Salil Bhate, MA (Cantab.), MMath., MS, PhD

D.O.B.: 2nd May 1992 Nationality: UK sbhate@broadinstitute.org salilb1.github.io

EMPLOYMENT Broad Institute of MIT and Harvard, 2021 - present:

Eric and Wendy Schmidt Center Postdoctoral Fellow

Mentors: Juan Caicedo and Caroline Uhler.

EDUCATION

Stanford University, 2015 - 2021: MS and PhD in Bioengineering:

MS in Bioengineering (synthetic biology, machine learning and statistics).

PhD Supervisor: Garry Nolan.

Thesis: Towards semantic representations of tissue organization from high-parameter imaging data.

University of Cambridge, 2014: MMath (with distinction):

Part III Mathematical Tripos. Thesis: Statistical properties of geometric flows and equidistribution. Supervisor: Prof. Vlad Markovic FRS.

University of Cambridge, 2013: BA Mathematics (first class honours):

Mathematical Tripos: top 10% in each year.

FIRST-AUTHOR biorxiv preprint, Aug. 2022: Deciphering causal genomic templates of complex **PUBLICATIONS** molecular phenotypes.

S.S. Bhate* et al. (corresponding author)

Cell Systems, Oct. 2021: Tissue schematics map the specialization of immune tissue motifs and their appropriation by tumors.

S.S. Bhate*, G. L. Barlow* et al.

European Journal of Immunology, Jan. 2021: Highly multiplexed tissue imaging using repeated oligonucleotide exchange reaction.

J. Kennedy-Darling*, S.S. Bhate*, J. Hickey* et al.

Cell, Aug. 2020: Coordinated cellular neighborhoods orchestrate antitumoral immunity at the colorectal cancer invasive front.

C. Schürch*, S.S. Bhate*, G. Barlow*, D. Phillips* et al.

OTHER.

Science Advances, in press: Immunotherapy of glioblastoma explants induces PUBLICATIONS interferon- γ responses and spatial immune cell rearrangements in tumor center, but not periphery

T. Shekarian et al.

BMC Bioinformatics, Jan. 2022: CellSeg: a robust, pre-trained nucleus segmentation and pixel quantification software for highly multiplexed fluorescence images M. Li et al.

Nature Communications, Nov. 2021: Immune cell topography predicts response to PD-1 blockade in cutaneous T cell lymphoma.

D. Phillips et al.

Nature Communications, June 2021: Subcellular localization of drug distribution by super-resolution ion beam imaging.

X. Rovira-Clave et al.

ResearchSquare preprint Feb. 2021: A tissue atlas of ulcerative colitis to guide TNF inhibitor therapy

A. Mayer et al.

Journal of Clinical Investigation, Jan. 2021: Landscape of coordinated immune responses to H1N1 challenge in humans. Z. Rahil et al.

Cell Stem Cell, Oct. 2018: Systematic identification of factors driving cell-fate

conversion using CRISPR activation screens. Y. Liu et al.

Cell, Aug. 2018: Deep profiling of mouse splenic architecture with CODEX multiplexed imaging.

Y.Goltsev et al.

BioBricks Foundation RFC 106, Mar 2015: A Standard Type IIS Syntax for Plants.

V. Rutten et al.

ACTIVITIES

Models, Inference and Algorithms, Broad Institute, Mar. 2022.

Neural Information Processing Systems, Los Angeles, Dec. 2017

Poster in computational biology workshop: "Unsupervised representation learning to interrogate cellular behaviours in 46-parameter imaging data"

<u>Poster</u> in computational biology workshop: "A multi-modal neural network for learning cis and trans regulation of stress response in yeast"

Keystone Single Cell 'Omics, Stockholm, May 2017

Poster: "Single-cell reference maps of tissue architecture using multiparameter imaging and unsupervised representation learning with neural networks" (SBSA Travel Award, \$1000)

AACR-CIMT-EATI-CRI International Cancer Immunotherapy Conference, New York, Sep. 2016

Awarded short talk: "Automatic identification of cellular niches using multiparameter imaging and deep learning"

(SBSA Travel Award, \$1000, BioX Oral Presentation travel award \$500)

Plants Workshop, iGEM Giant Jamboree, Nov. 2014, Boston: Conducted technical session on *M. polymorpha* in the plants for iGEM workshop.

Gold Medal, iGEM Nov. 2014, Boston: competed as member of University of Cambridge-JIC team

AWARDS AND

Bruce and Elizabeth Dunlevie Bio-X Stanford Interdisciplinary Graduate SCHOLARSHIPS Fellowship, 2016: full tuition and stipend support at elevated rate, 3 years.

> Clark Bioengineering Fellowship, 2015: full tuition and stipend support, 2 years. Cambridge PMC Bursary, 2014: £2000 grant for interdisciplinary collaboration. Senior Scholarship Prize, (2011, 2012, 2013, 2014) for excellent performance in Parts IA, IB, II and III of the Math Tripos.

Best A-Level Performance, King Edward VI Grammar School, 2010.

TEACHING

Ethics of Biotechniology, Stanford, Spring 2018: BioE131 teaching assistant. Hoefer award for mentorship of student writers.

Synthetic Biology lab, Stanford, Fall 2017: BioE44 teaching assistant

EXPERIENCE

Intern, Atum Technologies, April - June 2015

Protein engineering. Designed and performed high-throughput screen to optimize function of transposase variants in S. cerevisiae, finding several variants now in production.

Research assistant, Haseloff lab, Sep. 2014 - March 2015

Plant synthetic biology. Optimized constructs and protocols for electroporation of M. polymorpha spores.

Software engineering intern, FIS technologies, June - Sep. 2013

Implemented error logging tool for credit reporting software.

SKILLS AND INTERESTS

Programming: C++, Python, R, C#, (modelling/machine-learning/software development). Pytorch/Tensorflow. Basic: SQL, Java.

Language skills: Marathi (Fluent). German, Classical Greek, Latin, French (A-level) Other: Hindustani classical singing. Cycling and the outdoors.