Gregory Moyerbrailean

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Education

2012 PhD in Molecular Biology	and Genetics (C	Computational Biolog	av), Center	tor Molecular Medicine
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- 2016 and Genomics, Wayne State University, Detroit MI
 Advisors: Roger Pique-Regi and Francesca Luca

2006 – 2010 BS, Microbiology (Cell and Molecular Biology), Michigan State University, East Lansing, MI

Research Experience

Aug 2012 Center for Molecular Medicine and Genomics, Wayne State University

 present ° Computational analysis of non-coding genetic variation, differential gene expression, allelespecific gene expression, and phenotypic variation using high-throughput sequencing data

Nov 2011 Department of Physiology, Wayne State University

Aug 2012 ° Microbial community analysis from 16S deep-sequencing

Jul 2010 Department of Microbiology and Molecular Genetics, Michigan State University

- Sep 2011
 Molecular analysis of the beneficial effects of probiotics on bone health
 - ° Software development to assist targeted mutagenesis via high-efficiency recombineering

Nov 2008 Department of Plant Biology, Michigan State University

- Jul 2010 ° Data analysis of student performance for a broad NSF-funded project to reform introductory biology
 - ° Collection, organization, and storage of sensitive data relating to the NSF-funded Faculty Institutes for Reforming Scientific Teaching (FIRST) IV project

Sep 2007 Department of Microbiology and Molecular Genetics, Michigan State University

- Aug 2008 ° Molecular characterization of a putative thymidine-7-hydroxylase found in *Trypanosoma brucei*

Teaching Experience

Winter 2010 Teaching Assistant, Computer Concepts and Competencies (CSE101) at Michigan State University

Fall 2009 **Teaching Assistant**, Biology of Plants (PLB203) at Michigan State University **Teaching Assistant**, Computer Concepts and Competencies (CSE101) at Michigan State University

Professional Experience

Jul 2008 Laboratory Technician, MSU Diagnostic Center for Population and Animal Health, East Lansing, MI.

- Aug 2009 Tasks included tissue handling and preparations, cleanup and disposal of hazardous materials, working with a CERNER client database, and following Standard Operating Procedures

Intern, BrainCells, Inc, La Jolla, CA. Regular tasks included dishwashing, updating inventory, making solutions, and weighing out compounds. Extra tasks included coating cell culture plates, cutting tissue sections with a microtome, and antibody staining of tissue samples.

Publications

- 2016 Kato I, AA Vasquez, GA Moyerbrailean, S Land, J Sun, HS Lin, JL Ram. (2016). Oral microbiome and history of smoking and colorectal cancer. Journal of Epidemiological Research, 2(2), 92-101.
- 2016 Moyerbrailean GA, CT Harvey, C Kalita, X Wen, F Luca, and R Pique-Regi. (2016). Which genetic variants in DNase I sensitivity regions are functional? PLoS Genetics, 12(2), e1005875.

- 2015 Moyerbrailean GA, GO Davis, CT Harvey, D Watza, X Wen, R Pique-Regi, and F Luca. (2015). A high-throughput RNA-seq approach to profile transcriptional responses. Scientific Reports, 5:14976, DOI: 10.1038/srep14976
- 2014 Harvey CT, **GA Moyerbrailean**, GO Davis, X Wen, F Luca, and R Pique-Regi. (2014). <u>QuASAR:</u> <u>Quantitative Allele Specific Analysis of Reads.</u> *Bioinformatics*, 31(8), 1235-1242.
- Fujimoto M, **GA Moyerbrailean**, S Noman, JP Gizicki, ML Ram, J Henqinet, PA Green, JL Ram. (2014). <u>Application of Ion Torrent sequencing to the assessment of the effect of alkali ballast water treatment on microbial community diversity. *PLoS ONE*, *9*(9):e107534.</u>
- Bray Speth E, JL Momsen, **GA Moyerbrailean**, D Ebert-May, TM Long, SA Wyse and D Linton. (2010). 1, 2, 3, 4: Infusing quantitative literacy into introductory biology. *CBE-LSE*, 9(3), 323-32.

Select Presentations

- Nov 2015 **Moyerbrailean GA***, AL Richards, D Kurtz, CA Kalita, GO Davis, CT Harvey, A Alazizi, D Watza, Y Sorokin, N Hauff, X Zhou, X Wen, R Pique-Regi, F Luca. High-throughput allele-specific expression across 250 environmental conditions. Oral presentation at RECOMB/ISCB, Philadelphia, PA.
- Nov 2014 **Moyerbrailean GA***, CT Harvey, CA Kalita, X Wen, Luca F and R Pique-Regi. Which genetic variants in DNase I hypersensitive regions are functional? Oral presentation at RECOMB/ISCB, San Diego, CA.
- Jul 2014 Moyerbrailean GA*, CT Harvey, CA Kalita, X Wen, Luca F and R Pique-Regi. Empirical and computational survey of non-coding variants that affect transcription factor binding. Poster presentation at ISMB RegGenSIG, Boston, MA.
- Jun 2014 Luca F, **GA Moyerbrailean***, CT Harvey, CA Kalita, X Wen and R Pique-Regi. Molecular and Evolutionary Features of Regulatory Variants. Poster presentation at SMBE, San Juan, Puerto Rico.
- Nov 2013 Luca F, **GA Moyerbrailean**, CT Harvey and Pique-Regi R*. Analysis of non-coding genetic variants impact on gene regulation and complex traits. Oral presentation at RECOMB/ISCB, Toronto, ON, Canada.
- May 2013 Moyerbrailean GA*, F Luca and R Pique-Regi. Analysis of genetic variants and their effects on complex traits. Poster presentation at BOG, Cold Spring Harbor, NY.

Computational Skills

Programming Python, R, Java, C++, HTML, CSS, Javascript (incl. jQuery) Familiar with Perl, Ruby, SQL (MySQL and PostgreSQL), C/C#, PHP

OS Mac OSX, Linux, Microsoft Windows

Type Setting LaTeX, Microsoft Office, Google Docs

IDE Vim, emacs, Eclipse, Microsoft Visual Studio