### Author's Accepted Manuscript

Controlling cell-free metabolism through physiochemical perturbations

Ashty S. Karim, Jacob T. Heggestad, Samantha A. Crowe, Michael C. Jewett



 PII:
 S1096-7176(17)30172-6

 DOI:
 https://doi.org/10.1016/j.ymben.2017.11.005

 Reference:
 YMBEN1314

To appear in: Metabolic Engineering

Received date:26 May 2017Revised date:3 November 2017Accepted date:11 November 2017

Cite this article as: Ashty S. Karim, Jacob T. Heggestad, Samantha A. Crowe and Michael C. Jewett, Controlling cell-free metabolism through physiochemical p e r t u r b a t i o n s , *Metabolic* Engineering, https://doi.org/10.1016/j.ymben.2017.11.005

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting galley proof before it is published in its final citable form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.

# Controlling cell-free metabolism through physiochemical perturbations

Ashty S. Karim<sup>a,b,c</sup>, Jacob T. Heggestad<sup>a,b,c</sup>, Samantha A. Crowe<sup>a,b,c</sup>, Michael C. Jewett<sup>a,b,c,d,e\*</sup>

<sup>a</sup>Department of Chemical and Biological Engineering

<sup>b</sup>Chemistry of Life Processes Institute

<sup>c</sup>Center for Synthetic Biology, Northwestern University, Evanston, IL 60208, USA,

<sup>d</sup>Robert H. Lurie Comprehensive Cancer Center and <sup>e</sup>Simpson Querrey Institute, Northwestern

University, Chicago, IL 60611, USA

<sup>\*</sup>**Correspondence**: Department of Chemical and Biological Engineering, Northwestern University, Evanston, IL 60208, USA, m-jewett@northwestern.edu, **Tel.**: 1 847 467 5007; f**ax:** 1 847 491 3728

#### Postal Address for all authors:

2145 Sheridan Road, Tech E-136, Evanston, IL 60208, USA

#### Abstract:

Building biosynthetic pathways and engineering metabolic reactions in cells can be timeconsuming due to complexities in cellular metabolism. These complexities often convolute the combinatorial testing of biosynthetic pathway designs needed to define an optimal biosynthetic system. To simplify the optimization of biosynthetic systems, we recently reported a new cellfree framework for pathway construction and testing. In this framework, multiple crude-cell extracts are selectively enriched with individual pathway enzymes, which are then mixed to construct full biosynthetic pathways on the time scale of a day. This rapid approach to building pathways aids in the study of metabolic pathway performance by providing a unique freedom of Download English Version:

## https://daneshyari.com/en/article/6494129

Download Persian Version:

https://daneshyari.com/article/6494129

Daneshyari.com