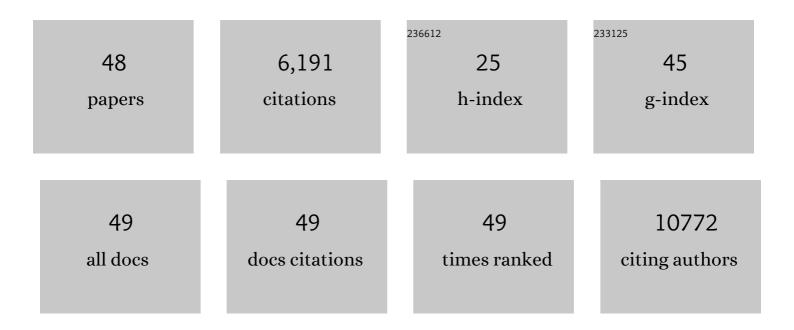
Balasubramanian Narasimhan

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	The changing impact of cytomegalovirus among hematopoietic cell transplant recipients during the past decade: A single institutional cohort study. Transplant Infectious Disease, 2022, 24, .	0.7	1
2	Accuracy of Electronic Medical Record Follow-Up Data for Estimating the Survival Time of Patients With Cancer. JCO Clinical Cancer Informatics, 2022, , .	1.0	3
3	Fast numerical optimization for genome sequencing data in population biobanks. Bioinformatics, 2021, 37, 4148-4155.	1.8	9
4	Penalized regression for leftâ€ŧruncated and right ensored survival data. Statistics in Medicine, 2021, 40, 5487-5500.	0.8	21
5	An open repository of real-time COVID-19 indicators. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	27
6	Letermovir Prophylaxis Decreases Burden of Cytomegalovirus (CMV) in Patients at High Risk for CMV Disease Following Hematopoietic Cell Transplant. Biology of Blood and Marrow Transplantation, 2020, 26, 1963-1970.	2.0	26
7	An Interview with Bradley Efron. International Statistical Review, 2020, 88, S2.	1.1	0
8	Efficacy of motivational-interviewing and guided opioid tapering support for patients undergoing orthopedic surgery (MI-Opioid Taper): A prospective, assessor-blind, randomized controlled pilot trial. EClinicalMedicine, 2020, 28, 100596.	3.2	19
9	Defining the features and duration of antibody responses to SARS-CoV-2 infection associated with disease severity and outcome. Science Immunology, 2020, 5, .	5.6	404
10	The Automatic Construction of Bootstrap Confidence Intervals. Journal of Computational and Graphical Statistics, 2020, 29, 608-619.	0.9	24
11	Derivation and validation of a prognostic score for neonatal mortality in Ethiopia: a case-control study. BMC Pediatrics, 2020, 20, 238.	0.7	12
12	deconvolveR : A G-Modeling Program for Deconvolution and Empirical Bayes Estimation. Journal of Statistical Software, 2020, 94, .	1.8	6
13	CVXR : An <i>R</i> Package for Disciplined Convex Optimization. Journal of Statistical Software, 2020, 94, .	1.8	41
14	Components of genetic associations across 2,138 phenotypes in the UK Biobank highlight adipocyte biology. Nature Communications, 2019, 10, 4064.	5.8	48
15	Imputation of Mixed Data With Multilevel Singular Value Decomposition. Journal of Computational and Graphical Statistics, 2019, 28, 552-566.	0.9	17
16	A scalable discrete-time survival model for neural networks. PeerJ, 2019, 7, e6257.	0.9	102
17	Big data modeling to predict platelet usage and minimize wastage in a tertiary care system. Proceedings of the United States of America, 2017, 114, 11368-11373.	3.3	54
18	Software for Distributed Computation on Medical Databases: A Demonstration Project. Journal of Statistical Software, 2017, 77, .	1.8	7

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19	A new approach to designing phase I-II cancer trials for cytotoxic chemotherapies. Statistics in Medicine, 2014, 33, 2718-2735.	0.8	7
20	Variability in CRP, regulatory T cells and effector T cells over time in gynaecological cancer patients: a study of potential oscillatory behaviour and correlations. Journal of Translational Medicine, 2014, 12, 179.	1.8	14
21	Minimal residual disease quantification using consensus primers and high-throughput IGH sequencing predicts post-transplant relapse in chronic lymphocytic leukemia. Leukemia, 2013, 27, 1659-1665.	3.3	123
22	Prophylactic rituximab after allogeneic transplantation decreases B-cell alloimmunity with low chronic GVHD incidence. Blood, 2012, 119, 6145-6154.	0.6	107
23	Effects of Exercise Training in Patients With Abdominal Aortic Aneurysm. Journal of Cardiopulmonary Rehabilitation and Prevention, 2010, 30, 374-383.	1.2	49
24	Sex differences in the prevalence of peripheral artery disease in patients undergoing coronary catheterization. Vascular Medicine, 2010, 15, 443-450.	0.8	32
25	An <i>R</i> Package for Assessing Drug Synergism/Antagonism. Journal of Statistical Software, 2010, 34,	1.8	14
26	Lymphatic invasion in cutaneous melanoma is associated with sentinel lymph node metastasis. Journal of Cutaneous Pathology, 2009, 36, 772-780.	0.7	79
27	The Relationship Between Human Papillomavirus Status and Other Molecular Prognostic Markers in Head and Neck Squamous Cell Carcinomas. International Journal of Radiation Oncology Biology Physics, 2009, 74, 553-561.	0.4	195
28	Characterizing the admixed African ancestry of African Americans. Genome Biology, 2009, 10, R141.	13.9	145
29	In Vivo 1H Magnetic Resonance Spectroscopy of Lactate in Patients With Stage IV Head and Neck Squamous Cell Carcinoma. International Journal of Radiation Oncology Biology Physics, 2008, 71, 1151-1157.	0.4	26
30	Response to Letter Regarding Article, "β2-Microglobulin as a Biomarker in Peripheral Arterial Disease: Proteomic Profiling and Clinical Studies― Circulation, 2008, 117, .	1.6	0
31	Associations Among Multiple Markers and Complex Disease: Models, Algorithms, and Applications. Advances in Genetics, 2008, 60, 437-464.	0.8	1
32	Estrogen Receptor–Negative Invasive Breast Cancer: Imaging Features of Tumors with and without Human Epidermal Growth Factor Receptor Type 2 Overexpression. Radiology, 2008, 246, 367-375.	3.6	135
33	β2-Microglobulin as a Biomarker in Peripheral Arterial Disease. Circulation, 2007, 116, 1396-1403.	1.6	172
34	Heritability of left ventricular mass in Japanese families living in Hawaii: the SAPPHIRe Study. Journal of Hypertension, 2007, 25, 985-992.	0.3	15
35	CD10 expression in peripheral T-cell lymphomas complicated by a proliferation of large B-cells. Modern Pathology, 2006, 19, 337-343.	2.9	21
36	Mouse Strain–Specific Differences in Vascular Wall Gene Expression and Their Relationship to Vascular Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2005, 25, 302-308.	1.1	39

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37	Signature patterns of gene expression in mouse atherosclerosis and their correlation to human coronary disease. Physiological Genomics, 2005, 22, 213-226.	1.0	73
38	A method for calling gains and losses in array CGH data. Biostatistics, 2005, 6, 45-58.	0.9	337
39	Lisp-StattoJavatoR. Journal of Statistical Software, 2005, 13, .	1.8	3
40	Tree-structured supervised learning and the genetics of hypertension. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 10529-10534.	3.3	42
41	Sample classification from protein mass spectrometry, by 'peak probability contrasts'. Bioinformatics, 2004, 20, 3034-3044.	1.8	196
42	Class Prediction by Nearest Shrunken Centroids, with Applications to DNA Microarrays. Statistical Science, 2003, 18, 104.	1.6	321
43	Diagnosis of multiple cancer types by shrunken centroids of gene expression. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 6567-6572.	3.3	2,632
44	Development of an internet-based protocol to facilitate randomized clinical trials in pediatric surgery. Journal of Pediatric Surgery, 2002, 37, 990-994.	0.8	21
45	Supervised Learning from Microarray Data. , 2002, , 67-77.		4
46	The Inference of Antigen Selection on Ig Genes. Journal of Immunology, 2000, 165, 5122-5126.	0.4	164
47	Bone Mineral Acquisition in Healthy Asian, Hispanic, Black, and Caucasian Youth: A Longitudinal Study ¹ . Journal of Clinical Endocrinology and Metabolism, 1999, 84, 4702-4712.	1.8	401
48	Dynamic Display of Changing Posterior in Bayesian Survival Analysis: The Software. Journal of Statistical Software, 1999, 4, .	1.8	2