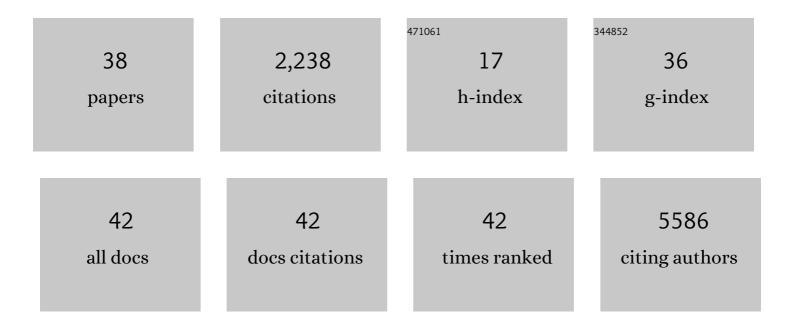
## Melanie E Garrett

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8283478/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Comprehensive functional genomic resource and integrative model for the human brain. Science, 2018, 362, .	6.0	618
2	The genetic architecture of the human cerebral cortex. Science, 2020, 367, .	6.0	450
3	Factors associated with survival in a contemporary adult sickle cell disease cohort. American Journal of Hematology, 2014, 89, 530-535.	2.0	235
4	<i>MYH9</i> and <i>APOL1</i> are both associated with sickle cell disease nephropathy. British Journal of Haematology, 2011, 155, 386-394.	1.2	139
5	Epigenome-wide meta-analysis of PTSD across 10 military and civilian cohorts identifies methylation changes in AHRR. Nature Communications, 2020, 11, 5965.	5.8	84
6	Genome-wide association study of posttraumatic stress disorder in a cohort of Iraq–Afghanistan era veterans. Journal of Affective Disorders, 2015, 184, 225-234.	2.0	81
7	Alloimmunization in sickle cell disease: changing antibody specificities and association with chronic pain and decreased survival. Transfusion, 2015, 55, 1378-1387.	0.8	75
8	An epigenome-wide association study of posttraumatic stress disorder in US veterans implicates several new DNA methylation loci. Clinical Epigenetics, 2020, 12, 46.	1.8	64
9	Genetic correlations and genome-wide associations of cortical structure in general population samples of 22,824 adults. Nature Communications, 2020, 11, 4796.	5.8	61
10	Cell-type-specific effects of genetic variation on chromatin accessibility during human neuronal differentiation. Nature Neuroscience, 2021, 24, 941-953.	7.1	47
11	In vivo Modeling Implicates APOL1 in Nephropathy: Evidence for Dominant Negative Effects and Epistasis under Anemic Stress. PLoS Genetics, 2015, 11, e1005349.	1.5	45
12	An Examination of the Association between 5-HTTLPR, Combat Exposure, and PTSD Diagnosis among U.S. Veterans. PLoS ONE, 2015, 10, e0119998.	1.1	29
13	Genetic determinants of telomere length from 109,122 ancestrally diverse whole-genome sequences in TOPMed. Cell Genomics, 2022, 2, 100084.	3.0	29
14	Clinical and metabolomic risk factors associated with rapid renal function decline in sickle cell disease. American Journal of Hematology, 2018, 93, 1451-1460.	2.0	28
15	DNA Copy Number Variants of Known Glaucoma Genes in Relation to Primary Open-Angle Glaucoma. Investigative Ophthalmology and Visual Science, 2014, 55, 8251-8258.	3.3	27
16	Clonal hematopoiesis in sickle cell disease. Journal of Clinical Investigation, 2022, 132, .	3.9	26
17	Pregnancy continuation and organizational religious activity following prenatal diagnosis of a lethal fetal defect are associated with improved psychological outcome. Prenatal Diagnosis, 2015, 35, 761-768.	1.1	19
18	Rare and de novo coding variants in chromodomain genes in Chiari I malformation. American Journal of Human Genetics. 2021. 108. 100-114.	2.6	17

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19	A common functional <i>PIEZO1</i> deletion allele associates with red blood cell density in sickle cell disease patients. American Journal of Hematology, 2018, 93, E362-E365.	2.0	15
20	Potential causal role of l-glutamine in sickle cell disease painful crises: A Mendelian randomization analysis. Blood Cells, Molecules, and Diseases, 2021, 86, 102504.	0.6	14
21	Systematic Functional Testing of Rare Variants: Contributions of <i>CFI</i> to Age-Related Macular Degeneration. , 2017, 58, 1570.		13
22	The association of single-nucleotide polymorphisms in theÂoxytocin receptor and G protein–coupled receptor kinase 6 ( GRK6 ) genes with oxytocin dosing requirements and labor outcomes. American Journal of Obstetrics and Gynecology, 2017, 217, 367.e1-367.e9.	0.7	12
23	Rare functional genetic variants in COL7A1, COL6A5, COL1A2 and COL5A2 frequently occur in Chiari Malformation Type 1. PLoS ONE, 2021, 16, e0251289.	1.1	12
24	Gene Expression Analysis in Three Posttraumatic Stress Disorder Cohorts Implicates Inflammation and Innate Immunity Pathways and Uncovers Shared Genetic Risk With Major Depressive Disorder. Frontiers in Neuroscience, 2021, 15, 678548.	1.4	12
25	Rapid decline in estimated glomerular filtration rate in sickle cell anemia: results of a multicenter pooled analysis. Haematologica, 2021, 106, 1749-1753.	1.7	11
26	Joint eQTL assessment of whole blood and dura mater tissue from individuals with Chiari type I malformation. BMC Genomics, 2015, 16, 11.	1.2	10
27	Thrombospondinâ€I gene polymorphism is associated with estimated pulmonary artery pressure in patients with sickle cell anemia. American Journal of Hematology, 2017, 92, E31-E34.	2.0	10
28	Examining Individual and Synergistic Contributions of PTSD and Genetics to Blood Pressure: A Trans-Ethnic Meta-Analysis. Frontiers in Neuroscience, 2021, 15, 678503.	1.4	10
29	Sex dependent glial-specific changes in the chromatin accessibility landscape in late-onset Alzheimer's disease brains. Molecular Neurodegeneration, 2021, 16, 58.	4.4	10
30	Effect of genetic variation in the nicotinic receptor genes on risk for posttraumatic stress disorder. Psychiatry Research, 2015, 229, 326-331.	1.7	6
31	Further evidence for a role of the ADRB2 gene in risk for posttraumatic stress disorder. Journal of Psychiatric Research, 2017, 84, 59-61.	1.5	5
32	Trauma and posttraumatic stress disorder modulate polygenic predictors of hippocampal and amygdala volume. Translational Psychiatry, 2021, 11, 637.	2.4	4
33	RNA sequencing of isolated cell populations expressing human APOL1 G2 risk variant reveals molecular correlates of sickle cell nephropathy in zebrafish podocytes. PLoS ONE, 2019, 14, e0217042.	1.1	3
34	Genome-Wide Studies in Sickle Cell Anemia Show Associations Between SNPs in the Olfactory Receptor Gene Cluster and Fetal Hemoglobin Concentration Blood, 2009, 114, 821-821.	0.6	2
35	Polymorphisms in TNFα Are Associated with Cerebrovascular Events in Sickle Cell Disease Blood, 2009, 114, 1540-1540.	0.6	1
36	Genes Associated with Survival in Adult Sickle Cell Disease. Blood, 2014, 124, 2719-2719.	0.6	1

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37	Prolonged Survival despite High Disease Burden in Elderly (≥55) Patients with Hb SS or Hb Sβ0 Thalassemia. Blood, 2008, 112, 710-710.	0.6	ο
38	Genome-Wide Evaluation of Epistasis with APOL1 Risk Variants in Sickle Cell Disease Nephropathy. Blood, 2015, 126, 3401-3401.	0.6	0