Mar RodrÃ-guez-Girondo

List of Publications by Year in descending order

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1040056 839539 19 369 9 18 citations h-index g-index papers 21 21 21 849 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Longevity defined as top 10% survivors and beyond is transmitted as a quantitative genetic trait. Nature Communications, 2019, 10, 35.	12.8	62
2	Validation of the BOADICEA model and a 313-variant polygenic risk score for breast cancer risk prediction in a Dutch prospective cohort. Genetics in Medicine, 2020, 22, 1803-1811.	2.4	49
3	A new approach to epigenome-wide discovery of non-invasive methylation biomarkers for colorectal cancer screening in circulating cell-free DNA using pooled samples. Clinical Epigenetics, 2018, 10, 53.	4.1	44
4	Effects of Family History on Relative and Absolute Risks for Colorectal Cancer: A Systematic Review and Meta-Analysis. Clinical Gastroenterology and Hepatology, 2019, 17, 2657-2667.e9.	4.4	42
5	Addition of a 161-SNP polygenic risk score to family history-based risk prediction: impact on clinical management in non- <i>BRCA1/2</i> breast cancer families. Journal of Medical Genetics, 2019, 56, 581-589.	3.2	35
6	Longevity Around the Turn of the 20th Century: Life-Long Sustained Survival Advantage for Parents of Today's Nonagenarians. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2018, 73, 1295-1302.	3.6	22
7	Molecular signatures of tumor progression in myxoid liposarcoma identified by N-glycan mass spectrometry imaging. Laboratory Investigation, 2020, 100, 1252-1261.	3.7	20
8	Universal Immunohistochemistry for Lynch Syndrome: A Systematic Review and Meta-analysis of 58,580 Colorectal Carcinomas. Clinical Gastroenterology and Hepatology, 2022, 20, e496-e507.	4.4	14
9	Longevity Relatives Count score identifies heritable longevity carriers and suggests case improvement in genetic studies. Aging Cell, 2020, 19, e13139.	6.7	13
10	An alternative approach to establishing unbiased colorectal cancer risk estimation in Lynch syndrome. Genetics in Medicine, 2019, 21, 2706-2712.	2.4	11
11	Methylation Assessment for the Prediction of Malignancy in Mediastinal Adenopathies Obtained by Endobronchial Ultrasound-Guided Transbronchial Needle Aspiration in Patients with Lung Cancer. Cancers, 2019, 11, 1408.	3.7	10
12	Survival analysis with delayed entry in selected families with application to human longevity. Statistical Methods in Medical Research, 2018, 27, 933-954.	1.5	8
13	Association between a 46-SNP Polygenic Risk Score and melanoma risk in Dutch patients with familial melanoma. Journal of Medical Genetics, 2021, 58, 760-766.	3.2	8
14	Comparison of Bisulfite Pyrosequencing and Methylation-Specific qPCR for Methylation Assessment. International Journal of Molecular Sciences, 2020, 21, 9242.	4.1	8
15	Methods for testing the Markov condition in the illness-death model: a comparative study. Statistics in Medicine, 2016, 35, 3549-3562.	1.6	7
16	The Apparent Genetic Anticipation in PMS2-Associated Lynch Syndrome Families Is Explained by Birth-cohort Effect. Cancer Epidemiology Biomarkers and Prevention, 2019, 28, 1010-1014.	2.5	6
17	Improving stability of prediction models based on correlated omics data by using network approaches. PLoS ONE, 2018, 13, e0192853.	2.5	6
18	Improved selection of participants in genetic longevity studies: family scores revisited. BMC Medical Research Methodology, 2021, 21, 7.	3.1	1

#	Article	lF	CITATIONS
19	The Hold me Tight Program for Couples Facing Huntington's Disease. Journal of Huntington's Disease, 2022, 11, 203-215.	1.9	1