

# TAEHOOON HA

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## Education

### Weill Cornell Medicine

*M.S., Biostatistics*

New York, NY

2018 – Present

- Advisor: Xi Kathy Zhou, PhD

### Duke University

*M.S., Business Analytics*

Durham, NC

2017 – 2018

- Capstone project: Duke University Hospital (Duke Health)

### Sungkyunkwan University

*B.B.A. with an emphasis on Quantitative Methods*

Seoul, Korea

2009 – 2017

- Dean's list with distinction
- Study-abroad: School of Arts and Sciences at the University of Pennsylvania (2014)
- Military service: Log analyst at Radar Operation and Computer Emergency Readiness Team (CERT), Republic of Korea Marine Corps (2011-2013)

### Independent Math Coursework

Online

- Intended (Spring 2020):
  - Differential Equations (University of North Dakota)
  - Real Analysis (University of Illinois at Urbana-Champaign)
- In progress (Fall 2019): Multivariable Calculus (University of North Dakota)
- Completed:
  - Calculus I, Calculus II, and Principles of Statistics (Brigham Young University)
  - Linear Algebra (University of North Dakota)

## Research Experience

### Johns Hopkins Bloomberg School of Public Health

Baltimore, MD

*Voluntary Researcher* (Advisor: Bongsoo Park, PhD)

(Remote)

04/2019 – Present

#### Transcriptome and epigenome atlas for air pollution PM<sub>2.5</sub>

- Construct data pipeline of both EdgeR and DESeq2 to identify differentially expressed genes associated with the exposure to ultra-fine dust, PM<sub>2.5</sub>
- Check the distribution by Relative Log Expression (RLE) and Principal Component Analysis (PCA) plots
- Normalize each sample and removed unwanted variances and filtered top 5,000 genes by p-values from EdgeR and DESeq2
- Perform DE (Differential Expression) analysis using EdgeR and Gene Ontology (GO) Term analysis on six different brain sub-areas and liver cells.
- Conduct pathway analysis using Ingenuity Pathway Analysis (IPA) software
- Prepare for publication “*Brain transcriptome map of air pollution PM<sub>2.5</sub>*”

### Weill Cornell Medicine

*Research Assistant* (Advisor: Xi Kathy Zhou, PhD)

New York, NY

04/2019 – Present

### Methodology: Bayesian model averaging

- Assist the advisor with developing a new statistical method using Bayesian model averaging to identify differentially expressed genes associated with one or more phenotypes, as well as their interactions.
- Develop R package ‘*BMAseq*’ using Bayesian model averaging to analyze observational gene-expression data
- Apply the Bayesian model averaging method to large scale public RNA-seq gene expression data
- Apply the Bayesian model averaging method (“BMA-seq”) to observational metabolomics data to improve differentially expressed (DE) metabolites identification in high dimensional setting
- Prepare for publication “*Bayesian model averaging approach for RNA-seq counts data (BMA-seq) and its application*”

### Application: Collaboration with Andrew J. Dannenberg, MD group

- Provide statistical consulting support to clinical and lab research projects using R
- Collaborate with Andrew J. Dannenberg, MD group to identify the link along the obesity→inflammation→Aromatase pathway in excess adipose tissue of mice and humans
- Update and customize R package ‘BTKR’ which includes multiple functions that implement some commonly used biostatistics analysis methods for a simple summary of data
- Prepare for publication “*Body composition is a determinant of breast adipose inflammation and Aromatase levels, including in normal-sized women*”

### JB Lab & Clinic

Research Scientist

Seoul, Korea

(Remote)

08/2018 – Present

- Identified the association of sodium intake and hypertension, metabolic syndrome, and ARB treatment effect from a 10K+ hypertension patients data acquired from a K-MetS study

### Weill Cornell Medicine

Leading Project Researcher

New York, NY

#### The Effect of Right Heart Catheterization (RHC) During the First 24 Hours on 30-day Mortality of Critically Ill Patients in Intensive Care Units

05/2019 – 08/2019

[Taehoon Ha, Jingjing Qi]

- Applied data adaptation method to estimate the propensity score
- Generated causal model estimands using multiple methods (random forest, GLM, elastic net, LASSO, and gradient boosting with SuperLearner)
- Conducted 5-fold cross-validation to evaluate each estimand’s performance and coefficient in the ensemble estimator

#### Rate of General Anesthesia Use for Cesarean Delivery Among Anesthesiologists with and without Fellowship Training in Obstetric Anesthesia

01/2019 – 05/2019

[Taehoon Ha, Yu Cai, Siyang Pei, Xuewei Quan]

- Fitted the logistic regression model to identify whether the fellowship-trained anesthesiologists are more or less likely to provide general anesthesia for non-routine C-section as compared to non-fellowship trained anesthesiologists
- Identified the significant difference between general and neuraxial anesthesia and the odds of using general anesthesia is 0.471 times lower in attending fellowship training anesthesiologists compared the non-fellowship training anesthesiologists

<b>Cost-Effective Optimization of Model-Based Prediction of Cardiovascular Disease (CVD) [Taehoon Ha]</b> <ul style="list-style-type: none"> <li>• Aimed to increase the accuracy of CVD diagnosis using a model-based approach</li> <li>• Determined the demographic factors and medical tests that help predict the likelihood of heart disease using proportional odds model and logistics regression</li> <li>• Identified a combination of necessary medical tests that help predict the probability of heart disease in a cost-effective manner and reduced the exam cost by \$110.17</li> </ul>	03/2019 – 05/2019
<b>Neuroendocrine Prostate Cancer (NEPC) [Taehoon Ha, Diane Li, Lingchen Lou]</b> <ul style="list-style-type: none"> <li>• Identified that there is no difference in the proportions of 3-month progression status between patients in the NEPC and non-NEPC group treated with alisertib</li> <li>• Determined that there is no association between 3-month progression-free survival for all clinical characteristics and gene abnormalities</li> <li>• Identified an association between PSA level and prior systemic therapies between NEPC and non-NEPC groups treated with alisertib</li> </ul>	11/2018 – 12/2018
<b>Major Risk Factors of Low Birth Weight Babies [Taehoon Ha]</b> <ul style="list-style-type: none"> <li>• Determined that low birth weight is associated with the mother's premature labor history, race, smoking status, weight of the last menstrual period, and history of hypertension</li> <li>• Fitted multiple generalized linear models with the model selection based on deviance and p-values</li> </ul>	10/2018 – 12/2018
<b>Study design of Tommy John Surgery for MLB Pitchers by Statcast Measurements [Taehoon Ha]</b> <ul style="list-style-type: none"> <li>• Identified the primary injury factors of Ulnar Collateral Ligament Reconstruction for Major League Baseball pitchers by Statcast measures</li> <li>• Designed a study to identify major factors of Ulnar Collateral Ligament Reconstruction (a.k.a. Tommy John Surgery) for Major League Baseball Pitchers by Statcast measurements</li> </ul>	09/2018 – 10/2018
<b>Policrew Research Group</b> <i>Co-founder and Head Research Scientist (Data)</i>	06/2015 – 06/2017
<b>Leadership</b> <ul style="list-style-type: none"> <li>• Managed all research projects, earning a reputation for excellent understanding of trends and insight of youth</li> <li>• Developed strong and long-term relationships with donors and Partner Research Organizations, beyond the scope of engagement</li> </ul>	
<b>Problem-solving</b> <ul style="list-style-type: none"> <li>• Conducted network analysis to publish annual youth employment and health trend report</li> <li>• Defined and evaluated the key performance indicators of the Korean government's current youth unemployment rate and healthcare policies</li> </ul>	

## Teaching

<b>Design and Analysis of Biomedical Studies (Master's level)</b> <i>Teaching Associate for Prof. Xi Kathy Zhou, Weill Cornell Medicine</i>	Spring 2020
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- Lead lab sessions for 21 Master's candidate students to teach biostatistics methods with R
- Review and grade weekly homework and provide guidance on lab assignments
- Hold regular office hours regarding questions on course materials, assignments, and academic concerns

### **Categorical and Censored Data Analysis (Master's level)**

Fall 2019

*Teaching Associate for Prof. Oleksandr Savenkov, Weill Cornell Medicine*

- Lead lab sessions for 32 Master's candidate students to teach biostatistics methods with R
- Review and grade weekly homework and provide guidance on lab assignments
- Hold regular office hours regarding questions on course materials, assignments, and academic concerns

### *Publications*

#### **In Preparation**

- Hanhan Wang, Lingsong Meng, **Taehoon Ha**, Xi Kathy Zhou. *A Bayesian model averaging approach for RNA-seq counts data (BMA-seq) and its application*
- B Park, S Kim, **Taehoon Ha**, JE Park, V Vinayachandran, KD Hansen, B Paul, S Rajagopalan, and S Biswal. *Brain transcriptome map of air pollution PM<sub>2.5</sub>*

### *Awards*

<b>Academic Excellence (Over 4.0 cumulative GPA), Weill Cornell Medicine</b>	07/2019
<b>Exchange Student Scholarship at the University of Pennsylvania (Travel Funding), Sungkyunkwan University</b>	03/2015
<b>Academic Excellence Scholarship, Sungkyunkwan University</b>	01/2015
<b>College Scholarship (Academic Excellence), Sungkyunkwan University</b>	03/2014
<b>Pursuit of Excellence Scholarship, Sungkyunkwan University</b>	01/2014
<b>Pursuit of Excellence Scholarship, Sungkyunkwan University</b>	01/2010
<b>Pursuit of Excellence Scholarship, Sungkyunkwan University</b>	07/2009

### *Invited Talks*

<b>How to organize, manage, share, collaborate and process bio/healthcare data, Konkuk University Hospital</b>	12/2018
<b>RNA-seq data analysis 101, JB Lab &amp; Clinic</b>	08/2018

### *Relevant Coursework*

#### **Weill Cornell Medicine**

- Biostatistics I, Biostatistics II – Regression, Statistical Learning, Causal Inference, Foundations Biomedical Applications, Study Design, Categorical Censored Data Analysis, Statistical Programming, Data Management

#### **Duke University**

- Applied Probability (Math), Data Infrastructure (SQL), Decision Analytics and Modeling (R), Data Visualization (Tableau), Data Science for Business (Python), Pricing (R), Empirical Economic Analysis: Advanced Econometrics (R), Operations Analytics (R)

#### **Sungkyunkwan University**

- Computer Data Processing and Computer Word Processing (Java, SQL), Marketing Research Methodology (SAS), Consumer Behavior (SAS), Introduction to Social Science Research (SAS)

#### **Independent Math Coursework**

- Intended (Spring 2020):
  - Differential Equations (University of North Dakota)
  - Real Analysis (University of Illinois at Urbana-Champaign)
- In progress (Fall 2019): Multivariable Calculus (University of North Dakota)
- Completed:
  - Calculus I, Calculus II, and Principles of Statistics (Brigham Young University)
  - Linear Algebra (University of North Dakota)

### *Service*

#### **Prudential Foundation**

2015

##### *Project CD34: Stem Cell Donation Trend Visualization*

- Integrated and constructed the stem cell donation trend databases in Korea
- Generated real-time dashboard to help the board's decision-making process

#### **S-ONE, Sungkyunkwan University**

2013 – 2014

##### *Data Analysis & Visualization Education Session Chief*

- Led lab sessions for 40 students to teach statistical methods and data visualization tools when analyzing social science data

#### **Samsung Dream Class Program**

2009 – 2010

##### *High School Mathematics Tutor*

- Taught high school and pre-college math to 20 low-income students

### *Technical Skills*

- **Programming:** R, SAS, Python, Java

- **Database:** MySQL, SQL Server, Microsoft Access
- **Document:** LaTeX, Markdown
- **Cloud / Parallel computing:** Google Cloud Platform, Amazon Web Services, Microsoft Azure, Sun Grid Engine (SGE)
- **Visualization:** Tableau Software

### *Membership*

- American Statistical Association (ASA)
- Korean Statistical Society (KSS)
- American Heart Association (AHA)
- American Association for the Advancement of Science (AAAS)