% (w/v) acetamide were added in a 10 mL vial. Then 0.5 mL of sample was added to this assay solution and incubated for 90 min at 25 °C. Finally, the absorbance was determined at 515 nm using a UV–vis spectrophotometer. The formate standard calibration curve was prepared against the appropriate concentrations of formate (Figures S2–S4). 500 µL of sample was used for the colorimetric assay.

3.12. GC for Determination Concentration of Methanol

A Hewlett Packard HP6890 gas chromatograph (Hewlett-Packard Company, Palo Alto, CA, USA) (GC) equipped with a FID (250 °C) and a Restek XTI-5 column (30 m × 0.25 mm i.d., film thickness 0.25 mm) was used for methanol concentration by using ethyl acetate as an internal standard. The carrier gas was N₂ with a flow rate of 0.4 mL min⁻¹. The injector temperature was 150 °C and the injection volume was 1 mL. Methanol GC chromatograms were calibrated with 0.01–1 mM methanol solution in 0.1 M pH 7.0 phosphate buffer.

3.13. ¹H Nuclear Magnetic Resonance Measurements

Spectra were recorded at 298 K on a Bruker av-600 MHz spectrometer (Bruker, Bilka, MA, USA) operating at 600 MHz. Deuterium oxide was as solvent.

4. Conclusions

In summary, we demonstrated that NADH degradation is unavoidable due to the presence of acid gas during enzymatic CO₂ conversion, which impedes the CO₂ conversion and overestimates the conversion when the absorbance of NADH is used as a method of detection. The mechanism of NADH degradation was investigated by UV, NMR, and DFT calculation and all methods suggested that the occurrence of acid in the reaction mixture is the main contributing factor to the degradation observed. By selecting neutral–basic ionic liquids and adjusting the concentration of ionic liquids in the buffer, stabilization of the cofactor (NADH) can be achieved, along with enabling a higher CO₂ concentration in the buffer. Finally, CO₂ conversion was more than twice as compared with the conversion reached by the enzymatic reaction in a phosphate buffer (traditional buffer). This study is a significant contribution to the use of enzymes like formate dehydrogenase in ionic liquids and paves the way for improving biocatalysts using ionic liquids.

Supplementary Materials: The following are available online at http://www.mdpi.com/2073-4344/8/8/304/s1, Figure S1. Standard calibration curve for NADH method. Figure S2. Standard calibration of formate for colorimetric method in buffer. Figure S3. Standard calibration curve of formate for colorimetric method in BmimBF₄. Figure S4. Standard calibration curve of formate for colorimetric method in BmimDCA. Figure S5. Arrhenius plots for degradation of NADH in BmimBF₄ as temperature increases. Figure S6. NADH degrades in phosphate buffer whose pH in range from 7.0 to 4.5. Figure S7. NADH degrades in contained-BmimBF₄ reaction. Figure S8. NADH degrades in ILs. Figure S9. Two possible pathway (red and black) for NADH degradation mechanism proposed by the Norman and co-worker. Figure S10. ¹H NMR (600 MHz, D₂O, 298K) (δ ppm). (a) BmimBF₄; (b) NADH; (c), NAD⁺; (d) a mixture of NADH and BmimBF₄ (molar ratio = 1:1); D₂O was used as the solvent. Table S1 Structure for the NADH and its derivatives. Table S2. Summary of chemicals NMR data.

Author Contributions: Z.Z., B.-h.X. and J.L. conceived and designed the experiment, performed the experiments, and analysed the data. M.P., N.V.S. and S.Z. wrote the paper and reviewed drafts of the paper. H.H. and Y.Z. conceived and designed the simulation experiment, analysed the data and reviewed drafts of the paper.

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