### **CURRICULUM VITAE**

#### **CONTACT INFORMATION**

Name: David M. Reif Office: Keystone 2018

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Morrisville, NC 27560

Fmail: david.reif@nih.gov

NIH https://www.niehs.nih.gov/research/atniehs/labs/ptb/staff/reif/index.cfm Websites:

> Reif Lab http://reif-lab.org/ ToxPi https://toxpi.org/

Google Scholar <a href="http://scholar.google.com/citations?user=0853ilkAAAAJ">http://scholar.google.com/citations?user=0853ilkAAAAJ</a>

ORCID 0000-0001-7815-6767

### **EDUCATION AND TRAINING**

2006-2008 U.S. Environmental Protection Agency (Research Triangle Park, NC)

Post-doc in Computational Toxicology

Vanderbilt University (Nashville, TN) 2002-2006

Ph.D. in Human Genetics

Vanderbilt University (Nashville, TN) 2003-2005

M.S. in Applied Statistics

College of William & Mary (Williamsburg, VA) 1998-2002

Monroe Scholar Honors Program

B.S. in Biology (Major) & Finance (Minor)

#### POSITIONS AND PROFESSIONAL EXPERIENCE

2022-Current Branch Chief & Senior Scientist,

Predictive Toxicology Branch, Division of Translational Toxicology,

National Institute of Environmental Health Sciences, RTP, NC

2020-2022 Professor.

> Department of Biological Sciences (Primary Appointment), Bioinformatics Research Center (Resident Member),

Department of Statistics (Associate Faculty),

NC Agromedicine Institute (Member),

Chancellor's Faculty Excellence Program (Bioinformatics Cluster),

Director, Bioinformatics Consulting and Service Center,

Director, Environmental Health Bioinformatics Training Program

Director, Data Management and Analysis Core (DMAC), Superfund Center for

Environmental and Health Effects of PFAS,

Co-Director, Integrated Health Science Facility Core (IHSFC), Center for Human Health

and the Environment

North Carolina State University, Raleigh, NC

2013-2020 Associate Professor,

Department of Biological Sciences (Primary Appointment),

Bioinformatics Research Center (Resident Member),

Department of Statistics (Associate Faculty),

NC Agromedicine Institute (Member),

Chancellor's Faculty Excellence Program (Bioinformatics Cluster),

Director, Bioinformatics Consulting and Service Center,

Lead. Bioinformatics Team. Center for Human Health and the Environment

North Carolina State University, Raleigh, NC

Affiliate Member, 2013-current

	Center for Pharmacogenomics and Individualized Therapy, University of North Carolina, Chapel Hill, NC
2008-2012	Principal Investigator (Statistician),
	National Center for Computational Toxicology,
	U.S. Environmental Protection Agency, Research Triangle Park, NC
2009-2012	Adjunct Assistant Professor,
	Department of Statistics,
	North Carolina State University, Raleigh, NC
2008	Visiting Scholar,
	Department of Statistics,
	North Carolina State University, Raleigh, NC
2006-2008	Federal post-doc (Biologist),
	National Center for Computational Toxicology,
	U.S. Environmental Protection Agency, Research Triangle Park, NC
	(advisor: Elaine Cohen Hubal)
2002-2006	Graduate Research Assistant,
	Center for Human Genetics Research,
	Vanderbilt University, Nashville, TN
	(advisors: Jason Moore and Jonathan Haines)
1999-2001	Research Assistant,
	Department of Biology,
	College of William & Mary, Williamsburg, VA
	(advisor: Patty Zwollo)
2001	Summer Associate,
	Navigant Consulting, Washington, DC

# SELECTED HONORS, AWARDS, AND APPOINTMENTS

2021-current	Member, Toxic Substances Control Act (TSCA) Science Advisory Committee on Chemicals (SACC), U.S. Environmental Protection Agency
2020-current	Co-Chair, Data Working Group, U.S. FDA Botanical Safety Consortium
2020-current	Advisory Committee, Pacific Northwest Center for Translational Environmental Health Research, Oregon State University
2020-current	Advisory Board, College of Sciences Mentorship Program, North Carolina State
	University
2019	Invited Expert, "Implementing a Class Approach to Hazard Assessment of
	Organohalogen Flame Retardants", National Academy of Sciences (NAS)
2018	Committee Member, "Committee on a Scoping Plan to Assess the Hazards of
	Organohalogen Flame Retardants", National Academy of Sciences (NAS)
2018	Planning Committee & Session Moderator, "Informing Environmental Health Decisions
	Through Data Integration", National Academy of Sciences (NAS)
2018	NIEHS Paper of the Month, "Elucidating Gene-by-Environment (GxE) Interactions
	Associated with Differential Susceptibility to Chemical Exposure"
2017	Working Group, IARC Monographs on the Evaluation of Carcinogenic Risks to Humans
	"Benzene", World Health Organization (WHO) International Agency for Research on
	Cancer (IARC)
2015	Working Group, IARC Monographs on the Evaluation of Carcinogenic Risks to Humans
2010	"Some Organochlorine Insecticides and Some Chlorphenoxy Herbicides", World
	Health Organization (WHO) International Agency for Research on Cancer (IARC)
2014	Committee Member & Chapter Lead, "Predictive Toxicology Approaches for Military
2011	Assessments of Acute Exposures", National Academy of Sciences (NAS)
2014	STAA Level II Award, for the Expocast project, Office of Research and Development,
2014	U.S. EPA
2013	STAA Level I Award, for Advancing Chemical Safety Assessment via the Development
2010	of Predictive Reproductive and Developmental Toxicity Models, Office of Research and
	Development, U.S. EPA
	Development, U.S. LFA

Page 2 (of 22)

David M. Reif

2012	Impact Award, for Comptox Chemical Toxicity Databases, Office of Research and
2012	Development, U.S. EPA Honor Award (Bronze Medal), for Development of Reproductive, Developmental, Vascular Disruption and Cancer Predictive Models, Office of Research and
2012	Development, U.S. EPA STAA Level III Award, for the Analysis of Eight Oil Spill Dispersants Using Rapid, In Vitro Tests for Endocrine and Other Biological Activity,
2012	Office of Research and Development, U.S. EPA STAA Level III Award, for the Endocrine Profiling and Prioritization of Environmental Chemicals Using ToxCast Data,
2011	Office of Research and Development, U.S. EPA Presidential Early Career Award for Scientists and Engineers (PECASE), Executive Office of the President  *The first time an EPA scientist won a PECASE, which is "The highest honor"
2011	bestowed by the United States government on science and engineering professionals in the early stages of their independent research careers"  Science Achievement Award in Health Sciences,
2011	Office of Research and Development, U.S. EPA  *An Agency-level award selected by an external peer panel to recognize significant advances in health sciences and impact on regulatory programs
2011	EPA Gold Coin Award, Office of the Administrator, U.S. EPA
2011	*A special award to recognize the rapid scientific response to the Deep Water Horizon oil spill emergency Honor Award (Bronze Medal), for the Mechanistic Indicators of Childhood Asthma
2011	(MICA) study, Office of Research and Development, U.S. EPA STAA Level II Award, for the Toxicity Reference Database (ToxRefDB),
2011	Office of Research and Development, U.S. EPA Quality Step Increase (QSI) Award,
2010	Office of Research and Development, U.S. EPA OTS Award,
2010	Office of Public Affairs, U.S. EPA S (Superior Accomplishment) Award,
2010	National Center for Computational Toxicology, U.S. EPA S (Superior Accomplishment) Award,
	National Health and Environmental Effects Research Laboratory, U.S. EPA
2009	OTS Award, National Center for Computational Toxicology, U.S. EPA
2007	OTS Award, Human Studies Division, U.S. EPA
2005	International Travel Grant, Vanderbilt University
2003-2005	NIH Training Grant in Human Genetics, Vanderbilt University
2002	Phi Sigma Biology Honors Fraternity, College of William & Mary
2001	Omicron Delta Kappa Leadership Fraternity, College of William & Mary
2001	Monroe Scholarship Supplemental Award for International Research, College of William & Mary
2000-2001	Howard Hughes Medical Institute Undergraduate Research Grant, College of William & Mary
1998-2002	Monroe Scholar, College of William and Mary

Page 3 (of 22)

David M. Reif

### **RESEARCH INTERESTS (KEYWORDS)**

Predictive Toxicology; Translational Toxicology; Bioinformatics; Data Integration; Computational Toxicology; Environmental Health Sciences; Statistical Genetics; Epidemiology (Genetic, Veterinary, Agricultural, Environmental); Translational Research; Computational Modeling; Environmental Exposure; Risk Assessment; Gene-Environment Interactions; Visual Analytics and Statistical Graphics; Aquatic Model Organisms (Zebrafish); High Throughput Screening; Bioassay Development; Software Development; Environmental Statistics; New Approach Methodologies (NAMs); Geographic Information Systems (GIS)

## PROFESSIONAL SOCIETIES (ACTIVE)

Society of Toxicology (SOT), Full Member, 2008 – current Federation of American Scientists (FAS), Board of Sponsors, 2012 – current International Society of Exposure Science (ISES), Member, 2016 – current American Society of Human Genetics (ASHG), Member, 2017 – current

#### **EDITORIAL AND REVIEWER SERVICE**

#### **Publications**

Editorial Board, Frontiers in Toxicology, 2019 – current Associate Editor, Environmental Health Perspectives (EHP), 2016 – current Editorial Board, Journal of Exposure Science and Environmental Epidemiology (JESEE), 2016 – current

Associate Editor, *BioData Mining*, 2018 – current Managing Editor, *BioData Mining*, 2015 – 18 Journal Reviewer for:

Bioinformatics; Biotechniques; BMC Bioinformatics; Environmental Health Perspectives; Environmental Pollution; Environmental Toxicology and Chemistry; Frontiers in Environmental Science; Genetic Epidemiology; Human Genetics; IEEE/ACM Transactions on Computational Biology and Bioinformatics; Journal of Agricultural, Biological, and Environmental Statistics; Journal of Exposure Science and Environmental Epidemiology; Journal of Infectious Disease; Journal of Statistical Software; Journal of the American Statistical Association; Medical Science Monitor; Neuropsychiatric Genetics; Nucleic Acids Research; Pharmacogenomics; PLoS Genetics; PLoS One; Toxicological Sciences

#### Grants

Grant Reviewer, Superfund Research Program (P42),

National Institutes of Health

Grant Reviewer, Medical Research Council (MRC),

Research Councils UK

Study Section *ad hoc* Member, Systemic Injury and Environmental Exposure (SIEE), National Institutes of Health

Study Section *ad hoc* Member, Digestive, Kidney and Urological Systems (DKUS), National Institutes of Health

Study Section *ad hoc* Member, Bioengineering Sciences and Technologies (BST), National Institutes of Health

Grant Reviewer, National Center for Environmental Research (NCER),

U.S. Environmental Protection Agency

Study Section *ad hoc* Member, Biomedical Computing and Health Informatics (BCHI), National Institutes of Health

Grant Reviewer, Division of Information and Intelligent Systems (IIS), National Science Foundation

Page 4 (of 22) David M. Reif

# Professional and Fellowships

External Tenure Evaluator,

City University of New Year (CUNY)

University of Indiana

Promotion Committee,

National Institutes of Environmental Health Sciences

Graduate Fellowship Reviewer, Science to Achieve Results (STAR),

U.S. Environmental Protection Agency

# Translational and Regulatory

Expert Reviewer,

CalEPA's Proposed Toxicological Priority (ToxPi) Framework

Office of Environmental Health Hazard Assessment (OEHHA), State of California

Peer Reviewer,

National Academy of Sciences (NAS)

Peer Reviewer,

International Agency for Research on Cancer (IARC)

### **TEACHING**

### Full Semester Courses

2019-current	Course Co-Director & Lecturer, Computational Environmental Health Sciences (BIO 592),
	North Carolina State University, Raleigh, NC, USA [Spring Session]
2014-current	Course Director & Lecturer,
	Introduction to Bioinformatics (GN 427),
	North Carolina State University, Raleigh, NC, USA [Fall Session]
2011	Lab Course Director & Lecturer,
	Statistical Genetics Practicum (STAT 489, STAT 498),
	North Carolina State University, Raleigh, NC, USA [Summer Session]
2010	Lab Course Director & Lecturer,
	Statistical Genetics Practicum (STAT 489, STAT 498),
	North Carolina State University, Raleigh, NC, USA [Summer Session]
2010	Course Director & Lecturer,
	Introduction to R (STAT 610),
	North Carolina State University, Raleigh, NC, USA [Spring Session]
2008	Course Director & Lecturer,
	Introduction to R (STAT 610),
	North Carolina State University, Raleigh, NC, USA [Spring Session]
2006	Teaching Assistant & Guest Lecturer,
_000	Statistics for Biomedical Researchers (IGP 304),
	Vanderbilt University, Nashville, TN, USA [Spring Session].
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# Short Courses

2010	Introduction to D and Dispetationies (with Kan Disp)
2019	Introduction to R and Biostatistics (with Ken Rice),
	Winter Institute in Statistical Genetics,
	NYU - Abu Dhabi, Abu Dhabi, United Arab Emirates [Short Course]
2016	Bioinformatics (with Nadia Singh and Dahlia Nielsen),
	Research Initiative for Scientific Enhancement (RISE) program,
	Fayetteville State University, Fayetteville, NC, USA [Short Course]
2015-2016	Practical Bioinformatics: Introduction to R,
	North Carolina State University, Raleigh, NC, USA [Short Course].

Page 5 (of 22)

David M. Reif

### Lectures

2021-current Computational Toxicology and Exposure Science (ENVR 890), University of North Carolina at Chapel Hill, Chapel Hill, NC, USA. 2016-current Responsible Conduct in Science (GN 850 / TOX 820) North Carolina State University, Raleigh, NC, USA. 2015-2017 Biochemical and Molecular Toxicology (TOXC / ENVR 442), University of North Carolina at Chapel Hill, Chapel Hill, NC, USA. Principles of Toxicology (TOX 710), 2015-current North Carolina State University, Raleigh, NC, USA. Advanced Toxicology (TOXC / ENVR 707), 2015-2018 University of North Carolina at Chapel Hill, Chapel Hill, NC, USA. 2013 Genetic Algorithms (CSC 530), North Carolina State University, Raleigh, NC, USA. 2011 Analytics Methods and Applications (Institute for Advanced Analytics), North Carolina State University, Raleigh, NC, USA. 2006 General Biology I & II, Nashville State Technical College, Nashville, TN, USA.

# Research & Academic Program Participation and Direction

2019-2021 Associate Director

Environmental Health Bioinformatics (T32 Training Grant) Program

North Carolina State University

2015-current Training Faculty

> Environmental Science and Engineering University of North Carolina at Chapel Hill

Training Faculty 2014-current

**Toxicology** 

North Carolina State University

2013-current Training Faculty

Genetics

North Carolina State University

2013-current **Training Faculty** 

**Statistics** 

North Carolina State University

Training Faculty 2013-current

**Bioinformatics** 

North Carolina State University

2013-current Training Faculty

Functional Genomics North Carolina State University

Faculty Mentor & Computational Lab Director

2011-2012 Computation for Undergraduates in Statistics Program

National Science Foundation (PI: Sujit Ghosh)

2010-2011 Faculty Mentor & Computational Lab Director

Computation for Undergraduates in Statistics Program

National Science Foundation (PI: Sujit Ghosh)

#### MENTORING AND MANAGEMENT

#### Graduate Students Mentored (As Primary Supervisor)

2021-current Jessie Chappel (joint with Erin Baker, UNC Chapel Hill)

Bioinformatics PhD (expected 2023)

Page 6 (of 22) David M. Reif North Carolina State University

2020-current Nnamdi Osakwe

Bioinformatics PhD (expected 2023) North Carolina State University

2020-current Jonathan Fleming

Bioinformatics PhD (expected 2023) North Carolina State University

2019-current Preethi Thunga

Bioinformatics PhD (expected 2022) North Carolina State University

2019-2022 Dylan Wallis

Toxicology PhD (graduated) North Carolina State University

2019-2021 Melody Hancock

Bioinformatics MS (graduated; currently Bioinformatics PhD student)

North Carolina State University

2017-current Aldo Carmona-Baez (joint with Reade Roberts)

Genetics PhD (expected 2020) North Carolina State University

2016-2019 Marissa Kosnik

Toxicology PhD (graduated) North Carolina State University

2015-2019 Kimberly To

Bioinformatics PhD (graduated) North Carolina State University

2014-2018 Kyle Roell

Bioinformatics PhD (graduated) North Carolina State University

2014-2017 Michele Balik-Meisner

Bioinformatics PhD (graduated) North Carolina State University

2013-2016 Guozhu Zhang

Bioinformatics PhD (graduated) North Carolina State University

2013 Ankita Kalantri

Department of Computer Science MS (graduated)

North Carolina State University

### Postdoctoral Scholars Mentored (As Primary Supervisor)

2019-2022 Adrian Green, PhD

Bioinformatics Research Center North Carolina State University

2012 Lisa Truong, PhD

National Center for Computational Toxicology

U.S. Environmental Protection Agency

2012 Cory Strope, PhD

National Center for Computational Toxicology

U.S. Environmental Protection Agency

# Scientific Staff Management (As Direct Supervisor)

2022-current Scientists and Staff,

Predictive Toxicology Branch, Division of Translational Toxicology

National Institute of Environmental Health Sciences

2018-2022 Alison Dickey (PhD, Research Associate)

Page 7 (of 22)

David M. Reif

	Bioinformatics Consulting and Service Core North Carolina State University
2018-2021	Elizabeth Scholl (PhD, Research Scholar) Bioinformatics Consulting and Service Core
	North Carolina State University
2014-2022	Skylar Marvel (PhD, Research Associate)
	Department of Biological Sciences
	North Carolina State University
2015-2016	Galen Collier (PhD, Research Scholar)
	Department of Biological Sciences
	North Carolina State University
2012	Sean Watford (BS, Research Fellow)
	National Center for Computational Toxicology US EPA
2011-2012	Dayne Filer (BS, Research Fellow)
	National Center for Computational Toxicology US EPA
2011-2012	Parth Kothiya (MS, Research Fellow)
	National Center for Computational Toxicology US EPA

### Graduate Student Committees (\*Chair or Co-Chair)

\* Guozhu Zhang, Bioinformatics PhD; North Carolina State University

\* Michele Balik-Meisner, Bioinformatics PhD; North Carolina State University

Rachel Spreng, Bioinformatics, PhD; North Carolina State University

Ryan Lougee, Toxicology, MS; North Carolina State University

\* Kyle Roell, Bioinformatics PhD; North Carolina State University

\* Kimberly To, Bioinformatics PhD; North Carolina State University

Sarah Wisotsky, Bioinformatics, PhD; North Carolina State University

Hsieh (Larry) Wu, Bioinformatics, MS; North Carolina State University

Maller (Edity) vvd, Bisinormatics, Nic, North Carolina State Criversity

William Kohlway, Bioinformatics PhD; North Carolina State University

Ander Wilson, Statistics PhD; North Carolina State University

Ravi Mathur, Bioinformatics PhD; North Carolina State University

Baljinder Kaur, Crop Sciences PhD; North Carolina State University

Sean Watford, Environmental Science and Engineering PhD; University of North Carolina

\* Tao Jiang, Bioinformatics PhD; North Carolina State University

Patrick Perkins, Bioinformatics PhD; North Carolina State University

Erin Peterson Genetics PhD; North Carolina State University

\* Jun Ma, Bioinformatics PhD; North Carolina State University

Brandon Baker, Genetics PhD; North Carolina State University

Hayden Brochu, Bioinformatics MS; North Carolina State University

Bethany Cook, Chemistry MS; North Carolina State University

\* Aldo Carmona-Baez, Genetics PhD; North Carolina State University

Shuping Ryan, Bioinformatics MS; North Carolina State University

\* Marissa Kosnik, Toxicology PhD; North Carolina State University

Desiree Unselt, Genetics PhD; North Carolina State University

Yaxu Wang, Bioinformatics PhD; North Carolina State University

Drake Phelps, Veterinary School PhD; North Carolina State University

Matthew Nethery, Functional Genomics PhD; North Carolina State University

Brian Ting, Statistics PhD; North Carolina State University

\* Melody Hancock, Bioinformatics MS; North Carolina State University

Mark Simmers, Toxicology PhD; North Carolina State University

Sagi Guillerra, Toxicology PhD; North Carolina State University

Melanie Odenkirk, Chemistry PhD; North Carolina State University

\* Preethi Thunga, Bioinformatics PhD; North Carolina State University

Page 8 (of 22) David M. Reif

\* Dylan Wallis, Toxicology PhD; North Carolina State University

- \* Jonathon Fleming, Bioinformatics PhD; North Carolina State University
- \* Nnamdi Osakwe, Bioinformatics PhD; North Carolina State University

Yueyang Huang, Statistics PhD; North Carolina State University

Thomas Howard, Bioinformatics MS: North Carolina State University

Jacob Freudenberg, Bioinformatics MS; North Carolina State University

William Marinello, Toxicology PhD; North Carolina State University

Michael Doyle, Chemistry PhD; North Carolina State University

Hannah Starnes, Toxicology PhD; North Carolina State University

Amanda Brucker, Statistics PhD; North Carolina State University

Megan Dillon, Comparative Biomedical Sciences PhD; North Carolina State University

Melody Hancock, Bioinformatics PhD; North Carolina State University

#### **FUNDING AND BUDGET MANAGEMENT**

# External (Competitive Grant) Research Support

NIH/NIEHS R01 R01ES033243

07/01/2022 - 04/30/2027

Characterizing gene-environment interactions that affect individual susceptibility to an expanding chemical exposome

Role: PI

NIH/NIEHS P42 Diversity Supplement

01/01/2022 - 12/31/2023

Gene-environment interactions causing differential susceptibility to chemical stressors in an expanding chemical exposome

Role: PI of Supplement

NIH/NIEHS P30 ES-025128

04/1/2015 - 03/31/2026

Center for Human Health and the Environment (CHHE) Role: Co-Director of IHSFC & Co-I (PI: Jane Hoppin, NCSU)

NIH/NIEHS U01 Supplement

10/01/2021 - 08/31/2022

Integrative Machine Learning for Synthesis of Cross-Consortium ENM Data

Role: PI of Supplement

NIH/NIEHS U01 ES-027294

09/01/2016 - 08/31/2022

Multidimensional in vivo Assessments of Engineered Nanomaterials and Biological Interactions

Role: PI of Sub-contract (PI: Robert Tanguay, Oregon State University)

EPA R83948101 01/01/2019 – 12/31/2022

System toxicological approaches to define and predict the toxicity of Per and Polyfluoroalkyl Substances

Role: PI of Sub-contract (PI: Robert Tanguay, Oregon State University)

NIH/NCI R01 CA-161608 04/01/2019 – 03/31/2023

Genetic Etiology of Cancer Drug Response

Role: PI

NIH/EPA P42 ES-031009 04/1/2020 – 03/31/2024

Superfund Center for Environmental and Health Effects of PFAS Role: Director of DMAC & Co-I (PI: Carolyn Mattingly, NCSU)

NIH RIVER R35 ES-031709

04/1/2021 - 03/31/2029

Discovering Chemical Activity Networks - Predicting Bioactivity Based on Structure

Role: PI of Sub-contract (PI: Robert Tanguay, Oregon State University)

### Completed Research Support

Page 9 (of 22) David M. Reif

NIH R01 ES-19604 10/01/2014 – 09/30/2015

Integrating Big Data and curated literature to advance discoveries about disease

Role: Co-Investigator (PI: Carolyn Mattingly, NCSU)

University Global Partnership Network (UPGN)

07/01/2016 -- 06/30/2017

An international alliance for Population, Wellbeing and Environment Research

Role: Co-Investigator (PI: Jane Hoppin, North Carolina State University)

NIH P42 ES-005948 10/01/2014 – 09/30/2017

Elucidating Risks: From Exposure and Mechanism to Outcome

Role: Co-Investigator (PI: James Swenberg, University of North Carolina at Chapel Hill)

EPA EPA-G2014-STAR-E1

06/01/2015 - 06/30/2018

System toxicological approaches to define flame retardant adverse outcome pathways

Role: PI of Sub-contract (PI: Robert Tanguay, Oregon State University)

EPA G2013-STAR-L1

09/01/2014 - 08/31/2018

Cardiotoxicity Adverse Outcome Pathway: organotypic culture model and in vitro/in vivo extrapolation for high-throughput hazard, dose-response and variability assessments

Role: Co-Investigator (PIs: Ivan Rusyn and David Threadgill, Texas A&M University; Fred Wright, NCSU)

NIH R01 ES-023788 01/01/2015 – 12/31/2019

Advancing mechanism-based studies with cross-species chemical-phenotype data

Role: Co-Investigator (PI: Carolyn Mattingly, NCSU)

NIH/NICEATM Contract

07/01/2018 - 06/30/2019

Integrated Data Analysis for Systematic Evaluation of the Application of Zebrafish in Toxicology

Role: PI

CalEPA Contract 09/01/2018 – 02/28/2019

Bioinformatic and cheminformatic modeling of perfluorinated compounds

Role: PI

Texas A&M Superfund Research Program, Pilot Project

10/31/2018 - 10/31/2019

Translation of multi-stream data into interactive visual profiles

Role: PI

NIH/NIEHS R56 ES-0300007

06/15/2019 - 06/14/2021

Gene-environment interactions causing differential susceptibility to chemical stressors in high-throughput data

Role: PI

NIH/NIEHS R15

06/01/2018 - 03/31/2021

Prenatal Supplementation Reduces the Severity of Toxicant-Induced Birth Defects

Role: Co-Investigator (PI: Krista McCoy, East Carolina University)

Federal Budget and Contracts Management

Predictive Toxicology Branch,

2022 - current

Division of Translational Toxicology,

National Institute of Environmental Health Sciences

Cooperative Research and Development Agreement (CRADA)

2012 - 2013

L'Oréal – U.S. Environmental Protection Agency

Role: PI

Page 10 (of 22)

David M. Reif

Presidential Early Career Award for Scientists and Engineers (PECASE) 2011 – 2012

Office of The White House

Role: PI

ToxCast *in vitro* assay contracts (multiple external vendors) 2009 – 2012

U.S. Environmental Protection Agency

Role: Contract Officer Representative (COR) / Task Order Contract Officer (TOCOR)

### **DISTRIBUTED SOFTWARE**

toxpiR: Open-source R package for ToxPi

(Public software and Github documentation)

https://cran.r-project.org/package=toxpiR

https://toxpi.github.io/toxpiR/

ToxPi\*GIS Toolkit: Software suite to create customized ToxPi interactive layers within ArcGIS

(Public software toolkit and Github documentation)

https://github.com/Jonathon-Fleming/ToxPi-GIS

COVID19 Pandemic Vulnerability Index (PVI) Dashboard: Web application for analytics of data streams

related to COVID-19

(Public web application and open source data repository)

https://covid19pvi.niehs.nih.gov

ToxPi\*GIS: Dashboard for ToxPi built atop geographic information system (GIS) layers

(Public web application)

https://toxpi.org/covid-19/map/

ToxPi GUI: Stand-alone Java application for building ToxPi models

(Free download)

https://toxpi.org/

MDR R: R implementation of Multifactor Dimensionality Reduction (MDR) modeling and internal validation (Free download)

http://cran.r-project.org/web/packages/MDR/index.html

zfish GUI: Graphical User Interface (GUI) implementation of analysis and graphics pipeline for high-

throughput, zebrafish experimental data

(Shiny package for collaborators)

zfish DB: Web-enabled browser interface for access to results and meta-analysis for high-throughput,

zebrafish experimental data from multiple labs

(Password-limited to collaborators)

## EXTENSION, SERVICE, COMMUNITY OUTREACH, AND MEDIA

2021 Interviewee, "At-home COVID-19 tests offer quick results but can go uncounted in NC

totals", ABC11 News

2021 Interviewee, "Biostatistics & Computational Biology in COVID-19", COVIDCalls (Apple

Podcasts)

Interviewee, "How do vaccines impact an area's risk for COVID-19?", ABC11 News
Member, Carcinogenicity Gene Signature Development, HESI eSTAR Consortium
Member, Carcinogenicity Gene Signature Development, HESI eSTAR Consortium
Advisory Committee, Pacific Northwest Center for Translational Environmental Health

Research, Oregon State University

Page 11 (of 22)

David M. Reif

2020-current 2020-current	Advisory Board, College of Sciences Mentorship Program, North Carolina State University Co-Chair, Data Working Group, U.S. Food and Drug Administration (FDA) Botanical Safety Consortium
2020	Panelist, DR2 Work Group SARS-CoV-2/COVID-19 Environmental Health Research Needs Panel (NIEHS)
2019 2019	Interviewee, "Why Anti-vax Doctors Are Ordering 23andMe Tests", <i>The Atlantic</i> Interviewee, "Daily Sports Feed – David Reif", ESPN Radio
2018-current	Faculty Workloads and Rewards Program, NC State University [committee member]
2017 2016	Undergraduate Education Seminar Series, Society of Toxicology (SOT) [speaker] Biomedical Career Symposium, National Institutes of Environmental Health Sciences (NIEHS) [invited speaker]
2016	Bioinformatics short course (with Nadia Singh and Dahlia Nielsen), Fayetteville State University, Research Initiative for Scientific Enhancement (RISE) program
2014	NCSU Genetics Club student-invited speaker
2013-2015	Search Committee, "Bioinformatics Faculty Cluster Hire", North Carolina State University
2008-2012	Program Committee, "Bioinformatics, Computational, and Systems Biology", Genetic and Evolutionary Computation Conference (GECCO)
2011	Guest/Interviewee, Radio In Vivo, WCOM-FM, Carrboro, NC
2011	Guest/Interviewee, Federal News Radio, Washington, DC
2009-2011	Chair, NCCT Seminar Series, Research Triangle Park, NC, USA
2010	EPA Volunteer/Presenter, US Science & Engineering Festival on the National Mall, Washington, DC
2009	"Gene-Environment Interactions", Science Café presented by the North Carolina Museum of Science, Raleigh, NC, USA [public seminar]
2009	"Lessons on Modern Toxicology: How Darwin Saw It Coming", EPA <i>Greenversations</i> weblog
2008-2009	Career Panel, Durham Public Schools "Stay in School" program, Durham, NC, USA
2008	"Human Genetics", EPA-Shaw University Research Associateship Program, Chapel Hill, NC, USA [public seminar]
2007	"Genotyping technology and analysis in cancer research", Association for Biomedical Research, Chapel Hill, NC, USA [public seminar]
2006-2008	EPA Speaker's Bureau, Multiple presentations in high schools on research and careers in science, Cary, NC; Chapel Hill, NC; Durham, NC; Raleigh, NC; Mishawaka, IN

# PRESENTATIONS AND INVITED WORKSHOPS

2021	Genetics and Genomics Initiative (GGI) Seminar Series, "Characterizing Gene- Environment Interactions in an Expanding Chemical Exposome" [invited seminar]
2020	Society of Toxicology (SOT) Risk Assessment Specialty Section Webinar, "Fighting COVID-19 using data science" [invited seminar]
2020	UC Berkeley Superfund, "The Problem of PFAS Contamination: How Can We Make Rapid Progress to Address it?", Berkeley, CA, USA [invited seminar and session chair]
2019	International Neurotoxicology Association, "Systems neurotoxicology: From in vivo towards predictive in vitro models", Dusseldorf, GERMANY, [seminar]
2018	Texas A&M Superfund External Advisory Board Meeting, "ToxPi-GIS: Visual analytics for integrating hazard and geospatial data", College Station, TX, USA [presentation]
2018	ToxCast Seminar, "Behavioral assessment using zebrafish", Durham, NC, USA [invited seminar]
2018	Fish and Amphibian Embryo as Alternative Models in Toxicology and Teratology, "Leveraging high-throughput screening data to find evidence for susceptibility differences in genetically diverse zebrafish", Paris, FRANCE [invited seminar]
2018	Biology Seminar Series, East Carolina University, "Searching for GxE in high-throughput data", Greenville, NC, USA [invited seminar]
2018	Principles of Data Visualization Workshop, "Data Visualization: Effective Communication Across Disciplines", Raleigh, NC, USA [invited seminar].

Page 12 (of 22)

David M. Reif

2017	OpenTox USA Meeting, "Implications of Population Genetic Diversity for Safety Assessment", Durham, NC, USA [invited seminar]
2017	Environmental Mutagenesis and Genomics Society (EMGS), "Gene-Environment Interactions Associated with Differential Susceptibility to Chemical Exposure", Raleigh, NC, USA [invited seminar]
2017	International Neurotoxicology Association, "Analytical methods to distill useful information from high-throughput screening of chemicals", Florianopolis, BRAZIL [seminar]
2017	National Institutes of Health Workshop, "SEAZIT: Overview of zebrafish screening data", Durham, NC, USA [invited seminar]
2017	University of Wollongong, "Bioinformatics for Linking Social and Environmental Determinants of Health", Wollongong, AUSTRALIA [invited seminar]
2017	Tox Forum, "Integrating multiple sources of high-dimensional data to quantify effects of genetic variability in chemical risk assessment", Washington, DC, USA [invited seminar]
2016	Division of Extramural Research and Training (DERT) Data Science Retreat, "Applied Data Science for the Basic Sciences", NIEHS, RTP, NC, USA [keynote seminar]
2016	Aggregate Exposure Pathway: A conceptual framework to support exposure science research and complete the source-to-outcome continuum for risk assessment. U.S. EPA, Durham, North Carolina, USA [invited expert]
2016	New Approach Methodologies in Regulatory Science. European Chemicals Agency (ECHA), Helsinki, FINLAND [invited workshop]
2016	Society of Toxicology Annual Meeting, "Beyond Benchmark Dose: Advancing Probabilistic and Bayesian Approaches in Hazard Characterization", New Orleans, LA, USA [seminar]
2015	North Carolina Department of Health and Human Services (NCDHHS), "Computational Toxicology Resources for Public Health Applications", Raleigh, NC, USA [seminar]
2015	Data Science Seminar Series, "Big Data in Environmental Bioinformatics", National Institute of Environmental Health Sciences (NIEHS), RTP, NC, USA [seminar]
2015	"Environmental Bioinformatics", Oregon State University, Corvallis, OR, USA [seminar]
2015	PopMed Forum, "Big Data and High-Throughput Screening in Environmental Toxicology", College of Veterinary Medicine, North Carolina State University, Raleigh, NC, USA [seminar]
2014	Workshop on fish and amphibian embryos as alternative models in toxicology and teratology, "Rapid identification and characterization of chemical bioactivity using an embryonic zebrafish system", Paris, FRANCE [seminar]
2014	"Big Data in Environmental Toxicology", University of Kentucky, Lexington, KY, USA [seminar]
2014	A Collaborative Workshop on Aquatic Models and 21st Century Toxicology, "Rapid identification and characterization of neuromodulator chemicals using an embryonic zebrafish system", NIH/NCSU, Raleigh, NC, USA [seminar]
2014	Workshop on Big Data in Genetics and Toxicogenomics, "ToxPi and other visualization tools for toxicological data", NC Biotechnology Center, Durham, NC, USA [seminar]
2013	Triangle Statistical Genetics Consortium, "High-throughput characterization of behavioral and developmental zebrafish phenotypes in response to chemical exposure: Setting the stage for GxE studies", Duke University, Durham, NC, USA [seminar]
2013	Laboratory for Analytical Sciences (LAS), "Visualization for Interdisciplinary Communication", North Carolina State University, Raleigh, NC, USA [seminar]
2012	Application of Toxicity Testing in the 21 <sup>st</sup> Century – Beyond Environmental Chemicals, Johns Hopkins Center for Alternatives to Animal Testing (CAAT), Ranco, ITALY [invited participant/presenter]
2012	Building for Better Decisions: Multi-Scale Integration of Human Health and Environmental Data, Society of Toxicology, Research Triangle Park, NC, USA [invited participant/presenter]
2011	International Chemical Regulation REACH Workshop Asia, Hangzhou, CHINA [seminar]
2011	Special Awards Seminar, U.S. EPA Office of Science Coordination and Policy, Washington, DC, USA [seminar]
2011	Chemical Prioritization Community of Practice, U.S. NIH & U.S. EPA, Research Triangle Park, NC, USA [seminar]

Page 13 (of 22)

David M. Reif

2011	NIH Workshop on the Role of Environmental Chemicals in the Development of Diabetes and Obesity, NIH/NIEHS, Durham, NC, USA [invited participant/presenter]
2010	Office of Chemical Safety and Pollution Prevention Seminar Series, U.S. EPA, Washinton, DC, USA [seminar]
2010	Advancing the Next Generation (NexGen) of Risk Assessment: The Prototypes, National Center for Environmental Assessment, Research Triangle Park, NC, USA [seminar]
2010	Center for Environmental Health Studies, Dartmouth College, Hanover, NH, USA [seminar]
2010	Biomolecular Screening Seminar Series, U.S. NIEHS/National Toxicology Program, Research Triangle Park, NC, USA [seminar]
2009	Chemical Prioritization Community of Practice, U.S. NIH & U.S. EPA, Research Triangle Park, NC, USA [seminar]
2009	ToxCast Data Analysis Summit, U.S. EPA, Research Triangle Park, NC, USA [seminar]
2008	Genetics Department Seminar Series, North Carolina State University, Raleigh, NC, USA [seminar]
2008	Visiting Pulmonary Scholar Seminar Series, University of North Carolina, Chapel Hill, NC, USA [seminar]
2008	International Society of Exposure Analysis-International Society of Environmental Epidemiology (ISEA-ISEE) Joint Annual Meeting, Pasadena, CA, USA [platform presentation]
2008	Human Studies Division Seminar Series, U.S. EPA, Chapel Hill, NC, USA [seminar]
2007	Therapeutic Applications of Computational Biology and Chemistry (TACBAC), Wellcome Trust Conference Centre, Cambridge, UK [seminar]
2007	Community Based Risk Assessment Workshop, "Statistical and Mathematical Modeling for Community-Based Risk Assessment", National Center for Environmental Research, Research Triangle Park, NC, USA [session chair]
2006	Vanderbilt University Genetics Interest Group, Center for Human Genetics Research, Nashville, TN, USA [seminar]
2005 2003	Pacific Symposium on Biocomputing (PSB), Lihue, HI, USA [platform presentation] Genetic and Evolutionary Computation Conference (GECCO), Chicago, IL, USA [platform presentation]

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## Peer-Reviewed Publications (link to Google Scholar profile)

- 1. Thunga P, Truong L, Rericha Y, Du J, Morshead M, Tanguay RL, Reif DM. Utilizing a Population-Genetic Framework to Test for Gene-Environment Interactions between Zebrafish Behavior and Chemical Exposure. Toxics. 2022 Dec 9;10(12):769. doi: 10.3390/toxics10120769. PMID: 36548602; PMCID: PMC9781692.
- 2. Watson ATD, Carmona Baez A, Jima D, Reif D, Ding J, Roberts R, Kullman SW. TCDD alters essential transcriptional regulators of osteogenic differentiation in multipotent mesenchymal stem cells. Toxicol Sci. 2022 Nov 12:kfac120. doi: 10.1093/toxsci/kfac120. Epub ahead of print. PMID: 36370075.
- 3. Carberry CK, Koval LE, Payton A, Hartwell H, Ho Kim Y, Smith GJ, Reif DM, Jaspers I, Ian Gilmour M, Rager JE. Wildfires and extracellular vesicles: Exosomal MicroRNAs as mediators of cross-tissue cardiopulmonary responses to biomass smoke. Environ Int. 2022 Sep;167:107419. doi: 10.1016/j.envint.2022.107419. Epub 2022 Jul 16. PMID: 35863239; PMCID: PMC9389917.
- 4. Wolkin A, Collier S, House JS, Reif D, Motsinger-Reif A, Duca L, Sharpe JD. Comparison of National Vulnerability Indices Used by the Centers for Disease Control and Prevention for the COVID-19 Response. Public Health Rep. 2022 Jul-Aug;137(4):803-812. doi: 10.1177/00333549221090262. Epub 2022 May 5. Erratum in: Public Health Rep. 2022 Sep-Oct;137(5):1041. PMID: 35514159; PMCID: PMC9257512.
- 5. Roell K, Koval LE, Boyles R, Patlewicz G, Ring C, Rider CV, Ward-Caviness C, Reif DM, Jaspers I, Fry RC, Rager JE. Development of the InTelligence And Machine LEarning (TAME) Toolkit for Introductory Data Science, Chemical-Biological Analyses, Predictive Modeling, and Database Mining

Page 14 (of 22)

David M. Reif

- for Environmental Health Research. Front Toxicol. 2022 Jun 22;4:893924. doi: 10.3389/ftox.2022.893924. PMID: 35812168; PMCID: PMC9257219.
- 6. Wallis DJ, La Du J, Thunga P, Elson D, Truong L, Kolluri SK, Tanguay RL, Reif DM. Leveraging a High-Throughput Screening Method to Identify Mechanisms of Individual Susceptibility Differences in a Genetically Diverse Zebrafish Model. Front Toxicol. 2022 Apr 29;4:846221. doi: 10.3389/ftox.2022.846221. PMID: 35573279; PMCID: PMC9098949.
- 7. Thessen AE, Marvel S, Achenbach JC, Fischer S, Haendel MA, Hayward K, Klüver N, Könemann S, Legradi J, Lein P, Leong C, Mylroie JE, Padilla S, Perone D, Planchart A, Prieto RM, Muriana A, Quevedo C, Reif D, Ryan K, Stinckens E, Truong L, Vergauwen L, Vom Berg C, Wilbanks M, Yaghoobi B, Hamm J. Implementation of Zebrafish Ontologies for Toxicology Screening. Front Toxicol. 2022 Mar 11;4:817999. doi: 10.3389/ftox.2022.817999. PMID: 35387429; PMCID: PMC8979167.
- 8. Fleming J, Marvel SW, Supak S, Motsinger-Reif AA, Reif DM. ToxPi\*GIS Toolkit: creating, viewing, and sharing integrative visualizations for geospatial data using ArcGIS. J Expo Sci Environ Epidemiol. 2022 Apr 26:1–8. doi: 10.1038/s41370-022-00433-w. Epub ahead of print. PMID: 35474345; PMCID: PMC9039976.
- 9. Shankar P, Garcia GR, La Du JK, Sullivan CM, Dunham CL, Goodale BC, Waters KM, Stanisheuski S, Maier CS, Thunga P, Reif DM, Tanguay RL. The Ahr2-Dependent wfikkn1 Gene Influences Zebrafish Transcriptome, Proteome, and Behavior. Toxicol Sci. 2022 May 26;187(2):325-344. doi: 10.1093/toxsci/kfac037. PMID: 35377459; PMCID: PMC9308396.
- 10. Truong L, Rericha Y, Thunga P, Marvel S, Wallis D, Simonich MT, Field JA, Cao D, Reif DM, Tanguay RL. Systematic developmental toxicity assessment of a structurally diverse library of PFAS in zebrafish. J Hazard Mater. 2022 Jun 5;431:128615. doi: 10.1016/j.jhazmat.2022.128615. Epub 2022 Mar 2. PMID: 35263707: PMCID: PMC8970529.
- 11. Kirkwood KI, Fleming J, Nguyen H, Reif DM, Baker ES, Belcher SM. Utilizing Pine Needles to Temporally and Spatially Profile Per- and Polyfluoroalkyl Substances (PFAS). Environ Sci Technol. 2022 Mar 15;56(6):3441-3451. doi: 10.1021/acs.est.1c06483. Epub 2022 Feb 17. PMID: 35175744; PMCID: PMC9199521.
- 12. Hubal EAC, DeLuca NM, Mullikin A, Slover R, Little JC, Reif DM. Demonstrating a systems approach for integrating disparate data streams to inform decisions on children's environmental health. BMC Public Health. 2022 Feb 15;22(1):313. doi: 10.1186/s12889-022-12682-3. PMID: 35168583; PMCID: PMC8845296.
- 13. Akhtari FS, Green AJ, Small GW, Havener TM, House JS, Roell KR, Reif DM, McLeod HL, Wiltshire T, Motsinger-Reif AA. High-throughput screening and genome-wide analyses of 44 anticancer drugs in the 1000 Genomes cell lines reveals an association of the NQO1 gene with the response of multiple anticancer drugs. *PLoS Genet*. 2021 Aug 26;17(8):e1009732.
- Green AJ, Mohlenkamp MJ, Das J, Chaudhari M, Truong L, Tanguay RL, Reif DM. Leveraging highthroughput screening data, deep neural networks, and conditional generative adversarial networks to advance predictive toxicology. *PLoS Comput Biol.* 2021 July 02; 17(7): e1009135.
- 15. Thunga P, Truong L, Tanguay RL, Reif DM. Concurrent evaluation of mortality and behavioral responses: A fast and efficient testing approach for high-throughput chemical hazard identification. *Frontiers in Toxicology.* 2021 Jun 15; 3.
- Odenkirk MT, Reif DM, Baker ES. Multiomic big data analysis challenges: Increasing confidence in the interpretation of artificial intelligence assessments. *Anal Chem.* 2021 June 08; 93(22): 7763-7773.
- 17. Green AJ, Anchang B, Akhtari FS, Reif DM, Motsinger-Reif A. Extending the lymphoblastoid cell line model for drug combination pharmacogenomics. *Pharmacogenomics*. 2021 June 01; 22(9): 543-551.
- 18. Marvel SW, House JS, Wheeler M, Song K, Zhou Y, Wright FA, Chiu WA, Rusyn I, Motsinger-Reif A, Reif DM. The COVID-19 pandemic vulnerability index (PVI) dashboard: Monitoring county-level vulnerability using visualization, statistical modeling, and machine learning. *Environ Health Perspect.* 2021 jan; 129(1): 17701.
- 19. Gilchrist PO, Alexander AB, Green AJ, Sanders FE, Hooker AQ, Reif DM. Development of a pandemic awareness STEM outreach curriculum: Utilizing a computational thinking taxonomy framework. *Education Sciences*. 2021; 11(3).

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- 21. Cohen Hubal Elaine A, Reif DM, Slover R, Mullikin A, Little JC. Children's environmental health: A systems approach for anticipating impacts from chemicals. *International journal of environmental research and public health.* 2020 nov; 17(22): 8337.
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Page 16 (of 22)

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Page 19 (of 22)

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Page 22 (of 22)

David M. Reif