

Supplementary List

Figure S1. Lineweaver-Burk plots in presence of other four alkylimidazolium chlorides

(a) [Bmim]Cl, (b) [Hmim]Cl, (c) [Omim]Cl, (d) [Dmim]Cl

Figure S2. Binding of syringaldazine, 2,6-DMP and guaiacol with the T₁ Cu active pocket of *Mth* laccase

(a) syringaldazine, (b) 2,6-DMP, (c) guaiacol

Figure S3. Scavenging of 2,6-DMP oxidative radicals in presence of [Bmim]Cl

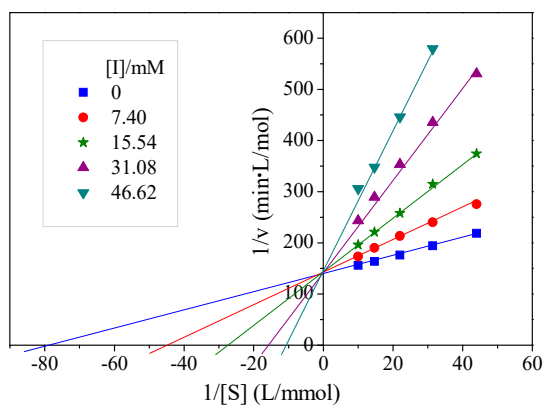
Figure S4. Change of media pH by alkylimidazolium ILs

(Note: Buffer concentration 30 mM)

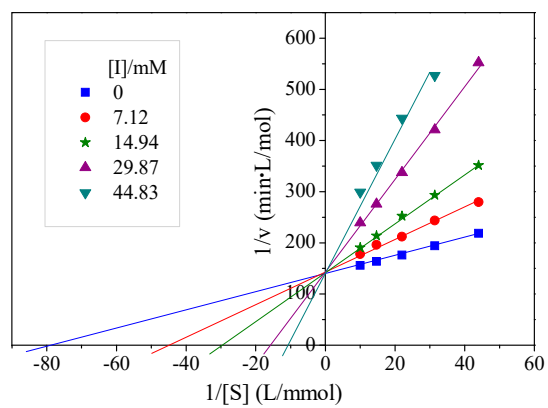
Figure S5. Homology modeling of *Myceliophthora thermophila* laccase

(a) Amino acids sequences, (b) 3D model, (c) QMEAN Z-score, (d) Ramachandran Plot

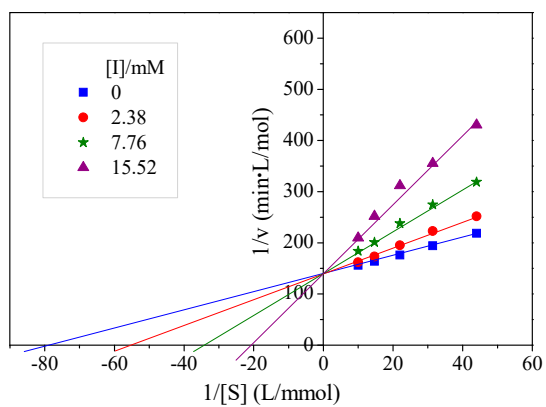
Figure S6. Plots of $\ln[A_0]-\ln[A_1]$ vs. radical scavenging time



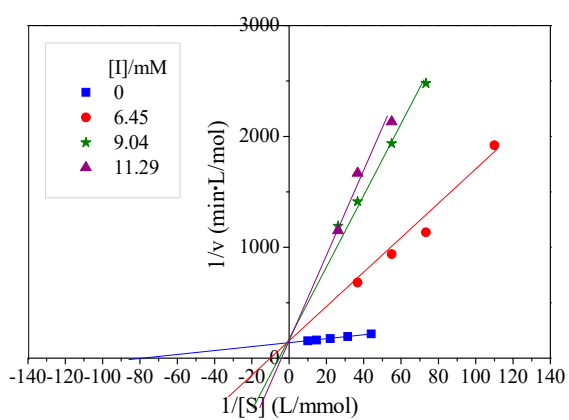
(a) [Bmim]Cl



(b) [Hmim]Cl

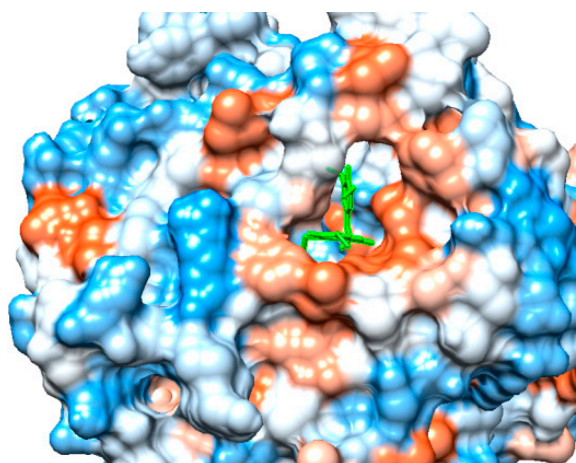


(c) [Omim]Cl

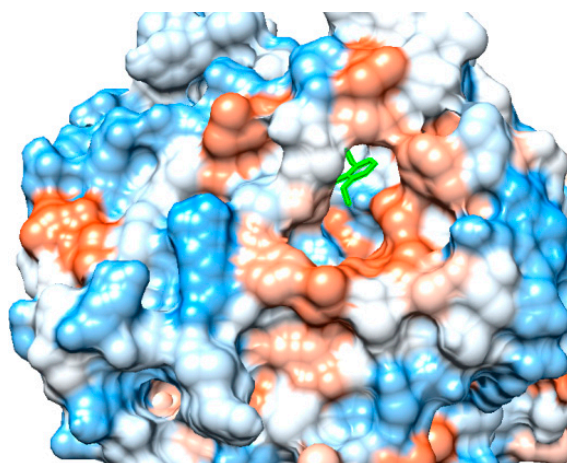


(d) [Dmim]Cl

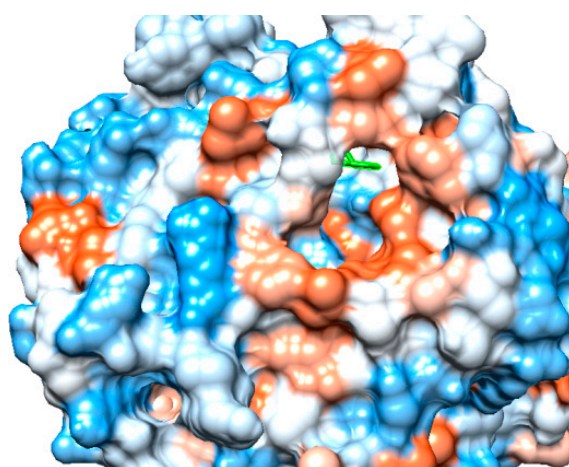
Figure S1 Lineweaver-Burk plots in presence of other four alkyimidazolium chlorides.



(a) Syringaldazine



(b) 2,6-DMP



(c) Guaiacol

Figure S2. Binding of syringaldazine, 2,6-DMP and guaiacol with the T₁ Cu active pocket of *Mth* laccase

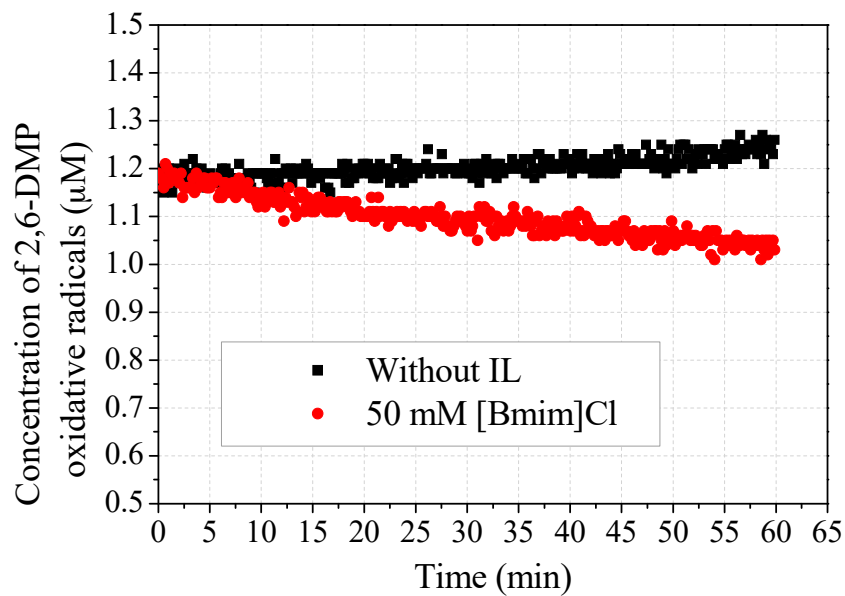


Figure S3. Scavenging of 2,6-DMP oxidative radicals in presence of [Bmim]Cl.

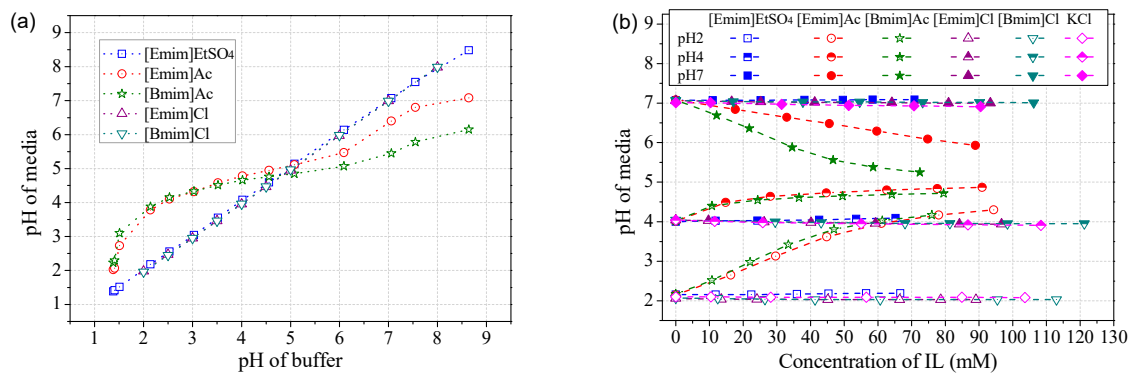


Figure S4. Change of media pH by different alkyimidazolium ILs (Note: Buffer concentration 30 mM).

Myceliophthora thermophila
(*Mth*) laccase

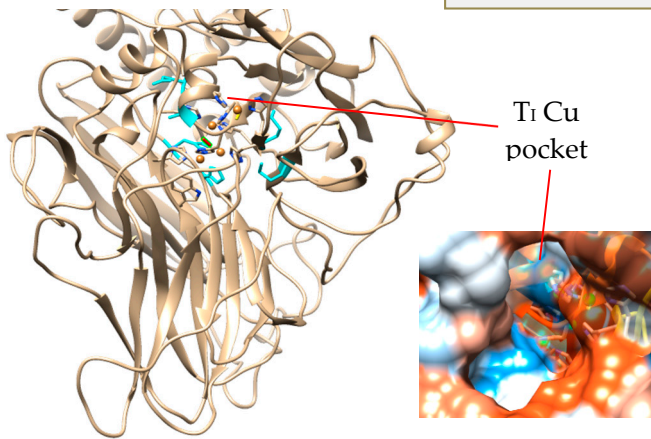
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Melanocarpus albomyces
(*Mal*) laccase

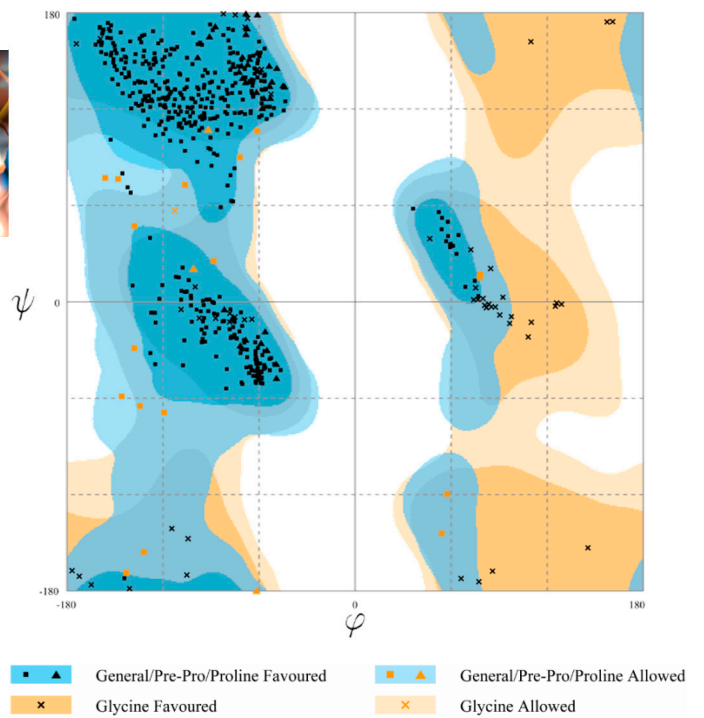
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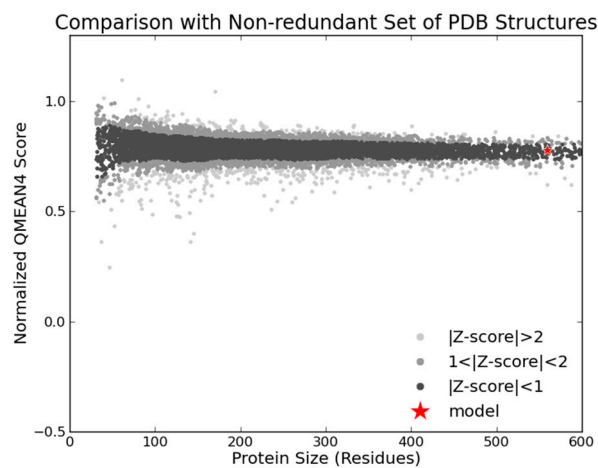


(b) 3D model

(a) Amino acids sequences



(d) Ramachandran Plot



(c) QMEAN Z-score

Figure S5. Homology modeling of *Myceliophthora thermophila* laccase

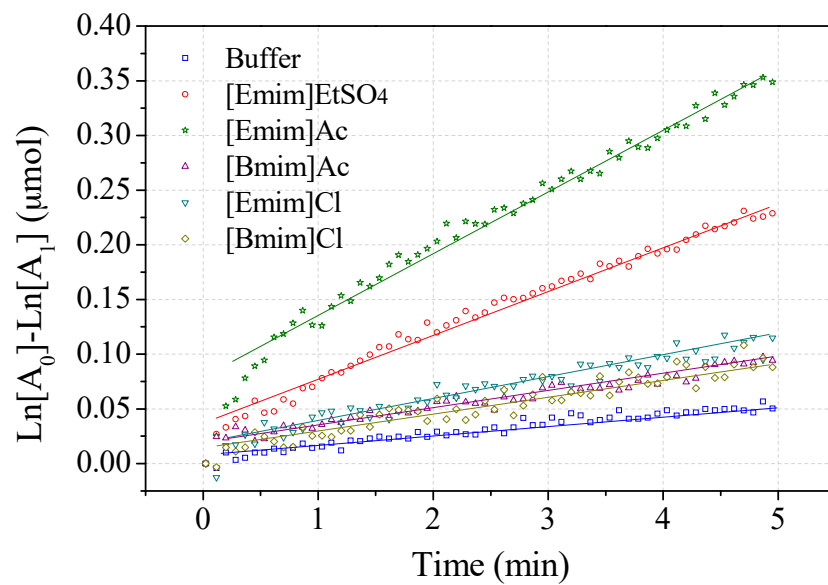


Figure S6. Plots of $\text{Ln}[A_0]-\text{Ln}[A_1]$ vs. radical scavenging time