## Supporting Information

for

## Role of Imidazolium Cation on Structure and Activity of Candida

## antarctica Lipase B Enzyme in Ionic Liquids

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Table S1. Simulation setup

| Solvent | Components | Molecular Volume $\left(\AA^{3}\right)$ | Volume <br> Ratio <br> $\mathrm{V}_{\mathrm{a}} / \mathrm{V}_{\mathrm{c}}$ | Volume of Simulation Box ( $\AA^{3}$ ) | $\begin{gathered} \hline \text { Density } \\ \text { from } \\ \text { MD } \\ (\mathrm{g} / \mathrm{ml}) \\ \hline \end{gathered}$ | Density from experiment (g/ml) |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| [Emim][TfO] | [Emim] ${ }^{+}$ | 234.25 | 0.71 | 192,151.21 | 1.39 | $1.39^{1}$ |
|  | [TfO] ${ }^{-}$ | 165.70 |  | $( \pm 121.28)$ |  |  |
| [Bmim][TfO] | [Bmim] ${ }^{+}$ | 296.77 | 0.56 | 205,552.22 | 1.32 | $1.30^{2}$ |
|  | [TfO] ${ }^{-}$ | 165.70 |  | $( \pm 109.61)$ |  |  |
| [Hmim][TfO] | $\left[^{\text {Hmim] }}{ }^{+}\right.$ | 360.29 | 0.46 | 251,349.19 | 1.25 | $1.20{ }^{1}$ |
|  | [TfO] ${ }^{-}$ | 165.70 |  | ( $\pm 204.87$ ) |  |  |
| [Omim][TfO] | [Omim] $^{+}$ | 423.70 | 0.39 | 277,985.43 | 1.21 | $1.19^{3}$ |
|  | [TfO] ${ }^{-}$ | 165.70 |  | $( \pm 83.18)$ |  |  |



Figure S1. Root mean square deviation (RMSD) of CALB solvated in [Emim][TfO] (black), [Bmim][TfO] (Red), [Hmim][TfO] (Dark green) and [Omim][TfO] (Blue)


Figure S2. Radial distribution function of carbon atom in imidazolium cation. [Emim] ${ }^{+}$(black), $[\mathrm{Bmim}]^{+}$(red), $[\mathrm{Hmim}]^{+}$(green), and [Omim $]^{+}$(blue). Second solvation shell (7 $\dot{\mathrm{A}}$ ) is used for estimation of coordination number.

Table S2. Reaction rate of butyl acetate synthesis reaction using CALB in ILs

| Solvents | Reaction Rate <br> $(\mathrm{mol} / \mathrm{L} \cdot \mathrm{hr})$ |
| :---: | :---: |
| $[$ Emim $][\mathrm{TfO}]$ | 0.20 |
| $[$ Bmim $][\mathrm{TfO}]$ | 0.85 |
| $[\mathrm{Hmim}][\mathrm{TfO}]$ | 0.77 |
| $[\mathrm{Omim}][\mathrm{TfO}]$ | 0.74 |

Table S3. Secondary structure of $\alpha-10$ helix (residue 285 to 287) in ILs

| CALB in | Residue | $3-10$ <br> Helix | Alpha <br> Helix | Turn |
| :---: | :---: | :---: | :---: | :---: |
|  | 285 ILE | 1.469 | 83.668 | 13.756 |
|  | 286 VAL | 6.258 | 62.206 | 25.974 |
|  | 287 ALA | 6.312 | 27.957 | 42.023 |
|  | 285 ILE | 0.043 | 99.887 | 0.370 |
| $[B m i m][T f O]$ | 286 VAL | 0.037 | 99.196 | 0.764 |
|  | 287 ALA | 0.086 | 80.785 | 11.383 |
|  | 285 ILE | 0.697 | 99.141 | 0.162 |
| $[H m i m][T f O]$ | 286 VAL | 0.025 | 98.581 | 1.394 |
|  | 287 ALA | 0.019 | 90.414 | 5.123 |
|  | 285 ILE | 0.021 | 95.418 | 4.540 |
| $[$ Omim][TfO] | 286 VAL | 0.127 | 94.087 | 5.744 |
|  | 287 ALA | 0.127 | 70.626 | 17.115 |

Each percentage value was calculated in consecutive frame of 30 ns simulations

## References

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(3) S. H. Ha, N. L. Mai and Y.M. Koo, Process. Biochem., 45, 1899 (2010).

