Tim H H Coorens

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5184604/publications.pdf

Version: 2024-02-01

30 papers 3,447 citations

331670 21 h-index 477307 29 g-index

44 all docs 44 docs citations

times ranked

44

3412 citing authors

#	Article	IF	CITATIONS
1	Single-cell transcriptomics reveals a distinct developmental state of KMT2A-rearranged infant B-cell acute lymphoblastic leukemia. Nature Medicine, 2022, 28, 743-751.	30.7	35
2	Somatic mutation rates scale with lifespan across mammals. Nature, 2022, 604, 517-524.	27.8	211
3	Genetic and chemotherapeutic influences on germline hypermutation. Nature, 2022, 605, 503-508.	27.8	43
4	Mutational landscape of normal epithelial cells in Lynch Syndrome patients. Nature Communications, 2022, 13, 2710.	12.8	19
5	Clonal dynamics of haematopoiesis across the human lifespan. Nature, 2022, 606, 343-350.	27.8	160
6	Inherited MUTYH mutations cause elevated somatic mutation rates and distinctive mutational signatures in normal human cells. Nature Communications, 2022, 13, .	12.8	30
7	Reliable detection of somatic mutations in solid tissues by laser-capture microdissection and low-input DNA sequencing. Nature Protocols, 2021, 16, 841-871.	12.0	82
8	Inherent mosaicism and extensive mutation of human placentas. Nature, 2021, 592, 80-85.	27.8	126
9	Somatic mutations and single-cell transcriptomes reveal the root of malignant rhabdoid tumours. Nature Communications, 2021, 12, 1407.	12.8	41
10	Somatic mutation landscapes at single-molecule resolution. Nature, 2021, 593, 405-410.	27.8	254
11	Clonal hematopoiesis and therapy-related myeloid neoplasms following neuroblastoma treatment. Blood, 2021, 137, 2992-2997.	1.4	19
12	Lineage tracing of human development through somatic mutations. Nature, 2021, 595, 85-90.	27.8	79
13	Single cell derived mRNA signals across human kidney tumors. Nature Communications, 2021, 12, 3896.	12.8	27
14	A single cell characterisation of human embryogenesis identifies pluripotency transitions and putative anterior hypoblast centre. Nature Communications, 2021, 12, 3679.	12.8	63
15	Extensive phylogenies of human development inferred from somatic mutations. Nature, 2021, 597, 387-392.	27.8	87
16	An in vitro stem cell model of human epiblast and yolk sac interaction. ELife, 2021, 10, .	6.0	24
17	The mutational landscape of human somatic and germline cells. Nature, 2021, 597, 381-386.	27.8	180
18	Increased somatic mutation burdens in normal human cells due to defective DNA polymerases. Nature Genetics, 2021, 53, 1434-1442.	21.4	85

#	Article	IF	CITATIONS
19	Convergent somatic mutations in metabolism genes in chronic liver disease. Nature, 2021, 598, 473-478.	27.8	87
20	Two of a kind: transmissible Schwann cell cancers in the endangered Tasmanian devil (Sarcophilus) Tj ETQq0 0 C	gBT/Ove	erlock 10 Tf 5
21	Somatic Evolution in Non-neoplastic IBD-Affected Colon. Cell, 2020, 182, 672-684.e11.	28.9	122
22	Lineage-Independent Tumors in Bilateral Neuroblastoma. New England Journal of Medicine, 2020, 383, 1860-1865.	27.0	23
23	Tobacco smoking and somatic mutations in human bronchial epithelium. Nature, 2020, 578, 266-272.	27.8	336
24	The mutational landscape of normal human endometrial epithelium. Nature, 2020, 580, 640-646.	27.8	338
25	Extensive heterogeneity in somatic mutation and selection in the human bladder. Science, 2020, 370, 75-82.	12.6	195
26	Embryonal precursors of Wilms tumor. Science, 2019, 366, 1247-1251.	12.6	101
27	The landscape of somatic mutation in normal colorectal epithelial cells. Nature, 2019, 574, 532-537.	27.8	468
28	Abstract 970: The mutational landscape of normal human endometrial epithelium., 2019,,.		4
29	The Origins and Vulnerabilities of Two Transmissible Cancers in Tasmanian Devils. Cancer Cell, 2018, 33, 607-619.e15.	16.8	88
30	Notwithstanding Circumstantial Alibis, Cytotoxic T Cells Can Be Major Killers of HIV-1-Infected Cells. Journal of Virology, 2016, 90, 7066-7083.	3.4	18