Supporting Information for

Insights into Catalytic Mechanism of Unsaturated Glucuronyl Hydrolase of *Bacillus sp.* GL1

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Supporting Information

1. Average distances between the carboxylate group of D88 and HO2/HO3 of ∆GlcA unit.

2. Minimum energy pathway and CPR optimized energetic profile for pathway I using SCC-DFTB/MM approach.

3. Minimum energy pathways and CPR optimized energetic profiles for pathway II using SCC-DFTB/MM approach.

4. Overlay representations of EI from pathway □ and from pathway □.

5. Snapshot of ES with No of H193 unprotonated.

6. Full citation for Ref. 24

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7. Full citation for Ref. 30

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Figures S1. Average distances between the carboxylate group of D88 and HO2 of Δ GlcA (left panel) or HO3 of Δ GlcA (right panel).



Fig S2. Minimum energy path for the mechanism I calculated using the SCC-DFTB/MM method, in which $d_1 = d_{\text{OD2}(\text{D149})\cdots\text{HD2}} - d_{\text{C4}\cdots\text{HD2}}, d_2 = d_{\text{C5}\cdots\text{Ow}}$.



Fig S3. CPR-refined energetic profiles using the SCC-DFTB/MM approach for the hydration reaction undergoing mechanism I.



Fig S4. Minimum energy path for the mechanism II along the putative reaction coordinates calculated using the SCC-DFTB/MM method, in which $d_3 = d_{\text{OD2}(\text{D149})\cdots\text{HD2}} - d_{\text{C4}\cdots\text{HD2}}$, $d_4 = d_{\text{O2}\cdots\text{C1}}$, $d_5 = d_{\text{O2}\cdots\text{C1}}$, $d_6 = d_{\text{OD2}(\text{D149})\cdots\text{H}_2} - d_{\text{Ow}\cdots\text{H}_2}$ and $d_7 = d_{\text{C5}\cdots\text{Ow}}$.



Fig S5. CPR-refined energetic profiles using the SCC-DFTB/MM approach for each step in the hydration reaction undergoing mechanism II.



Fig S6. Overlay representations of EI from mechanism \Box (carbon in green) and from mechanism \Box (carbon in magenta).



Figure S7. Snapshot of ES with N δ of H193 unprotonated.