# **Burak Ogan Mancarci**

2790 W 33rd Ave V6N 2G1 Vancouver, BC Canada ogan.mancarci@alumni.ubc.ca github.com/oganm Tel: +1 778 995 1591

111	Ca	1	$\sim$ 1	
,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	L-0		U	

2013-2018

PhD, Bioinformatics; University of British Columbia (Vancouver, Canada)

(expected)

Thesis title: Identification/validation of cell type marker genes of the brain and their

use in estimating cell type proportions in brain samples.

2009-2013

BSc, Molecular Biology And Genetics; Bilkent University (Ankara, Turkey)

## Research Experience

2014 - present

PhD Student at UBC Micheal Smith Laboratories - Supervisor: Dr. Paul Pavlidis

- Identification/validation of cell type marker genes of the brain and their use in estimating cell type proportions in brain samples (github.com/PavlidisLab/neuroExpressoAnalysis).
- Development of a web application to visualize gene expression in brain cell types (neuroexpresso.org).

2013

Rotation at BC Children's Hospital - Supervisor: Dr. Wyeth Wasserman

- Analysis of CAGE data for detection of microRNA transcription start sites.
- 2013

Rotation at Simon Fraser University - Supervisor: Dr. Fiona Brinkman

Analysis of antisense transcription in genomic islands.

2012

Summer internship at University of Zurich - Supervisor: Dr. Barbara Tschirren

 Selective mating of Japanese quails and computational analysis of various properties of quail and tit eggs.

2011

Summer internship at Harvard Medical School - Supervisor: Dr. George Daley

• Reprogramming of murine and human cells via viral vectors.

# **Teaching Experience**

2018

Teaching Assistant for Statistical Methods for High Dimensional Biology

course (STAT 540) at UBC - Instructors: Sara Mostafavi, Paul Pavlidis

2016

Teaching Assitant for Exploratory Data Analysis course (STAT 545) at UBC -

Instructor: Dr. Jenny Brian

2015 Teaching Assitant for Exploratory Data Analysis course (STAT 545) at UBC -

Instructor: Dr. Jenny Brian

2015 Instructional Skills Workshop at UBC - iswnetwork.ca

## Awards and Scholarships

2016 1st place in HackSeq hackaton - hackseq.com

2015 3rd place in SportsHack hackaton - sportshackweekend.org/ca/2015

2013-2014 Canadian Institutes of Health Research Training Program Scholarship

2009-2013 Bilkent 50% Scholarship

#### **Publications**

Toker, L., **Mancarci, B.O.**, Tripathy, S., and Pavlidis, P. (2018). Transcriptomic evidence for alterations in astrocytes and parvalbumin interneurons in bipolar disorder and schizophrenia subjects. Biological Psychiatry.

Tripathy, S., Toker, L., Bomkamp, C., **Mancarci, O.**, Belmadani, M., and Pavlidis, P. (2018). A critical assessment of single-cell transcriptomes sampled following patch-clamp electrophysiology. BioRxiv 298133.

**Mancarci, B.O.**, Toker, L., Tripathy, S.J., Li, B., Rocco, B., Sibille, E., and Pavlidis, P. (2017). Cross-Laboratory Analysis of Brain Cell Type Transcriptomes with Applications to Interpretation of Bulk Tissue Data. ENeuro 4.

Tripathy, S.J., Toker, L., Li, B., Crichlow, C.-L., Tebaykin, D., **Mancarci, B.O.**, and Pavlidis, P. (2017). Transcriptomic correlates of neuron electrophysiological diversity. PLOS Computational Biology 13, e1005814.

Horvath, G.A., Demos, M., Shyr, C., Matthews, A., Zhang, L., Race, S., Stockler-Ipsiroglu, S., Van Allen, M.I., **Mancarci, O.**, Toker, L., et al. (2016). Secondary neurotransmitter deficiencies in epilepsy caused by voltage-gated sodium channelopathies: A potential treatment target? Mol. Genet. Metab. 117, 42–48.

Onder, T.T., Kara, N., Cherry, A., Sinha, A.U., Zhu, N., Bernt, K.M., Cahan, P., **Mancarci, B.O.**, Unternaehrer, J., Gupta, P.B., et al. (2012). Chromatin-modifying enzymes as modulators of reprogramming. Nature 483, 598–602.

#### Software

**NeuroExpresso:** A web application for visualization of gene expression data in brain cell types. Available at **neuroexpresso.org**.

markerGeneProfile: An R package for calculation of marker gene profiles as described in Mancarci et al. (2017). Available at github.com/PavlidisLab/markerGeneProfile.

**VASCO:** A web application for visualization of gene expression data from single cell RNA sequencing experiments. Developed for HackSeq hackaton. Available at **hackseq.github.io/vasco**.

**Impact Replays:** A web application for visualization play-by-play data from football games. Developed for SportsHack hackaton. Available at **daattali.com/shiny/cfl**.

**eggstractor:** A GUI based software to automate extraction of shape and pigmentation related information from quail eggs. Developed for 2012 summer internship under Barbara Tschirren. Available at **github.com/oganm/eggstractor**.

**Misc. Bioinformatics related R packages:** I have been maintaining various R packages aimed at making life easier for myslef and other bioinformaticians. See **github.com/topics/ogan-bio** for a full list.

### Presentations

**Society for Neuroscience Annual Meeting 2017: Mancarci, B.O.**, Toker, L., Li, B., Rocco, B., Tripathy, S., Sibille, E., and Pavlidis, P. NeuroExpresso: A brain cell type specific gene expression database composed of pooled microarray and single cell RNA sequencing data.

**25th Annual International Conference on Intelligent Systems for Molecular Biology: Mancarci, B.O.**, Toker, L., Li, B., Rocco, B., Tripathy, S., Sibille, E., and Pavlidis, P. (2017). NeuroExpresso: Cross laboratory database of brain cell type specific gene expression.

**Society for Neuroscience Annual Meeting 2016:** Toker, L., **Mancarci, B.O.**, Tripathy, S., and Pavlidis, P. (2016). Deciphering the cell-type specific component in the pathophysiology of brain-related disorders.

**Society for Neuroscience Annual Meeting 2016:** Tripathy, S., Tebaykin, D., **Mancarci, O.**, Toker, L., and Pavlidis, P. (2016). Transcriptomic correlates of brain-wide electrophysiological diversity.

**Society for Neuroscience Annual Meeting 2016: Mancarci, O.**, Toker, L., and Pavlidis, P. (2016). Comparison of single cell and pooled cell expression data from mouse and human brain.

**24th Annual International Conference on Intelligent Systems for Molecular Biology: Mancarci, O.**, Toker, L., Li, B., Tripathy, S., and Pavlidis, P. (2016). Identification of novel markers for mammalian brain cell types.

**Organization of Computational Neurosciences Conference 2015:** Tripathy, S.J., Tebaykin, D., Li, B., **Mancarci, O.**, Toker, L., and Pavlidis, P. (2015). Large-scale analysis of brain-wide electrophysiological diversity reveals novel characterization of mammalian neuron types. BMC Neurosci 16, O4.

**23rd Annual International Conference on Intelligent Systems for Molecular Biology: Mancarcı, O.**, Toker, L., Tripathy, S., Pavlidis, P., Mancarcı, O., Toker, L., Tripathy, S., and Pavlidis, P. (2015). A comprehensive database of cell-type specific marker genes for the mammalian brain. F1000Research 4.

**23rd Annual International Conference on Intelligent Systems for Molecular Biology:** Toker, L., **Mancarci, O.**, Tripathy, S., and Pavlidis, P. (2015). A transcriptomics approach for revealing cell-type proportion changes in psychiatric disorders.