

# Abigail L Savage

## List of Publications by Year in descending order

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Version: 2024-02-01

24  
papers

403  
citations

933264

10  
h-index

839398

18  
g-index

27  
all docs

27  
docs citations

27  
times ranked

574  
citing authors

#	ARTICLE	IF	CITATIONS
1	Innovative approaches for treatment of osteosarcoma. <i>Experimental Biology and Medicine</i> , 2022, 247, 310-316.	1.1	18
2	Characterisation of the Function of a SINE-VNTR-Alu Retrotransposon to Modulate Isoform Expression at the MAPT Locus. <i>Frontiers in Molecular Neuroscience</i> , 2022, 15, 815695.	1.4	7
3	Locus specific reduction of L1 expression in the cortices of individuals with amyotrophic lateral sclerosis. <i>Molecular Brain</i> , 2022, 15, 25.	1.3	2
4	Longitudinal intronic RNA-Seq analysis of Parkinson's disease patients reveals disease-specific nascent transcription. <i>Experimental Biology and Medicine</i> , 2022, 247, 945-957.	1.1	5
5	Mechanisms of disease-associated SINE-VNTR-Alus. <i>Experimental Biology and Medicine</i> , 2022, 247, 756-764.	1.1	7
6	At the dawn of the transcriptomic medicine. <i>Experimental Biology and Medicine</i> , 2021, 246, 286-292.	1.1	7
7	Transcript Variants of Genes Involved in Neurodegeneration Are Differentially Regulated by the APOE and MAPT Haplotypes. <i>Genes</i> , 2021, 12, 423.	1.0	7
8	The TOMM40 523 polymorphism in disease risk and age of symptom onset in two independent cohorts of Parkinson's disease. <i>Scientific Reports</i> , 2021, 11, 6363.	1.6	6
9	Variable number tandem repeats – Their emerging role in sickness and health. <i>Experimental Biology and Medicine</i> , 2021, 246, 1368-1376.	1.1	11
10	Reference SVA insertion polymorphisms are associated with Parkinson's Disease progression and differential gene expression. <i>Npj Parkinson's Disease</i> , 2021, 7, 44.	2.5	22
11	Expression Quantitative Trait Loci (eQTLs) Associated with Retrotransposons Demonstrate their Modulatory Effect on the Transcriptome. <i>International Journal of Molecular Sciences</i> , 2021, 22, 6319.	1.8	10
12	TOMM40 523 poly-T repeat length is a determinant of longitudinal cognitive decline in Parkinson's disease. <i>Npj Parkinson's Disease</i> , 2021, 7, 56.	2.5	2
13	An Increased Burden of Highly Active Retrotransposition Competent L1s Is Associated with Parkinson's Disease Risk and Progression in the PPMI Cohort. <i>International Journal of Molecular Sciences</i> , 2020, 21, 6562.	1.8	18
14	Frequency and methylation status of selected retrotransposition competent L1 loci in amyotrophic lateral sclerosis. <i>Molecular Brain</i> , 2020, 13, 154.	1.3	7
15	Disease-modifying effects of an SCAF4 structural variant in a predominantly SOD1 ALS cohort. <i>Neurology: Genetics</i> , 2020, 6, e470.	0.9	9
16	Single Nucleotide Polymorphisms Associated With Gut Homeostasis Influence Risk and Age-at-Onset of Parkinson's Disease. <i>Frontiers in Aging Neuroscience</i> , 2020, 12, 603849.	1.7	16
17	Non-coding genetic variation shaping mental health. <i>Current Opinion in Psychology</i> , 2019, 27, 18-24.	2.5	14
18	The Role of SINE-VNTR-Alu (SVA) Retrotransposons in Shaping the Human Genome. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5977.	1.8	22

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19	Retrotransposons in the development and progression of amyotrophic lateral sclerosis. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2019, 90, 284-293.	0.9	29
20	The RNA processing factors THRAP3 and BCLAF1 promote the DNA damage response through selective mRNA splicing and nuclear export. <i>Nucleic Acids Research</i> , 2017, 45, 12816-12833.	6.5	79
21	Molecular signatures of mood stabilisers highlight the role of the transcription factor REST/NRSF. <i>Journal of Affective Disorders</i> , 2015, 172, 63-73.	2.0	10
22	An Evaluation of a SVA Retrotransposon in the FUS Promoter as a Transcriptional Regulator and Its Association to ALS. <i>PLoS ONE</i> , 2014, 9, e90833.	1.1	32
23	Characterisation of the potential function of SVA retrotransposons to modulate gene expression patterns. <i>BMC Evolutionary Biology</i> , 2013, 13, 101.	3.2	55
24	Polymorphic variation as a driver of differential neuropeptide gene expression. <i>Neuropeptides</i> , 2013, 47, 395-400.	0.9	8