

Angela K Hodges

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

31 papers	2,102 citations	15 h-index	41 g-index
41 ext. papers	3,442 ext. citations	12.2 avg, IF	3.15 L-index

#	Paper	IF	Citations
31	Dysregulated expression levels of APOE4 in peripheral blood are associated with brain atrophy and amyloid- β deposition in Alzheimer's disease. <i>Alzheimer's Research and Therapy</i> , 2021 , 13, 183	9	1
30	Altered nuclear architecture in blood cells from Huntington's disease patients. <i>Neurological Sciences</i> , 2021 , 1	3.5	0
29	Genome sequencing analysis identifies new loci associated with Lewy body dementia and provides insights into its genetic architecture. <i>Nature Genetics</i> , 2021 , 53, 294-303	36.3	31
28	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	36.3	40
27	Dysregulated Fc gamma receptor-mediated phagocytosis pathway in Alzheimer's disease: network-based gene expression analysis. <i>Neurobiology of Aging</i> , 2020 , 88, 24-32	5.6	6
26	Association of blood-based transcriptional risk scores with biomarkers for Alzheimer disease. <i>Neurology: Genetics</i> , 2020 , 6, e517	3.8	3
25	An epigenome-wide association study of Alzheimer's disease blood highlights robust DNA hypermethylation in the HOXB6 gene. <i>Neurobiology of Aging</i> , 2020 , 95, 26-45	5.6	17
24	Genome-wide transcriptome analysis identifies novel dysregulated genes implicated in Alzheimer's pathology. <i>Alzheimer's and Dementia</i> , 2020 , 16, 1213-1223	1.2	8
23	Integrated lipidomics and proteomics network analysis highlights lipid and immunity pathways associated with Alzheimer's disease. <i>Translational Neurodegeneration</i> , 2020 , 9, 36	10.3	15
22	Transcriptomic analysis of probable asymptomatic and symptomatic alzheimer brains. <i>Brain, Behavior, and Immunity</i> , 2019 , 80, 644-656	16.6	27
21	ALSGeneScanner: a pipeline for the analysis and interpretation of DNA sequencing data of ALS patients. <i>Amyotrophic Lateral Sclerosis and Frontotemporal Degeneration</i> , 2019 , 20, 207-215	3.6	6
20	Regional mitochondrial DNA and cell-type changes in post-mortem brains of non-diabetic Alzheimer's disease are not present in diabetic Alzheimer's disease. <i>Scientific Reports</i> , 2019 , 9, 11386	4.9	10
19	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates APOE4, tau, immunity and lipid processing. <i>Nature Genetics</i> , 2019 , 51, 414-430	36.3	917
18	Plasma levels of soluble TREM2 and neurofilament light chain in TREM2 rare variant carriers. <i>Alzheimer's Research and Therapy</i> , 2019 , 11, 94	9	15
17	Loss of Trem2 in microglia leads to widespread disruption of cell coexpression networks in mouse brain. <i>Neurobiology of Aging</i> , 2018 , 69, 151-166	5.6	17
16	Mitochondrial genes are altered in blood early in Alzheimer's disease. <i>Neurobiology of Aging</i> , 2017 , 53, 36-47	5.6	78
15	Rare coding variants in PLCG2, ABI3, and TREM2 implicate microglial-mediated innate immunity in Alzheimer's disease. <i>Nature Genetics</i> , 2017 , 49, 1373-1384	36.3	508

14	[P2098]: TSPO IMMUNOSTAINING IN AD CASES WITH/WITHOUT AN AD-ASSOCIATED TREM2 VARIANT 2017 , 13, P644-P645		
13	[P1020]: SCREENING FDA-APPROVED COMPOUNDS IN A TREM2 CELL MODEL OF ALZHEIMER'S DISEASE 2017 , 13, P240		
12	[O20306]: ASSESSING TREM2 FUNCTION IN ALZHEIMER'S DISEASE WITH RNA-SEQ 2017 , 13, P590		
11	A Pathway Based Classification Method for Analyzing Gene Expression for Alzheimer's Disease Diagnosis. <i>Journal of Alzheimer's Disease</i> , 2016 , 49, 659-69	4.3	26
10	IC-P-072: Gene Expression Of ABCA7 Dysregulated in Peripheral Blood is Associated With Decreased Metabolic Activity in Hippocampus 2016 , 12, P56-P58		
9	IC-P-074: Genome-Wide Meta-Analysis of Transcriptome Profiling Identifies Novel Dysregulated Genes Implicated in Alzheimer's Disease 2016 , 12, P58-P60		
8	P3-087: Gene Expression of ABCA7 Dysregulated in Peripheral Blood is Associated With Decreased Metabolic Activity in Hippocampus 2016 , 12, P851-P853		
7	O2-06-02: Genome-Wide Meta-Analysis of Transcriptome Profiling Identifies Novel Dysregulated Genes Implicated in Alzheimer's Disease 2016 , 12, P238-P239		
6	P1-002: AD-Associated TREM2 Variants Lead to Some Subpopulations of Microglia to be Less Abundant But More Activated 2016 , 12, P397-P397		
5	P4-321: Using Gene Expression and Genetics to Predict Amyloid Burden Before Dementia 2016 , 12, P1157-P1157		
4	A novel multi-tissue RNA diagnostic of healthy ageing relates to cognitive health status. <i>Genome Biology</i> , 2015 , 16, 185	18.3	112
3	No Evidence to Suggest that the Use of Acetylcholinesterase Inhibitors Confounds the Results of Two Blood-Based Biomarker Studies in Alzheimer's Disease. <i>Journal of Alzheimer's Disease</i> , 2015 , 47, 741-50	4.3	2
2	Alzheimer's disease biomarker discovery using SOMAscan multiplexed protein technology. <i>Alzheimer's and Dementia</i> , 2014 , 10, 724-34	1.2	133
1	Mitochondrial dysfunction and immune activation are detectable in early Alzheimer's disease blood. <i>Journal of Alzheimer's Disease</i> , 2012 , 30, 685-710	4.3	104