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# Supporting Information

# Ionozyme: ionic liquids as solvent and stabilizer for efficient bioactivation of CO<sub>2</sub>

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Table S2. Full name and structure of IL cations

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### **Supplementary Figures**



**Figure S1.** Phylogenetic analysis of the FDHs. Phylogenetic tree of the top one ranking FDH protein sequences in 53 unique PPR groups and the well characterized FDH protein sequences.



Figure S2. Peptides in the region of NADH binding sites and active site for PPR group 1.



Figure S3. Sequences alignment of the predicted FDHs after the PPR prediction and phylogenetic analysis. The possible NADH



binding sites and active sites are boxed in green and red color, respectively.

**Figure S4.** SDS-PAGE profile for the selected FDHs. Lane 1 M (marker), lane 2 CK (control check), lane 3 FDHFu from fungal sp. No.11243, lane 4 FDHPa from *Paracoccus* sp., lane 5 FDHSn from *Starkeya novella*.



**Figure S5.** Biocatalytic performance of the recombinant FDHs. Enzyme activities of FDHCb and FDHPa with or without DTT for CO<sub>2</sub> conversion at pH 7.0, 25 °C.



**Figure S6.** Effect of pH and temperature on enzyme activity of FDHPa. (A) Relative activity as a function of pH. (B) Relative activity observed as a function of temperature. Enzyme assays were carried out in the presence of NaHCO<sub>3</sub> (50 mM). The optimum pH for recombinant FDHPa activity was determined by incubating the reaction mixture at 25 °C, while the optimum temperature was at pH 7.0 with PB (100 mM). Activities at the optimal temperature and pH were defined as 100%. Each value represented the mean of triplicate measurements.



**Figure S7.** Screening of ILs. (A) Prediction of CO<sub>2</sub> solubility by COSMO-RS model. (B) Relative activity of FDHPa in 20 different kinds of biocompatible ILs (0.05%). Activity in [TNGH][PhO] were defined as 100%. Each value represented the mean of triplicate measurements.



**Figure S8.** σ-Profiles prediction by COSMO-RS. σ-Profiles of [TMGH][Pyrr] (black line), [CH][Gly] (red line), [TMGH][Phe] (blue line), [TMGH][Im] (Cyan line), [CH][Ole] (Olive line) and CO<sub>2</sub> (yellow line).



**Figure S9.** Optimization of reaction conditions for the ionozymes. (A) FDHPa concentration effect. (B) ILs ([CH][Glu], [CH][Arg] and [CH][Lys]) concentration effect. (C) NADH concentration effect on ionozyme activities. Enzyme assays were carried out under standard conditions in the presence of CO<sub>2</sub>. Activities at the optimal concentration were defined as 100%. Each value represented the mean of triplicate measurements.



**Figure S10.** Analysis of determining conditions on catalytic activity of FDHPa and FDHPa-ILs. (A) Kinetic analysis based on Michaelis–Menten equation. (B) Far-UV CD spectra. (C) Thermal stability.



Figure S11. Root mean square deviation (RMSD) of FDHPa and FDHPa-ILs. All stimulations were carried out at 25 °C.



Figure S12. Structural comparison of FDHPa (red) and FDHCb (blue).



Figure S13. Calibration Curve. Absorbance of FDHPa at 562 nm as a function of concentration.



Figure S14. Calibration Curve. Absorbance of NADH at 340 nm as a function of concentration.



**Figure S15.** Proposed hydride transfer process from NADH to CO<sub>2</sub> in ionozymes. The binding models of NADH and CO<sub>2</sub> in the active sites of FDHPa-[CH][Pro] and FDHPa-[TMGH][PhO].

## **Supplementary Tables**

lonozymes	K <sub>m</sub> (mM)	K <sub>cat</sub> (s <sup>-1</sup> )	V <sub>max</sub> (U mg <sup>-1</sup> )	$K_{cat}/K_{m}$ (s <sup>-1</sup> mM <sup>-1</sup> )
FDHPa	20.14	0.1839	1148	0.0091
FDHPa-[CH][Pro]	40.61	0.2244	1484	0.0055
FDHPa-[TMGH][PhO]	25.35	0.2724	1812	0.0107
FDHPa-[CH][Pro]-[TMGH][PhO]	15.74	0.2595	1782	0.0165

 Table S1. Kinetic analysis of ionozymes based on Michaelis–Menten equation.

#### Table S2. Full name and structure of IL cations.

No.	Full name	Abbreviation	Structure	COSMO sigma surface
1	Choline	СН	N <sup>+</sup> OH	
2	Tetramethylgunidinium	TMGH	$H_2N^+$ -N	

Table S3.	. Full	name	and	structure	of	IL	anions.
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No.	Full name	Abbreviation	Structure	COSMO sigma surface
1	∟-glycinate	Gly	$H_2N \int_0^0$	
2	∟-alaninate	Ala	O ↓↓O NH₂	8
3	∟-valinate	Val	↓ ↓ o- NH₂	<b>é</b>
4	∟-leucinate	Leu	↓ ↓ ↓ O <sup>-</sup>	
5	∟-isoleucinate	lso		٠
6	∟-phenylalaninate	Phe	<sup>№</sup> H <sub>2</sub> 0	
7	∟-prolinate	Pro	√ N O <sup>−</sup>	
8	∟-lysinate	Lys	$H_2N$ $M_2$ $O^-$	
9	∟-argininate	Arg		<b>***</b>
10	∟-histidinate	His		<b>()</b>
11	∟-tryptophan	Try	······································	
12	∟-serinate	Ser		
13	∟-tyrosinate	Tyr	HO NH <sub>2</sub> O	
14	∟-cysteinate	Cys		
15	Oleic acid	Ole	,,,,,,,,	
16	Dihydrogen phosphate	H <sub>2</sub> PO <sub>4</sub>	H <sub>2</sub> PO <sub>4</sub>	4
17	∟-glutamate	Glu		<b>(</b> )
18	Pyrrole	Pyrr	<b>N</b>	
19	Imidazole	Im		0
20	Phenol	PhO	o c	