

Boying Gong

boyinggong@berkeley.edu ◇ (510) 708-9381

EDUCATION	University of California, Berkeley <i>Ph.D. in Biostatistics</i> ; GPA: 4.0/4.0 University of California, Berkeley <i>M.A. in Statistics</i> ; GPA: 3.96/4.0 Zhejiang University <i>B.S. in Statistics, Minor in Finance</i> ; GPA: 3.92/4.0	Expected Aug. 2021 2017 2015
EXPERIENCE	Barclays Capital, New York <i>Statistical Modeling and Development Intern</i> <ul style="list-style-type: none">Evaluated the efficiency of the pricing and reward system in equity financing. Investigated models on optimizing funding efficiency. Department of Statistics, UC Berkeley <i>Graduate Student Researcher. Advisor: Elizabeth Purdom</i> <ul style="list-style-type: none">Developed statistical methods and software for the analysis of high-dimensional genomic data. Focused on multimodal machine learning for integrating sequencing datasets and region identification from spatio-temporal epigenetic data. <i>Administrative Researcher</i> <ul style="list-style-type: none">Co-developed the curriculum for a new course, <i>Statistical Methods for Data Science</i>, in Fall 2016 and taught the course in Spring 2017.	June – Aug. 2019 2017 – Present 2016 – 2017
PUBLICATIONS	◇ <i>Methodology</i> <ul style="list-style-type: none">[1] Boying Gong, Yun Zhou, Elizabeth Purdom. “Cobolt: Joint analysis of multimodal single-cell sequencing data.” <i>Preprint</i>, 2021.<ul style="list-style-type: none">Keywords: multimodal machine learning, variational autoencoder[2] Yun Zhou, Boying Gong, Tao Jiang, Ting Xu, Haiyan Huang. “Remote homologue representation in random heteropolymer with stochastic variational methods.” <i>Submitted to Journal of American Statistical Association</i>, 2021.<ul style="list-style-type: none">Keywords: variational inference, sequential data, expectation–maximization[3] Boying Gong, Elizabeth Purdom. “MethCP: Differentially methylated region detection with change point models.” <i>Journal of Computational Biology</i>, 2019.<ul style="list-style-type: none">Top 5% paper RECOMB 2019Keywords: change point detection, time-course data, differential analysis ◇ <i>Application</i> <ul style="list-style-type: none">[4] Boying Gong, et al. “Single-cell RNA-seq reveals age-associated changes in olfactory stem cell regeneration.” <i>Prepare for submission</i>, 2021.[5] Brann, David H., . . . , Boying Gong et al. “Non-neuronal expression of SARS-CoV-2 entry genes in the olfactory system suggests mechanisms underlying COVID-19-associated anosmia.” <i>Science Advances</i>, 2020.<ul style="list-style-type: none">Uncovered the target cells for the loss of smell caused by Covid-19. Covered by <i>NY Times</i>, <i>WSJ</i>, and other news outlets.[6] Lim, Marc, Vishnu Dharmaraj, Boying Gong, et al. “Estimating tumor vascular permeability of nanoparticles using an accessible diffusive flux model.” <i>ACS Biomaterials Science and Engineering</i>, 2020.	
SOFTWARE PACKAGES	MethCP : An R package for differentially methylated region detection (GitHub) Cobolt : A Python package for jointly analyzing multimodal single-cell datasets (Github)	
SELECTED HONORS	Block Grant Fellowship Elizabeth Scott Memorial Award <i>Awarded annually to one M.A. student showing the greatest promise in statistical research.</i>	UC Berkeley, 2018 UC Berkeley, 2017
SKILLS	Programming Proficient in R. Intermediate experience with Python and Git. Languages Native speaker of Mandarin. Fluent in English. Beginner in French.	