## Juliette Randerson-Moor

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7325586/publications.pdf Version: 2024-02-01

	933447		1199594	
12	1,452	10	12	
papers	citations	h-index	g-index	
12	12	12	3851	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	Ulcerated melanoma: Systems biology evidence of inflammatory imbalance towards proâ€ŧumourigenicity. Pigment Cell and Melanoma Research, 2022, 35, 252-267.	3.3	4
2	MX 2 is a novel regulator of cell cycle in melanoma cells. Pigment Cell and Melanoma Research, 2020, 33, 446-457.	3.3	11
3	Genome-wide association meta-analyses combining multiple risk phenotypes provide insights into the genetic architecture of cutaneous melanoma susceptibility. Nature Genetics, 2020, 52, 494-504.	21.4	138
4	High-Resolution Copy Number Patterns From Clinically Relevant FFPE Material. Scientific Reports, 2019, 9, 8908.	3.3	6
5	β-Catenin–mediated immune evasion pathway frequently operates in primary cutaneous melanomas. Journal of Clinical Investigation, 2018, 128, 2048-2063.	8.2	71
6	Genome-wide meta-analysis identifies five new susceptibility loci for cutaneous malignant melanoma. Nature Genetics, 2015, 47, 987-995.	21.4	218
7	The Effect on Melanoma Risk of Genes Previously Associated With Telomere Length. Journal of the National Cancer Institute, 2014, 106, .	6.3	109
8	A variant in FTO shows association with melanoma risk not due to BMI. Nature Genetics, 2013, 45, 428-432.	21.4	111
9	Genome-wide association study identifies three new melanoma susceptibility loci. Nature Genetics, 2011, 43, 1108-1113.	21.4	230
10	Gene Expression Profiling of Paraffin-Embedded Primary Melanoma Using the DASL Assay Identifies Increased Osteopontin Expression as Predictive of Reduced Relapse-Free Survival. Clinical Cancer Research, 2009, 15, 6939-6946.	7.0	93
11	Genome-wide association study identifies three loci associated with melanoma risk. Nature Genetics, 2009, 41, 920-925.	21.4	422
12	The Relationship Between the Epidermal Growth Factor (EGF) 5′UTR Variant A61G and Melanoma/Nevus Susceptibility. Journal of Investigative Dermatology, 2004, 123, 755-759.	0.7	39