

# Nicholas Giangreco

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<b>Email:</b>	nick.giangreco@gmail.com
<b>Links:</b>	Github LinkedIn
<b>Website:</b>	nickg.bio
<b>Programming Languages:</b>	Frequent: R, Python, bash, css, html, sql Previously: Java
<b>Formatting:</b>	Markdown, Rmarkdown, Microsoft Office Suite

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## EDUCATION

**2021 (expected) PhD, Systems Biology;** Columbia University, New York City, NY

PhD advisor: Nicholas Tatonetti

Translational Bioinformatics, 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

- 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*)

## CONFERENCES AND HACKATHONS

- 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

## 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 wide range of topics such as AI & Society, AI & Healthcare, and AI economic impact.
    - \* 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 a group of mostly wet-lab biologists learning data science topics using the R programming language
  - Leading project management and learn strategies for doing bioinformatics projects.