Andrew T Magis

List of Publications by Year in descending order

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62 papers

4,430 citations

185998 28 h-index 61 g-index

79 all docs

79 docs citations

79 times ranked 7320 citing authors

#	Article	IF	CITATIONS
1	Integrated analysis of plasma and single immune cells uncovers metabolic changes in individuals with COVID-19. Nature Biotechnology, 2022, 40, 110-120.	9.4	81
2	Multiple early factors anticipate post-acute COVID-19 sequelae. Cell, 2022, 185, 881-895.e20.	13.5	605
3	Personal Dense Dynamic Data Clouds Connect Systems Biomedicine to Scientific Wellness. Methods in Molecular Biology, 2022, 2486, 315-334.	0.4	1
4	Manifestations of Alzheimer's disease genetic risk in the blood are evident in a multiomic analysis in healthy adults aged 18 to 90. Scientific Reports, 2022, 12, 6117.	1.6	12
5	Heterogeneity in statin responses explained by variation in the human gut microbiome. Med, 2022, 3, 388-405.e6.	2.2	21
6	Gut microbiome pattern reflects healthy ageing and predicts survival in humans. Nature Metabolism, 2021, 3, 274-286.	5.1	278
7	Characteristics and Factors Associated With Coronavirus Disease 2019 Infection, Hospitalization, and Mortality Across Race and Ethnicity. Clinical Infectious Diseases, 2021, 73, 2193-2204.	2.9	41
8	DUOX2 variants associate with preclinical disturbances in microbiota-immune homeostasis and increased inflammatory bowel disease risk. Journal of Clinical Investigation, 2021, 131, .	3.9	35
9	The geometry of clinical labs and wellness states from deeply phenotyped humans. Nature Communications, 2021, 12, 3578.	5.8	19
10	Baseline Gut Metagenomic Functional Gene Signature Associated with Variable Weight Loss Responses following a Healthy Lifestyle Intervention in Humans. MSystems, 2021, 6, e0096421.	1.7	19
11	Discovery of Small Molecule Bak Activator for Lung Cancer Therapy. Theranostics, 2021, 11, 8500-8516.	4.6	19
12	Health and disease markers correlate with gut microbiome composition across thousands of people. Nature Communications, 2020, 11 , 5206 .	5.8	378
13	Plasma levels of soluble ACE2are associated with sex, Metabolic Syndrome, and its biomarkers in a large cohort, pointing to a possible mechanism for increased severity in COVID-19. Critical Care, 2020, 24, 452.	2.5	59
14	Multi-Omics Resolves a Sharp Disease-State Shift between Mild and Moderate COVID-19. Cell, 2020, 183, 1479-1495.e20.	13.5	449
15	Multiomic blood correlates of genetic risk identify presymptomatic disease alterations. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 21813-21820.	3.3	22
16	Longitudinal analysis reveals transition barriers between dominant ecological states in the gut microbiome. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 13839-13845.	3.3	47
17	Untargeted longitudinal analysis of a wellness cohort identifies markers of metastatic cancer years prior to diagnosis. Scientific Reports, 2020, 10, 16275.	1.6	12
18	Multi-Omic Biological Age Estimation and Its Correlation With Wellness and Disease Phenotypes: A Longitudinal Study of 3,558 Individuals. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2019, 74, S52-S60.	1.7	56

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19	Mcl-1 Interacts with Akt to Promote Lung Cancer Progression. Cancer Research, 2019, 79, 6126-6138.	0.4	25
20	Blood metabolome predicts gut microbiome \hat{l}_{\pm} -diversity in humans. Nature Biotechnology, 2019, 37, 1217-1228.	9.4	213
21	Genetic Predisposition Impacts Clinical Changes in a Lifestyle Coaching Program. Scientific Reports, 2019, 9, 6805.	1.6	48
22	Small Molecule KRAS Agonist for Mutant KRAS Cancer Therapy. Molecular Cancer, 2019, 18, 85.	7.9	41
23	Reply to "Precision medicine in the clouds". Nature Biotechnology, 2018, 36, 680-682.	9.4	2
24	A Multi-omic Association Study of Trimethylamine N-Oxide. Cell Reports, 2018, 24, 935-946.	2.9	115
25	Modulation of Bax and mTOR for Cancer Therapeutics. Cancer Research, 2017, 77, 3001-3012.	0.4	24
26	Genomic, Transcriptional, and Phenotypic Analysis of the Glucose Derepressed <i>Clostridium beijerinckii</i> Mutant Exhibiting Acid Crash Phenotype. Biotechnology Journal, 2017, 12, 1700182.	1.8	14
27	A wellness study of 108 individuals using personal, dense, dynamic data clouds. Nature Biotechnology, 2017, 35, 747-756.	9.4	340
28	Targeting Mcl-1 enhances DNA replication stress sensitivity to cancer therapy. Journal of Clinical Investigation, 2017, 128, 500-516.	3.9	48
29	Sulfatide Preserves Insulin Crystals Not by Being Integrated in the Lattice but by Stabilizing Their Surface. Journal of Diabetes Research, 2016, 2016, 1-4.	1.0	8
30	Small-Molecule Bcl2 BH4 Antagonist for Lung Cancer Therapy. Cancer Cell, 2015, 27, 852-863.	7.7	108
31	SNAPR: A Bioinformatics Pipeline for Efficient and Accurate RNA-Seq Alignment and Analysis. IEEE Life Sciences Letters, 2015, 1, 22-25.	1.2	15
32	Detection of $1\hat{l}_{\pm}$,25-Dihydroxyvitamin D-Regulated miRNAs in Zebrafish by Whole Transcriptome Sequencing. Zebrafish, 2014, 11, 207-218.	0.5	14
33	Small-molecule Bax agonists for cancer therapy. Nature Communications, 2014, 5, 4935.	5.8	110
34	Design, synthesis, and biological evaluation of novel FAK scaffold inhibitors targeting the FAK–VEGFR3 protein–protein interaction. European Journal of Medicinal Chemistry, 2014, 80, 154-166.	2.6	26
35	Measuring the Effect of Inter-Study Variability on Estimating Prediction Error. PLoS ONE, 2014, 9, e110840.	1.1	19
36	AUREA: an open-source software system for accurate and user-friendly identification of relative expression molecular signatures. BMC Bioinformatics, 2013, 14, 78.	1.2	7

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37	Disruption of focal adhesion kinase and p53 interaction with small molecule compound R2 reactivated p53 and blocked tumor growth. BMC Cancer, 2013, 13, 342.	1.1	49
38	Novel Small-Molecule Inhibitors of Bcl-XL to Treat Lung Cancer. Cancer Research, 2013, 73, 5485-5496.	0.4	62
39	Multi-study Integration of Brain Cancer Transcriptomes Reveals Organ-Level Molecular Signatures. PLoS Computational Biology, 2013, 9, e1003148.	1.5	16
40	A Small-molecule Inhibitor, 5'-O-Tritylthymidine, Targets FAK and Mdm-2 Interaction, and Blocks Breast and Colon Tumorigenesis in vivo. Anti-Cancer Agents in Medicinal Chemistry, 2013, 13, 532-545.	0.9	27
41	Mitoxantrone Targets the ATP-binding Site of FAK, Binds the FAK Kinase Domain and Decreases FAK, Pyk-2, c-Src, and IGF-1R In Vitro Kinase Activities. Anti-Cancer Agents in Medicinal Chemistry, 2013, 13, 546-554.	0.9	20
42	Inhibiting the Interaction of cMET and IGF-1R with FAK Effectively Reduces Growth of Pancreatic Cancer Cells in vitro and in vivo. Anti-Cancer Agents in Medicinal Chemistry, 2013, 13, 595-602.	0.9	25
43	Research Resource: Whole Transcriptome RNA Sequencing Detects Multiple 1α,25-Dihydroxyvitamin D3-Sensitive Metabolic Pathways in Developing Zebrafish. Molecular Endocrinology, 2012, 26, 1630-1642.	3.7	41
44	Kinetic Characterization of Newly Discovered Inhibitors of Various Constructs of Human T-Cell Leukemia Virus-1 (HTLV-1) Protease and Their Effect on HTLV-1-Infected Cells. Antiviral Therapy, 2012, 17, 883-892.	0.6	6
45	The top-scoring â€~N' algorithm: a generalized relative expression classification method from small numbers of biomolecules. BMC Bioinformatics, 2012, 13, 227.	1.2	24
46	A small molecule focal adhesion kinase (FAK) inhibitor, targeting Y397 site: 1-(2-hydroxyethyl) -3, 5, 7-triaza-1-azoniatricyclo [3.3.1.13,7]decane; bromide effectively inhibits FAK autophosphorylation activity and decreases cancer cell viability, clonogenicity and tumor growth in vivo. Carcinogenesis, 2012, 33, 1004-1013.	1.3	73
47	A shift in the salt bridge interaction of residues D620 and E621 mediates the constitutive activation of Jak2-H538Q/K539L. Molecular and Cellular Biochemistry, 2012, 367, 125-140.	1.4	5
48	A46, a benzothiophene-derived compound, suppresses Jak2-mediated pathologic cell growth. Experimental Hematology, 2012, 40, 22-34.	0.2	0
49	Crystallization and preliminary X-ray analysis of the human long myosin light-chain kinase 1-specific domain IgCAM3. Acta Crystallographica Section F: Structural Biology Communications, 2011, 67, 221-223.	0.7	4
50	Noise Contributions in an Inducible Genetic Switch: A Whole-Cell Simulation Study. PLoS Computational Biology, 2011, 7, e1002010.	1.5	85
51	Graphics processing unit implementations of relative expression analysis algorithms enable dramatic computational speedup. Bioinformatics, 2011, 27, 872-873.	1.8	11
52	Structure-Function Correlation of G6, a Novel Small Molecule Inhibitor of Jak2. Journal of Biological Chemistry, 2010, 285, 31399-31407.	1.6	11
53	The Constitutive Activation of Jak2-V617F Is Mediated by a π Stacking Mechanism Involving Phenylalanines 595 and 617. Biochemistry, 2010, 49, 9972-9984.	1.2	36
54	A novel small molecule inhibitor of FAK decreases growth of human pancreatic cancer. Cell Cycle, 2009, 8, 2435-2443.	1.3	150

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55	Identification of Enoxacin as an Inhibitor of Osteoclast Formation and Bone Resorption by Structure-Based Virtual Screening. Journal of Medicinal Chemistry, 2009, 52, 5144-5151.	2.9	46
56	Targeting of the protein interaction site between FAK and IGF-1R. Biochemical and Biophysical Research Communications, 2009, 388, 301-305.	1.0	30
57	Small Molecule Chloropyramine Hydrochloride (C4) Targets the Binding Site of Focal Adhesion Kinase and Vascular Endothelial Growth Factor Receptor 3 and Suppresses Breast Cancer Growth in Vivo. Journal of Medicinal Chemistry, 2009, 52, 4716-4724.	2.9	67
58	Crystallization of the focal adhesion kinase targeting (FAT) domain in a primitive orthorhombic space group. Acta Crystallographica Section F: Structural Biology Communications, 2008, 64, 564-566.	0.7	2
59	A Bony Fish Immunological Receptor of the NITR Multigene Family Mediates Allogeneic Recognition. Immunity, 2008, 29, 228-237.	6.6	42
60	A Small Molecule Inhibitor, 1,2,4,5-Benzenetetraamine Tetrahydrochloride, Targeting the Y397 Site of Focal Adhesion Kinase Decreases Tumor Growth. Journal of Medicinal Chemistry, 2008, 51, 7405-7416.	2.9	204
61	Z3, a novel Jak2 tyrosine kinase small-molecule inhibitor that suppresses Jak2-mediated pathologic cell growth. Molecular Cancer Therapeutics, 2008, 7, 2308-2318.	1.9	22
62	Crystallization and X-ray diffraction analysis of a novel immune-type receptor fromIctalurus punctatusand phasing by selenium anomalous dispersion methods. Acta Crystallographica Section F: Structural Biology Communications, 2007, 63, 1035-1037.	0.7	3