Kévin Contrepois

List of Publications by Year in descending order

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136740 5,241 52 32 citations papers

51 h-index g-index 63 63 63 8629 docs citations times ranked citing authors all docs

182168

#	Article	IF	CITATIONS
1	The NASA Twins Study: A multidimensional analysis of a year-long human spaceflight. Science, 2019, 364,	6.0	576
2	Mass spectrometry-based metabolomics: a guide for annotation, quantification and best reporting practices. Nature Methods, 2021, 18, 747-756.	9.0	403
3	Longitudinal multi-omics of host–microbe dynamics in prediabetes. Nature, 2019, 569, 663-671.	13.7	391
4	The Human Tumor Atlas Network: Charting Tumor Transitions across Space and Time at Single-Cell Resolution. Cell, 2020, 181, 236-249.	13.5	334
5	A longitudinal big data approach for precision health. Nature Medicine, 2019, 25, 792-804.	15.2	329
6	Macrophage de novo NAD+ synthesis specifies immune function in aging and inflammation. Nature Immunology, 2019, 20, 50-63.	7.0	304
7	Molecular Choreography of Acute Exercise. Cell, 2020, 181, 1112-1130.e16.	13.5	261
8	Physiological blood–brain transport is impaired with age by a shift in transcytosis. Nature, 2020, 583, 425-430.	13.7	243
9	Personal aging markers and ageotypes revealed by deep longitudinal profiling. Nature Medicine, 2020, 26, 83-90.	15.2	225
10	Optimized Analytical Procedures for the Untargeted Metabolomic Profiling of Human Urine and Plasma by Combining Hydrophilic Interaction (HILIC) and Reverse-Phase Liquid Chromatography (RPLC)–Mass Spectrometry*. Molecular and Cellular Proteomics, 2015, 14, 1684-1695.	2.5	183
11	Integrative Personal Omics Profiles during Periods of Weight Gain and Loss. Cell Systems, 2018, 6, 157-170.e8.	2.9	183
12	Multiomics modeling of the immunome, transcriptome, microbiome, proteome and metabolome adaptations during human pregnancy. Bioinformatics, 2019, 35, 95-103.	1.8	162
13	Exercise plasma boosts memory and dampens brain inflammation via clusterin. Nature, 2021, 600, 494-499.	13.7	156
14	Metabolic Dynamics and Prediction of Gestational Age and Time to Delivery in Pregnant Women. Cell, 2020, 181, 1680-1692.e15.	13.5	154
15	Histone variant H2A.J accumulates in senescent cells and promotes inflammatory gene expression. Nature Communications, 2017, 8, 14995.	5.8	131
16	An exercise-inducible metabolite that suppresses feeding and obesity. Nature, 2022, 606, 785-790.	13.7	96
17	Altered Cardiac Energetics and Mitochondrial Dysfunction in Hypertrophic Cardiomyopathy. Circulation, 2021, 144, 1714-1731.	1.6	90
18	Integrated trajectories of the maternal metabolome, proteome, and immunome predict labor onset. Science Translational Medicine, 2021, 13, .	5.8	82

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19	Cross-Platform Comparison of Untargeted and Targeted Lipidomics Approaches on Aging Mouse Plasma. Scientific Reports, 2018, 8, 17747.	1.6	81
20	Biallelic Mutations in ATP5F1D, which Encodes a Subunit of ATP Synthase, Cause a Metabolic Disorder. American Journal of Human Genetics, 2018, 102, 494-504.	2.6	59
21	Multi-omic profiling of primary mouse neutrophils predicts a pattern of sex- and age-related functional regulation. Nature Aging, 2021, 1, 715-733.	5.3	55
22	Multiomics Characterization of Preterm Birth in Low- and Middle-Income Countries. JAMA Network Open, 2020, 3, e2029655.	2.8	53
23	The MEK5–ERK5 Kinase Axis Controls Lipid Metabolism in Small-Cell Lung Cancer. Cancer Research, 2020, 80, 1293-1303.	0.4	49
24	Systematic Identification of Regulators of Oxidative Stress Reveals Non-canonical Roles for Peroxisomal Import and the Pentose Phosphate Pathway. Cell Reports, 2020, 30, 1417-1433.e7.	2.9	49
25	Global, distinctive, and personal changes in molecular and microbial profiles by specific fibers in humans. Cell Host and Microbe, 2022, 30, 848-862.e7.	5.1	48
26	Multiomic immune clockworks of pregnancy. Seminars in Immunopathology, 2020, 42, 397-412.	2.8	47
27	Ultra-High Performance Liquid Chromatographyâ^'Mass Spectrometry for the Fast Profiling of Histone Post-Translational Modifications. Journal of Proteome Research, 2010, 9, 5501-5509.	1.8	43
28	Cross-Laboratory Standardization of Preclinical Lipidomics Using Differential Mobility Spectrometry and Multiple Reaction Monitoring. Analytical Chemistry, 2021, 93, 16369-16378.	3.2	40
29	Global metabolic profiling to model biological processes of aging in twins. Aging Cell, 2020, 19, e13073.	3.0	38
30	Profiling of ARDS pulmonary edema fluid identifies a metabolically distinct subset. American Journal of Physiology - Lung Cellular and Molecular Physiology, 2017, 312, L703-L709.	1.3	36
31	Deep longitudinal multiomics profiling reveals two biological seasonal patterns in California. Nature Communications, 2020, 11, 4933.	5.8	36
32	Deacetylation of H4-K16Ac and heterochromatin assembly in senescence. Epigenetics and Chromatin, 2012, 5, 15.	1.8	35
33	Nat1 Deficiency Is Associated with Mitochondrial Dysfunction and Exercise Intolerance in Mice. Cell Reports, 2016, 17, 527-540.	2.9	35
34	ALDH1A3 Coordinates Metabolism With Gene Regulation in Pulmonary Arterial Hypertension. Circulation, 2021, 143, 2074-2090.	1.6	34
35	A scalable, secure, and interoperable platform for deep data-driven health management. Nature Communications, 2021, 12, 5757.	5.8	27
36	Can Metabolic Profiles Be Used as a Phenotypic Readout of the Genome to Enhance Precision Medicine?. Clinical Chemistry, 2016, 62, 676-678.	1.5	21

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37	Proteomic signatures predict preeclampsia in individual cohorts but not across cohorts – implications for clinical biomarker studies. Journal of Maternal-Fetal and Neonatal Medicine, 2022, 35, 5621-5628.	0.7	20
38	Towards personalized medicine in maternal and child health: integrating biologic and social determinants. Pediatric Research, 2021, 89, 252-258.	1.1	19
39	metID: an R package for automatable compound annotation for LCâ^'MS-based data. Bioinformatics, 2022, 38, 568-569.	1.8	15
40	Incremental value of diastolic stress test in identifying subclinical heart failure in patients with diabetes mellitus. European Heart Journal Cardiovascular Imaging, 2020, 21, 876-884.	0.5	12
41	Plasma Metabolites in Early Sepsis Identify Distinct Clusters Defined by Plasma Lipids. , 2021, 3, e0478.		10
42	Endogenous Retroviral Elements Generate Pathologic Neutrophils in Pulmonary Arterial Hypertension. American Journal of Respiratory and Critical Care Medicine, 2022, 206, 1019-1034.	2.5	10
43	Robust identification of temporal biomarkers in longitudinal omics studies. Bioinformatics, 2022, 38, 3802-3811.	1.8	10
44	The Right Heart Network and Risk Stratification in Pulmonary Arterial Hypertension. Chest, 2022, 161, 1347-1359.	0.4	9
45	Impact of acute lymphoblastic leukemia induction therapy: findings from metabolomics on non-fasted plasma samples from a biorepository. Metabolomics, 2021, 17, 64.	1.4	7
46	Cross-Platform Evaluation of Commercially Targeted and Untargeted Metabolomics Approaches to Optimize the Investigation of Psychiatric Disease. Metabolites, 2021, 11, 609.	1.3	6
47	Peripheral Oxygen Extraction and Exercise Limitation in Asymptomatic Patients with Diabetes Mellitus. American Journal of Cardiology, 2021, 149, 132-139.	0.7	4
48	In-depth triacylglycerol profiling using MS3 Q-Trap mass spectrometry. Analytica Chimica Acta, 2021, 1184, 339023.	2.6	4
49	Prediction of gestational age using urinary metabolites in term and preterm pregnancies. Scientific Reports, 2022, 12, 8033.	1.6	4
50	H2B Type 1-K Accumulates in Senescent Fibroblasts with Persistent DNA Damage along with Methylated and Phosphorylated Forms of HMGA1. Proteomes, 2021, 9, 30.	1.7	3
51	Temporal changes in soluble angiotensin-converting enzyme 2 associated with metabolic health, body composition, and proteome dynamics during a weight loss diet intervention: a randomized trial with implications for the COVID-19 pandemic. American Journal of Clinical Nutrition, 2021, 114, 1655-1665.	2.2	3
52	Design and Methods of the Validating Injury to the Renal Transplant Using Urinary Signatures (VIRTUUS) Study in Children. Transplantation Direct, 2021, 7, e791.	0.8	3