## Angela K Hodges

## List of Publications by Citations

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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.

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

#	Paper	IF	Citations
31	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates All tau, immunity and lipid processing. <i>Nature Genetics</i> , <b>2019</b> , 51, 414-430	36.3	917
30	Rare coding variants in PLCG2, ABI3, and TREM2 implicate microglial-mediated innate immunity in Alzheimer's disease. <i>Nature Genetics</i> , <b>2017</b> , 49, 1373-1384	36.3	508
29	Alzheimer's disease biomarker discovery using SOMAscan multiplexed protein technology. <i>Alzheimer</i> and Dementia, <b>2014</b> , 10, 724-34	1.2	133
28	A novel multi-tissue RNA diagnostic of healthy ageing relates to cognitive health status. <i>Genome Biology</i> , <b>2015</b> , 16, 185	18.3	112
27	Mitochondrial dysfunction and immune activation are detectable in early Alzheimer's disease blood. <i>Journal of Alzheimer</i> Disease, <b>2012</b> , 30, 685-710	4.3	104
26	Mitochondrial genes are altered in blood early in Alzheimer's disease. <i>Neurobiology of Aging</i> , <b>2017</b> , 53, 36-47	5.6	78
25	A genome-wide association study with 1,126,563 individuals identifies new risk loci for Alzheimer's disease. <i>Nature Genetics</i> , <b>2021</b> , 53, 1276-1282	36.3	40
24	Genome sequencing analysis identifies new loci associated with Lewy body dementia and provides insights into its genetic architecture. <i>Nature Genetics</i> , <b>2021</b> , 53, 294-303	36.3	31
23	Transcriptomic analysis of probable asymptomatic and symptomatic alzheimer brains. <i>Brain, Behavior, and Immunity,</i> <b>2019</b> , 80, 644-656	16.6	27
22	A Pathway Based Classification Method for Analyzing Gene Expression for Alzheimer's Disease Diagnosis. <i>Journal of Alzheimer</i> Disease, <b>2016</b> , 49, 659-69	4.3	26
21	Loss of Trem2 in microglia leads to widespread disruption of cell coexpression networks in mouse brain. <i>Neurobiology of Aging</i> , <b>2018</b> , 69, 151-166	5.6	17
20	An epigenome-wide association study of Alzheimer's disease blood highlights robust DNA hypermethylation in the HOXB6 gene. <i>Neurobiology of Aging</i> , <b>2020</b> , 95, 26-45	5.6	17
19	Integrated lipidomics and proteomics network analysis highlights lipid and immunity pathways associated with Alzheimer's disease. <i>Translational Neurodegeneration</i> , <b>2020</b> , 9, 36	10.3	15
18	Plasma levels of soluble TREM2 and neurofilament light chain in TREM2 rare variant carriers. <i>Alzheimer Research and Therapy</i> , <b>2019</b> , 11, 94	9	15
17	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> , <b>2019</b> , 9, 11386	4.9	10
16	Genome-wide transcriptome analysis identifies novel dysregulated genes implicated in Alzheimer's pathology. <i>Alzheimer</i> and <i>Dementia</i> , <b>2020</b> , 16, 1213-1223	1.2	8
15	ALSgeneScanner: a pipeline for the analysis and interpretation of DNA sequencing data of ALS patients. <i>Amyotrophic Lateral Sclerosis and Frontotemporal Degeneration</i> , <b>2019</b> , 20, 207-215	3.6	6

## LIST OF PUBLICATIONS

14	Dysregulated Fc gamma receptor-mediated phagocytosis pathway in Alzheimer's disease: network-based gene expression analysis. <i>Neurobiology of Aging</i> , <b>2020</b> , 88, 24-32	5.6	6
13	Association of blood-based transcriptional risk scores with biomarkers for Alzheimer disease. <i>Neurology: Genetics</i> , <b>2020</b> , 6, e517	3.8	3
12	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</i> Disease, <b>2015</b> , 47, 741-50	4.3	2
11	Dysregulated expression levels of APH1B in peripheral blood are associated with brain atrophy and amyloid-Ideposition in Alzheimer's disease. <i>Alzheimer Research and Therapy</i> , <b>2021</b> , 13, 183	9	1
10	Altered nuclear architecture in blood cells from Huntington's disease patients. <i>Neurological Sciences</i> , <b>2021</b> , 1	3.5	O
9	[P2 <b>0</b> 98]: TSPO IMMUNOSTAINING IN AD CASES WITH/WITHOUT AN AD-ASSOCIATED TREM2 VARIANT <b>2017</b> , 13, P644-P645		
8	[P1020]: SCREENING FDA-APPROVED COMPOUNDS IN A TREM2 CELL MODEL OF ALZHEIMER'S DISEASE <b>2017</b> , 13, P240		
7	[O2l13l06]: ASSESSING TREM2 FUNCTION IN ALZHEIMER's DISEASE WITH RNA-SEQ <b>2017</b> , 13, P590		
6	IC-P-072: Gene Expression Of ABCA7 Dysregulated in Peripheral Blood is Associated With Decreased Metabolic Activity in Hippocampus <b>2016</b> , 12, P56-P58		
5	IC-P-074: Genome-Wide Meta-Analysis of Transcriptome Profiling Identifies Novel Dysregulated Genes Implicated in Alzheimer Disease <b>2016</b> , 12, P58-P60		
4	P3-087: Gene Expression of ABCA7 Dysregulated in Peripheral Blood is Associated With Decreased Metabolic Activity in Hippocampus <b>2016</b> , 12, P851-P853		
3	O2-06-02: Genome-Wide Meta-Analysis of Transcriptome Profiling Identifies Novel Dysregulated Genes Implicated in Alzheimer's Disease <b>2016</b> , 12, P238-P239		
2	P1-002: AD-Associated TREM2 Variants Lead to Some Subpopulations of Microglia to be Less Abundant But More Activated <b>2016</b> , 12, P397-P397		
1	P4-321: Using Gene Expression and Genetics to Predict Amyloid Burden Before Dementia <b>2016</b> , 12, P1	157-P1	157