

Title: Metabolomic and Proteomic Analysis of *Zymomonas mobilis* During Nitrogen Fixation Reveals Metabolic Remodeling of Biofuel Producing Pathways

Authors: Julia I. Martien,^{1,2*} (martien@wisc.edu), Edna A. Trujillo,^{3,4} David M. Stevenson,^{1,2} Joshua J. Coon,^{1,3,4,5}, and **Daniel Amador-Noguez**,^{1,2}

Institutions: ¹DOE Great Lakes Bioenergy Research Center, University of Wisconsin-Madison; ²Department of Bacteriology, University of Wisconsin-Madison; ³Department of Chemistry, University of Wisconsin-Madison; ⁴National Center for Quantitative Biology of Complex Systems, University of Wisconsin-Madison; and ⁵Morgridge Institute for Research, Madison, WI

Project Goals: Characterize native metabolic regulation of biofuel producing pathways (such as ethanol production and isoprenoid synthesis) in order to inform genetic engineering of *Z. mobilis* for improved biofuel production.

Abstract text: *Zymomonas mobilis* is a promising biofuel producer capable of rapid glucose consumption and ethanol production. Recently, it was demonstrated that *Z. mobilis* can fix N₂ as a sole nitrogen source (1). Under N₂ fixing conditions, *Z. mobilis* exhibited a higher specific rate of ethanol production than when NH₄⁺ was supplied in the media (1, 2). In order to better understand the metabolic remodeling that occurs during N₂ fixation, we performed metabolomics, proteomics, and thermodynamic analysis of *Z. mobilis* under conditions of N₂ fixation compared to replete NH₄⁺ availability. We also performed metabolomic and proteomic analysis during the dynamic shift to N₂ fixing conditions (NH₄⁺ downshift) and during the shift to NH₄⁺ replete conditions (NH₄⁺ upshift). We found that intracellular concentrations of intermediates of the Entner-Doudoroff (ED) glycolytic pathway were depleted during N₂ fixation. Protein levels of zinc-dependent alcohol dehydrogenase (encoded by *adhA*, ZMO1236) increased by 10-fold during the shift to N₂ fixation, helping to explain the previously observed increase in specific ethanol production. Positional stable isotope labeling revealed that labeled forms indicative of reverse flux were more abundant under NH₄⁺ replete conditions for all five labeled schemes tested, implying increased thermodynamic favorability of the ED pathway during N₂ fixation. We also observed severe depletion of intermediates of the methylerythritol 4-phosphate (MEP) pathway during N₂ fixation, which was accompanied by decreased protein abundance of deoxyxylulose 5-phosphate synthase (DXS), the first enzyme of the MEP pathway. Unexpectedly, we found that intracellular arginine levels were over 3-fold higher during N₂ fixation and decreased by over 3-fold within 10 minutes of NH₄⁺ addition. Based on an overall depletion in intermediates of arginine biosynthesis during N₂ fixation and dynamic changes in protein abundance of a group IV pyridoxal-dependent decarboxylase, encoded in an operon with a deoxyhypusine synthase-like gene, we hypothesize that polyamine synthesis from arginine plays an important role in *Z. mobilis* physiology during changes in NH₄⁺ availability. This study has expanded our fundamental understanding of nitrogen metabolism in *Z. mobilis*, identified DXS protein abundance as a native control-point for MEP pathway activity, and demonstrated that metabolic remodeling during N₂ fixation results in increased thermodynamic favorability of

the ED pathway *in vivo*. These results will help to inform future efforts for metabolic engineering in *Z. mobilis* to increase biofuel production.

References/Publications

1. Kremer TA, LaSarre B, Posto AL, McKinlay JB. 2015. N₂ gas is an effective fertilizer for bioethanol production by *Zymomonas mobilis*. *Proc Natl Acad Sci U S A* 112:2222–6.
2. Palamae S, Choorit W, Chatsungnoen T, Chisti Y. 2020. Simultaneous nitrogen fixation and ethanol production by *Zymomonas mobilis*. *J Biotechnol*.

Funding statement: This material is based upon work supported by the Great Lakes Bioenergy Research Center, U.S. Department of Energy, Office of Science, Office of Biological and Environmental Research under Award Numbers DE-SC0018409 and DE-FC02-07ER64494