

# Nicos Angelopoulos

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5988148/publications.pdf>

Version: 2024-02-01

27  
papers

1,881  
citations

687363

13  
h-index

642732

23  
g-index

29  
all docs

29  
docs citations

29  
times ranked

4362  
citing authors

#	ARTICLE	IF	CITATIONS
1	Classification and Personalized Prognosis in Myeloproliferative Neoplasms. <i>New England Journal of Medicine</i> , 2018, 379, 1416-1430.	27.0	442
2	The vaginal microbiome during pregnancy and the postpartum period in a European population. <i>Scientific Reports</i> , 2015, 5, 8988.	3.3	415
3	Timing the Landmark Events in the Evolution of Clear Cell Renal Cell Cancer: TRACERx Renal. <i>Cell</i> , 2018, 173, 611-623.e17.	28.9	398
4	Genomic landscape and chronological reconstruction of driver events in multiple myeloma. <i>Nature Communications</i> , 2019, 10, 3835.	12.8	183
5	Timing the initiation of multiple myeloma. <i>Nature Communications</i> , 2020, 11, 1917.	12.8	99
6	Molecular Evolution of <i>IDH</i> Wild-Type Glioblastomas Treated With Standard of Care Affects Survival and Design of Precision Medicine Trials: A Report From the EORTC 1542 Study. <i>Journal of Clinical Oncology</i> , 2020, 38, 81-99.	1.6	84
7	Revealing the Impact of Structural Variants in Multiple Myeloma. <i>Blood Cancer Discovery</i> , 2020, 1, 258-273.	5.0	81
8	Targeting of EGFR by a combination of antibodies mediates unconventional EGFR trafficking and degradation. <i>Scientific Reports</i> , 2020, 10, 663.	3.3	23
9	LMTK3 Represses Tumor Suppressor-like Genes through Chromatin Remodeling in Breast Cancer. <i>Cell Reports</i> , 2015, 12, 837-849.	6.4	21
10	mmsig: a fitting approach to accurately identify somatic mutational signatures in hematological malignancies. <i>Communications Biology</i> , 2021, 4, 424.	4.4	21
11	ATG9A loss confers resistance to trastuzumab via c-Cbl mediated Her2 degradation. <i>Oncotarget</i> , 2016, 7, 27599-27612.	1.8	21
12	Characterization of the Tyrosine Kinase-Regulated Proteome in Breast Cancer by Combined use of RNA interference (RNAi) and Stable Isotope Labeling with Amino Acids in Cell Culture (SILAC) Quantitative Proteomics. <i>Molecular and Cellular Proteomics</i> , 2015, 14, 2479-2492.	3.8	17
13	Bayesian learning of Bayesian networks with informative priors. <i>Annals of Mathematics and Artificial Intelligence</i> , 2008, 54, 53-98.	1.3	15
14	Architecture of a mediator for a bioinformatics database federation. <i>IEEE Transactions on Information Technology in Biomedicine</i> , 2002, 6, 116-122.	3.2	13
15	Bayesian Model Averaging for Ligand Discovery. <i>Journal of Chemical Information and Modeling</i> , 2009, 49, 1547-1557.	5.4	13
16	Proteomic profile of KSR1-regulated signalling in response to genotoxic agents in breast cancer. <i>Breast Cancer Research and Treatment</i> , 2015, 151, 555-568.	2.5	10
17	Distributional logic programming for Bayesian knowledge representation. <i>International Journal of Approximate Reasoning</i> , 2017, 80, 52-66.	3.3	5
18	Probabilistic Finite Domains: A Brief Overview. <i>Lecture Notes in Computer Science</i> , 2002, , 475-475.	1.3	5

#	ARTICLE	IF	CITATIONS
19	Bayesian networks elucidate complex genomic landscapes in cancer. <i>Communications Biology</i> , 2022, 5, 306.	4.4	5
20	Proteome-wide dataset supporting functional study of tyrosine kinases in breast cancer. <i>Data in Brief</i> , 2016, 7, 740-746.	1.0	3
21	Advances in integrative statistics for logic programming. <i>International Journal of Approximate Reasoning</i> , 2016, 78, 103-115.	3.3	2
22	Selective Chemical Intervention in the Proteome of <i>Caenorhabditis elegans</i> . <i>Journal of Proteome Research</i> , 2010, 9, 6060-6070.	3.7	1
23	Accessing biological data as Prolog facts. , 2017, , .		1
24	clp(pdf(y)): Constraints for Probabilistic Reasoning in Logic Programming. <i>Lecture Notes in Computer Science</i> , 2003, , 784-788.	1.3	1
25	Advances in Big Data Bio Analytics. <i>Electronic Proceedings in Theoretical Computer Science</i> , EPTCS, 0, 306, 309-322.	0.8	1
26	Extending the CLP Engine for Reasoning under Uncertainty. <i>Lecture Notes in Computer Science</i> , 2003, , 365-373.	1.3	0
27	Probabilistic Space Partitioning in Constraint Logic Programming. <i>Lecture Notes in Computer Science</i> , 2004, , 48-62.	1.3	0