

# Julia M Sealock

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6867566/publications.pdf>

Version: 2024-02-01

12  
papers

2,651  
citations

840776

11  
h-index

1058476

14  
g-index

20  
all docs

20  
docs citations

20  
times ranked

5392  
citing authors

#	ARTICLE	IF	CITATIONS
1	Proposing a Sex-Adjusted Sodium-Adjusted MELD Score for Liver Transplant Allocation. <i>JAMA Surgery</i> , 2022, 157, 618.	4.3	16
2	Genetic risk for major depressive disorder and loneliness in sex-specific associations with coronary artery disease. <i>Molecular Psychiatry</i> , 2021, 26, 4254-4264.	7.9	26
3	Clinical laboratory test-wide association scan of polygenic scores identifies biomarkers of complex disease. <i>Genome Medicine</i> , 2021, 13, 6.	8.2	49
4	Pancreatic adenocarcinoma in liver transplant recipients: a case series. <i>Annals of Pancreatic Cancer</i> , 2021, 4, 0-0.	1.2	1
5	Elevated Polygenic Burden for Autism Spectrum Disorder Is Associated With the Broad Autism Phenotype in Mothers of Individuals With Autism Spectrum Disorder. <i>Biological Psychiatry</i> , 2021, 89, 476-485.	1.3	32
6	A genome-wide association study with 1,126,563 individuals identifies new risk loci for Alzheimer's disease. <i>Nature Genetics</i> , 2021, 53, 1276-1282.	21.4	430
7	Use of the PsycheMERGE Network to Investigate the Association Between Depression Polygenic Scores and White Blood Cell Count. <i>JAMA Psychiatry</i> , 2021, 78, 1365.	11.0	31
8	Genome-wide association study of problematic opioid prescription use in 132,113 23andMe research participants of European ancestry. <i>Molecular Psychiatry</i> , 2021, 26, 6209-6217.	7.9	26
9	Genome-wide meta-analysis of problematic alcohol use in 435,563 individuals yields insights into biology and relationships with other traits. <i>Nature Neuroscience</i> , 2020, 23, 809-818.	14.8	242
10	Phenome-wide investigation of health outcomes associated with genetic predisposition to loneliness. <i>Human Molecular Genetics</i> , 2019, 28, 3853-3865.	2.9	62
11	Genome-wide meta-analysis identifies new loci and functional pathways influencing Alzheimer's disease risk. <i>Nature Genetics</i> , 2019, 51, 404-413.	21.4	1,625
12	Sex-specific genetic predictors of Alzheimer's disease biomarkers. <i>Acta Neuropathologica</i> , 2018, 136, 857-872.	7.7	87