

Andy G Lynch

List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

1,644
citations

623574

14
h-index

887953

17
g-index

21
all docs

21
docs citations

21
times ranked

4376
citing authors

#	ARTICLE	IF	CITATIONS
1	Microbiomes of Urine and the Prostate Are Linked to Human Prostate Cancer Risk Groups. <i>European Urology Oncology</i> , 2022, 5, 412-419.	2.6	21
2	Protocol for an interdisciplinary cross-sectional study investigating the social, biological and community-level drivers of antimicrobial resistance (AMR): Holistic Approach to Unravel Antibacterial Resistance in East Africa (HATUA). <i>BMJ Open</i> , 2021, 11, e041418.	0.8	24
3	CamGFR v2: A New Model for Estimating the Glomerular Filtration Rate from Standardized or Non-standardized Creatinine in Patients with Cancer. <i>Clinical Cancer Research</i> , 2021, 27, 1381-1390.	3.2	9
4	Resolving the immune landscape of human prostate at a single-cell level in health and cancer. <i>Cell Reports</i> , 2021, 37, 110132.	2.9	40
5	Different laterality indexes are poorly correlated with one another but consistently show the tendency of males and females to be more left- and right-lateralized, respectively. <i>Royal Society Open Science</i> , 2020, 7, 191700.	1.1	19
6	Whole-genome sequencing of a sporadic primary immunodeficiency cohort. <i>Nature</i> , 2020, 583, 90-95.	13.7	148
7	Multicenter Validation of the CamGFR Model for Estimated Glomerular Filtration Rate. <i>JNCI Cancer Spectrum</i> , 2019, 3, pkz068.	1.4	6
8	Deciphering the genomic, epigenomic, and transcriptomic landscapes of pre-invasive lung cancer lesions. <i>Nature Medicine</i> , 2019, 25, 517-525.	15.2	178
9	Sequencing of prostate cancers identifies new cancer genes, routes of progression and drug targets. <i>Nature Genetics</i> , 2018, 50, 682-692.	9.4	182
10	Timing the Landmark Events in the Evolution of Clear Cell Renal Cell Cancer: TRACERx Renal. <i>Cell</i> , 2018, 173, 611-623.e17.	13.5	398
11	Telomerecat: A ploidy-agnostic method for estimating telomere length from whole genome sequencing data. <i>Scientific Reports</i> , 2018, 8, 1300.	1.6	48
12	DESNT: A Poor Prognosis Category of Human Prostate Cancer. <i>European Urology Focus</i> , 2018, 4, 842-850.	1.6	30
13	Appraising the relevance of DNA copy number loss and gain in prostate cancer using whole genome DNA sequence data. <i>PLoS Genetics</i> , 2017, 13, e1007001.	1.5	34
14	New Model for Estimating Glomerular Filtration Rate in Patients With Cancer. <i>Journal of Clinical Oncology</i> , 2017, 35, 2798-2805.	0.8	78
15	Choline Kinase Alpha as an Androgen Receptor Chaperone and Prostate Cancer Therapeutic Target. <i>Journal of the National Cancer Institute</i> , 2016, 108, djv371.	3.0	37
16	Analysis of the genetic phylogeny of multifocal prostate cancer identifies multiple independent clonal expansions in neoplastic and morphologically normal prostate tissue. <i>Nature Genetics</i> , 2015, 47, 367-372.	9.4	380