

Zeynep H GÃ¼mÃ¼s

List of Publications by Year in descending order

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Version: 2024-02-01

65
papers

6,659
citations

201674

27
h-index

182427

51
g-index

86
all docs

86
docs citations

86
times ranked

16256
citing authors

#	ARTICLE	IF	CITATIONS
1	A Brave New World: Virtual Reality and Augmented Reality in Systems Biology. <i>Frontiers in Bioinformatics</i> , 2022, 2, .	2.1	2
2	Germline Pathogenic Variants Impact Clinicopathology of Advanced Lung Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 1450-1459.	2.5	10
3	SNPs at SMG7 associated with time from biochemical recurrence to prostate cancer death. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, , .	2.5	1
4	Brain aging is faithfully modelled in organotypic brain slices and accelerated by prions. <i>Communications Biology</i> , 2022, 5, .	4.4	1
5	Functional Common and Rare <i>ERBB2</i> Germline Variants Cooperate in Familial and Sporadic Cancer Susceptibility. <i>Cancer Prevention Research</i> , 2021, 14, 441-454.	1.5	0
6	Generation of TRAIL-resistant cell line models reveals distinct adaptive mechanisms for acquired resistance and re-sensitization. <i>Oncogene</i> , 2021, 40, 3201-3216.	5.9	5
7	Novel ultra-rare exonic variants identified in a founder population implicate cadherins in schizophrenia. <i>Neuron</i> , 2021, 109, 1465-1478.e4.	8.1	21
8	A proteogenomic portrait of lung squamous cell carcinoma. <i>Cell</i> , 2021, 184, 4348-4371.e40.	28.9	170
9	Integrated Proteogenomic Characterization across Major Histological Types of Pediatric Brain Cancer. <i>Cell</i> , 2020, 183, 1962-1985.e31.	28.9	177
10	Inherited Rare, Deleterious Variants in ATM Increase Lung Adenocarcinoma Risk. <i>Journal of Thoracic Oncology</i> , 2020, 15, 1871-1879.	1.1	24
11	Comprehensive Immunoprofiling of Pediatric Zika Reveals Key Role for Monocytes in the Acute Phase and No Effect of Prior Dengue Virus Infection. <i>Cell Reports</i> , 2020, 31, 107569.	6.4	43
12	ProNetView ^{ccRCC} : A Web-Based Portal to Interactively Explore Clear Cell Renal Cell Carcinoma Proteogenomics Networks. <i>Proteomics</i> , 2020, 20, e2000043.	2.2	6
13	Ancestry-specific predisposing germline variants in cancer. <i>Genome Medicine</i> , 2020, 12, 51.	8.2	35
14	Immunology of COVID-19: Current State of the Science. <i>Immunity</i> , 2020, 52, 910-941.	14.3	1,387
15	Proteogenomic Characterization Reveals Therapeutic Vulnerabilities in Lung Adenocarcinoma. <i>Cell</i> , 2020, 182, 200-225.e35.	28.9	410
16	Comparative effects of oncogenic mutations G12C, G12V, G13D, and Q61H on local conformations and dynamics of K-Ras. <i>Computational and Structural Biotechnology Journal</i> , 2020, 18, 1000-1011.	4.1	33
17	Oncogenic G12D mutation alters local conformations and dynamics of K-Ras. <i>Scientific Reports</i> , 2019, 9, 11730.	3.3	46
18	Integrated Proteogenomic Characterization of Clear Cell Renal Cell Carcinoma. <i>Cell</i> , 2019, 179, 964-983.e31.	28.9	430

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19	Dynamical Rearrangement of Human Epidermal Growth Factor Receptor 2 upon Antibody Binding: Effects on the Dimerization. <i>Biomolecules</i> , 2019, 9, 706.	4.0	6
20	ImmuneRegulation: a web-based tool for identifying human immune regulatory elements. <i>Nucleic Acids Research</i> , 2019, 47, W142-W150.	14.5	4
21	Identification of SERPINE1 as a Regulator of Glioblastoma Cell Dispersal with Transcriptome Profiling. <i>Cancers</i> , 2019, 11, 1651.	3.7	43
22	Rare, Pathogenic Germline Variants in <i>Fanconi Anemia</i> Genes Increase Risk for Squamous Lung Cancer. <i>Clinical Cancer Research</i> , 2019, 25, 1517-1525.	7.0	31
23	Ancestral reconstruction reveals mechanisms of ERK regulatory evolution. <i>ELife</i> , 2019, 8, .	6.0	24
24	Exploring Biological Networks in 3D, Stereoscopic 3D, and Immersive 3D with iCAVE. <i>Current Protocols in Bioinformatics</i> , 2018, 61, 8.27.1-8.27.26.	25.8	6
25	High-depth whole genome sequencing of an Ashkenazi Jewish reference panel: enhancing sensitivity, accuracy, and imputation. <i>Human Genetics</i> , 2018, 137, 343-355.	3.8	24
26	Identifying Causality in Mutant KRas Residue Pairs from Molecular Dynamics Data Analysis. <i>Biophysical Journal</i> , 2018, 114, 397a.	0.5	1
27	Comprehensive functional genomic resource and integrative model for the human brain. <i>Science</i> , 2018, 362, .	12.6	618
28	Landscape of Conditional eQTL in Dorsolateral Prefrontal Cortex and Co-localization with Schizophrenia GWAS. <i>American Journal of Human Genetics</i> , 2018, 102, 1169-1184.	6.2	128
29	Spoton: A Machine-Learning Approach for Hot-Spot Determination. <i>Biophysical Journal</i> , 2017, 112, 45a.	0.5	0
30	iCAVE: an open source tool for visualizing biomolecular networks in 3D, stereoscopic 3D and immersive 3D. <i>GigaScience</i> , 2017, 6, 1-13.	6.4	21
31	SpotOn: High Accuracy Identification of Protein-Protein Interface Hot-Spots. <i>Scientific Reports</i> , 2017, 7, 8007.	3.3	77
32	KDM2B, an H3K36-specific demethylase, regulates apoptotic response of GBM cells to TRAIL. <i>Cell Death and Disease</i> , 2017, 8, e2897-e2897.	6.3	26
33	Germline and Somatic Smoothened Mutations in Nonâ€“Small-Cell Lung Cancer Are Potentially Responsive to Hedgehog Inhibitor Vismodegib. <i>JCO Precision Oncology</i> , 2017, 1, 1-10.	3.0	3
34	Inhibition of colorectal cancer genomic copy number alterations and chromosomal fragile site tumor suppressor FHIT and WWOX deletions by DNA mismatch repair. <i>Oncotarget</i> , 2017, 8, 71574-71586.	1.8	6
35	Response of germline and somatic smoothened (SMO) mutations in non-small cell lung cancer (NSCLC) to hedgehog inhibitor vismodegib.. <i>Journal of Clinical Oncology</i> , 2017, 35, 9062-9062.	1.6	3
36	A Machine Learning Approach for Hot-Spot Detection at Protein-Protein Interfaces. <i>International Journal of Molecular Sciences</i> , 2016, 17, 1215.	4.1	46

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37	DRES-12. PROFILING OF DIFFERENT GBM CELL POPULATIONS WITH VARYING APOPTOTIC THRESHOLDS IDENTIFIES IGFBP-2 AS A NOVEL MEDIATOR OF TRAIL RESISTANCE. <i>Neuro-Oncology</i> , 2016, 18, vi54-vi54.	1.2	0
38	Gene expression elucidates functional impact of polygenic risk for schizophrenia. <i>Nature Neuroscience</i> , 2016, 19, 1442-1453.	14.8	952
39	Intrinsic K-Ras dynamics: A novel molecular dynamics data analysis method shows causality between residue pair motions. <i>Scientific Reports</i> , 2016, 6, 37012.	3.3	26
40	A Novel Genetic Variant in Long Non-coding RNA Gene NEXN-AS1 is Associated with Risk of Lung Cancer. <i>Scientific Reports</i> , 2016, 6, 34234.	3.3	48
41	ANGI-08. IDENTIFICATION OF NOVEL MOLECULAR PLAYERS OF GBM CELL DISPERSAL THROUGH AN IN VITRO PROFILING APPROACH. <i>Neuro-Oncology</i> , 2016, 18, vi16-vi16.	1.2	0
42	Comprehensive models of human primary and metastatic colorectal tumors in immunodeficient and immunocompetent mice by chemokine targeting. <i>Nature Biotechnology</i> , 2015, 33, 656-660.	17.5	30
43	EPIG-01 THE FUNCTION OF CHROMATIN-MODIFYING ENZYMES IN GBM CELL APOPTOSIS. <i>Neuro-Oncology</i> , 2015, 17, v86.1-v86.	1.2	0
44	CBIO-08 IGFBP2 IS A NOVEL MOLECULAR DETERMINANT IN TRAIL-RESISTANT SUBPOPULATIONS OF GBM CELL LINES. <i>Neuro-Oncology</i> , 2015, 17, v56.3-v56.	1.2	0
45	miR-1269 promotes metastasis and forms a positive feedback loop with TGF- β 2. <i>Nature Communications</i> , 2015, 6, 6879.	12.8	110
46	Integrative Annotation of Variants from 1092 Humans: Application to Cancer Genomics. <i>Science</i> , 2013, 342, 1235587.	12.6	341
47	A microRNA miR-34a-Regulated Bimodal Switch Targets Notch in Colon Cancer Stem Cells. <i>Cell Stem Cell</i> , 2013, 12, 602-615.	11.1	325
48	Rapid Temporal Changes in the Expression of a Set of Neuromodulatory Genes During Alcohol Withdrawal in the Dorsal Vagal Complex: Molecular Evidence of Homeostatic Disturbance. <i>Alcoholism: Clinical and Experimental Research</i> , 2012, 36, 1688-1700.	2.4	32
49	miR-23a Promotes the Transition from Indolent to Invasive Colorectal Cancer. <i>Cancer Discovery</i> , 2012, 2, 540-553.	9.4	132
50	Resistance/Response Molecular Signature for Oral Tongue Squamous Cell Carcinoma. <i>Disease Markers</i> , 2012, 32, 51-64.	1.3	28
51	Chemokine CXCL12-induced signaling suppresses colon cancer invasion and metastasis. <i>Journal of Clinical Investigation</i> , 2012, 122, 3184-3196.	8.2	67
52	Resistance/response molecular signature for oral tongue squamous cell carcinoma. <i>Disease Markers</i> , 2012, 32, 51-64.	1.3	11
53	Abstract 4911: Molecular investigation of the FTI/Taxol synergy using an integrated framework of transcriptome analysis. , 2011, , .		0
54	Effects of Cigarette Smoke on the Human Oral Mucosal Transcriptome. <i>Cancer Prevention Research</i> , 2010, 3, 266-278.	1.5	146

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55	Quantification and Analysis of Combination Drug Synergy in High-Throughput Transcriptome Studies. , 2010, , .		0
56	Abstract 115: An integrated framework of transcriptome analysis leads to mechanistic insights into the FTI/Taxol synergy. , 2010, , .		0
57	Towards a quantitative representation of the cell signaling mechanisms of hallucinogens: Measurement and mathematical modeling of 5-HT1A and 5-HT2A receptor-mediated ERK1/2 activation. Neuropharmacology, 2009, 56, 213-225.	4.1	25
58	Effects of Tobacco Smoke on Gene Expression and Cellular Pathways in a Cellular Model of Oral Leukoplakia. Cancer Prevention Research, 2008, 1, 100-111.	1.5	54
59	Bilevel Programming: Applications in Engineering. , 2008, , 243-248.		0
60	Mixed Integer Nonlinear Bilevel Programming: Deterministic Global Optimization. , 2008, , 2225-2230.		0
61	Global optimization of mixed-integer bilevel programming problems. Computational Management Science, 2005, 2, 181-212.	1.3	109
62	Nonlinear bilevel programming: A deterministic global optimization framework. Computer Aided Chemical Engineering, 2001, 9, 393-400.	0.5	0
63	Global Optimization in Design under Uncertainty:Â Feasibility Test and Flexibility Index Problems. Industrial & Engineering Chemistry Research, 2001, 40, 4267-4282.	3.7	136
64	Global Optimization of Nonlinear Bilevel Programming Problems. Journal of Global Optimization, 2001, 20, 1-31.	1.8	125
65	Reactive distillation column design with vapor/liquid/liquid equilibria. Computers and Chemical Engineering, 1997, 21, S983-S988.	3.8	22