# Héctor García Martín, Ph. D.

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Quantitative modeler with strong mathematical background experienced in integrating diverse data sets to produce actionable items using machine learning and mechanistic modeling

# **EDUCATION**

Physics Ph. D.
University of Illinois at Urbana-Champaign

October 2004 Urbana, IL USA

Physics B. Sc. University of the Basque Country June 1999 Bilbao, Spain

## WORK EXPERIENCE

# Deputy Vice President, Biofuels and Bioproducts Division

Joint BioEnergy Institute (www.jbei.org)

2018-Present Berkeley, CA USA

Director, Quantitative Metabolic Modeling

Joint BioEnergy Institute (www.jbei.org)

2011-Present Berkeley, CA USA

# Computational Biologist Research Scientist Engineer

Lawrence Berkeley National Laboratory (www.lbl.gov/)

2009-Present

- Berkeley, CA USA
- Used *mechanistic models* and metabolomics data to improve biofuel production in bioengineered strains.
- Used machine learning and proteomics data to improve biofuel production in bioengineered strains.
- Developed an online tool for large-scale standardized data storage and visualization.
- Developed a *microfluidics* chip for performing automated CRISPR-based strain modification.

## **Deputy Director, Host Engineering**

Joint BioEnergy Institute

2008-2011

Berkeley, CA USA

## **Computer Project Scientist**

Lawrence Berkeley National Laboratory

2007-2009

Berkeley, CA USA

- Developed *quantitative predictive models* for microbial metabolism.
- Developed *novel methods to measure metabolic fluxes* using <sup>13</sup>C tracing experiments, genome-scale models and nonlinear optimization techniques.

# **Computational Biologist Post Doctoral Fellow**

DOE Joint Genome Institute (Lawrence Berkeley National Laboratory)

2005-2007 Berkeley, CA USA

- Used metagenomics to create *first comprehensive metabolic map* of phosphorus removing microbial communities.
- Researched wood degradation in termite guts through *metagenomic* techniques.

## **Graduate Research Assistant**

University of Illinois at Urbana-Champaign, Dept. of Physics

2000-2004

Urbana, IL USA

- Derived quantitative explanation for the *universal scaling law* known as Species Area Relationship.
- Used Path Integral Monte Carlo techniques to simulate Bose Einstein Condensates at very high rotation rates.

Thesis: http://guava.physics.uiuc.edu/people/Theses/Martin\_PhD\_thesis.pdf

## SELECTED INVITED TALKS

10<sup>th</sup> International Workshop on Bio-Design and Automation August 2018, Berkeley, USA

REDBIO 2016 conference June 2016, Lima, Peru

Society for Industrial Microbiology (SIMB) annual meeting August 2015, Philadelphia, USA

Cell factories and sustainability conference May 2015, Hillerod, Denmark

Metabolomics 2014 conference June 2014, Tsuruoka, Japan

RECOMB ISCB Conference on Regulatory and Systems Genomics

November 2013, Toronto, Canada

EMSL Biosciences Theme Advisory Panel May 2013, Richland, USA

2<sup>nd</sup> International conference on COBRA October 2012, Elsinore, Denmark

Industrial Biotechnology conference April 2012, Xian, China

## SELECTED PUBLICATIONS

Costello, Z. et al. "A machine learning approach to predict metabolic pathway dynamics from time-series multiomics data" Nature pj Systems Biology & Applications 4.1: 19 (2018)

Chubukov, V. et al. "Synthetic and systems biology for microbial production of commodity chemicals" Nature pi Systems Biology & Applications 2: 16009 (2016)

Garcia Martin, H. *et al.* "A method to constrain genome-scale models with <sup>13</sup>C labeling data" **PLoS Computational Biology 11(9):e1004363 (2015)** 

Warnecke, F., et al. "Metagenomic and functional analysis of hindgut microbiota of a wood feeding higher termite" Nature, 450(7169):560-5 (2007).

García Martín, H., *et al.* "Metagenomic analysis of phosphorus removing sludge communities" Nature Biotechnology, 24: 1263-9 (2006).

García Martín, H. and Goldenfeld, N. "On the origin and robustness of power-law species-area relationships in ecology" **Proceedings of the National Academy of Sciences USA (PNAS)**, 103(27):10310-5 (2006).

Other publications available at: <a href="http://www.researcherid.com/rid/B-5357-2009">http://www.researcherid.com/rid/B-5357-2009</a>

## **AWARDS AND HONORS**

Member of Congresswoman Barbara Lee's biotech advisory committee

US permanent resident as Oustanding Researcher (EB-12)

Member of the Pacific Northwest National Lab's advisory committee for the "Microbes in Transition" initiative

Renato Bobone Award to the Outstanding European Graduate Student in Physics (UIUC)

Excellence in Teaching award (UIUC)

Incomplete List of Teachers Ranked as Excellent (UIUC)

Phi Kappa Phi Honors Society

# **SKILLS**

Python, Matlab, Perl, GAMS, C/C++, UNIX, Linux