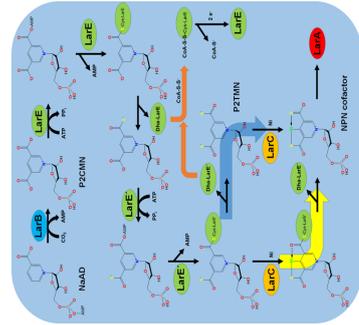
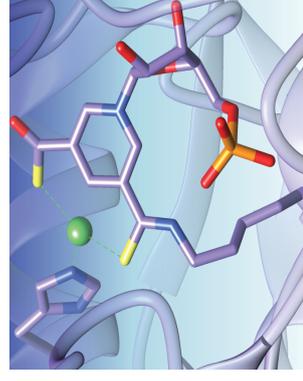


Lactate racemization, a story of so much more than just a nickel

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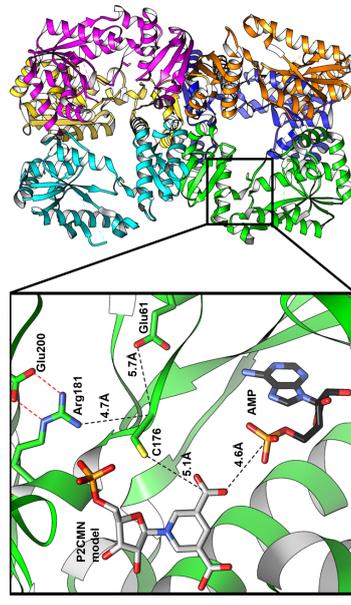


Lactate racemase **LarA**, the ninth discovered nickel-dependent enzyme, was shown to contain a newly identified vitamin B3 derived Ni cofactor (P2TMN-Ni = NPN cofactor). The cofactor is bound via Lys184 with the Ni being coordinated by His200. Synthesis of the cofactor involves three proteins **LarB**, **LarC**, and **LarE**: Cofactor biosynthesis begins with **LarB**, a carboxylase/hydrolase of NaAD. **LarE**, a new member of the PP-loop ATP Pyrophosphatase family, then inserts two sulfur atoms. Structural analysis, combined with structure-guided mutagenesis establishes **LarE** as a paradigm for sulfur transfer through sacrificing its catalytic cysteine residue, only the second sacrificial sulfur transferase to be described^{1,2}. Finally, **LarC** inserts a nickel atom to form a five-membered nickelacycle structure in which a stable nickel-carbon bond, as well as a nickel-sulfur bond is created (a metallacycle). **LarC** is therefore the first cyclometalase identified in nature. Structure-function characterization discovered that **LarC** requires cytidine triphosphate (CTP) hydrolysis³. **LarA** then utilizes the cofactor for lactate racemization. Again using functional and structural methods we provided compelling evidence, that this is accomplished via a proton-coupled hydride transfer mechanism⁴. Lactate racemization is involved in lactate metabolism and cell wall assembly but the cofactor may also be used for a wide range of other yet to be discovered reactions.

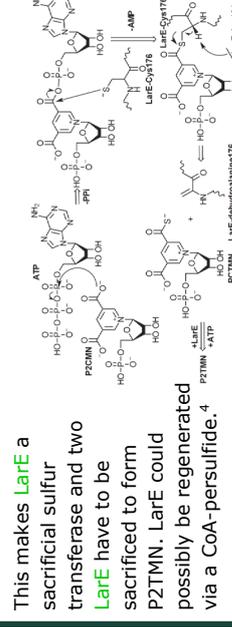


P2TMN-Ni cofactor in LarA

LarE a sacrificial sulfur transferase^{1,2}

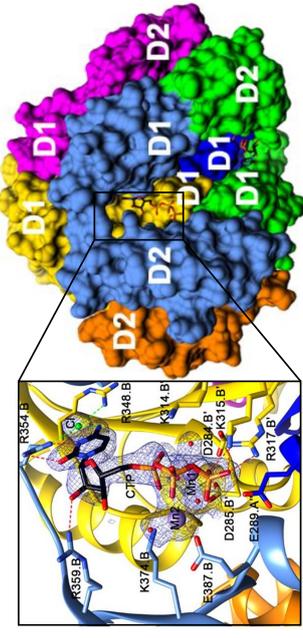


LarE forms a hexamer (6 monomers). It binds ATP-Mg to activate its substrate P2CMN via AMPylation (model based on nicotinamide mononucleotide bound structure - PDB 5UDR). The activated P2CMN is attacked by Cys176 and the sulfur is transferred from the protein to the cofactor.



This makes **LarE** a sacrificial sulfur transferase and two **LarE** have to be sacrificed to form P2TMN. **LarE** could possibly be regenerated via a CoA-persulfide⁴.

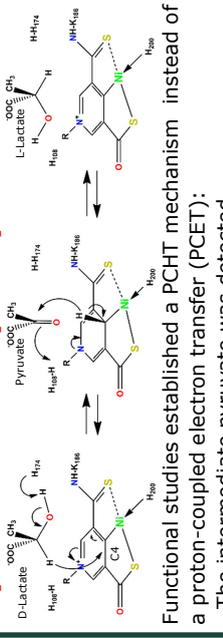
LarC a CTP-dependent cyclometalase³



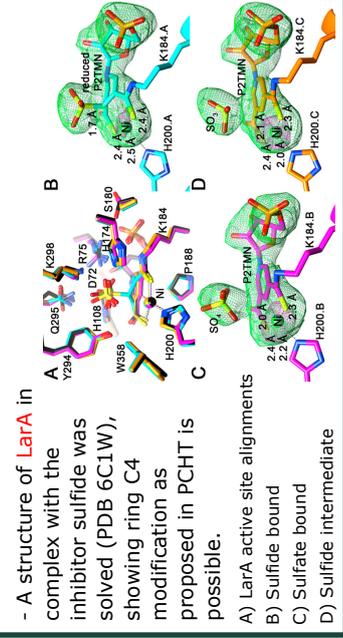
LarC contains a N-terminal C1 domain, responsible for P2TMN and Ni binding. The above structure (PDB 6BWQ) shows the C-terminal C2 domain with a novel CTP binding pocket. The C2 domain forms a hexamer (6 monomers) and each C2 monomer contains two domains D1 and D2. CTP is bound by the D1 and D2 domains of two separate monomers. No structure of C1 is available, therefore the mechanism of **LarC** to create the cofactor metal cycle (Ni-S and Ni-C bonds) using the C2 bound CTP remains unclear.

Future directions, beyond lactate racemase
10% bacteria have **LarA**, many might have different substrate
Additional 15% have only **LarBCE** and no **LarA**
-> undiscovered protein uses cofactor for unknown activity
Lactate and other racemase reaction utilization for industry
Role of these proteins in pathogenic bacteria - drug targets

LarA Operates by a Proton-Coupled Hydride Transfer (PCHT) Mechanism⁴



Functional studies established a PCHT mechanism instead of a proton-coupled electron transfer (PCET):
- The intermediate pyruvate was detected
- No Ni EPR signal was detected (PCHT proposes inactive Ni²⁺ instead of PCET involves EPR active Ni³⁺)
- A normal substrate Substrate Kinetic Isotope Effect of kH/kD of 3.1 is consistent with direct cleavage of the C-H bond (PCHT) instead of C-C bond cleavage in (PCET).



- A structure of **LarA** in complex with the inhibitor sulfide was solved (PDB 6C1W), showing ring C4 modification as proposed in PCHT is possible.
A) LarA active site alignments
B) Sulfide bound
C) Sulfate bound
D) Sulfide intermediate

1. Fellner, M., Desguin, B. et al. (2017). Structural insights into the catalytic mechanism of a sacrificial sulfur transferase LarE from Lactobacillus plantarum. *Biochemistry*, 56(11), 1492-1502.
2. Fellner, M., Desguin, B. et al. (2018). Active Site Cysteine Residue of the Sacrificial Sulfur Transferase LarE from Lactobacillus plantarum. *Biochemistry*, 57(11), 1702-1711.
3. Rankin, J. A. et al. (2018). CTP-dependent Hydride Transfer Mechanism. *Biochemistry*, 57, 3244-3251.
4. Rankin, J. A. et al. (2018). Lactate Racemase Nickel-Pincer Cofactor Operates by a Proton-Coupled Hydride Transfer Mechanism. *Biochemistry*, 57, 3244-3251.