Carlos Martinez

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Education

Georgia Institute of Technology

Master of Science Electrical & Computer Engineering Aug. 2016 – May 2018 3.6 GPA **New York University**

Bachelor of Science Mathematics Sept. 2011 – Dec. 2015 3.7 GPA; Magna Cum Laude

Minors: Computer Science, Electrical Engineering,

Nuclear Science & Engineering

Employment

Assistant Research Technician

NYU Langone Medical Center, Cancer Institute May 2014 – July 2016; New York, New York

- Researcher acting in a bioinformatic and statistical capacity for the Kirchhoff Lab specializing in cancer genetics
- Implement genomic data processing pipelines for next-generation sequencing using e.g. GATK, SAMtools, BWA, Cufflinks, VEP
- Perform statistical association tests for variant-disease interactions, expression-disease interactions, and survival analyses
- Conduct exploratory analyses with visualizations and descriptive statistics to discover novel trends in high-risk populations
- Utilize a high-performance computing cluster running on Sun Grid Engine to achieve the above computations
- Secondary responsibilities include grant and manuscript preparation, project presentations, and digital data storage

(Head) Teaching Assistant – Georgia Institute of Technology

Head TA: May 2017 - Dec. 2017; TA: Aug. 2016 - May 2017; Atlanta, Georgia

- Lead MATLAB-based laboratory sections of ECE2026 (intro digital signal processing) and host review sessions
- As head TA: coordinate and train other TAs to conduct labs and grade homework

Tutor – NYU Tandon School of Engineering, TRIO Scholars Program

September 2013 – December 2015; Brooklyn, New York

- One-on-one and small group tutor for introductory-to-advanced level math and physics courses.
- Prepare individualized review plans per student

Publications

M. Vogelsang, **C. N. Martinez**, J. Rendleman, A. Bapodra, K. Malecek, A. Romanchuk, E. Kazlow, R. L. Shapiro, R. S. Berman, M. Krogsgaard, et al., "The expression quantitative trait loci in immune pathways and their effect on cutaneous melanoma prognosis," Clinical Cancer Research, 2016.

R. Ferguson, M. Vogelsang, E. Ucisik-Akkaya, K. Rai, R. Pilarski, **C. N. Martinez**, J. Rendleman, E. Kazlow, K. Nagdimov, I. Osman, et al., "Genetic markers of pigmentation are novel risk loci for uveal melanoma," Scientific reports, vol. 6, p. 31191, 2016.

Conference Poster Sessions

American Society of Clinical Oncology (ASCO) 2017 Annual Meeting

R. Ferguson, D. Simpson, C. N. Martinez, M. Vogelsang, E. Kazlow, U. Moran, J. S. Weber, R. J. Sullivan, K. Flaherty, A. C. Pavlick, A. Ribas, I. Osman, and T. Kirchhoff, "Expression quantitative trait loci (eqtls) as germline determinants of melanoma immunotherapy response.," Journal of Clinical Oncology, vol. 35, no. 15 suppl, pp. 3017–3017, 2017.

D. Simpson, R. Ferguson, C. N. Martinez, E. Kazlow, U. Moran, A. Heguy, D. Hanniford, E. Hernando, I. Osman, and T. Kirchhoff, "Mutation burden as a potential prognostic marker of melanoma progression and survival.," Journal of Clinical Oncology, vol. 35, no. 15 suppl, pp. 9567–9567, 2017.

E. Kazlow, R. Ferguson, D. Simpson, C. N. Martinez, M. Vogelsang, U. Moran, Y. Lee, I. Osman, D. Polsky, and T. Kirchhoff, "Novel germline risk loci in familial melanoma (fm).," Journal of Clinical Oncology, vol. 35, no. 15 suppl, pp. 1535–1535, 2017.

ASCO 2016 Annual Meeting

(Presented) D. Hanniford, C. N. Martinez, I. Dolgalev, M. W. Lattanzi, E. V.-S. de Miera, E. M. Robinson, C. Goldman, A. Heguy, T. Kirchhoff, I. Osman, and E. Hernando, "Targeted next-generation sequencing of melanoma patient samples to reveal mutations in non-protein coding regions of targetable oncogenes.," Journal of Clinical Oncology, vol. 34, no. 15 suppl, pp. 9559–9559, 2016.

T. Kirchhoff, E. Ucisik-Akkaya, M. Vogelsang, K. Rai, R. Pilarski, **C. N. Martinez**, R. Ferguson, E. Kazlow, I. Osman, F. H. Davidorf, C. M. Cebulla, and M. Abdel-Rahman, "The identification of novel genetic risk loci in uveal melanoma.," Journal of Clinical Oncology, vol. 34, no. 15 suppl, pp. 1543–1543, 2016.

S. A. Weiss, **C. N. Martinez**, E. V.-S. de Miera, I. Dolgalev, R. L. Shapiro, A. Heguy, E. Hernando, T. Kirchhoff, and I. Osman, "Genomic characterization of acral lentiginous melanoma: Identification of altered metabolism as a potential therapeutic target.," Journal of Clinical Oncology, vol. 34, no. 15 suppl, pp. 9524–9524, 2016.

Technical Skills

Core:

- Statistical data analysis and numerical computation: NumPy, pandas, R, MATLAB, Excel
- Machine learning libraries: Scikit, Keras, Tensorflow
- Interactive and static visualizations with Dash, plot.ly, matplotlib, seaborn, ggplot2
- General programming: Python, C++, shell scripting
- Querying, joining, aggregating SQL databases (familiar with PostgreSQL, MySQL, SQLite)
- Equally comfortable in Windows or UNIX-based OS
- Basic cloud computing with AWS EC2 instances and S3 storage
- Basic web development and deployment via Django and Heroku
- Professional communication in technical formats (manuscripts, reviews, poster presentations, slide decks)

Bioinformatics / Genomics:

- Genomic pipeline development & analysis for next-generation sequencing data with various software including: GATK,
 SAMtools, BWA, Cufflinks, ANNOVAR, Variant Effect Predictor
- Custom track management for use on UCSC Genome Browser
- Querying and investigating public datasets such as TCGA, Ensembl, 1000 Genomes Project

Academics & Certificates

Gates Millennium Scholar

2011 - 2018

Full-ride scholarship awarded by the Bill & Melinda Gates Foundation to minority high school students exhibiting academic promise and commitment to community service.

Deep Learning Specialization

March 2018

Deep Learning, a 5-course specialization by deeplearning.ai on Coursera

Selected Projects

City Perimeter Detection - cmartinez.io/perimeter-detection/

Automatically detect the perimeters of cities from aerial images. A CNN detects roads in the image followed by image processing to expand the road network into a contour of the city. Tools used include **Keras with Tensorflow** for implementing a CNN, **AWS** for data acquisition, and **OpenCV** for image processing.

Toeing the Party Manifold – cmartinez.io/toeing-party-manifold/

Interactive 2- and 3D visualization of various dimensionality reduction techniques applied to US congressional voting data. Made with **Dash** and deployed using **Heroku**.