

# Boris Hayete

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3602873/publications.pdf>

Version: 2024-02-01

19  
papers

4,517  
citations

1162367

8  
h-index

1125271

13  
g-index

20  
all docs

20  
docs citations

20  
times ranked

7071  
citing authors

#	ARTICLE	IF	CITATIONS
1	Multiple Myeloma DREAM Challenge reveals epigenetic regulator PHF19 as marker of aggressive disease. <i>Leukemia</i> , 2020, 34, 1866-1874.	3.3	36
2	Crowdsourced High-Risk Classifiers for Multiple Myeloma Patients Commonly Identify PHF19 As a Robust Progression Biomarker. <i>Blood</i> , 2019, 134, 4370-4370.	0.6	0
3	Systems biology and inÂvitro validation identifies family with sequence similarity 129 member AA(FAM129A) as an asthma steroid response modulator. <i>Journal of Allergy and Clinical Immunology</i> , 2018, 142, 1479-1488.e12.	1.5	15
4	Preclinical Validation Studies Support Causal Machine Learning Based Identification of Novel Drug Targets for High-Risk Multiple Myeloma. <i>Blood</i> , 2018, 132, 3210-3210.	0.6	3
5	Causal modeling of CALGB 80405 (Alliance) to identify network drivers of metastatic colorectal cancer (CRC).. <i>Journal of Clinical Oncology</i> , 2018, 36, 3570-3570.	0.8	0
6	Abstract 5299: Machine learning approach to personalized medicine in breast cancer patients: Development of data-driven, personalized, causal modeling through identification and understanding of optimal treatments for predicting better disease outcomes. , 2018, , .		1
7	Large-scale identification of clinical and genetic predictors of motor progression in patients with newly diagnosed Parkinson's disease: a longitudinal cohort study and validation. <i>Lancet Neurology</i> , The, 2017, 16, 908-916.	4.9	132
8	A Bayesian mathematical model of motor and cognitive outcomes in Parkinsonâ€™s disease. <i>PLoS ONE</i> , 2017, 12, e0178982.	1.1	11
9	Statistical modeling of CALGB 80405 (Alliance) to identify influential factors in metastatic colorectal cancer (CRC) dependent on primary (1o) tumor side.. <i>Journal of Clinical Oncology</i> , 2017, 35, 3528-3528.	0.8	1
10	Inferring Gene Networks for Strains of <i>Dehalococcoides</i> Highlights Conserved Relationships between Genes Encoding Core Catabolic and Cell-Wall Structural Proteins. <i>PLoS ONE</i> , 2016, 11, e0166234.	1.1	4
11	Bayesian Network Models of Multiple Myeloma: Drivers of High Risk and Durable Response. <i>Blood</i> , 2016, 128, 4406-4406.	0.6	5
12	Investigation of Mechanisms of Response in Multiple Myeloma Via Bayesian Causal Inference: An Early Analysis of the Compass Study Data. <i>Blood</i> , 2015, 126, 1794-1794.	0.6	0
13	Abstract 5663: Confirmation of peroxiredoxin II as a driver gene for doxorubicin sensitivity identified from drug-induced expression profiling of the NCI-60 cell lines using Reverse Engineering (REFS) network models. , 2012, , .		0
14	Gyrase inhibitors induce an oxidative damage cellular death pathway in <i>Escherichia coli</i> . <i>Molecular Systems Biology</i> , 2007, 3, 91.	3.2	397
15	Large-Scale Mapping and Validation of <i>Escherichia coli</i> Transcriptional Regulation from a Compendium of Expression Profiles. <i>PLoS Biology</i> , 2007, 5, e8.	2.6	1,308
16	Many Microbe Microarrays Database: uniformly normalized Affymetrix compendia with structured experimental metadata. <i>Nucleic Acids Research</i> , 2007, 36, D866-D870.	6.5	227
17	Size matters: network inference tackles the genome scale. <i>Molecular Systems Biology</i> , 2007, 3, 77.	3.2	26
18	A Common Mechanism of Cellular Death Induced by Bactericidal Antibiotics. <i>Cell</i> , 2007, 130, 797-810.	13.5	2,334

#	ARTICLE	IF	CITATIONS
19	GOTREES: PREDICTING GO ASSOCIATIONS FROM PROTEIN DOMAIN COMPOSITION USING DECISION TREES. , 2004, , .		15