

QM-Cluster Model Study of Enzymatic Reaction  
between Cytochrome P450 Enzyme GcoA and Various Substrates  
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Recently, a promising *Amycolatopsis* cytochrome P450-reductase pair -- a two component P450 class composed of GcoA and GcoB has been discovered.<sup>1</sup> GcoA was found to efficiently demethylate several lignin-derived monomers,<sup>2</sup> suggesting a broad substrate scope for lignin degradation and valorization. The key step of the O-demethylation of guaiacol by GcoA was studied by our group with Density Functional Theory using various sized quantum mechanical (QM)-cluster models.<sup>3</sup> The computed free energy of activation is in good agreement with experimentally measured  $k_{cat}$  of demethylation of guaiacol by GcoA ( $6.8 \pm 0.5 \text{ s}^{-1}$  at 25°C). Hence, the enzymatic reactions of seven different substrates catalyzed by GcoA were studied by a similar approach, starting from Molecular Dynamics simulations of GcoA in complex with each substrate. QM-cluster models were generated by the toolkit *Residue Interaction Network-based Residue Selector* (RINRUS). This study will provide details for better understanding enzymatic O-demethylation of lignins to form catechol derivatives by GcoA and shine a light on structure-activity relationship models for future ligand design and enzyme bioengineering.

Reference:

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