

# Mark T W Ebbert

## 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.

44 papers	2,831 citations	22 h-index	51 g-index
51 ext. papers	3,449 ext. citations	9 avg, IF	4.31 L-index

#	Paper	IF	Citations
44	Culture shock: microglial heterogeneity, activation, and disrupted single-cell microglial networks in vitro.. <i>Molecular Neurodegeneration</i> , <b>2022</b> , 17, 26	19	1
43	Long-read targeted sequencing uncovers clinicopathological associations for C9orf72-linked diseases. <i>Brain</i> , <b>2021</b> , 144, 1082-1088	11.2	2
42	Systematic analysis of dark and camouflaged genes reveals disease-relevant genes hiding in plain sight. <i>Genome Biology</i> , <b>2019</b> , 20, 97	18.3	68
41	Microglial translational profiling reveals a convergent APOE pathway from aging, amyloid, and tau. <i>Journal of Experimental Medicine</i> , <b>2018</b> , 215, 2235-2245	16.6	85
40	Long-read sequencing across the C9orf72 'GGGGCC' repeat expansion: implications for clinical use and genetic discovery efforts in human disease. <i>Molecular Neurodegeneration</i> , <b>2018</b> , 13, 46	19	66
39	An Epigenetic Spin to ALS and FTD. <i>Advances in Neurobiology</i> , <b>2018</b> , 20, 1-29	2.1	4
38	Corticobasal degeneration with TDP-43 pathology presenting with progressive supranuclear palsy syndrome: a distinct clinicopathologic subtype. <i>Acta Neuropathologica</i> , <b>2018</b> , 136, 389-404	14.3	27
37	Common DNA Variants Accurately Rank an Individual of Extreme Height. <i>International Journal of Genomics</i> , <b>2018</b> , 2018, 5121540	2.5	3
36	Poly(GR) impairs protein translation and stress granule dynamics in C9orf72-associated frontotemporal dementia and amyotrophic lateral sclerosis. <i>Nature Medicine</i> , <b>2018</b> , 24, 1136-1142	50.5	149
35	Evaluation of the Abbott RealTime HCV genotype II plus RUO (PLUS) assay with reference to core and NS5B sequencing. <i>Journal of Clinical Virology</i> , <b>2017</b> , 90, 26-31	14.5	10
34	Linkage, whole genome sequence, and biological data implicate variants in RAB10 in Alzheimer's disease resilience. <i>Genome Medicine</i> , <b>2017</b> , 9, 100	14.4	40
33	Conserved DNA methylation combined with differential frontal cortex and cerebellar expression distinguishes C9orf72-associated and sporadic ALS, and implicates SERPINA1 in disease. <i>Acta Neuropathologica</i> , <b>2017</b> , 134, 715-728	14.3	18
32	Repetitive element transcripts are elevated in the brain of C9orf72 ALS/FTLD patients. <i>Human Molecular Genetics</i> , <b>2017</b> , 26, 3421-3431	5.6	63
31	Interaction between variants in CLU and MS4A4E modulates Alzheimer's disease risk. <i>Alzheimer's and Dementia</i> , <b>2016</b> , 12, 121-129	1.2	15
30	Presenilin E318G variant and Alzheimer's disease risk: the Cache County study. <i>BMC Genomics</i> , <b>2016</b> , 17 Suppl 3, 438	4.5	8
29	A novel approach for multi-SNP GWAS and its application in Alzheimer's disease. <i>BMC Bioinformatics</i> , <b>2016</b> , 17 Suppl 7, 268	3.6	8
28	Evaluating the necessity of PCR duplicate removal from next-generation sequencing data and a comparison of approaches. <i>BMC Bioinformatics</i> , <b>2016</b> , 17 Suppl 7, 239	3.6	65

27	<p> Variants in ACPP are associated with cerebrospinal fluid Prostatic Acid Phosphatase levels. <i>BMC Genomics</i>, <b>2016</b>, 17 Suppl 3, 439 </p>	4.5	1
26	<p> Variants in CCL16 are associated with blood plasma and cerebrospinal fluid CCL16 protein levels. <i>BMC Genomics</i>, <b>2016</b>, 17 Suppl 3, 437 </p>	4.5	1
25	<p> Genome-wide association study of prolactin levels in blood plasma and cerebrospinal fluid. <i>BMC Genomics</i>, <b>2016</b>, 17 Suppl 3, 436 </p>	4.5	2
24	<p> P3-082: Assessment of the Genetic Variance of Late-Onset Alzheimer's Disease <b>2016</b>, 12, P849-P850 </p>		
23	<p> Liver kinase B1 inhibits the expression of inflammation-related genes postcontraction in skeletal muscle. <i>Journal of Applied Physiology</i>, <b>2016</b>, 120, 876-88 </p>	3.7	8
22	<p> Population-based analysis of cholesteryl ester transfer protein identifies association between I405V and cognitive decline: the Cache County Study. <i>Neurobiology of Aging</i>, <b>2015</b>, 36, 547.e1-3 </p>	5.6	5
21	<p> Bridging the gap between statistical and biological epistasis in Alzheimer's disease. <i>BioMed Research International</i>, <b>2015</b>, 2015, 870123 </p>	3	22
20	<p> Genetic discoveries in AD using CSF amyloid and tau. <i>Current Genetic Medicine Reports</i>, <b>2014</b>, 2, 23-29 </p>	2.2	7
19	<p> Variant Tool Chest: an improved tool to analyze and manipulate variant call format (VCF) files. <i>BMC Bioinformatics</i>, <b>2014</b>, 15 Suppl 7, S12 </p>	3.6	9
18	<p> Intrinsic subtypes from PAM50 gene expression assay in a population-based breast cancer cohort: differences by age, race, and tumor characteristics. <i>Cancer Epidemiology Biomarkers and Prevention</i>, <b>2014</b>, 23, 714-24 </p>	4	84
17	<p> Population-based analysis of Alzheimer's disease risk alleles implicates genetic interactions. <i>Biological Psychiatry</i>, <b>2014</b>, 75, 732-7 </p>	7.9	43
16	<p> Assessment of TREM2 rs75932628 association with Alzheimer's disease in a population-based sample: the Cache County Study. <i>Neurobiology of Aging</i>, <b>2013</b>, 34, 2889.e11-3 </p>	5.6	41
15	<p> A phase II study of UCN-01 in combination with irinotecan in patients with metastatic triple negative breast cancer. <i>Breast Cancer Research and Treatment</i>, <b>2013</b>, 137, 483-92 </p>	4.4	82
14	<p> Application of a new informatics tool for contamination screening in the HIV sequencing laboratory. <i>Journal of Clinical Virology</i>, <b>2013</b>, 57, 249-53 </p>	14.5	5
13	<p> PAM50 proliferation score as a predictor of weekly paclitaxel benefit in breast cancer. <i>Breast Cancer Research and Treatment</i>, <b>2013</b>, 138, 457-66 </p>	4.4	78
12	<p> Genetics of Alzheimer's disease. <i>BioMed Research International</i>, <b>2013</b>, 2013, 254954 </p>	3	59
11	<p> TRIM29 functions as a tumor suppressor in nontumorigenic breast cells and invasive ER+ breast cancer. <i>American Journal of Pathology</i>, <b>2012</b>, 180, 839-47 </p>	5.8	38
10	<p> PAM50 breast cancer subtyping by RT-qPCR and concordance with standard clinical molecular markers. <i>BMC Medical Genomics</i>, <b>2012</b>, 5, 44 </p>	3.7	199

9	Agreement in risk prediction between the 21-gene recurrence score assay (Oncotype DX <sup>®</sup> ) and the PAM50 breast cancer intrinsic Classifier <sup>®</sup> in early-stage estrogen receptor-positive breast cancer. <i>Oncologist</i> , <b>2012</b> , 17, 492-8	5.7	63
8	Responsiveness of intrinsic subtypes to adjuvant anthracycline substitution in the NCIC.CTG MA.5 randomized trial. <i>Clinical Cancer Research</i> , <b>2012</b> , 18, 2402-12	12.9	117
7	Characterization of uncertainty in the classification of multivariate assays: application to PAM50 centroid-based genomic predictors for breast cancer treatment plans. <i>Journal of Clinical Bioinformatics</i> , <b>2011</b> , 1, 37		15
6	Tumor grafts derived from women with breast cancer authentically reflect tumor pathology, growth, metastasis and disease outcomes. <i>Nature Medicine</i> , <b>2011</b> , 17, 1514-20	50.5	695
5	A comparison of PAM50 intrinsic subtyping with immunohistochemistry and clinical prognostic factors in tamoxifen-treated estrogen receptor-positive breast cancer. <i>Clinical Cancer Research</i> , <b>2010</b> , 16, 5222-32	12.9	546
4	Evolutionary pressure on mitochondrial cytochrome b is consistent with a role of Cytb17T affecting longevity during caloric restriction. <i>PLoS ONE</i> , <b>2009</b> , 4, e5836	3.7	29
3	DNA reference alignment benchmarks based on tertiary structure of encoded proteins. <i>Bioinformatics</i> , <b>2007</b> , 23, 2648-9	7.2	42
2	Pharmacogenomics: analysing SNPs in the CYP2D6 gene using amino acid properties. <i>International Journal of Bioinformatics Research and Applications</i> , <b>2007</b> , 3, 471-9	0.9	2
1	Long-read sequencing across the C9orf72 CCGGCC repeat expansion: implications for clinical use and genetic discovery efforts in human disease		1