

CONTACT INFORMATION	Kidney Disease Initiative and Klarman Cell Observatory Broad Institute 5407G, 415 Main Street Cambridge, MA 02412 USA	twitter: @ayshwaryas website: ayshwaryas.github.io email: subraman@broadinstitute.org
AREAS OF EXPERTISE	Computational Biology (Kidney disease, Cancer, Inflammatory Bowel Disease, RNA Biology), Genomic Data Analysis (Single-cell and Bulk-RNAseq, Metagenomics, Exome and single-nucleus DNA sequencing), Machine Learning, Probabilistic modeling, Phylogenetics, Applied Statistics	
EDUCATION	<p>2013 <b>Ph.D., Biological Sciences</b> Carnegie Mellon University, Pittsburgh, PA USA <i>Doctoral Advisor:</i> Russell Schwartz, Ph.D. <i>Dissertation:</i> Inferring tumor evolution using computational phylogenetics</p> <p>2007 <b>M.Sc. (Hons), Biological Sciences (Undergraduate degree)</b> Birla Institute of Technology and Science (BITS–Pilani), Rajasthan, India <i>CGPA 9.21/10, Major GPA 10/10, with Distinction</i> <i>Undergraduate Honors Thesis:</i> A mathematical model for phototactic responses in <i>Halobacterium salinarium</i>, Max Planck Institute for Complex Technical Systems, Germany.</p>	
CURRENT APPOINTMENT	<b>Computational Scientist</b> , Cambridge MA <i>Mentors:</i> Anna Greka, M.D., Ph.D. & Aviv Regev, Ph.D. <i>Research Summary:</i> Single-cell portraits of disease and normal states using human data, mouse and organoid models Kidney Disease Initiative and Klarman Cell Observatory Broad Institute, Cambridge, MA 02142	2017–Present

PUBLICATIONS **Pre-prints**

- [1] Korthauer K<sup>†</sup>, Kimes PK<sup>†</sup>, Duvallet C<sup>‡</sup>, Reyes A<sup>‡</sup>, **Subramanian A<sup>‡</sup>**, Teng M, Shukla C, Alm EJ, Irizarry RA, Hicks SC. (2018). A Practical Guide to Methods Controlling False Discoveries in Computational Biology. *bioRxiv*. doi: <https://doi.org/10.1101/458786>

**Peer-reviewed Journal Articles**

6. Mehta RS, Abu-Ali GS, Drew DA, Lloyd-Price J, **Subramanian A**, Lochhead P, Joshi AD, Ivey KL, Khalili H, Brown GT, DuLong C, Song M, Nguyen LH, Mallick H, Rimm E, Izard J, Huttenhower C, Chan AT. Stability of the human faecal microbiome in a cohort of adult men. *Nat Microbiol*. 2018 Mar;3(3):347-355. PMID: 29335554
5. **Subramanian A** and Schwartz R. Reference-free inference of tumor phylogenies from single-cell sequencing data. *BMC Genomics*. 2015;16 Suppl 11:S7. PMID: 26576947
4. **Subramanian A**, Shackney S, Schwartz R. Novel multi-sample scheme for inferring phylogenetic markers from whole genome tumor profiles. *IEEE/ACM Trans Comput Biol Bioinform*. 2013 Apr 23. PMID: 24407301
3. **Subramanian A**, Shackney S, Schwartz R. Inference of tumor phylogenies from genomic assays on heterogeneous samples. *J Biomed Biotechnol*. 2012; 2012:797812. PMID: 22654484
2. Tolliver D, Tsourakakis C, **Subramanian A**, Shackney S, Schwartz R. Robust unmixing of tumor states in array comparative genomic hybridization data. *Bioinformatics*. 2010 Jun 15; 26(12): i106-14. PMID: 20529894
1. Adithi M, Kandam M, Ramkumar HL, **Subramanian A**, Venkatesan N, Krishnakumar S. Retinoblastoma: expression of HLA-G. *Ocul Immunol Inflamm*. 2006 Aug; 14(4): 207-13. PMID: 16911982

<sup>†</sup>CD, AR, AS contributed equally and are listed alphabetically

## Peer-reviewed full-length Conference Papers

4. **Subramanian A** and Schwartz R, "Reference-free inference of tumor phylogenies from single-cell sequencing data," 2014 IEEE 4th International Conference on Computational Advances in Bio and Medical Sciences (ICCABS), Miami, FL, 2014, pp. 1-1. *Oral Presentation*
3. **Subramanian A**, Shackney S, Schwartz R. Novel multi-sample scheme for inferring phylogenetic markers from whole genome tumor profiles. Proceedings of the 8th International Symposium on Bioinformatics Research and Applications (ISBRA) 2012, Dallas, TX, USA, May 21-23, 2012. Bioinformatics Research and Applications. Lecture Notes in Computer Science Volume 7292, 2012, pp 250-262. *Oral Presentation*.
2. **Subramanian A**, Shackney S, Schwartz R. Inference of tumor phylogenies from genomic assays on heterogeneous samples. ACM-BCB, Chicago, IL, August 01 - 03, 2011. BCB '11 Proceedings of the 2nd ACM Conference on Bioinformatics, Computational Biology and Biomedicine (ACM-BCB). pp 172-181. *Oral Presentation*.
1. Tolliver D, Tsourakakis C, **Subramanian A**, Shackney S, Schwartz R. Robust unmixing of tumor states in array comparative genomic hybridization data. Intelligent Systems for Molecular biology (ISMB) 2010. *Oral Presentation*.

## Book Chapter

1. **Subramanian A**, Shackney S, Schwartz R. "Tumor phylogenetics in the Next Generation Sequencing era: Strategies and Challenges." Applications of Next Generation Sequencing in Cancer Research. 2013

## Papers in progress

1. **Subramanian A**, et al. Single-cell portraits of human kidney disease.*in preparation*
2. **Subramanian A**, Schwartz R and Lee AV. Genomic variation among MCF-7 cancer cell lines grown in different laboratories: a phylogenetic analysis.*in preparation*
3. **Subramanian A**, Eric Franzosa, Dirk Gevers, Ramnik Xavier, Huttenhower C. Computational prediction of novel secreted bioactive microbial gene products from microbial shotgun sequencing studies.*in preparation*
4. **Subramanian A**, Irizarry RA. Creating a transcript expression barcode using RNA-Seq Data.*in preparation*

## Selected Poster Presentations

7. Mallick H, Tickle TL, McIver LJ, Weingart G, Paulson JN, Ma S, Ren B, Schwager E, **Subramanian A**, Franzosa EA, Bravo HC, Huttenhower C. Multivariable Association in Population-scale Meta'omic Surveys (Oral Presentation at the Intelligent Systems for Molecular Biology (ISMB), Chicago, IL, July 2018).
6. **Subramanian A**, Heidi-Sidhom E, Emani M, Zhou Y, Zhang F, Xiao L, Vernon K, Dionne D, Nguyen L, Cuoco M, Rozenblatt-Rosen O, Regev A, Greka A. 2018 Klarman-Cell Observatory Meeting.
5. **Subramanian A**, Francoza E, Vlamakis H, Xavier R and Huttenhower C. de novo functional genomic annotation of the gut microbiome in inflammatory bowel disease. 2015 Annual Broad Institute retreat.
4. **Subramanian A**, and Schwartz R. Reference-free inference of tumor phylogenies from single-cell sequencing data. 2015 Annual Program in Quantitative Genetics (PQG) symposium.
3. **Subramanian A**, Mehta RS, Dulong Casey et al. Diet-linked gut microbial risk factors in Colorectal Carcinogenesis. STARR Cancer Consortium Annual Retreat 2015.
2. **Subramanian A**, Shackney S, Schwartz R. Inference of robust tumor phylogenetic markers from multi-sample data. GLBIO (Great Lakes Bioinformatics Conference) 2012.

1. **Subramanian A**, Shackney S, Schwartz R. Phylogenetic Methods for inferring tumor progression pathways from aCGH profiles of mixed cell populations. Poster, Intelligent Systems in Molecular Biology (ISMB) 2010.

## Conference Proceedings

5. Chang Y-L, Harre N, Rossetti M, **Subramanian A**, Kostic A, Huttenhower C, Xavier R, Stappenbeck T, Simpson K W., Sartor R. B, Wu G D., Lewis J, Bushman F D., McGovern D, Salzman N, Borneman J, Braun J. Su1873 Identification of IBD-Related Microbial Metabolites Affecting Human Th17 Differentiation. *Gastroenterology* Volume 150, Issue 4, Supplement 1, April 2016, Page S576.
4. Chowdhury SA, **Subramanian A**, Schaffer AA, Shackney SE, Wangsa D, Heselmeyer-Haddad K, Ried T, Schwartz RS. Inferring evolutionary models of tumor progression from single-cell heterogeneity data.[abstract]. In: Proceedings of the 105th Annual Meeting of the American Association for Cancer Research; 2014 Apr 5-9; San Diego, CA. Philadelphia (PA): AACR; Cancer Res 2014;74(19 Suppl):Abstract nr 5338. doi:10.1158/1538-7445.AM2014-533.
3. **Subramanian A**, Shackney S, Schwartz R. Inference of tumor phylogenetic markers from large copy number datasets.[abstract]. In: Proceedings of the 104th Annual Meeting of the American Association for Cancer Research; 2013 Apr 6-10; Washington, DC. Philadelphia (PA): AACR; Cancer Res 2013;73(8 Suppl):Abstract nr 5133. doi:10.1158/1538-7445.AM2013-5133
2. **Subramanian A**, Shackney S, Schwartz R. Inference of robust tumor phylogenetic markers from multi-sample data.[abstract]. In: Proceedings of the 103rd Annual Meeting of the American Association for Cancer Research; 2012 Mar 31-Apr 4; Chicago, IL. Philadelphia (PA): AACR; Cancer Res 2012;72(8 Suppl):Abstract nr 3964. doi:1538-7445.AM2012-3964
1. **Subramanian A**, Shackney S, Schwartz R. Towards novel marker discovery from phylogenetic analysis of heterogeneous tumor samples.[abstract]. In: Proceedings of the 102nd Annual Meeting of the American Association for Cancer Research; 2011 Apr 2-6; Orlando, FL. Philadelphia (PA): AACR; Cancer Res 2011;71(8 Suppl):Abstract nr 44. doi:10.1158/1538-7445.AM2011-44

## Ph.D. Dissertation

1. **Subramanian A**, “Inferring tumor evolution using computational phylogenetics” (2013). *Dissertations*. 275.

## GRANTS AND FELLOWSHIP

- |      |   |
|------|---|
| 2016 | <i>Principal Investigator</i> Broadnext10 Round 3 Catalytic Steps Trainee Award by the Broad Institute, (\$40,000 direct costs)             |
| 2006 | Undergraduate Summer Science Fellowship Award by the Indian Academy of Science (IAS)  |
| 2006 | Undergraduate Summer Science Fellowship Award by the Jawaharlal Nehru Center for Advanced Scientific Research (JNCASR) ( <i>declined</i> ). |
| 2001 | Ford Environment and Conservation grant awarded to only 3 high-school science projects, Ford Motors, USA.                                   |

## AWARDS AND HONORS

- 2017 American Statistical Association (ASA) Travel award to attend the Women in Statistics and Data Science (WSDS) conference, La Jolla, CA
- 2017 RStudio Diversity Scholarship to attend the RStudio International Conference at Orlando, FL
- 2016 UseR! Diversity Scholarship to attend UseR!, the largest conference for users of the statistical programming language R
- 2015 Stellar Abstract Award for poster at the Harvard Program in Quantitative Genetics (PQG) Annual Conference
- 2015 Travel Award for oral presentation at the Human Microbiome Data (HMD) workshop by the Statistical and Applied Mathematical Sciences Institute (SAMSI)
- 2014 NSF Travel Fellowship for the 4th IEEE International Conference on Computational Advances in Bio and Medical Sciences (IEEE/ICCABS)
- 2012 Travel Fellowship for oral presentation at the International Symposium on Bioinformatics Research and Applications (ISBRA)
- 2011 Only Invited Student Panelist, ACM Women in Bioinformatics, ACM-BCB
- 2011 Travel Fellowship by the Department of Biological Sciences, Carnegie Mellon University
- 2011 Travel Fellowship for oral presentation at the ACM Conference on Bioinformatics, Computational Biology, and Health Informatics (ACM-BCB)
- 2010 Global Champion, [Hult Global Case Challenge for social innovation and entrepreneurship](#)
- 2010 2nd Place, [Carnegie Mellon Open Innovation Competition](#)
- 2003 National Certificate of Merit in Physics and Biology (Top 0.1% All India Senior Secondary Certificate Examination (AISSCE))
- 2003 Score of 485/500 (Known highest score in STEM in 2003 = 487/500) in AISSCE Grade 12 examinations

## TALKS

### Invited Talks

- [1] An introduction to single-cell transcriptomics. Broad Institute Cancer Program Bootcamp. 2018 May. Cambridge, MA.
- [2] Inferring Insights from Omics Data with Applications in Tumor Evolution and Inflammatory Bowel Disease. Department of Bioinformatics and Computational Biology, Genentech. 2017 June 8. South San Francisco, USA
- [3] Computational Methods for Inferring Insights from omics Data with Applications in Inflammatory Bowel Disease and Cancer. Inflammation and Immunology Research Unit, Pfizer. 2017 May 17. Cambridge, MA
- [4] Defining transcriptional activity states by leveraging massive, public RNAseq datasets. Bioinformatics Meeting, Division of Immunology, Harvard Medical School. 2017 May 11. Boston, MA.
- [5] *de novo* functional genomic annotation of the gut microbiome in inflammatory bowel disease. Bioinformatics Meeting, Division of Immunology, Harvard Medical School. 2016 May 12. Boston, MA.
- [6] Computational Screens for identifying bioactive microbial compounds in Inflammatory Bowel Disease. Novartis Institute for Biomedical Research (NIBR). 2015 July 13. Cambridge, MA.
- [7] Reference-free inference of tumor phylogenies from single-cell sequencing data. Workshop on Computational Advances for Next Generation Sequencing (CANGS). 2014 June 06. Miami Beach, FL
- [8] Inferring tumor progression using computational phylogenetics. MIT-Computer Science and Artificial Intelligence Laboratory (CSAIL). 2013 August 22. Boston, MA.

### Contributed Talks

- [1] Domain Knowledge as a Key Enabler of Impactful Data Analyses in Single-Cell Genomics. Women in Statistics and Data Science Conference. 2018 October 18. Cincinnati, OH.

- [2] Defining transcriptional activity states by leveraging massive, public RNAseq datasets. [Women in Statistics and Data Science Conference](#). 2017 October 19. La Jolla, CA.
- [3] Computational screens for novel gut microbial bioactive compounds. [Discovering Patterns in Human Microbiome Data \(HMD\) Workshop](#). Statistical and Applied Mathematical Sciences Institute (SAMSI) 2015 March 17. Research Triangle Park, NC.

TEACHING  
EXPERIENCE

- 2017 **Teaching Fellow**, [Introduction to Omics Research](#)  
Prepared course assessment questions and answered online discussion forum. Harvard Catalyst, The Harvard Clinical and Translational Science Center, Harvard University, MA, USA
- 2016 **Teaching Fellow**, [BIO 261/ CS107-E Introduction to Data Science](#)  
Contributed to and graded HWs, advised final projects, assisted in-class, held office hours for online students. Course by Rafael Irizarry at Harvard T. H. Chan School of Public Health & Harvard Extension School
- 2010 **Teaching Assistant**, [03712 Computational Methods for Biological Modeling and Simulation](#).  
Graded exams and HWs, held office hours. Course by Russell Schwartz at Carnegie Mellon.
- 2009 **Teaching Assistant**, [03711 Computational Molecular Biology and Genomics](#).  
Graded exams and HWs, held office hours. Course by Dannie Durand at Carnegie Mellon.
- 2009 **Teaching Assistant**, [03710 Computational Biology](#).  
Created and graded quizzes, held recitations and office hours, graded HWs and exams. Course by Robert Murphy at Carnegie Mellon.
- 2006 **Professional Assistant**, [BIO C461 Recombinant DNA Technology](#)  
Course by Ashis K. Das at Birla Institute of Technology & Science (BITS-Pilani)

TUTORIALS  
AND SHORT  
COURSES

- [1] co-Instructor. [Analysis of single-cell RNA-seq data](#), Physalia courses. February 2019, Berlin, Germany.
- [2] co-Instructor. [Single Cell Data Analysis Lab](#), Center for Excellence in Genomics (CEGS) Workshop. 12 September 2018. Cambridge, USA
- [3] co-Instructor. [Single cell RNA-seq toolkit](#), tutorial at the 25th Intelligent Systems in Molecular Biology (ISMB) Conference. 6 July 2018. Chicago, USA
- [4] Computational Lead. [CyTOF Analysis Workshop](#). 5 December 2017. Dana-Farber Cancer Institute cBio Center. Boston, USA.

PROFESSIONAL  
EXPERIENCE

- 05/14-10/17 **Postdoctoral Research Associate**  
Department of Biostatistics, Harvard T. H. Chan School of Public Health, Boston, MA, Department of Biostatistics and Computational Biology
- 11/15-10/17 PI: Dr. Rafael Irizarry
- 05/14-10/15 PI: Dr. Curtis Huttenhower
- 01/14-05/14 **Bridge Postdoctoral Research Associate**
- 08/07-12/13 **Graduate Research and Teaching Assistant**. Department of Biological Sciences, Carnegie Mellon University, Pittsburgh
- Summer 2011 **Summer Research Intern** (Mentor: Dr. Peter Haverty).  
Department of Bioinformatics and Computational Biology, Genentech, South San Francisco
- 01/07-06/07 **Undergraduate Honors Thesis Advisee** (Mentor: Dr. Wolfgang Marwan)  
Molecular Network Analysis Group, Max Planck Institute for Dynamics of Complex Technical Systems, Magdeburg, Germany
- 2005-2007 **Undergraduate Research Assistant** (Mentor: Dr. A.K. Das). Birla Institute of Technology & Science (BITS Pilani), India
- Summer 2006 **Indian Academy of Science Summer Research Fellow** (Mentor: Dr. M.S. Shaila). Indian Institute of Science (IISc), Bangalore, India
- Summer 2005 **Undergraduate Summer Research Intern** (Mentor: Dr. Krishnakumar Subramanian). Sankara Nethralaya Eye Hospitals, Chennai, India
- 2002 **High School Term project**. Childs Trust Hospital, Human Genetics Department, Chennai, India

TECHNICAL  
SKILLS

**Programming:** Stan, R, Python, Git, Perl, LaTeX, Matlab, Shell scripting, Java (basic), C(basic)  
**Operating Systems:** Mac OS X, Unix, Linux  
**Relevant Graduate coursework:** 10701 Machine Learning, 10705 Intermediate Statistics, 10702 Statistical Machine Learning, 15211 Data Structures and Algorithms, 03712 Computational Methods for Biological Modeling and Simulation, 03210-A3 Information Noise and Entropy in the Brain

SERVICE

**Program Committee**

2017 International Conference on Machine Learning (ICML) Workshop on Computational Biology

**Grant reviewing**

2016 Medical Research Council (MRC) UK  
2015 Florida Department of Health

**Ad hoc Peer-Review**

Journals PLOS Computational Biology, Nature, BMC Genomics  
Conferences Annual International Conference on Research in Computational Molecular Biology (RECOMB), Intelligent Systems in Molecular Biology (ISMB), International Symposium on Bioinformatics Research and Applications (ISBRA), Workshop on Algorithms in Bioinformatics (WABI), IEEE International Conference on Bioinformatics and Biomedicine (BIBM)

**Research Mentoring**

2011–2012 Titus Banerjee, Carnegie Mellon University

**STEM Outreach**

2016, 2017 Career exploration day for high school students, Broad Institute & MassBioEd  
2012 Mentor, “Creative Tech Nights” outreach program for Pittsburgh middle school girls, Carnegie Mellon Women@SCS (School of Computer Science)  
2010–13 Co-president, Scientists and Engineers for America (SEA) student group for Science and Technology policy-making education for graduate students

**Community**

06/14–present Volunteer, Haley House Shelter for Homeless, Boston, MA  
2017 Charles River Watershed Association Earth Day cleanup  
2016 Started the “Rally for Jimmy Fund” employee personal electronics upcycling program at the Department of Biostatistics & Computational Biology, DFCI  
2014, 2016 Volunteer, Perkins School for the Blind, Watertown, MA  
03/08–12/12 Volunteer Teacher (Human Values Character Education Program), Allegheny County Shuman Juvenile Detention Center, Pittsburgh, PA  
07/10–12/13 Volunteer, Bethlehem Haven, Pittsburgh, PA  
07/10–02/14 Volunteer, Alice Davis Personal Care Home, Braddock, PA  
2009 Founding member, CMU Panama Global Business Brigade, Pittsburgh PA