# **Nicholas Attila Kovacs**

#### **BIOINFORMATICIST & DATA SCIENTIST**

🛘 (248) 895-2704 | 🗷 NAttilaKovacs@gmail.com | 😭 NicholasAKovacs.com | 🖸 NicholasAKovacs | 🛅 NicholasAKovacs

## Expertise \_\_\_

- · Computational biologist seeking to apply intellectual curiosity, creativity, and problem solving-skills as a consultant at BCG
  - Productive computational scientist with 139 citations from the past 6 years
  - Spearheaded the transition of a theoretical model to a computational project resulting in two, 1<sup>st</sup>-author scientific publications with 3<sup>rd</sup> in preparation.
  - Coauthor of 7 multidisciplinary scientific publications with experimental and computational scientists and engineers
  - Managed 8 senior undergraduates per semester for 6 semesters as a teaching assistant
  - Independently wrote and awarded NSF grant to conduct summer research in Taiwan
  - Presented research as oral and poster presentations at conferences in Japan, Vietnam, and US
  - Research mentor and project manager of undergraduate for one year

## Education \_\_\_\_\_

Ph.D. Bioinformatics GPA: 3.5

Aug 2013 - Dec 2018

Atlanta, GA

GEORGIA INSTITUTE OF TECHNOLOGY

B.S. Biochemistry and Molecular Biology/Biotechnology GPA: 3.3

Aug 2008 - May 2012 East Lansing, MI

MICHIGAN STATE UNIVERSITY

## Experience \_\_\_\_

**Graduate Research Assistant**Georgia Institute of Technology

Aug 2013 - Current

Atlanta, GA

**Dissertation**: Data Mining the Atomic Structure of the Ribosome to Unravel the History of Protein Folding

- **Summary**: Applied structural bioinformatics methods and machine learning algorithms such as clustering to atomic coordinate datasets from over 100 biomolecules composed of 150,000-200,000 atoms to unravel the interralatedness and origin of life.
- **Results**: Two 1<sup>st</sup>-author research articles published. 3<sup>rd</sup> in preparation.
- Funding: NASA Astrobiology Institute
- Collaboration: Provided computational analysis for coworkers projects; resulted in coauthor of 2 experimental and 4 computational research articles.
- **Communication**: Independently wrote and awarded \$7,000 NSF grant to support summer research in Taiwan. Oral and poster presentations at 7 domestic and international scientific conferences.
- Mentoring: Awarded \$2,500 conference and travel funding for mentoring undergraduate student.
- Courses: 9 courses in biochemisty, computational biology, statistics, and computer science. Concept-to-Market business short-course completed.

### **Atomic Interaction Network Analysis of the Ribosome**

Fall 2017

COURSE PROJECT FOR CS 7280 - NETWORK SCIENCE

Atlanta, GA

- Collaborated with a team member to apply course concepts and algorithms to 3 atomic interaction networks of the biomolecule, the ribosome, each composed of more than 100,000 edges between approx. 50,000 nodes.
- Results: Predicted RNA and protein folding domains within the ribosome by applying community detection algorithms.

#### **Graduate Teaching Assistant**

Aug 2013 - Dec 2016

Atlanta, GA

GEORGIA INSTITUTE OF TECHNOLOGY

- Biophysical Chemistry Lab(CHEM 4582) 6 semesters Instructed ~8 undergraduate students on experimental and computational protocols.
- Macromolecular Structure(CHEM 6572) 2 semesters Directed ~25 graduate students on the use of computational modelling programs.
- Survey of Biochemistry(CHEM 3511) 1 semester Guided ~40 undergraduate students to solve homework problems in weekly recitation.

## Analysis and Interpretation of NGS Data from CDC

Aug 2014 - May 2015

 ${\tt Course\,Project\,for\,BIOL\,8803B\,-\,Programming\,for\,Bioinformatics\,and\,BIOL\,7210\,-\,Computational\,Genomics}$ 

Atlanta, GA

- · Worked in multidisciplinary teams of biologists and computer scientists to identify pathogens from DNA sequences provided by the CDC.
- Analyzed 97 NGS single-end and paired-end reads of *Neisseria meningitidis*, *Haempophilus influenza*, and *Haemophuilus haemolyticus* generated from GAII or Illumina HiSeq/MiSeq instruments.
- **Results**: Developed a typing-tool that identifies the organism and its serotype/serogroup from DNA sequence file inputs and constructed a genome browser of 53 annotated genomes to view annotated genomes.

# Skills & Interests

**Computational Acumen** Python, R, SQL, Bash, Perl, MATLAB, Javascript, Git, Spark, Hadoop, Tableau, Adobe **Computational Sciences** Structural Bioinformatics, Next-Generation Sequencing, Molecular Dynamics, Drug Docking

**Data Science** Machine Learning, Network Science, Statistics, Software Engineering **Experimental Sciences** Molecular Biology, Biochemistry, Organic Chemistry, Analytical Chemistry

**Personal Interests** Bodyweight fitness, snowboarding, guitar, world travel, bachata dance, Korean-culture