

# QM/MM Molecular Dynamics of Polycaprolactone enzymatic synthesis

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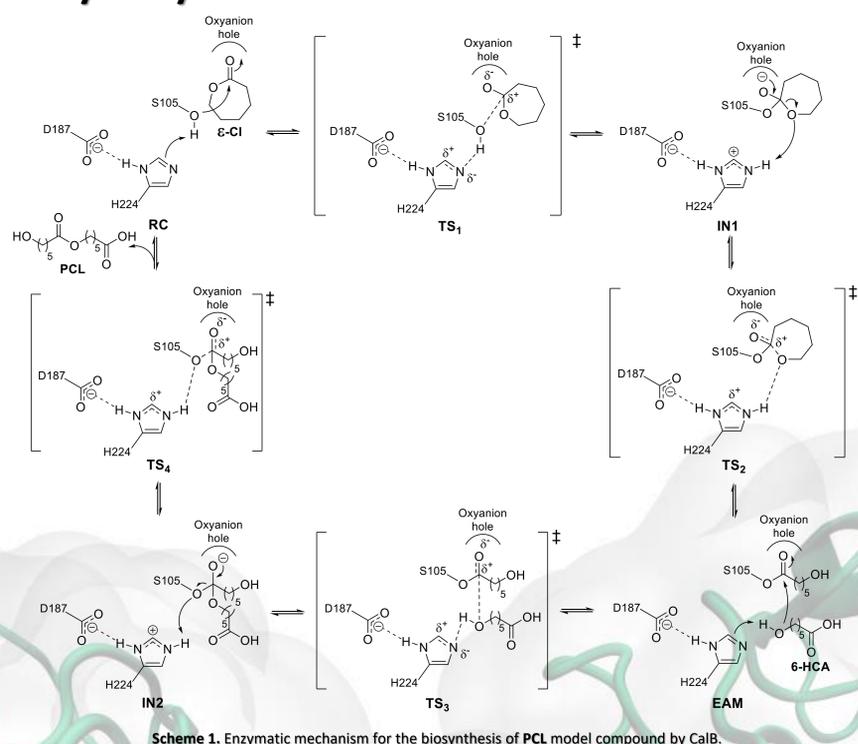


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## Abstract

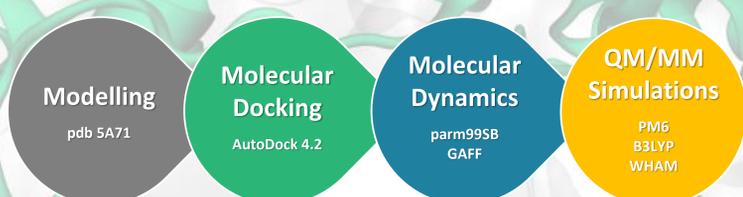
Bioresorbable polymers have gained much attention in the biomedical field, since these materials are non-toxic, biodegradable and biocompatible.<sup>1</sup> Poly( $\epsilon$ -caprolactone) is one of those polymers that, its slow degradation rate has been exploited for tissue engineering, drug-delivery systems and scaffold technologies.<sup>2</sup> The enzyme *Candida antarctica* lipase B (CalB) efficiently catalyzes the ring-opening polymerization (ROP) of  $\epsilon$ -caprolactone to products suitable for biomedical applications.<sup>3</sup> Here, we investigated the catalytic cycle for ROP of  $\epsilon$ -caprolactone using Quantum Mechanics/Molecular Mechanics Molecular Dynamics (QM/MM MD) calculations.<sup>4</sup>

## Catalytic Cycle



CalB, which contains the well described catalytic triad S105, H224 and D187<sup>5</sup> is able to catalyze the synthesis of poly( $\epsilon$ -caprolactone) (PCL).<sup>3</sup> The backbone amides of residues T40 and Q106 and the side-chain hydroxyl of T40 stabilize the negative charge developed in the tetrahedral intermediates (INT-1 and INT-2), which are formed by nucleophilic attack by S105 in  $\epsilon$ -caprolactone ( $\epsilon$ -Cl) and a molecule of 6-hydroxycaproic acid (6-HCA) to the EAM. After product release, the enzyme is ready for another turnover to increase PCL polymer chain.

## Workflow



## References

- <sup>1</sup> Wcislek A. et al., *Polymers* **2018**, 10(6), p1-18
- <sup>2</sup> Malikkammadov E. et al., *J. Biomater. Sci. Polym. Ed.* **2018**, 29, p863-93
- <sup>3</sup> Zhao H. et al., *RSC Adv.* **2017**, 7, p48639-48
- <sup>4</sup> Carvalho A.T.P. et al., *J. Mol. Graph. Model.* **2014**, 54, p67-79
- <sup>5</sup> der Mee L. et al., *Macromolecules* **2006**, 39(15), p5021-27
- <sup>6</sup> Figueiredo P.R. et al., *Front. Mol. Biosci.* **2020**, 6, p109
- <sup>7</sup> Figueiredo P.R. et al., *ChemCatChem* **2020**, 12, p4845

## Discussion

The reaction starts with a nucleophilic attack by the S105 side-chain to the carbonyl group of  $\epsilon$ -Cl forming INT-1 which, after ring-opening assisted by H224 (Fig 1), generates the EAM structure (Fig 2). The TS<sub>1</sub> is  $6.0 \pm 0.1$  kcal/mol above the reactant complex and the TS<sub>2</sub> overall  $\Delta G^\ddagger$  is 9.5 kcal/mol (with the PMF corrected with B3LYP).

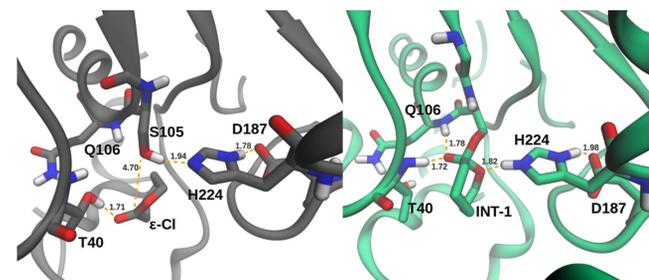


Fig 1. Representation of the enzyme active site RC with  $\epsilon$ -Cl (grey) and the INT-1 complex (dark green).

The alcohol moiety of 6-hydroxycaproic acid 6-HCA performs a nucleophilic attack to the EAM structure (Fig 2). The TS<sub>3</sub> has a free energy barrier of  $7.7 \pm 0.2$  kcal/mol, leading to the formation of the second intermediate structure (INT-2, Fig 2).

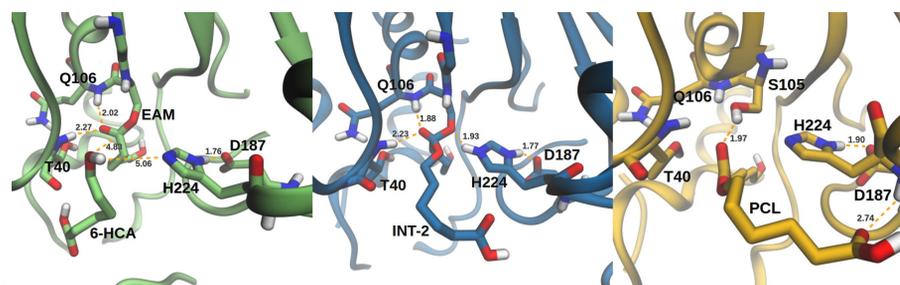


Fig 2. Representation of the EAM complex with 6-HCA (light green), INT-2 complex (blue) and RC with PCL model compound (dark yellow).

The INT-2 complex evolves to the release of a PCL model compound molecule ( $\Delta G^\ddagger$  of 4.1 kcal/mol, Fig 2), regenerating the free enzyme RC (Fig 1).

We were able to fully characterise the catalytic mechanism for this enzyme and for a PCL model compound synthesis.<sup>6</sup> Our results show that the rate-determining step point out to the acylation step and corresponds to the formation of the EAM structure, which is in good agreement with the experimental data.<sup>5</sup>

## Final remarks

Poly( $\epsilon$ -caprolactone) has collected much attention among other biodegradable polymers to be applied in tissue engineering, drug-delivery systems and scaffold technologies due to its biodegradability, biocompatibility and tailorability. Our QM/MM MD calculations for the production of PCL with this enzyme suggest that the rate-determining step is the formation of the EAM structure with an overall free energy of 9.5 kcal/mol. We have compared this mechanism with the one for the thermophilic carboxylesterase AfEST and found that the orientation of the Histidine/Aspartate residues in the INT-1 and connected transition states (TS<sub>1</sub> and TS<sub>2</sub>) differs between both enzymes, providing an explanation for the different catalytic efficiency for this reaction.<sup>6,7</sup>

## Acknowledgements

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