Jordan A. Berg

CONTACT INFORMATION	Department of Biochemistry, University of Utah 15 N Medical Drive East Rm. 4100 Salt Lake City, UT 84112		ORCID: 0000-0002-5096-0558 Website: https://j-berg.github.io/ Email: jordan.berg@biochem.utah.edu	
RESEARCH INTERESTS	Cancer Metabolism, Biological Networks, Pattern Recognition, Software Development			
EDUCATION	2016-present	Ph.D. in Biochemistry, University of Utah		
	2010-2016	B.S. in Molecular Biology, Brigham Young University		
RESEARCH EXPERIENCE	2016-present	Graduate Research Assistant, Jared Rutter Lab, University of Utah		
	2013-2016	Undergraduate Research Assistant, Julianne Grose Lab, Brigham Young University		
SERVICE	2020-present	Codechecker, CODECHECK organization		
	2018-present	Web Designer/Social Media Outreach Officer, University of Utah SACNAS Chapter		
	2018	Lead Recruitment Host, Molecular Biology Graduate Program, University of Utah		
	2018	Volunteer, Adventure Scientists		
	2014-2016	Volunteer, Provo Youth Mentoring, Provo,	UT	
	Reviewer (ad hoc): Bioinformatics, NAR Genomics and Bioinformatics, Comput Struct Biotechnol J, F1000 Research, and Journal of Emerging Investigators. See Publons record.			
Honors, Awards, and Fellowships	2020-2022	F99 Predoctoral Fellow, National Cancer In	stitute	
	2018-2020	T32 Graduate Trainee, National Institute o eases	of Diabetes and Digestive and Kidney Dis-	
	2016	Outstanding Research Award, Department Brigham Young University	of Microbiology and Molecular Biology,	
	2014	iGem Silver Medal, iGEM World Jamboree,	Boston, MA	
SELECTED PUBLICATIONS	& indicates co-corresponding authors			
	Berg JA ^{&} , George I, Zhou Y, Hicks KG, Wang B, Rutter J ^{&} . Electrum contextualizes enzyme-metabolite interactions and predicts allosteric regulatory mecha-			

Electrum contextualizes enzyme-metabolite interactions and predicts allosteric regulatory mechanisms.

[working draft] [software]

Berg JA $^{\&}$, Zhou Y, Waller TC, Ouyang Y, Nowinski SM, Van Ry T, George I, Cox JE, Wang B, Rutter J $^{\&}$.

Gazing into the Metaboverse: Automated exploration and contextualization of metabolic data. *bioRxiv*. (2020) https://doi.org/10.1101/2020.06.25.171850. [working draft] [software]

 $\textbf{Berg JA}\&, \ \text{Belyeu JR}, \ \text{Morgan JT}, \ \text{Ouyang Y}, \ \text{Bott AJ}, \ \text{Quinlan AR}, \ \text{Gertz J}, \ \text{Rutter J}\&.$

XPRESSyourself: Enhancing, Standardizing, and Automating Ribosome Profiling Computational Analyses Yields Improved Insight into Data.

PLoS Comp. Biol. (2020) https://doi.org/10.1371/journal.pcbi.1007625. [software]

Berg JA, Merrill BD, Breakwell DP, Grose JH, Hope S.

A PCR-based method for distinguishing between two common beehive bacteria, *Paenibacillus larvae* and *Brevibacillus laterosporus*.

Journal of Appl Environ Microbiol. (2018) https://doi.org/10.1128/AEM.01886-18. [software]

Berg JA, Merrill BD, Crockett JT, Esplin KP, Evans MR, Heaton KE, Hilton JA, Hyde JR, McBride MS, Schouten JT, Simister AR, Thurgood TL, Ward AT, Breakwell DP, Burnett SH, Grose JH. Characterization of five novel *Brevibacillus* bacteriophages and genomic comparison of *Brevibacillus* phages.

PLoS ONE. (2016) https://doi.org/10.1371/journal.pone.0156838.

OTHER PUBLICATIONS

Hicks KG, Cluntun AA, Schubert HL, Hackett SR, **Berg JA**, Leonard PG, Ajalla Aleixo MA, Blevins A, Barta P, Tilley S, Fogarty S, Ahn H-C, Allen KN, Atsushi M, Block SD, Ding J, Dreveny I, Gasper C, Ho Q, Palladino MJ, Prajapati S, Sun PK, Tittmann K, Tolan DR, Unterlass J, VanDemark AP, Vander Heiden MG, Webb B, Yun C-H, Zhap PK, Hill CP, Nonato MC, Muller FL, Gottschling DE, Cox JE, Rutter J.

Protein-Metabolite Interactomics Reveals Novel Regulation of Carbohydrate Metabolism. *bioRxiv*. (2021) https://doi.org/10.1101/2021.08.28.458030.

Winter JM, Fresenius H, Keys HR, Cunningham CN, Ryan J, Sirohi D, Berg JA, Tripp S, Barta P, Agarwal N, Letai A, Sabatini D, Wohlever M, Rutter J.

Co-deletion of ATAD1 with PTEN primes cells for BIM-mediated apoptosis.

bioRxiv. (2021) https://doi.org/10.1101/2021.07.01.450781.

Nuebel E, Morgan JT, Fogarty S, Winter JM, Lettlova S, **Berg JA**, Chen Y-C, Kidwell CU, Maschek JA, Clowers KJ, Argyriou C, Chen L, Wittig I, Cox JE, Roh-Johnson M, Braverman N, Steinberg SJ, Gygi SP, Rutter J.

The biochemical basis of mitochondrial dysfunction in Zellweger Spectrum Disorder. *EMBO Reports*. (2021) https://doi.org/10.15252/embr.202051991. [code notebook]

Cluntun AA, Badolia R, Lettlova S, Parnell KM, Shankar TS, Diakos NA, Olson KA, Taleb I, Tatum SM, **Berg JA**, Cunningham CN, Van Ry T, Bott AJ, Krokidi AT, Fogarty S, Skedros S, Swiatek WI, Yu X, Luo B, Merx S, Navankasattusas S, Cox JE, Ducker GS, Holland WL, McKellar SH, Rutter J, Drakos SG.

The Pyruvate-Lactate Axis Modulates Cardiac Hypertrophy and Heart Failure. *Cell Metabolism*. (2020) https://doi.org/10.1016/j.cmet.2020.12.003. [code notebook]

Nowinski SM, Solmonson A, Rusin SF, Maschek JA, Bensard CL, Fogarty S, Jeong M, Lettlova S, **Berg JA**, Morgan JT, Ouyang Y, Naylor BC, Paulo JA, Funai K, Cox JE, Gygi SP, Winge DR, Deberardinis RJ. Rutter J.

Mitochondrial fatty acid synthesis coordinates oxidative metabolism in mammalian mitochondria. *eLife*. (2020) https://doi.org/10.7554/eLife.58041. [code notebook]

Waller TC, Berg JA, Lex A, Chapman BE, Rutter J.

Compartment and Hub Definitions Tune Metabolic Networks for Metabolic Interpretations. *GigaScience*. (2020) https://doi.org/10.1093/gigascience/giz137. [software]

Hughes CE, Coody TK, Jeong M, **Berg JA**, Winge DR, Hughes AL. Amino acid toxicity drives age-related mitochondrial decline by altering iron metabolism. *Cell*. (2020) https://doi.org/10.1016/j.cell.2019.12.035. [code notebook]

Bensard CL, Wisidigama DR, Olsen KA, **Berg JA**, Krah NM, Schell JC, Nowinski SM, Fogarty S, Bott AJ, Wei P, Dove KK, Tanner JM, Panic V, Cluntun A, Lettlova S, Earl CS, Namnath DF, Vázquez-Arregun K, Villanueva CJ, Tantin D, Murtaugh LC, Evason KJ, Ducker GS, Thummel CS, Rutter J.

Regulation of Tumor Initiation by the Mitochondrial Pyruvate Carrier. *Cell Metabolism*. (2019) https://doi.org/10.1016/j.cmet.2019.11.002. [code notebook]

Van Vranken JG, Nowinski SM, Clowers KJ, Jeong M, Ouyang Y, **Berg JA**, Gygi J, Gygi SP, Winge DR, Rutter JP.

ACP acylation is an acetyl-CoA-dependent modification required for electron transport chain assembly.

Molecular Cell. (2018) https://doi.org/10.1016/j.molcel.2018.06.039. [code notebook]

A complete list of publications can be found at Google Scholar.

ORAL PRESENTATIONS

Gazing into the Metaboverse — Automated exploration and contextualization of metabolic data CSHL Biological Data Science (2020; virtual) - Selected Talk

Program Slides

ISMB (2020; virtual) - Selected Short Talk

Program Slides

EMBL/EMBO Epigenetics Meets Metabolism (2019) - Selected Talk

Program Slides

XPRESSyourself: Enhancing, Standardizing, and Automating Ribosome Profiling Computational Analyses Yields Improved Insights

CSHL Genome Informatics (2019) - Lightning Talk

Program

Bioinformatics Seminar Series, Center for Computational Biology & Bioinformatics, UCSD (2019) - Invited Talk

Program Slides

Characterization and analysis of six novel Erwinia phages reveals relationship to Enterobacteriaceae family members

Tri-branch ASM meeting (2015) - Selected Talk

POSTER PRESENTATIONS

A novel metabolic signature in malignant human lung adenocarcinomas is revealed by contextual, reaction-resolution analysis of metabolic data

Keystone Symposia: Metabolic Decisions in Development and Disease (2021; virtual) Poster

Keystone Symposia: Tumor Metabolism and the Microenvironment (2021; virtual) *Poster*

Gazing into the Metaboverse — Automated exploration and contextualization of metabolic data Systems Approaches to Cancer Biology (2020; virtual)

Poster

XPRESSyourself: Enhancing, Standardizing, and Automating Ribosome Profiling Computational Analyses Yields Improved Insights

CSHL Genome Informatics (2019)

RECOMB (2018)

Ribosome profiling reveals translation-level regulation of peroxins in response to loss of peroxi-

somes

Frontiers in Metabolism (2018)

GENBANK PUBLICATIONS	A complete list	A complete listing of bacteriophage genome publications can be found on GenBank (60 total).		
TEACHING EXPERIENCE	2018, 2020	MBIOL 6200: Lit. & Problem Solving, University of Utah (3CR; 4 class hours/week; TA)		
	2015-2016	MMBIOL194B: Phage Hunters Genomics, Brigham Young University (2CR; 6 lab hours/week; TA)		
	2014-2015	MMBIOL194A: Phage Hunters Discovery, Brigham Young University (2CR; 6 lab hours/week: TA)		

TRAINEES

Lilly R. (High School Student) Ian George* (Undergraduate, U. of Utah) Claudia Charles (Undergraduate, U. of Utah) Nolan Beatty (Undergraduate, Brigham Young U.) Braden Brundage* (Undergraduate, Brigham Young U.) Alisa Buchanan (Undergraduate, Brigham Young U.) Minsey Choi* (Undergraduate, Brigham Young U.) Justin Crockett* (Undergraduate, Brigham Young U.) Kyle Esplin* (Undergraduate, Brigham Young U.) Marlee Evans* (Undergraduate, Brigham Young U.) Hannah Ferguson* (Undergraduate, Brigham Young U.) Karli Heaton* (Undergraduate, Brigham Young U.) Jared Hilton* (Undergraduate, Brigham Young U.) Emily Hurst (Undergraduate, Brigham Young U.) Jonathan Hyde* (Undergraduate, Brigham Young U.) Moon Hee I (Undergraduate, Brigham Young U.) Morgan McBride* (Undergraduate, Brigham Young U.) Sam Pollock (Undergraduate, Brigham Young U.) Micah Putnam* (Undergraduate, Brigham Young U.) Jordan Schouten* (Undergraduate, Brigham Young U.) Jeremy Severe (Undergraduate, Brigham Young U.) Austin Simister* (Undergraduate, Brigham Young U.) Philip (PJ) Tatlow* (Undergraduate, Brigham Young U.) Trever Thurgood* (Undergraduate, Brigham Young U.) Charles (CJ) Webb* (Undergraduate, Brigham Young U.)

An asterisk indicates those who presented their research at a conference or in a publication.

OTHER

Languages: English (native), Spanish (professional working proficiency), Mandarin Chinese (elementary proficiency)

Programming: Python, R, Javascript, HTML/CSS, ŁTEX, Bash, SLURM, C++, Julia

Citizenship: United States; Born: 1991

Last update: August 31, 2021