## Nicholas Giangreco

**Programming Languages:** 

Email: nick.giangreco@gmail.com
Links: Github Linkedin ORCID
Website: nickg.bio

Frequent: R, Python, bash, css, html, sql Previously: Java

Formatting: LaTeX, Markdown, Rmarkdown, Microsoft Office Suite

#### **EDUCATION**

2021 (expected) PhD Candidate, Systems Biology; Columbia University,

New York City, NY

PhD advisor: Nicholas Tatonetti

Topics: Translational Bioinformatics, Pharmacoepidemiology, Analysis of

Electronic Health Records

2010-2014 BS, Biochemistry; University of Rochester, Rochester, NY

Minor: Philosophy

#### WORK EXPERIENCE

# 2014-Present National Human Genome Research Institute; Bethesda, MD

Principal Investigator: Dr. Laura Elnitski

- Molecular events during tumorigenesis of murine ovarian tumor subtypes differing in deletion of tumor suppressor genes
- RNA sequencing, DNA methylation sequencing, and their integration.

Trainee 2014-2016

Special Volunteer 2016-Present

#### FELLOWSHIPS AND AWARDS

- Columbia Diversity Fellowship
- Department of Systems Biology Merit Fellowship
- Donald Charles Award, University of Rochester Department of Biology

- Fulbright Fellowship Alternate 2013-2014: Sweden, Molecular Modeling, "Novel Antibody-SpA Complex Modeling".
- Travel Award to 9th Student Council and ISMB/ECCB conference 2013 Berlin, Germany.

### PUBLICATIONS AND POSTERS

- Nicholas Giangreco, Emmanuel Zorn, Emily Chen, Paolo Colombo, Nicholas Tatonetti, and Barry Fine. Identification of Novel Primary Graft Dysfunction Biomarkers using Exosome Proteomics. American Heart Association Scientific Sessions 2017, poster.
- Sarah Kim-Hellmuth, Matthias Bechheim, Benno Puetz, Pejman Mohammadi, Yohann Nedelec, Nicholas Giangreco, Jessica Becker, Vera Kaiser, Nadine Fricker, Esther Beier, Peter Boor, Stephane Castel, Markus M. Noethen, Luis B. Barreiro, Joseph K. Pickrell, Bertram Mueller-Myhsok, Tuuli Lappalainen, Johannes Schumacher, Veit Hornung. Genetic regulatory effects modified by immune activation contribute to autoimmune disease associations Nature Communications, 8 (266): 1-10.
- Giangreco N., Petrykowska H., Scott A., Margolin G., Gotea V., Cho K. R., and Elnitski L. Inactivation of Arid1a drives aberrant epigenetic traits in a mouse model of Apc/Pten defective ovarian endometrioid tumors. (in preparation)
- Nicholas Giangreco, Tim Lezon. Alternative conformation prediction of Vibrio Cholerae concentrative nucleoside transporter. F1000Research 2012, poster.

#### CONFERENCES AND HACKATHONS

- American Heart Association Scientific Sessions 2017, poster presentation Giangreco et al. 2017.
- NCBI Hackathon @ New York Genome Center June 2017.
- NCBI Hackathon @ NCBI March 2017.
- CSHL Biological Data Science meeting October 2016.
- JHU DaSH Hackathon September 2015.
- ISMB/ECCB conference @ Berlin, Germany July 2013, poster presentation *Giangreco et al. 2012.* .

#### INVITED TALKS AND PANELS

- "AI, Life Sciences, and Big Data", AI Meetup in NYC, August 2017
- NIDDK Undergraduate Step-Up Judge, NIH, Bethesda MD, August 2015

#### COMMUNITY INVOLVEMENT

- NYC MeetUps
  - AI @ Columbia University Medical Center
    - \* Attend and engage in public discourse on a wide range of topics such as AI & Society, AI & Healthcare, and economic impact by AI.
    - \* Facilitate group engagement and organization by consult
  - Data Storytelling
    - \* Monthly lectures by data scientists and managers from private companies presenting stories and narratives using visualizations and interactive methods on diverse sources of data.
  - Statistical programming
    - \* Monthly seminars by data scientists, engineers and analysts using novel and statistically rigorous methods applied within academic and company research teams mainly using the R programming language.
- CUMC Data Science Study Group
  - Co-organizer of mostly wet-lab biologists learning data science topics using the R programming language
  - Leading project management and learn strategies for doing bioinformatics projects.