Oscar Harari

List of Publications by Year
in descending order

51 papers	3,799 citations	218677 26 h-index	254184 43 g-index
73 all docs	73 docs citations	73 times ranked	6449 citing authors

#	Article	IF	Citations
1	Murine roseolovirus does not accelerate amyloid-Î² pathology and human roseoloviruses are not over-represented in Alzheimer disease brains. Molecular Neurodegeneration, 2022, 17, 10.	10.8	9
2	Multi-ancestry GWAS reveals excitotoxicity associated with outcome after ischaemic stroke. Brain, 2022, 145, 2394-2406.	7.6	15
3	Circular RNA detection identifies circPSEN1 alterations in brain specific to autosomal dominant Alzheimer's disease. Acta Neuropathologica Communications, 2022, 10, 29.	5.2	11
4	Alzheimer's disease alters oligodendrocytic glycolytic and ketolytic gene expression. Alzheimer's and Dementia, 2021, 17, 1474-1486.	0.8	37
5	Meningeal lymphatics affect microglia responses and anti-Aî² immunotherapy. Nature, 2021, 593, 255-260.	27.8	179
6	Alzheimer's Disease Alters Oligodendrocytic Glycolytic and Ketolytic Gene Expression. FASEB Journal, 2021, 35, .	0.5	1
7	Genomic atlas of the proteome from brain, CSF and plasma prioritizes proteins implicated in neurological disorders. Nature Neuroscience, 2021, 24, 1302-1312.	14.8	105
8	Neuronal VCP loss of function recapitulates FTLD-TDP pathology. Cell Reports, 2021, 36, 109399.	6.4	25
9	Profiling the metabolic landscape of AD. Alzheimer's and Dementia, 2021, 17, e050086.	0.8	0
10	Genome-wide scan of Alzheimer disease cohort identifies genetic loci associated with human brain metabolite levels.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e051756.	0.8	0
11	Functional exploration of AGFG2, a novel player in the pathology of Alzheimer disease.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e054240.	0.8	1
12	LMNA-mediated nucleoskeleton dysregulation in Alzheimer disease.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e054396.	0.8	0
13	Single nuclei RNA-sequencing of GWAS loci variant carriers elucidates cell-types and transcriptional profile alterations associated with Alzheimer disease.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e054402.	0.8	0
14	Multi-omics approaches reveal a link between the MS4A gene loci, TREM2, and microglia function.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e054553.	0.8	0
15	Stem cell models of primary tauopathies reveal defects in synaptic function.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e054566.	0.8	0
16	The TMEM106B FTLD-protective variant, rs1990621, is also associated with increased neuronal proportion. Acta Neuropathologica, 2020, 139, 45-61.	7.7	51
17	Overlapping genetic architecture between Parkinson disease and melanoma. Acta Neuropathologica, 2020, 139, 347-364.	7.7	23
18	Functional genomic analyses uncover APOE-mediated regulationÂofÂbrain and cerebrospinal fluid beta-amyloid levels in Parkinson disease. Acta Neuropathologica Communications, 2020, 8, 196.	5.2	8

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19	TREM2 activation on microglia promotes myelin debris clearance and remyelination in a model of multiple sclerosis. Acta Neuropathologica, 2020, 140, 513-534.	7.7	186
20	Examination of the Effect of Rare Variants in TREM2, ABI3, and PLCG2 in LOAD Through Multiple Phenotypes. Journal of Alzheimer's Disease, 2020, 77, 1469-1482.	2.6	18
21	CCL23: A Chemokine Associated with Progression from Mild Cognitive Impairment to Alzheimerâ€™s Disease. Journal of Alzheimer's Disease, 2020, 73, 1585-1595.	2.6	25
22	A single-nuclei RNA sequencing study of Mendelian and sporadic AD in the human brain. Alzheimer's Research and Therapy, 2019, 11, 71.	6.2	131
23	The MS4A gene cluster is a key modulator of soluble TREM2 and Alzheimerâ€™s disease risk. Science Translational Medicine, 2019, 11, .	12.4	170
24	TREM2 brain transcript-specific studies in AD and TREM2 mutation carriers. Molecular Neurodegeneration, 2019, 14, 18.	10.8	58
25	Polygenic Risk Scores in Neurodegenerative Diseases: a Review. Current Genetic Medicine Reports, 2019, 7, 22-29.	1.9	23
26	Analysis of whole genome-transcriptomic organization in brain to identify genes associated with alcoholism. Translational Psychiatry, 2019, 9, 89.	4.8	66
27	An atlas of cortical circular RNA expression in Alzheimer disease brains demonstrates clinical and pathological associations. Nature Neuroscience, 2019, 22, 1903-1912.	14.8	242
28	Assessment of the Genetic Architecture of Alzheimerâ€™s Disease Risk in Rate of Memory Decline. Journal of Alzheimer's Disease, 2018, 62, 745-756.	2.6	45
29	Polygenic risk score of sporadic lateâ€onset Alzheimer's disease reveals a shared architecture with the familial and earlyâ€onset forms. Alzheimer's and Dementia, 2018, 14, 205-214.	0.8	109
30	P2â€105: NOMINATION OF NOVEL CANDIDATE GENES FOR FAMILIAL LATE ONSET ALZHEIMER DISEASE AFTER EVALUATION OF GENEâ€BASED FAMILYâ€BASED METHODS. Alzheimer's and Dementia, 2018, 14, P709.	0.8	0
31	Integrative system biology analyses of CRISPR-edited iPSC-derived neurons and human brains reveal deficiencies of presynaptic signaling in FTLD and PSP. Translational Psychiatry, 2018, 8, 265.	4.8	47
32	Evaluation of Gene-Based Family-Based Methods to Detect Novel Genes Associated With Familial Late Onset Alzheimer Disease. Frontiers in Neuroscience, 2018, 12, 209.	2.8	21
33	Genetic variants associated with Alzheimerâ€™s disease confer different cerebral cortex cell-type population structure. Genome Medicine, 2018, 10, 43.	8.2	62
34	Genome-wide association study identifies four novel loci associated with Alzheimerâ€™s endophenotypes and disease modifiers. Acta Neuropathologica, 2017, 133, 839-856.	7.7	199
35	CSF protein changes associated with hippocampal sclerosis risk gene variants highlight impact of GRN/PGRN. Experimental Gerontology, 2017, 90, 83-89.	2.8	7
36	A common haplotype lowers PU.1 expression in myeloid cells and delays onset of Alzheimer's disease. Nature Neuroscience, 2017, 20, 1052-1061.	14.8	330

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37	[O1â€“11â€“03]: CEREBROSPINAL FLUID ENDOPHENOTYPES PROVIDE INSIGHT INTO BIOLOGY UNDERLYING ALZHEIMER'S DISEASE. Alzheimer's and Dementia, 2017, 13, P218.	0.8	0
38	[O2â€“13â€“04]: CELLâ€TYPE PROFILING TO IDENTIFY THE TRANSCRIPTOMIC DOWNSTREAM EVENTS TRIGGERED BY EARLYâ€ONSET AUTOSOMAL DOMINANT AD MUTATIONS. Alzheimer's and Dementia, 2017, 13, P589.	0.8	1
39	Parkinson disease polygenic risk score is associated with Parkinson disease status and age at onset but not with alpha-synuclein cerebrospinal fluid levels. BMC Neurology, 2017, 17, 198.	1.8	55
40	Linkage, whole genome sequence, and biological data implicate variants in RAB10 in Alzheimerâ€™s disease resilience. Genome Medicine, 2017, 9, 100.	8.2	67
41	Analysis of neurodegenerative Mendelian genes in clinically diagnosed Alzheimer Disease. PLoS Genetics, 2017, 13, e1007045.	3.5	40
42	Paving the road for the study of epigenetics in neurodegenerative diseases. Acta Neuropathologica, 2016, 132, 483-485.	7.7	6
43	Genome-Wide Association Study of CSF Levels of 59 Alzheimer's Disease Candidate Proteins: Significant Associations with Proteins Involved in Amyloid Processing and Inflammation. PLoS Genetics, 2014, 10, e1004758.	3.5	109
44	Coding variants in TREM2 increase risk for Alzheimer's disease. Human Molecular Genetics, 2014, 23, 5838-5846.	2.9	263
45	Rare coding variants in the phospholipase D3 gene confer risk for Alzheimerâ€™s disease. Nature, 2014, 505, 550-554.	27.8	425
46	Genome-wide survival analysis of age at onset of alcohol dependence in extended high-risk COGA families. Drug and Alcohol Dependence, 2014, 142, 56-62.	3.2	29
47	Phosphorylated Tau-AÎ²42 Ratio as a Continuous Trait for Biomarker Discovery for Early-Stage Alzheimerâ€™s Disease in Multiplex Immunoassay Panels of Cerebrospinal Fluid. Biological Psychiatry, 2014, 75, 723-731.	1.3	72
48	Missense variant in TREML2 protects against Alzheimer's disease. Neurobiology of Aging, 2014, 35, 1510.e19-1510.e26.	3.1	110
49	O4-01-01: Association of genetic variants with cerebrospinal fluid protein levels of ACE, MMP3 and other proteins and risk for Alzheimer's disease. , 2013, 9, P677-P678.		1
50	GWAS of Cerebrospinal Fluid Tau Levels Identifies Risk Variants for Alzheimerâ€™s Disease. Neuron, 2013, 78, 256-268.	8.1	344
51	Pathway Analysis of Smoking Quantity in Multiple GWAS Identifies Cholinergic and Sensory Pathways. PLoS ONE, 2012, 7, e50913.	2.5	11