

# Burak Ogan Mancarci

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## Education

- 2013-2019 (expected)**      **PhD, Bioinformatics;** University of British Columbia (Vancouver, Canada)  
*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](https://github.com/PavlidisLab/neuroExpressoAnalysis)).
  - Development of a web application to visualize gene expression in brain cell types ([neuroexpresso.org](http://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 Assistant for Exploratory Data Analysis course (STAT 545) at UBC - Instructor: Dr. Jenny Brian
- 2015 Instructional Skills Workshop at UBC - [iswnetwork.ca](http://iswnetwork.ca)

## Awards and Scholarships

- 2016 1st place in HackSeq hackaton - [hackseq.com](http://hackseq.com)
- 2015 3rd place in SportsHack hackaton - [sportshackweekend.org/ca/2015](http://sportshackweekend.org/ca/2015)
- 2013-2014 Canadian Institutes of Health Research Training Program Scholarship
- 2009-2013 Bilkent 50% Scholarship

## Publications

- Tripathy, S.J., Toker, L., Bomkamp, C., **Mancarci, B.O.**, Belmadani, M., and Pavlidis, P. (2018). Assessing Transcriptome Quality in Patch-Seq Datasets. *Front. Mol. Neurosci.* 11.
- 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*.
- 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](http://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](https://github.com/PavlidisLab/markerGeneProfile).
- VASCO:** A web application for visualization of gene expression data from single cell RNA sequencing experiments. Developed for HackSeq 2016. Available at [hackseq.github.io/vasco](http://hackseq.github.io/vasco).
- Viral Voyager:** A web application for visualization and analysis of metavirome data collected for the Tara Oceans Project. Developed for HackSeq 2018. Available at [oganm.com/shiny/taracyc](http://oganm.com/shiny/taracyc).

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

**eggstractor:** 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](https://github.com/oganm/eggstractor).

**Misc. Bioinformatics related R packages:** I maintain R packages aimed at making life easier for myself and other bioinformaticians. See [github.com/topics/ogan-bio](https://github.com/topics/ogan-bio) for a full list.

**Misc. D&D related repositories:** I maintain R packages and web applications to be used while playing Dungeon's and Dragons, a popular tabletop roleplaying game. See [github.com/topics/ogan-dnd](https://github.com/topics/ogan-dnd) 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: Mancarci, O.,** Toker, L., Tripathy, S., Pavlidis, P., Mancarci, 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.