

**Curriculum Vitae  
Robert W. Corty**

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updated: February 10, 2022

**EDUCATION**

<b>residency</b>	Vanderbilt University Medical Center <b>specialty:</b> internal medicine	<b>June 2022 (expected)</b>
<b>MD</b>	University of North Carolina at Chapel Hill	<b>May 2020</b>
<b>PhD</b>	University of North Carolina at Chapel Hill curriculum: Bioinformatics and Computational Biology <b>dissertation:</b> Variance Heterogeneity in Genetic Mapping	<b>August 2018</b>
<b>AB</b>	Harvard University, Cambridge, Massachusetts concentration: Chemical and Physical Biology <b>thesis:</b> Studies on the Oligomerization of the Dengue Envelope Protein	<b>May 2010</b>

**STANDARDIZED TESTING**

USMLE Step 1: 231  
USMLE Step 2 CK: 251  
USMLE Step 2 CS: pass  
USMLE Step 3: 236

**RESEARCH**

**Independent** **2018 – 2020**

I investigated the effect of antibiotics on the risk for developing gemcitabine-associated toxicities by re-purposing the data from clinical trial of patients with stage IV pancreatic cancer.

**Supervised by Dr. William Valdar** **2013 – 2018**

**Dissertation project:** Variance Heterogeneity in Genetic Mapping

I investigated the use of the double generalized linear model in genetic mapping of experimental crosses. I found that this model has two advantages as compared to traditional approaches: (1) it re-weights observations based on their empirical precision and (2) it detects genetic loci that influence residual variance, so-called “vQTL”.

**Related project:** I derived a novel procedure for efficiently fitting a linear mixed model with arbitrary genetic covariance and heterogeneous residual variance, which has applications in genetic mapping.

**Supervised by Dr. Stephen Harrison 2007 – 2010**

I designed, cloned, expressed, purified, and refolded protein constructs to recapitulate various steps of process by which the dengue virus fuses its membrane with a host cell. This research was supported by HHMI and NIH and led to my senior thesis, which won the Thomas Temple Hoopes prize for outstanding undergraduate work.

**GRANT SUPPORT**

**University Cancer Research Fund MD/PhD Scholarship** **2019 – 2020**

**National Research Service Award:** F30MH108265 **2016 – 2019**  
title: Statistical Modeling of Genetic Effects on Behavior and its Variability

<b>UNC Big Data to Knowledge training grant:</b> T32LM012420	<b>2015 – 2016</b>
<b>UNC Bioinformatics and Computational Biology training grant:</b> T32GM067553	<b>2014 – 2014</b>
<b>UNC Medical Scientist Training Program:</b> T32GM008719	<b>2011 – 2012</b>

## PUBLICATIONS

1. **Corty, RW**, Langworthy BW, Fine JP, Buse JB, Sanoff HK, and Lund JL. Antibacterial use is associated with an increased risk of hematologic and gastrointestinal adverse events in patients treated with gemcitabine for stage IV pancreatic cancer. *Oncologist* 2020;25.
2. Mosedale M, Cai Y, Eaddy JS, **Corty, RW**, Nautiyal M, Watkins PB, and Valdar W. Identification of Candidate Risk Factor Genes for Human Idelalisib Toxicity Using a Collaborative Cross Approach. *Toxicological Sciences* 2019;172:265–78.
3. **Corty, RW** and Valdar W. QTL Mapping on a Background of Variance Heterogeneity. *G3: Genes, Genomes, Genetics* 2018;8:3767–82.
4. **Corty, RW** and Valdar W. vqtl: An R package for Mean-Variance QTL Mapping. *G3: Genes, Genomes, Genetics* 2018;8:3757–66.
5. **Corty, RW**, Kumar V, Tarantino LM, Takahashi JS, and Valdar W. Mean-Variance QTL Mapping Identifies Novel QTL for Circadian Activity and Exploratory Behavior in Mice. *G3: Genes, Genomes, Genetics* 2018;8:3783–90.
6. Mosedale M, Kim Y, Brock WJ, Roth SE, Wiltshire T, Eaddy JS, Keele GR, **Corty, RW**, Xie Y, Valdar W, et al. Editor's Highlight: Candidate Risk Factors and Mechanisms for Tolvaptan-Induced Liver Injury Are Identified Using a Collaborative Cross Approach. *Toxicological Sciences* 2017;156:438–54.
7. Green AK\*, **Corty, RW**, Wood WA, Meenaghan M, Reeder-Hayes KE, Basch E, Milowsky MI, and Dusetzina SB. Comparative effectiveness of mitoxantrone plus prednisone versus prednisone alone in metastatic castrate-resistant prostate cancer after docetaxel failure. *The Oncologist* 2015;20:516–22.
8. Green AK, Reeder-Hayes KE, **Corty, RW**, Basch E, Milowsky MI, Dusetzina SB, Bennett AV, and Wood WA. The project data sphere initiative: accelerating cancer research by sharing data. *The Oncologist* 2015;20:464–e20.
9. Matson BC, **Corty, RW**, Karpnich NO, Murtha AP, Valdar W, Grotegut CA, and Caron KM. Midregional pro-adrenomedullin plasma concentrations are blunted in severe preeclampsia. *Placenta* 2014;35:780–3.
10. Corty EW and **Corty, RW**. Setting sample size to ensure narrow confidence intervals for precise estimation of population values. *Nursing Research* 2011;60:148–53.

## ORAL PRESENTATIONS

<b>UNC Quality Improvement Symposium</b> title: Equalizing the Workload Among UNC Hospitalists	<b>2020</b>
<b>Dissertation Defense</b> title: Variance Heterogeneity in Genetic Mapping	<b>2018</b>
<b>MD-PhD Association for Clinical Research Outcomes</b> title: Patients taking gemcitabine for metastatic pancreatic cancer have an increased risk of gemcitabine toxicity while on incidental antibiotics.	<b>2018</b>

<b>MS3 Quality Improvement Project</b> title: Improving Hypertension Management at Moncure Community Health Center	<b>2018</b>
<b>UNC Bioinformatics Colloquium</b> title: Recognizing Idiosyncrasy in Environmental Variation to Improve QTL Mapping	<b>2017</b>
<b>Triangle Statistical Genetics Conference</b> title: Two Reasons to Model Phenotype Mean and Variance in QTL Mapping	<b>2016</b>
<b>MD-PhD student session at the Annual Molecular Psychiatry Association Meeting</b> title: Variance Variants	<b>2014</b>

#### UNIVERSITY SERVICE

MD-PhD admissions committee member	<b>2016 – 2019</b>
student representative for Bioinformatics and Computation Biology curriculum recruitment	<b>2016 – 2017</b>
team captain in MD-PhD program “vertical mentorship” system	<b>2015 – 2017</b>
John B. Graham society for medical student research, secretary	<b>2014 – 2016</b>
guest lecturer for high dimensional data analysis course	<b>2015</b>
teaching assistant for Introduction to Statistical Modeling	<b>2014</b>

#### COMMUNITY SERVICE

organizer and speaker, NAACP Historic Thousands on Jones Street	<b>2016 – 2017</b>
volunteer, Homestart residential facility for women and children	<b>2012 – 2014</b>
guest teacher, Brier Creek Elementary School wellness week	<b>2013</b>
intake coordinator, St. Petersburg Free Clinic	<b>2010 – 2011</b>

#### POSTER PRESENTATIONS

\* = presenter

<b>UNC Pediatrics Day of Scholarship, Chapel Hill, NC</b> title: Seize the diagnosis authors: Pamela Londres*, <b>Corty, RW</b> , and William Mills	<b>2019</b>
<b>Society of Toxicology Annual Meeting, Baltimore, MD</b> title: Identification of Genetic Risk Factors for Human Idelalisib Toxicity Using a Collaborative Cross Approach authors: Merrie Mosedale*, Yanwei Cai, J Scott Eaddy, <b>Corty, RW</b> , William Valdar, and Paul Watkins	<b>2019</b>
<b>UNC Quality Improvement Dinner</b> title: Optimizing Blood Pressure Recheck Protocol authors: Amir Ghodka*, Mercedes Yoder, Joan East, . . . , Nicholas Alexander, and <b>Corty, RW</b> .	<b>2018</b>
<b>UNC Quality Improvement Dinner</b> title: Improving Hypertension Management at Moncure Community Health Center	<b>2018</b>

authors: **Corty, RW\***, Angelia Flanagan, Joan East, Nicholas Alexander, and Jan Lee Santos

- American Society of Human Genetics** 2017  
title: Evaluating the role of genetic variants on blood cell count variability in the Jackson Heart Study  
authors: Jessica Shaw\*, **Corty, RW**, Laura Raffield, . . . , Leslie Lange, William Valdar, and Ethan Lange
- National MD-PhD Student Keystone Conference** 2017  
title: How and Why To Model Heteroscedasticity in Mouse Genetics and Personalized Medicine  
authors: **Corty, RW\***, Jessica Shaw, Laura Raffield, Leslie Lange, Ethan Lange, and William Valdar
- International College of Quantitative Genetics** 2016  
title: Jointly Modeling Phenotype Mean and Variance for Robust (m|v|mv)QTL Mapping  
authors: **Corty, RW\***, Janice S. Bailey, Lisa M. Tarantino, and William Valdar
- Society of Toxicology Annual Meeting** 2015  
title: Risk Factors for Tolvaptan-Induced Liver Injury Are Identified Using a Collaborative Cross Approach  
authors: Merrie Mosedale\*, Darol E. Dodd, **Corty, RW**, Yuying Xie, William Valdar, and Paul B. Watkins
- Gordon Research Conference on Quantitative Genetics** 2015  
title: Identification of QTL that Control Mean and Variability of Murine Anxiety Traits in a Pair of F2 Crosses  
authors: **Corty, RW\***, Robin Ervin, Lisa M Tarantino, and William Valdar
- Complex Trait Consortium** 2015  
title: Identification of QTL that Control Mean and Variability of Murine Anxiety Traits in a Pair of F2 Crosses  
authors: **Corty, RW\***, Robin Ervin, Lisa M Tarantino, and William Valdar
- UNC Genetics Department retreat** 2013  
title: Inbred Strain Association for Anxiety Behaviors in 45 Classical and Wild Mouse Strains  
authors: **Corty, RW\***, Robin Ervin, Lisa M Tarantino, and William Valdar

#### AWARDS

- NINDS travel award to attend MD-PhD student conference in Washington, D.C. 2015
- USDA scholarship to attend Gordon Research Conference on quantitative genetics in Barga, Italy 2015
- NIMH award to attend Molecular Psychiatry annual meeting in San Francisco, CA 2014
- Jackson Lab scholarship to attend short course in systems genetics in Bar Harbor, ME 2014
- Thomas Temple Hoopes prize for outstanding senior thesis 2010