PART I: General Information

Name: Hasan H. Otu DATE PREPARED: November 20, 2018

Office Address: University of Nebraska-Lincoln, Department of Electrical and Computer

Engineering, 209N SEC, Lincoln, NE 68588-0511, USA

Education:

Year	Degree	Institution
1996	B.S.	Electrical and Electronics Engineering, Bogazici
		University, Istanbul, Turkey
1997	M.S.	Electrical and Electronics Engineering, Bogazici
		University, Istanbul, Turkey
2002	Ph.D.	Electrical Engineering, University of Nebraska-
		Lincoln, USA

Academic Appointments:

Title	Institution
Teaching Asst.	Bogazici University, Department of Electrical and
	Electronics Engineering
Teaching Asst.	University of Nebraska-Lincoln, Department of
	Electrical Engineering
Instructor of EE	University of Nebraska-Lincoln, Department of
	Electrical Engineering
Research Fellow	Harvard Medical School, Boston, MA
Instructor of Medicine	Harvard Medical School, Boston, MA
Assistant Prof.	Istanbul Bilgi University Department of
	Bioengineering, Istanbul, Turkey
Professor	University of Nebraska-Lincoln, Department of
	Electrical Engineering (Tenure 2013)
	Teaching Asst. Teaching Asst. Instructor of EE Research Fellow Instructor of Medicine Assistant Prof.

Visiting and Other Academic Appointments:

Year	Title	Institution
2005-2008	Adjunct Instructor	Boston Uni. Bioinformatics Program, Boston, MA
2005-2006	Teaching Instructor	Northeastern University Department of Biology,
		Graduate Program in Bionformatics, Boston, MA
2006-2007	Assistant Prof.	Yeditepe University Department of Genetics and
		Bioengineering, Istanbul, Turkey
2008	Assistant Prof.	Sabanci University Department of Biological
		Sciences and Bioengineering, Istanbul, Turkey

Major Administrative Responsibilities:

Year	Title	Institution
2003-2007	Director	Bioinformatics Core, BIDMC Genomics Center
2004-2012	Associate Director	Dana-Farber/Harvard Cancer Center Proteomics
		Core
2006-2008	Steering Committee	Biotechnology Institute, Yeditepe University
2007-2012	Associate Director	Bioinformatics Core, BIDMC Genomics Center
2010-2013	Chair	Department of Bioengineering, Istanbul Bilgi U.
2016-	Advisory Committee	Department of Bioengineering, Istanbul Bilgi U.

Professional Societies:

Year	Role	Society
2004-	Member	American Association for the Advancement of
		Science
2004-	Member	International Society for Computational Biology

Editorial Boards and Program Committees:

<i>Year</i> 2000-	Role Ad hoc reviewer	Journal/Conference IEEE Transactions on Communications, Bioinformatics, Systematic Biology, Journal of Computational Chemistry, Journal of Molecular Modeling, Cancer Informatics, Journal of Molecular Evolution, BMC Bioinformatics, BMC Medical Genomics, Journal of Computational Biology, Molecular Simulation, The Computer Journal, African Journal of Biotechnology, European Journal of Human Genetics, Journal of Theoretical Biology, Acta Biotheoretica, Journal of Heredity, IEEE/ACM Transactions on Computational Biology and Bioinformatics, Anatolian Journal of Cardiology, Biometrical Journal, Data Compression Conference, Entropy, Electro-Information Technology Conference, Nature Scientific Reports, PLOS One, Statistical Bioscience, Current Bioinformatics
2010-2017	Associate Editor	EURASIP Journal on Bioinformatics and Systems Biology
2011	PC Co-chair	The 6 th International Symposium on Health Informatics and Bioinformatics, (HIBIT)
2012	Organizer	Workshop, "Bioinformatics Approaches for Analysis of High-throughput Biological Data", International Centre for Genetic Engineering and Biotechnology
2013-2017	Associate Editor	Advances in Biology
2014-2018	Editor in Chief	Journal of Bioinformatics, Computational and Systems Biology
2014-	Associate Editor	Journal of Bioinformatics, Proteomics and Imaging Analysis
2017	Member	Technical Program Committee Electro-Information Technology Conference

2017 Member Program Committee International Symposium on

Integrative Bioinformatics

2018- Associate Editor Advances in Bioinformatics

Awards and Honors:

Year Name of Award

- 1992 Turkish Government scholarship for undergraduate studies (Ranked 27th Nationwide)
- 1997 Turkish Oil Foundation monetary award for graduate studies
- 2001 Ranked 1st in Graduate Student Research Paper Competition, Department of Electrical Engineering, University of Nebraska-Lincoln
- 2001 Travel Award, 3rd Georgia Tech-Emory International Conference on Bioinformatics
- 2003 Cited by the National Physical Laboratory of the United Kingdom in its recommendations to the UK government as work that should be studied in order to meet future challenges in the bioinformatics area in NPL Report CMSC 23/03 Report to the National Measurement System Directorate, Department of Trade and Industry New Directions Software Issues in Bioinformatics
- 2003 Invited Presentation and Travel Award, International Stem Cell Conference, Singapore
- 2003 Best Study Award, "Transcriptional Profiling for Detection of a Gene Signature in Renal Cell Cancer" 55th Congress of the German Urological Society, Hamburg.
- 2004 Second Best Study Award, "Gene expression profiles in Renal Cell Cancer: Characterization of gene signatures in various histological subtypes and application of a metastatic signature" 56th Congress of the German Urological Society, Hamburg.
- 2008 Research Grant Award, Dubai Harvard Foundation for Medical Research
- 2017 Outstanding Paper, Second Place Award, The International Conference on Electro-Information Technology.

Part II: Research, Teaching, and Clinical Contributions

A. Narrative Report of Research, Teaching, and Clinical Contributions.

My career to date has involved research in Information Theory and Bioinformatics, which involves computational methods to organize and analyze biological data. My earlier work was in the areas of image compression and joint source/channel coding. With the availability of genomic sequences, my interest shifted into bioinformatics by trying to understand how information is organized in DNA sequences. Using Information Theory, I developed characterizations of this organization, with applications to fragment assembly and phylogeny reconstruction which led to two US patents.

My research at Harvard Medical School (HMS) focused on management and analysis of high-throughput biological data (HTBD) in the context of functional genomics and proteomics. These included, but were not limited to, transcription profiling (e.g. Affymetrix GeneChip system), proteomics (e.g. SELDI TOF, MALDI TOF/TOF), and genotyping (e.g. Affymetrix SNP arrays) efforts targeting questions in computational biology, systems biology, cancer, stem cells, heart disease, diabetes, and obesity. As the director of Bioinformatics Core at BIDMC Genomics Center, I managed the core to establish a state-of-the-art computer infrastructure and a research web portal (www.bidmcgenomics.org), which functioned as the front end for automated experiment ordering, data storage and analysis. As part of this portal, we designed databases for various data types including gene microarrays and protein chips. We developed embedded analysis tools and analytical methods for HTBD, which resulted in two stand alone computer programs. Specifically, I focused

on problems regarding data normalization, differential expression, clustering, functional group/pathway analysis and biomarker discovery.

Currently I have been working on analysis of HTBD within the context of networks, along the lines of systems biology. We use probabilistic graph models, e.g Bayesian Networks, to answer two questions: (i) what are the active known pathways given experimental data? (ii) what are the interaction networks based on observed data? Our goal is to incorporate existing biological knowledge in interpreting specific experimental results in the context of pathways. I have also recently led next-generation sequencing projects regarding whole genome sequencing and metagenomics.

In addition to research efforts, I have supervised and trained summer interns, IT personnel, BS, MS, and PhD students totaling around 30 people over the last 16 years. My teaching experience involves assisting in undergraduate Electrical Engineering (EE) courses at Bogazici University and University of Nebraska-Lincoln (UNL), teaching undergraduate/graduate EE, Bioinformatics, and Bioengineering courses at UNL, Northeastern University, Yeditepe University, Sabanci University, Acibadem University and Istanbul Bilgi University.

B. Funding Information

2002-2003	NIH/NIDDK; Project No: 5U24DK058739-03; "NIDDK Biotechnology Center";
2002 2003	Libermann (PI); \$595,856; Role: Investigator
2002-2007	NIH/NCI; Project No: PO1 CA92664-03; "Spatial and Temporal Regulation of
	Angiogenesis"; Dvorak (PI); \$141,237; Role: Investigator
2003-2005	NIH/NIAID; Project No: P01 AI041521; "Costimulation and Cytokines in
	Tolerance"; Turka (PI); \$1,459,884; Role: Investigator
2003-2005	NIH/NCI; Project No: 1R21 CA108303-01; "Proteomics and Biomarkers for
	Hepatocellular Cancer"; Afdahl (PI); \$100,000; Role: Investigator
2004-2009	NIH/NCI; Project No: P01 HL076540; "Endothelial Cell Phenotypes in Health and
	Disease", Aird (PI); \$80,000; Role: Investigator
2005-2007	NIH/NIAID; Project No: R21 CA107352-01; "Novel Approaches to Gene Profiling
	in Ovarian Cancer"; Libermann (PI); \$86,000; Role: Investigator
2006-2010	Michigan State University-NAY Project; Project No: MSU 95464; "Direct
	Dedifferentiation of Primary Somatic Cells", Cibelli (PI); Role: Consultant
2007-2010	King Abdulaziz City for Science and Technology; Project No: 26-64; "Camel
	Genome Project Phase I"; Al-Swailem (PI), Otu (Co-PI); \$519,281.
2009-2012	The Dubai Harvard Foundation for Medical Research; "Analysis of high-throughput
2011 2012	genomic data using an integrated approach; Otu (PI); \$182,000
2011-2013	Istanbul Bilgi University Research Fund; "Human Whole Genome Sequencing"
2011 2012	Otu (PI); \$25,000
2011-2013	The Scientific and Technological Research Council of Turkey; Project No: 111E042;
	"Bayesian Network Analysis of High Throughput Biological Data: A Systems
2016-2019	Biology Approach"; Otu (PI); \$85,000
2010-2019	NIH/NIA; Project No: R01AG051658; "Advancing the Understanding of Postoperative Delirium Mechanisms via Multi-Omics"; Marcantonio/Libermann
	(MPI's), Otu (Co-PI); ~\$2.3M
2018-2020	NIH/NLM; Project No: R21LM012759; "Identification and characterization of
2010-2020	interaction at lases in human"; Otu (PI); \$443,862
2018-2024	NIH/NIA; Project No: P01AG031720; "Delirium, Dementia, and the Vulnerable
2010 2021	Brain: An Integrative Approach"; Inouye (PI); "The role of inflammation in the

pathophysiology of delirium and its associated long term cognitive decline (Project 2)"; Marcantonio/Libermann (MPI's), Otu (Co-PI); ~\$13.6M

In Preparation/Pending:

Electrochemical Chip to Profile Circulating miRNA at Attomolar Level with Zero Background (Submitted as a BRG R01, Saraf – PI, Otu – Co-PI)

Pathway Connectivity Maps: Bisociation of Interaction Networks Using Network Pruning and Compressive Sensing (To be submitted as an R01, Otu – PI)

Somatic cell's acquisition of pluripotency in Zebrafish (To be resubmitted as an R01, Cibelli – PI, Otu – Co-PI)

C. Report of Current Research Activities

<u>Project</u>	Role
Gene Interaction Atlas Generation	Method Development/Supervision
Applications of Random Matrix Theory on Biological Networks	Method Development/Supervision
Predictive Models for Methylation Disposition of CpG Islands	Method Development/Supervision
Using Probabilistic Graph Representations for Multi-omic Data Integration	Method Development/Supervision
Comparative Analysis of Gastrointestinal Cancers using Network Theory	Method Development/Supervision
Correlation Network/Pathway Analysis using Network Pruning, Bisociation, and Compressive Sensing	Method Development/Supervision
Biomarker discovery in NASH disease	Method Development/ Data Analysis (w/Harvard Medical School)
TGFB3 involvement in cleft palate	Method Development/ Data Analysis (w/UNMC)
Effect of Simvastatin in Bone Regenaration Following Dental Grafts	Method Development/ Data Analysis (w/UNMC)
Exosome Proteomics	Method Development/ Data Analysis (w/Harvard Medical Scool)
Multi-omics of Delirium	Method Development/ Data Analysis/Leadership/Training (w/Harvard Medical School)
Mechanisms of Cellular Reprogramming	Method Development/Data Analysis (w/Michigan State University)
Alternative splicing in pancreatic cancer	Method Development/ Data Analysis (w/Harvard Medical School)
Proteomics of IBD	Method Development/ Data Analysis (w/Harvard Medical School)

D. Report of Teaching

Graduate and Undergraduate Courses

i.	Bogazici University, Istanbul, Turkey
1996	Department of Electrical and Electronics Engineering. EE 374 Communication Engineering (current listing). Lecturer. ~20 senior EE students. Teaching: 2 hrs/week. Preparation: 3hrs/week. Duration: Fall Semester.
1997	Department of Electrical and Electronics Engineering. EE 477 DigitalCommunication (current listing). Lecturer. ~20 senior EE students. Teaching: 2 hrs/week. Preparation: 3hrs/week. Duration: Spring Semester.
1997	Department of Electrical and Electronics Engineering. EE 210 Introduction to Electrical Engineering (current listing). Lecturer. ~50 sophomore/junior EE students. Teaching: 2 hrs/week. Preparation: 3hrs/week. Duration: Spring Semester.
ii.	University of Nebraska-Lincoln, Lincoln, NE USA
2000-2002	Department of Electrical Engineering ELEC 464/864 Digital Communication Systems. Core Faculty. ~10 senior Electrical Engineering students and ~10 Electrical Engineering graduate students. Teaching: 3hrs/week. Preparation: 5hrs/week. Duration:
2000-2002	Spring Semester (each year). Department of Electrical Engineering. ELEC 462/862 Communication Systems. Core Faculty. ~10 senior Electrical Engineering students and ~10 Electrical Engineering graduate students. Teaching: 3hrs/week. Preparation: 5hrs/week. Duration: Fall Semester (each year).
2013-	Department of Electrical and Computer Engineering. ELEC 450/850 Bioinformatics. Core Faculty. ~15 senior/graduate engineering students. Teaching: 3hrs/week. Preparation: 5hrs/week. Duration: Fall Semester (each year).
2013-	Department of Electrical and Computer Engineering. ELEC 498/898 Computational and Systems Biology. Core Faculty. ~15 senior/graduate engineering students. Teaching: 3hrs/week. Preparation: 5hrs/week. Duration: Spring Semester (each year).
2016-	Department of Electrical and Computer Engineering. ELEC 996 Bayesian Networks. Core Faculty. ~15 graduate engineering/sciences students, Teaching: 3hrs/week. Preparation: 5hrs/week. Duration: Spring Semester (each odd year).
2016-	Department of Electrical and Computer Engineering. ECEN 215 Electronics and Circuits I. Core Faculty. ~90 undergraduate engineering/sciences students, Teaching: 3hrs/week. Preparation: 5hrs/week. Duration: Fall Semester (each year).
iii.	Northeastern University, Boston, MA USA

2005	Biology Department, Graduate Program in Bioinformatics. BIO G385, Seminar in Bioinformatics. Core Faculty. ~5 Bioinformatics graduate students. Teaching: 2hrs/week. Preparation: 5hrs/week. Duration: Fall Semester.
iv.	Yeditepe University Istanbul, Turkey
2007	Department of Genetics and Bioengineering GBE 313, Experimental Bioengineering Lab. Core Faculty. 6 GBE undergraduate students. Teaching: 4hrs/week. Preparation: 5hrs/week. Duration: Fall Semester.
2007	Department of Genetics and Bioengineering GBE 311, Principles of Bioengineering. Core Faculty. 13 GBE undergraduate students. Teaching: 3hrs/week. Preparation: 5hrs/week. Duration: Spring Semester.
v.	Sabanci University Istanbul, Turkey
2008	Department of Biological Sciences and Bioengineering BIO 512, Advanced Computational Bilogy. Core Faculty. 3 graduate students; 9 participants. Teaching: 3hrs/week. Preparation: 7hrs/week. Duration: Spring Semester.
vi.	Acibadem University Istanbul, Turkey
2010	Medical School, MED 106, Medical Informatics. Core Faculty. 23 undergraduate students. Teaching: 5hrs/week. Preparation: 5hrs/week. Duration: Spring Semester.
vii.	Istanbul Bilgi University Istanbul, Turkey
2010	College of Engineering, ENG 179, Engineering in Society. Guest Lecturer. 12 undergraduate students. Teaching: 3hrs lecture on Human Genome Project. Duration: Fall Semester.
2011	College of Engineering, ENG 180, Engineering and Sciences. Adjunct Faculty. 43 undergraduate students. Teaching: 10hrs lecture on Introduction to Bioengineering. Duration: Spring Semester.
2011	College of Engineering, PHYS 101 Physics I. Adjunct Faculty. 43 undergraduate students. Teaching: Problem Sessions 2hrs/week. Duration: Spring Semester.
2011	College of Engineering, PHYS 100 Physics for Scientists and Engineers. Faculty. ~60 undergraduate students. Teaching: Lecture, 2 sections, 3 hrs/week per section. Problem Sessions / Laboratory, 3 sections, 2hrs/week per section. Duration: Fall Semester.
2012	College of Engineering, PHYS 100 Physics for Scientists and Engineers. Faculty. ~45 undergraduate students. Teaching: Lecture, 3 hrs/week. Duration: Spring Semester.
2012	College of Engineering, ENGR 230 Probability and Random Processes. Faculty. ~30 undergraduate students. Teaching: Lecture, 2 hrs/week. PS / Lab, 2hrs/week. Duration: Spring Semester.

2012	College of Engineering, PHYS 100 Physics for Scientists and
	Engineers. Faculty. ~65 undergraduate students. Teaching: Lecture, 3
	hrs/week. Duration: Fall Semester.
2012	College of Engineering, BIOE 341 Bioinformatics. Faculty. ~10
	undergraduate students. Teaching: Lecture, 2 hrs/week. PS / Lab,
	2hrs/week. Duration: Fall Semester.
2013	College of Engineering, BIOE 346 Microarrays. Faculty. 8
	undergraduate students. Teaching: Lecture, 3 hrs/week. Duration:
	Spring Semester.
2013	College of Engineering, BIOE 241 Fundamentals of Biostatistics and
	Experimental Design. Faculty. ~30 undergraduate students. Teaching:
	Lecture, 3 hrs/week. PS / Lab, 2hrs/week. Duration: Spring Semester.
2013	Experimental Design. Faculty. ~30 undergraduate students. Teaching:

Local Invited Teaching Presentations

2002	Characterization of DNA Sequences. BIDMC Genomics Center
	Invited Lecture. Attending: ~20 HMS Faculty, Post-doctoral Fellows
	and Residents. Presentation and Follow-up: 5 hrs. Preparation: 20 hrs.
2003	Networks. BIDMC Genomics Center Core Meeting. Attending ~15
	Post-doctoral Fellows and Residents. Presentation: 1 hr. Preparation:
	10 hrs.
2004	Bioinformatics Core at BIDMC Genomics Center. MIT CSBI
	BioMicro Center. ~50 Faculty, Post-doctoral fellows and graduate
	students. Presentation and Follow-up: 2 hrs. Preparation: 10 hrs.
2005	Progress of Challenges in Bioinformatics: From Sequence to Function
	to Networks. Boston University Bioinformatics Program. ~20
	Faculty, Post-doctoral fellows and graduate students. Presentation and
	Follow-up:5 hrs. Preparation: 20 hrs.

Advisees and trainees

2003	Jian Li	PhD student at Baylor College of Medicine
2003	Charles Bailey	Student at Tufts School of Veterinary
		Medicine
2003	Chris Porter	Children's Hospital IT department
2003	Osman Osman	Student at MIT EECS Dept.
2003-2006	Shakir A. Kolia	Research Associate at BIDMC Genomics
		Center Bioinformatics Core
2005-2006	Taehyun Park	Research Associate at BIDMC Genomics
		Center
2006-2008	Al-Arawi MS, Al-Kh	ider AY, Al-Muhaimeed AN, Al-Qahtani FH,
	Al-Manee MM, Al-Si	homrani BM (KACST Bioinformatics Group)
2007-2013	Senol Isci	PhD student at Bogazici University
		Biomedical Engineering Institute
2007-2009	Caner Akdemir	Undergraduate student at Yeditepe
		University Department of Computer
		Engineering and Department of Genetics and
		Bioengineering
2007-2011	Cem Meydan	PhD student at Sabanci University Department
		of Biological Sciences and Bioengineering

2007-2011	Aydin Albayrak	PhD student at Sabanci University Department of Biological Sciences and Bioengineering
2007-2011	Yasin Bakis	PhD student at Sabanci University Department of Biological Sciences and Bioengineering
2010-2013	Haluk Dogan	Teaching Assistant at Istanbul Bilgi University, Department of Bioengineering and MS student at Bogazici University Department of Computer Engineering
2011-2013	Umut Agyuz	MS Student, Bogazici University Institute of Biomedical Engineering
2011-2013	Melike Korucuoglu	MS Student, Bogazici University Department of Computer Engineering
2013-2016	Haluk Dogan	PhD student at University of Nebraska-Lincoln, Department of Electrical and Computer Engineering
2014-2016	Zeynep Hakguder	PhD student at University of Nebraska-Lincoln, Department of Electrical and Computer Engineering
2014-	Dicle Yalcin	PhD student at University of Nebraska-Lincoln, Department of Electrical and Computer Engineering
2016-	Sree Chanumolu	Postdoctoral researcher at University of Nebraska-Lincoln, Department of Electrical and Computer Engineering
2017-	Bridget Tripp	PhD student at University of Nebraska-Lincoln, Program in Complex Biosystems

Regional, national, or international contributions

1997	A Compression Algorithm that Preserves NDVI and NDWI Values.
	Conference Presentation. Asilomar Conference on Circuits, Systems
	and Computers. Monterey, California, USA.
1998	A Joint Source Channel Coder with Block Constraints. Conference
	Presentation. IEEE International Conference on Acoustics, Speech,
	and Signal Processing. Seattle, Washington, USA.
1999	Issues in Joint Source Channel Coding. Seminar. UNL EE Dept.
	Journal Club. Lincoln, NE USA.
2001	A New Approach to Sequence Assembly Using Divide and Conquer
	Algorithms. Conference Presentation. 3rd Georgia Tech-Emory
	International Conference on Bioinformatics. Atlanta, Georgia, USA.
2002	An Information-theoretic Sequence Distance Measure with
	applications to Phylogeny Analysis. Seminar. UNL EE Dept. Journal
	Club. Lincoln, NE USA.
2004	A Seminar in Bioinformatics: Looking for Familiar Faces in the
	Neighborhood. Invited Lecture. Bogazici University, Institute of
	Biomedical Engineering, Istanbul, Turkey.
2004	Challenges in Bioinformatics: DNA Sequence Analysis and Frontiers
	in Functional Genomics. Invited Lecture. Sabanci University, Faculty
	of Engineering and Natural Sciences, Istanbul, Turkey.

2004	From Sequence to Function: Issues in Computational Biology. Invited Lecture. Koc University, Department of Chemical and Biological
	Engineering, Istanbul, Turkey.
2006	Progress of Challenges in Bioinformatics: From Sequence to Function to Networks. Invited Lecture. Yeditepe University, Department of
	Genetics and Bioengineering, Istanbul, Turkey.
2007	Challenges in Bioinformatics: Invited Lecture. King Abdulaziz City for Science and Technology, Riyadh, KSA.
2007	Experimental Design and Analysis of High-Throughput Biological Data. Invited Lecture. Sabanci University, Faculty of Engineering and
	Natural Sciences, Istanbul, Turkey.
2007	DNA Sequence Analysis and Applications in Functional Genomics. Seminar. Bogazici University, Institute of Biomedical Engineering,
2007	Istanbul, Turkey.
2007	Algorithmic and practical approaches to issues in Bioinformatics.
2008	Seminar. Izmir Institute of Technology, Izmir, Turkey.
2008	Computational Approaches in DNA Sequence Analysis and Functional Genomics and Proteomics. Seminar. Bilgi University,
	Istanbul, Turkey.
2008	Computational Approaches in DNA Sequence Analysis and
2000	Functional Genomics and Proteomics. Seminar. Halic University,
	Istanbul, Turkey.
2009	Biomarker Discovery – Pregnancy Success. Seminar. Michigan State
	University, East Lansing, MI USA
2009	Analysis and Applications of High-throughput Biological Data.
	Seminar. University of Nebraska Medical Center, Lincoln/Omaha, NE USA
2009	Bioinformatic Approaches for High-throughput Biological Data
	Analysis. Seminar. Middle East Technical University, Ankara, Turkey
2010	Looking for Familiar Faces in the Old Neighborhood. Invited Lecture.
	Bogazici University, Department of Electrical and Electronics
	Engineering, Istanbul, Turkey.
2010	From Sequence to Function to Networks: Analysis Issues in
	Bioinformatics. Istanbul Technical University, Program in
	Biomedical Engineering, Istanbul, Turkey.
2010	Sequence, Function, and Networks based Analysis Issues in
	Bioinformatics. Istanbul University, Institute for Experimental
	Medicine, Istanbul, Turkey. Similar talk is given at N.K.U. Faculty of
	Engineering Corlu, Tekirdag, Kadir Has University, Fatih University,
	Bogazici University (Department of Computer Engineering), Bilgi
2010	University, Pakize Tarzi Laboratories, all in Istanbul, Turkey
2010	Algorithms in Bioinformatics, 9th National Medical Genetics Congress, Istanbul Turkey
2011	Contemporary Issues in and Applications of Computational Biology,
2011	Inonu University, School of Medicine, Malatya, Turkey
2011	Bioengineering Education in Turkey, Yildiz Technical University,
2011	Bioengineering Days.
2011	Bayesian Network based pathway analysis of microarray data,
	European Biotechnology Congress, Istanbul, Turkey

2012	Systems Biology, Bogazici University, Molecular Biology and
	Genetics Weekend, Istanbul, Turkey
2012	Bioinformatics, ITU Biotech, Istanbul, Turkey
2012	A Crash Course on Microarray Data Analysis, DONE Genetics and
	Bioinformatics, Istanbul, Turkey
2012	Bayesian Pathway Analysis, Sabanci University, Istanbul, Turkey
2012	HTBD Analysis within a BN Framework, Istanbul University,
	Institute for Experimental Medicine, Istanbul, Turkey.
2014	Pathway Analysis of Biological Data using Bayesian Networks.
	University of Nebraska Medical Center, Omaha, NE USA
2018	Keynote Speaker, UNL Plant Science Retreat Network Analysis of
	Multiomic Data Using Probabilistic Graph Representations

Description of major curriculum offerings, teaching cases or innovative educational programs developed

2007	Development of Undergraduate Curriculum at Yeditepe University,
	Department of Genetics and Bioengineering, Istanbul, Turkey.
2007	Development of Graduate Curriculum (both MS and PhD) at Yeditepe
	University, Bioengineering Institute, Istanbul, Turkey.
2010	Development of Undergraduate Curriculum at Istanbul Bilgi
	University, Department of Bioengineering, Istanbul, Turkey.
2013	Development of Bioinformatics Program at University of Nebraska-
	Lincoln, Department of Electrical and Computer Engineering.

Part III: Bibliography

Original Articles

- 1. Otu HH, Sayood K. "A joint source/channel coder with block constraints" *IEEE Transactions on Communications* 1999; 47 (11): 1615-1618.
- 2. Sayood K, Otu HH, Demir N. "Joint source/channel coding for variable length codes" *IEEE Transactions on Communications* 2000; 48 (5): 787-794.
- 3. Otu HH, Sayood K. "A divide and conquer approach to fragment assembly" *Bioinformatics* 2003; 19:22-29.
- 4. Otu HH*, Fortunel NO*, Ng HH*, Chen J, Mu X, Chevassut T, Li X, Joseph M, Bailey C, Hatzfeld JA, Usta F, Vega VB, Long PM, Liberman TA, Lim B. "Comment on 'Stemness: Transcriptional Profiling of Embryonic and Adult Stem Cells' and 'A Stem Cell Molecular Signature" *Science* 2003; 302: 393b.
- 5. Otu HH, Sayood K. "A new sequence distance measure for phylogenetic tree construction" *Bioinformatics* 2003; 19:2122-2130.
- 6. Bastola DR, Otu HH, Doukas SE, Sayood K, Hinrichs SH, Iwen PC. "Utilization of the relative complexity measure to construct a phylogenetic tree for fungi" *Mycological Research* 2004; 108(2):117-125. [This journal is called "Fungal Biology" as of Jan. 2010].
- 7. Voisine P, Ruel M, Khan TA, Bianchi C, Xu SH, Kohane I, Libermann TA, Otu HH, Saltiel AR, Sellke FW "Differences in gene expression profiles of diabetic and non-diabetic patients undergoing cardiopulmonary bypass and cardioplegic arrest" *Circulation* 2004; 110:II-280-286.

_

[•] These authors contributed equally to this work

- 8. von Stechow D, Zurakowski D, Pettit AR, Muller R, Gronowicz G, Otu HH, Libermann TA, Alexander JM "Differential transcriptional effects of PTH and estrogen during anabolic bone formation" *J. Cell. Biochem.* 2004; 93:476-490.
- 9. Aivado M, Spentzos D, Alterovitz G, Otu HH, Grall F, Porter C., Cho JY, Giagounidis AAN, Germing U, Ramoni M, Libermann TA "Optimization and evaluation of surface-enhanced laser desorption/ionization time-of-flight mass spectrometry (SELDI-TOF MS) with reversed-phase protein arrays for protein profiling" *Clinical Chemistry and Laboratory Medicine* 2005; 43(2), 133-140.
- 10. Jones J, Otu HH, Spentzos D, Kolia S, Inan M, Beecken WD, Fellbaum C, Gu X, Joseph M, Jonas D, Libermann TA. "Gene signatures of progression and metastasis in Renal Cell Cancer" *Clinical Cancer Research* 2005 11:5730-5739.
- 11. Spentzos D, Levine DA, Kolia S, Otu HH, Boyd J, Libermann TA, Cannistra SA. "Unique gene expression profile based upon pathologic response in epithelial ovarian cancer" *Journal of Clinical Oncology* 2005 23(31):7911-7918.
- 12. Wada Y, Otu HH, Wu S, Abid R, Okada H, Libermann TA, Kodama T, Shih S-C, Minami T, Aird WC. "Preconditioning of primary human endothelial cells with inflammatory mediators alters the "set point" of the cell" *FASEB Journal* 2005 19(13):1914-1916.
- 13. Ijiri K, Zerbini LF, Peng H, Correa RG, Lu B, Walsh N, Zhao Y, Taniguchi N, Huang XL, Otu HH, Hong W, Wang JF, Komiya S, Ducy P, Rahman MU, Flavell RA, Libermann TA, Goldring MB. "A novel role for GADD45β as a mediator of MMP-13 gene expression during chondrocyte terminal differentiation" *Journal of Biological Chemistry* 2005 280 (46): 38544-38555.
- 14. Ramnarain DB, Park S, Lee DY, Hatanpaa KJ, Scoggin SO, Otu HH, Libermann TA, Raisanen JM, Ashfaq R, Wong ET, Wu J, Elliott R, Habib AA. "Differential gene expression analysis reveals generation of an autocrine loop by a mutant EGFR in glioma cells" *Cancer Research* 2006 66(2): 867-874.
- 15. El Essawy B, Otu HH, Choy B, Xiao XZ, Libermann TA, Strom T. "Proteomic analysis of the allograft response" *Transplantation* 2006 82(2): 267-274.
- 16. Kocabas AM, Crosby J, Ross PJ, Otu HH, Beyhan Z, Can H, Leong TW, Rosa GJM, Halgren RG, Lim B, Fernandez E and Cibelli JB. "The transcriptome of human oocytes" *Proceedings of the National Academy of Sciences*, 2006 103: 14027-14032.
- 17. Steidl U, Rosenbauer F, Verhaak RGW, Gu X, Ebralidze A., Otu HH, Klippel S, Steidl C, Bruns I, Costa DB, Wagner K, Aivado M, Kobbe G, Valk PJ, Passegué E, Libermann TA, Delwel R, Tenen DG. "Essential role of Jun family transcription factors in PU.1 knockdown-induced leukemic stem cells" *Nature Genetics*, 2006 38(11):1269-77.
- 18. Abid R, Shih SC, Otu HH, Curiel DC, Spokes KC, Aird WC. "A novel class of vascular endothelial growth factor-responsive genes that require forkhead activity for expression" *Journal of Biological Chemistry* 2006 281(46):35544-53.
- 19. Zerbini LF, Czibere A, Wang Y, Correa RG, Otu HH, Joseph M, Takayasu Y, Silver M, Gu X, Ruchusatsawat K, Li L, Sarkar D, Zhou JR, Fisher PB, Libermann TA. "A novel pathway involving melanoma differentiation associated gene-7/interleukin-24 mediates nonsteroidal anti-inflammatory drug-induced apoptosis and growth arrest of cancer cells" *Cancer Res.* 2006 66(24):11922-31.
- 20. Aivado M, Spentzos D, Germing U, Alterowitz G, Meng XY, Grall F, Giagounidis AAN, Klement G, Steidl U, Otu HH, Czibere A, Prall WC, Iking-Konert C, Shayne M, Ramoni MF, Gattermann N, Haas R, Mitsiades CS, Fung ET, Libermann TA. "Serum proteome profiling detects myelodysplastic syndromes and identifies CXC chemokine ligands 4 and 7 as markers for advanced disease" *Proceedings of the National Academy of Sciences*, 2007 104(4):1307-12.

- 21. Otu HH, Can H, Spentzos D, Nelson RG, Hanson RL, Looker HC, Knowler WC, Monroy M, Libermann TA, Karumanchi SA, Thadhani R. "Prediction of diabetic nephropathy using urine proteomic profiling 10 years prior to development of nephropathy" *Diabetes Care*, 2007 30:638-643.
- 22. Otu HH, Noxerova K, Can H, Ho K, Nesbitt N, Libermann TA, Karp SJ. "Restoration of liver mass after injury requires proliferative and not embryonic transcriptional patterns" *Journal of Biological Chemistry*, 2007 282(15):11197-204.
- 23. Gu X, Zerbini LF, Otu HH, Joseph MG, Correa R, Libermann TA. "Reduced PDEF expression increases invasion and expression of mesenchymal genes in prostate cancer cells" *Cancer Research*, 2007 67(9):4219-26.
- 24. Kennedy AR, Pissios P, Otu HH, Xue B, Asakura K, Furukawa N, Marino FE, Liu F, Kahn B, Libermann T, Maratos-Flier E. "A high fat, ketogenic diet induces a unique metabolic state in mice" *American Journal of Physiology-Endocrinology and Metabolism*, 2007 292(6):E1724-39.
- 25. Ramlawi B, Otu HH, Rudolph JL, Mieno S, Kohane IS, Can H, Libermann TA, Marcantonio ER, Bianchi C, Sellke FW. "Genomic expression pathways associated with brain injury after cardiopulmonary bypass" *Journal of Thoracic and Cardiovascular Surgery*, 2007 134(4):996-1005.
- 26. Zhang X, Zhang L, Yang H, Huang X, Otu HH, Libermann T, DeWolf WC, Khosravi-Far R, Olumi AF. "c-Fos as a proapoptotic agent in TRAIL-induced apoptosis in prostate cancer cells" *Cancer Research*, 2007 67(19):9425-9434.
- 27. Ramlawi B, Otu HH, Mieno S, Boodhwani M, Sodha NR, Clements RT, Bianchi C, Sellke FW. "Oxidative stress and atrial fibrillation after cardiac surgery: a case-control study" *Ann. Thoracic Surgery*, 2007 84(4):1166 1173.
- 28. Zinkin NT, Grall F, Bhaskar KK, Otu HH, Spentzos D, Kalmowitz BD, Wells M, Guerrero M, Asara J, Libermann TA, Afdhal NH. "Serum proteomics and biomarkers in hepatocellular carcinoma and chronic liver disease" *Clinical Cancer Research*, 2008 14(2):470-477.
- 29. Jones J, Otu HH, Grall F, Spentzos D, Can H, Aivado M, Figlin RA, Belldegrun AS, Pantuck AJ, Libermann TA. "Proteomic identification of interleukin-2 therapy response in metastatic renal cell cancer" *Journal of Urology*, 2008 179(2):730-736.
- 30. Marselli L, Thorne J, Ahn JB, Omer A, Sgroi DC, Libermann T, Otu HH, Sharma A, Bonner-Weir S, Weir GC. "Gene expression of purified beta cell tissue obtained from human pancreas with laser capture microdissection" *The Journal of Clinical Endocrinology & Metabolism*, 2008 93(3):1046-1053.
- 31. Nikolova-Krstevski V, Bhasin M, Otu HH, Libermann T, Oettgen P. "Gene expression analysis of embryonic stem cells expressing VE-cadherin (CD144) during endothelial differentiation" *BMC Genomics*, 2008 9:240.
- 32. Ijiri K, Zerbini LFC, Peng H, Otu HH, Tsuchimochi K, Otero M, Walsh N, Bierbaum BE, Mattingly D, van Flandern G, Komiya S, Aigner T, Libermann TA, Goldring MB. "Differential expression of GADD45beta in normal and osteoarthritic cartilage: potential role in homeostasis of articular chondrocytes" *Arthritis & Rheumatism*, 2008 58(7):2075-87.
- 33. Otu HH*, Dusek JA*, Wohlhueter AL, Bhasin M, Zerbini LF, Joseph MG, Benson H, Libermann TA. "Genomic counter-stress changes induced by the relaxation response" *PLoS ONE*, 2008 3(7):e2576.
- 34. Russell DJ, Otu HH, Sayood K. "Grammar-based distance in progressive multiple sequence alignment" *BMC Bioinformatics*, 2008 9:306.
- 35. Haram KM, Peltier HJ, Lu B, Bhasin M, Otu HH, Choy B, Regan M, Libermann TA, Latham GJ, Sanda MG, Arredouani MS. "Gene expression profile of mouse prostate tumors

_

[·] These authors contributed equally to this work

- reveals dysregulations in major biological processes and identifies potential murine targets for preclinical development of human prostate cancer therapy" *The Prostate*, 2008 68(14):1517-30.
- 36. Prall WC, Czibere A, Grall F, Spentzos D, Steidl U, Giagounidis AA, Kuendgen A, Otu H, Rong A, Libermann TA, Germing U, Gattermann N, Haas R, Aivado M. "Differential gene expression of bone marrow-derived CD34+ cells is associated with survival of patients suffering from myelodysplastic syndrome" *Int. J. Hematol.* 2009 89(2):173-87.
- 37. Ramlawi B, Otu HH, Russo MJ, Novick RJ, Bianchi C, Sellke FW. "Aprotinin attenuates genomic expression variability following cardiac surgery" *J. Card. Surg.*, 2009 24, 772-780.
- 38. Al-Swailem AM, Shehata MM, Abu-Duhier FM, Al-Yamani EJ, Al-Busadah KA, Al-Arawi MS, Al-Khider AY, Al-Muhaimeed AN, Al-Qahtani FH, Al-Manee MM, Al-Shomrani BM, Al-Qhtani SM, Al-Harthi AS, Akdemir KC, Inan MS, Otu HH. "Sequencing, analysis, and annotation of expressed sequence tags for camelus dromedaries" *PLoS ONE*, 2010 5(5):e10720.
- 39. Bhasin M, Yuan L, Keskin DB, Otu HH, Libermann TA, Oettgen P. "Bioinformatic identification and characterization of human endothelial cell-restricted genes" *BMC Genomics*, 2010 11:342.
- 40. Ho KJ, Do N, Otu HH, Dib MJ, Ren X, Enjyoji K, Robson SC, Terwilliger EF, Karp SJ. "Tob1 is a constitutively expressed repressor of liver regeneration" *Journal of Experimental Medicine*, 2010 207(6):1197-208.
- 41. Kang J, Yoo J, Lee S, Tang W, Aguilar B, Swapnika R, Inho C, Otu HH, Shin JW, Dotto GP, Koh CJ, Detmar M, Hong, YK. "An exquisite cross-control mechanism among endothelial cell fate regulators directs the plasticity and heterogeneity of lymphatic endothelial cells" *Blood*, 2010 116(1):140-150.
- 42. Albayrak A, Otu HH, Sezerman OU. "Clustering of protein subfamilies into functional subtypes using relative complexity measure" *BMC Bioinformatics*, 2010 11(1):428.
- 43. Isci S, Jones J, Ozturk C, Otu HH. "Pathway analysis of high throughput biological data within a Bayesian Network framework" *Bioinformatics*, 2011 27(12):1667-1674.
- 44. Wang K, Otu HH, Chen Y, Lee Y, Latham K, Cibelli JB. "Reprogrammed transcriptome in rhesus-bovine interspecies somatic cell nuclear transfer embryos" *PLoS One*, 2011 6(7):e22197.
- 45. Bakis Y, Otu HH, Sezerman OU. "Inferring Phylogenies from Physico-Chemical Properties of DNA" *American Journal of Bioinformatics Research*, 2012 2(1): 1-6
- 46. Otu HH*, Iager AE*, Kocabas AM*, Ruppel P, Langerveld A, Schnarr P, Suarez M, Jarrett JC, Conaghan J, Rosa GJM, Fernández E, Rawlins RG, Cibelli JB, Crosby JA. "Identification of a novel gene signature in human cumulus cells predictive of an oocyte's pregnancy potential" *Fertility and Sterility*, 2013 99 (3): 745-752.
- 47. Meydan C, Otu HH, Sezerman OU. "Prediction of peptides binding to MHC class I and II alleles by temporal motif mining" *BMC Bioinformatics*, 2013 14(Suppl 2):S13.
- 48. Bakis Y, Otu HH, Tasci N, Meydan C, Bilgin N, Yuzbasioglu S, Sezerman OU. "Testing robustness of relative complexity measure method constructing robust phylogenetic trees for Galanthus L. Using the relative complexity measure" *BMC Bioinformatics*, 2013 14:20.
- 49. Coskun S, Otu HH, Awartani KA, Al-Alwan LA, Al-Hassan S, Al-Mayman H, Kaya N, Inan MS. "Gene expression profiling of granulosa cells from PCOS patients following varying doses of human chorionic gonadotropin" *J Assist Reprod Genet.*, 2013 30(3):341-352
- 50. Isci S, Dogan H, Ozturk C, Otu HH. "Bayesian Network Prior: Network Analysis of Biological Data Using External Knowledge" *Bioinformatics*, 2014 30(6):860-867.

-

[•] These authors contributed equally to this work

- 51. Dogan H, Can H, Otu HH. "Whole Genome Sequencing of a Turkish Individual" *PLoS One*, 2014 9(1): e85233.
- 52. Korucuoglu M, Isci S, Ozgur A, Otu HH. "Bayesian Pathway Analysis of Cancer Microarray Data" *PLoS One*, 2014 9(7): e102803.
- 53. Gonzalez-Muñoz E, Arboleda-Estudillo Y, Otu HH, Cibelli JB. "Histone chaperone ASF1A is required for maintenance of pluripotency and cellular reprogramming" *Science*, 2014 345(6198):822-825.
- 54. Yalcin D, Hakguder ZM, Otu HH. "Bioinformatics approaches to single-cell analysis in developmental biology" *Mol Hum Reprod.*, 2016 22(3):182-192.
- 55. Wang F, Kaplan JL, Gold BD, Bhasin MK, Ward NL, Kellermayer R, Kirschner BS, Heyman MB, Dowd SE, Cox SB, Dogan H, Steven B, Ferry GD, Cohen SA, Baldassano RN, Moran CJ, Garnett EA, Drake L, Otu HH, Mirny LA, Libermann TA, Winter HS, Korolev KS. "Detecting microbial dysbiosis associated with Pediatric Crohn's disease despite the high variability of the gut microbiota" *Cell Rep.*, 2016 14(4):945-55
- 56. Ozturk F, Sheldon E, Sharma J, Canturk KM, Otu HH, Nawshad A. "Nicotine Exposure During Pregnancy Results in Persistent Midline Epithelial Seam With Improper Palatal Fusion" *Nic. & Tob. Res.*, 2016 18(5):604-612.
- 57. Bhasin MK, Ndebele K, Bucur O, Yee EU, Otu HH, Plati J, Bullock A, Gu X, Castan E, Zhang P, Najarian R, Muraru MS, Miksad R, Khosravi-Far R, Libermann TA. "Meta-analysis of transcriptome data identifies a novel 5-gene pancreatic adenocarcinoma classifier" *Oncotarget*, 2016 7(17):23263-23281.
- 58. Tomov ML, Olmsted ZT, Dogan H, Gongorurler E, Tsompana M, Otu HH, Buck M, Chang EA, Cibelli J, Paluh JL. "Distinct and Shared Determinants of Cardiomyocyte Contractility in Multi-Lineage Competent Ethnically Diverse Human iPSCs" *Sci Rep.*, 2016 6:37637.
- 59. Dillon ST, Vasunilashorn SM, Ngo L, Otu HH, Inouye SK, Jones RN, Alsop DC, Kuchel GA, Metzger ED, Arnold SE, Marcantonio ER, A. Libermann TA. "Higher C-reactive Protein Levels Predict Postoperative Delirium in Older Patients Undergoing Major Elective Surgery: A Longitudinal Nested Case-Control Study" *Biological Psychiatry*, 2017 81(2): 145-153.
- 60. Vasunilashorn SM, Ngo LH, Chan NY, Zhou W, Dillon ST, Otu HH, Inouye SK, Wyrobnik I, Kuchel GA, McElhaney JE, Xie Z, Alsop DC, Jones RN, Libermann TA, Marcantonio ER. "Development of a Dynamic Multi-Protein Signature of Postoperative Delirium" *The Journal of Gerontology: Series A*, 2018 (in press).
- 61. Mueller SK, Nocera AL, Dillon ST, Gu X, Otu HH, Libermann TA, Bleier BS. "Non-invasive Exosomal Proteomic Biosignatures Including Cystatin-SN (CST 1), Peroxiredoxin-5 (PRDX5) and Glycoprotein-VI (GP6) Accurately Predict Chronic Rhinosinusitis with Nasal Polyps" *Allergy and Rhinology*, 2018 (in press).
- 62. Liu J, Chanumolu SK, Krei Z, Albahrani M, Akhtam A, Jia Z, Wang X, Wang D, Otu HH, Reinhardt RA, Nawshad A. "Identification of genes differentially expressed in simvastatin-induced alveolar bone formation" *Journal of Bone and Mineral Research Plus*, 2018 (in press).
- 63. Xie Z, Kuhns DB, Gu X, Otu HH, Libermann TA, Gallin JI, Parikh SM, Druey KM. "Neutrophil activation in Systemic Capillary Leak Syndrome (Clarkson Disease)" *American Journal of Hematology*, 2018 (in press).
- 64. Liu J, Chanumolu SK, White KM, Albahrani M, Otu HH, Nawshad A. "Comparative analysis of gene expression in TGF-\(\beta\)3 mutant mice with cleft palate" *BMC Genomics*, 2018 (submitted).
- 65. Fong TG, Chan N, Dillon ST, Zhou W, Tripp B, Ngo LH, Otu HH, Inouye SK, Vasunilashorn SM, Cooper Z, Xie Z, Marcantonio E, Libermann T. "Identification of

- plasma proteome signatures associated with surgery and post-operative outcomes using SOMAscan" *Annals of Surgery*, 2018 (submitted).
- 66. Gonzalez-Munoz E, Arboleda-Estudillo Y, Chanumolu SK, Otu HH, Cibelli JB. "Zebrafish macroH2A variants analysis reveals distinct embryo localization and function" *The EMBO Journal*, 2018 (submitted).

In Preparation

- 1. Chanumolu SK, Albahrani M, Otu HH. "FQStat: An automatically optimized parallel programming approach for statistical evaluation of sequencing files" *BMC Bioinformatics*, 2018.
- 2. Can H, Chanumolu SK, Otu HH, Cibelli JB. "Comparative analysis of the zebrafish and human oocyte transcriptome" *Stem Cells*, 2018.
- 3. Chanumolu SK, Albahrani M, Otu HH. "KEGG2DAG: Tools for obtaining directed acyclic graphs from network representations of pathways" *Bioinformatics*, 2019.
- 4. Shi Y, Chanumolu SK, Otu HH. "Bayesian Pathway Analysis of transcriptional regulation" *Bioinformatics*, 2019.
- 5. Yalcin D, Otu HH. "Understanding the CpG island distribution in model organisms using dbCGI" *Genome Research*, 2019.
- 6. Yalcin D, Otu HH. "Modeling the methylation propensity of CpG islands in humans" *Nucleic Acids Research*, 2019.
- 7. Tripp B, Otu HH. "Probabilistic graph representations of multi-omic data interactions" *Bioinformatics*, 2019.
- 8. Tripp B, Dillon ST, Zhou W, Ngo LH, Inouye SK, Vasunilashorn SM, Marcantonio E, Libermann TA, Otu HH. "Network analysis of multi-omic approaches in delirium" *TBD*, 2019.

Proceedings of Meetings

- 1. Otu HH, Sayood K. "A Compression Algorithm that Preserves NDVI and NDWI Values" Proceedings of Thirty-First Asilomar Conference on Circuits, Systems and Computers, pp. 205-208, 1997 November; Monterey, California, 1997.
- 2. Otu HH, Sayood K. "A Joint Source Channel Coder with Block Constraints" Proceedings of 1998 IEEE International Conference on Acoustics, Speech and Signal Processing (ICASSP), pp. 3461-3464,1998 May; Seattle, Washington.
- 3. Arnavut Z, Otu HH. "Compressing Color-mapped Images with Burrows-Wheeler Transformation" Proceedings of Signal Processing Conference IASTED, pp.185-189, 2001 July; Rhodes, Greece.
- 4. Jones J, Inan M, Otu HH, Cho JY, Gu X, Beecken W, Bailey C, Joseph M, Fellbaum C, Jonas D, Libermann TA. "Transcriptional Profiling for Detection of a Gene Signature in Renal Cell Cancer" 55th Congress of the German Urological Society, Der Urologe (A) Suppl 1: p. S19, 2003; Hamburg.
- 5. Jones J, Inan MS, Otu HH, Cho JY, Gu XS, Beecken WD, Joseph M, Fellbaum C, Libermann TA. "Gene Expression Analysis for Identification of a Gene Signature in Renal Cell Cancer", 56th Congress of the German Urological Society, 36th Annual Meeting of the American Society of Nephrology, Journal of the American Society of Nephrology, 14: Suppl S p.367A, November 2003; San Diego, California.
- 6. El Essawy B, Otu HH, Aivado M, Kim YS, Libermann T, Strom T. "Proteomic analysis of the allograft response" American Transplant Congress, American Journal of Transplantation, 4: Spl. 8 p.512, May 2004; Boston, Massachusetts.

- 7. Aivado M, Spentzos D, Germing U, Alterovitz G, Meng XY, Grall1 F, Giagounidis AAN, Klement G, Steidl U, Otu HH, Iking-Konert C, Czibere A, Prall WC, Shayne M, Ramoni MF, Gattermann N, Mitsiades CS, Haas R, Fung ET, Libermann TA. "Serum protein profiling with mass spectrometry for the diagnosis of Myelodysplastic Syndromes (MDS)" 46th Annual Meeting of the American Society of Hematology, Blood (104)11: Part1, p.649A, December 2004; San Diego, California.
- 8. Jones J, Inan M, Otu HH, Spentzos D, Gu X, Joseph M, Beecken W, Jonas D, Libermann TA. "Gene expression profiles in Renal Cell Cancer: Characterization of gene signatures in various histological subtypes and application of a metastatic signature" 56th Congress of the German Urological Society, Der Urologe (A) Suppl 1: p. S23, 2004; Hamburg.
- 9. Ijiri K, Peng H, Zerbini LFC, Walsh N, Wedig M, Gray M, Komiya S, Aigner T, Otu HH, Libermann TA, Goldring MB. "Growth arrest and dna damage-inducible gene 45β expression in normal and osteoarthiritic cartilage" 9th World Congress of the Osteoarthritis Research Society International, Osteoarthritis and Cartilage 12: Suppl. B, p. S8-S9, December 2004; Chicago, Illinois.
- 10. Otu HH, Kolia S, Jones J, Osman O, Libermann TA. "Significance analysis of clustering high-throughput biological data" Proceedings of IEEE Electro/Information Technology Conference, pp. 247-252, May 2005; Lincoln, Nebraska.
- 11. Jones J, Spentzos D, Otu H, Aivado M, Kolia S, Paullus J, Belldegrun AS, Figlin RA, Libermann TA, Pantuck AJ. "Proteomic identification of interleukin-2 therapy responders in metastatic renal cell cancer" Annual Meeting of the American Urological Association, Journal of Urology 173 (4): Suppl. S, pp. 171-172, May 2005; San Antonio, Texas.
- 12. Zinkin NT, Otu HH, Spentzos D, Aivado M, Wells M, Libermann TA, Afdhal NH. "Proteomic profiling of hepatocellular carcinoma with seldi-tof mass spectrometry in patients with chronic liver disease" 106th Annual Meeting of the American Gastroenterological Association, Gastroenterology 128 (4): Suppl. 2, p. A30 May 2005; Chicago, Illinois.
- 13. Zinkin NT, Otu HH, Spentzos D, Aivado M, Wells M, Kalmowitz BD, Bhaskar KK, Libermann TA, Afdhal NH. "A combined proteomic and serologic approach to diagnosis of hepatocellular carcinoma" 56th Annual Meeting of the American Association for the Study of Liver Diseases, Hepatology 42 (4): Suppl. 1, p. 239A, November 2005; San Fransisco, California.
- 14. Steidl U, Rosenbauer F, Verhaak RGW, Gu X, Otu HH, Kolia S, Owens BM, Klippel S, Wagner K, Aivado M, Passegué E, Libermann TA, Delwel R, Tenen DG. "Essential Role of Jun Family Transcription Factors in PU.1-induced Leukemic Stem Cell Transformation" 47th Annual Meeting of the American Society of Hematology, Blood 106 (11): pp. 139A-140A, December 2005; Atlanta, Georgia.
- 15. Abid MR, Shih SC, Otu HH, Spokes KC, Okada Y, Curiel DT, Minami T, Aird WC. "A novel class of VEGF-responsive genes that require forkhead aactivity for expression" 7th Annual Conference on Arteriosclerosis, Thrombosis and Vascular Biology, Arteriosclerosis Thrombosis and Vascular Biology 26 (5): p. E45, April 2006; Denver, Colorado.
- 16. Zhang XP, Yang HM, Zhang LA, Huang X, Otu H, Libermann T, Khosravi-Far R, DeWolf WC, Olumi AF. "c-FOS promotes trail-induced apoptosis by repressing c-FLIP(L)" Coatings Science International Conference, Journal of Urology 177(4): Suppl. S, p. 222, June 2006; Noordwijk, Netherlands.
- 17. Ramlawi B, Otu HH, Kohane IS, Bianchi C, Sellke FW. "Genomic expression pathways associated to brain injury after cardiopulmonary bypass" 79th Annual Scientific Session of the American-Heart-Association, Circulation 114 (18): Suppl. S, p. 402 November 2006; Chicago, Illinois.

- 18. Kang JJ, van den Akker NMS, Aguilar B, Tang WL, Kafka D, Lee SJ, Ramu S, Ganesan SK, Otu HH, van Vugt JMG, Shin JW, Dotto GP, Detmar M, Gittenberger-de Groot AC, Hong YK. "Dysregulated Notch signaling induces pathological arterialization of developing lymphatics in Down syndrome fetus" Experimental Biology 2007 Annual Meeting, Faseb Journal 21(5): p.A15, April 2007; Washington D.C.
- 19. Otu HH, Can H, Kaya N, Al-alwan L, Ozand P, Inan MS. "Computational analysis of transcriptional profiling in Dysmorphic Syndrome" Proceedings of IEEE 15th Signal Processing and Communication Applications Conference, pp. 75-78, June 2007; Eskisehir, Türkiye.
- 20. Marselli L, Sgroi DC, Thorne J, Dahiya S, Torri S, Omer A, Del Prato S, Libermann T, Otu HH, Sharma A, Bonner-Weir S, Marchetti P, Weir GC. "Evidence of inflammatory markers in beta cells of type 2 diabetic subjects" 43rd Annual Meeting of the European Association for the Study of Diabetes, Diabetologia 50: Suppl 1, pp. S178-S179, September 2007; Amsterdam, Netherlands.
- 21. Ramlawi B, Otu HH, Mieno S, Boodhwani M, Sodha NR, Clements RT, Bianchi C, Sellke FW. "Oxidative stress and postoperative atrial fibrillation after cardiac surgery" Canadian Cardiovascular Congress, Canadian Journal of Cardiology 23: p. 292C, October 2007; Ouebec, Canada.
- 22. Jones J, Otu HH, Spentzos D, Belldegrun AS, Libermann TA, Pantuck AJ. "Proteomic identification and validation of interleukin-2 therapy response in metastatic renal cell cancer" 103rd Annual Meeting of the American-Urological-Association, Journal of Urology 179(4): Suppl. S, pp. 35-36, May 2008; Orlando, Florida.
- 23. Ramlawi B, Otu HH, Emani S, Bianchi C, Sellke FW. "Tissue permeability associated with chemokine-class inflammatory response following cardiac surgery" 95th Annual Clinical Congress of the American-College-of-Surgeons, Journal of the American College of Surgeons 209(3): Suppl. S., pp. S29-S30, Ekim 2009; Chicago, Illinois.
- 24. Bakis Y, Sezerman OU, Otu HH. "Resampling techniques for non-alignment based sequence distance methods" Proceedings of the 16th National Biotechnology Conference S-D5: 235-237, 2009; Antalya, Turkey.
- 25. Meydan C, Sezerman U, Otu HH. "Prediction Of Peptides Binding To MHC Class I Alleles By Partial Periodic Pattern Mining" Proceedings of International Joint Conference on Bioinformatics, Systems Biology and Intelligent Computing, pp. 315-318, August 2009, Shanghai, PRC.
- 26. Isci S, Agyuz U, Ozturk C, Otu HH. "Detecting Gene Interactions within a Bayesian Network Framework Using External Knowledge" Proceedings of the 7th International Symposium on Health Informatics and Bioinformatics, (HIBIT 2012), pp. 82-87, April 2012, (appeared in IEEE Xplore).
- 27. Agyuz U, Isci S, Ozturk C, Ademoglu A, Otu HH. "A Dynamic Bayesian Framework to Learn Temporal Gene Interactions Using External Knowledge" Proceedings of the 8th International Symposium on Health Informatics and Bioinformatics, (HIBIT 2013), pp. 1-5, September 2013, DOI:10.1109/HIBIT.2013.6661680 (appeared in IEEE Xplore).
- 28. Yalcin D, Otu H.H. "CpG Island (CGI) Annotation Database and Analysis of CGIs in Human Genome" Proceedings of the Festival of Genomics; 2016 June 27-29; Boston, MA.
- 29. Yalcin D and Otu HH. "Comparative Analysis of Human and Mouse CpG Islands Using dbCGI", Proceedings of the Electro-Information Technology Conference 2017, Lincoln, NE.

Book Chapters

- 1. Otu HH and Libermann TA. "From Microarrays to Gene Networks". In: Appasani K., editor. Bioarrays: From Basics to Diagnostics, pp. 45-61. Humana Press Totowa, NJ. 2007 ISBN: 1588294765.
- 2. Thomson A, Hui NG, Robson P, Otu HH, Lim B. "Embryonic Stem cells as a model for Systems Biology". In: Rigoutsos I. and Stephanoplous D., Eds. Systems Biology 2, pp. 297-319. Oxford University Press. 2007 ISBN: 0195300807.
- 3. Otu HH. "Bioinformatics". In: Dundar M. and Bagis H., Eds. Modern Biotechnology and Applications, pp. 425-462. Erciyes University Press 2010 ISBN: 9789756478639.
- 4. Dogan H and Otu HH. "Objective Functions". In: Russel D., editor. Multiple Sequence Alignment Methods, pp. 45-58. Humana Press. 2013 ISBN: 1627036458.
- 5. Otu HH. "Bioinformatics". In: Dundar M. and Bagis H., Eds. Modern Biotechnology and Applications, 2nd Edition, pp. 425-462. Erciyes University Press 2017 ISBN: 978-60567442-0-4.

Thesis

- 1. Otu HH. Characterization and use of structure and complexity of DNA sequences [doctoral dissertation]. Department of Electrical Engineering, University of Nebraska-Lincoln, Lincoln, NE USA; 2002.
- 2. Otu HH. Constrained Joint Source/Channel Coder [M.S. thesis]. Department of Electrical and Electronics Engineering, Bogazici University, Istanbul, TURKEY; 1997.

Nonprint Materials

- 1. Otu HH. Fr-As, 2002. A computer program to calculate the contigs resulting from a shotgun sequencing data. Distributed as academic freeware.
- 2. Otu HH. OS Distance, 2002. A computer program to calculate the OS distance measure for a set of sequences. Distributed as academic freeware.
- 3. Otu HH and Kolia S. Annotation Database, 2003. A web-based tool to cross annotate more than 80 information facets regarding a gene or a protein. Available through free registration at www.bidmcgenomics.org.
- 4. Otu HH and Usta F. Statistical Analysis of Gene Expression (STAGE), 2003. A stand-alone computer program to analyze gene microarray data. Distributed as academic freeware.
- 5. Otu HH and Kolia S. and Osman O. Assigning Significance to Subclusters of Experimental SampleS (ASSESS), 2003. A stand-alone computer program to assign statistical significance to clustering of gene microarray data. Distributed as academic freeware.
- 6. Isci S and Otu HH. Bayesian pathway Analysis (BPA), 2011. A stand-alone computer program to find active pathways given High Throughput Biological Data. Distributed as academic freeware.
- 7. Isci S and Otu HH. Bayesian Network Prior (BNP), 2013. A stand-alone computer program to incorporate external knowledge to build interaction networks for given High Throughput Biological Data. Distributed as academic freeware.
- 8. Yalcin D and Out HH. Database of CpG Islands (dbCGI), 2017. A web portal acting as a database and tool for analysis of CGIs in different organisms. Freely available on the WWW.

Patents

1. Divide and Conquer System and Method of DNA Sequence Assembly. File Number: 20030224384.

- 2. System and Method for Sequence Distance Measure for Phylogenetic Tree Construction. File Number: 20070225918.
- 3. Human Transcriptome Corresponding To Human Oocytes And Use Of Said Genes Or The Corresponding Polypeptides To Trans-Differentiate Somatic Cells. File Number: 20090028835.
- 4. Genes differentially expressed by cumulus cells and assays using same to identify pregnancy competent oocytes. File Number: 20130053261/20140296104.
- 5. "Protein biomarkers for early detection of Pancreatic Cancer", provisional.
- 6. "Markers for the diagnosis and treatment of non-alcoholic steatohepatitis (NASH) and advanced liver fibrosis,". File Number: 62/586,8125

Abstracts (over 90 abstracts presented at International Conferences)