

Nilgün Ertekin-Taner

List of Publications by Year in descending order

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Version: 2024-02-01

127
papers

11,371
citations

44042

48
h-index

36008

97
g-index

166
all docs

166
docs citations

166
times ranked

14356
citing authors

#	ARTICLE	IF	CITATIONS
1	Common variants at MS4A4/MS4A6E, CD2AP, CD33 and EPHA1 are associated with late-onset Alzheimer's disease. <i>Nature Genetics</i> , 2011, 43, 436-441.	9.4	1,676
2	Alzheimer's disease: early alterations in brain DNA methylation at ANK1, BIN1, RHBDF2 and other loci. <i>Nature Neuroscience</i> , 2014, 17, 1156-1163.	7.1	800
3	Rare coding variants in PLCC2, ABI3, and TREM2 implicate microglial-mediated innate immunity in Alzheimer's disease. <i>Nature Genetics</i> , 2017, 49, 1373-1384.	9.4	783
4	Large-scale proteomic analysis of Alzheimer's disease brain and cerebrospinal fluid reveals early changes in energy metabolism associated with microglia and astrocyte activation. <i>Nature Medicine</i> , 2020, 26, 769-780.	15.2	547
5	Human whole genome genotype and transcriptome data for Alzheimer's and other neurodegenerative diseases. <i>Scientific Data</i> , 2016, 3, 160089.	2.4	361
6	Variants in the ATP-Binding Cassette Transporter (ABCA7), Apolipoprotein E ϵ 4, and the Risk of Late-Onset Alzheimer Disease in African Americans. <i>JAMA - Journal of the American Medical Association</i> , 2013, 309, 1483.	3.8	360
7	TREM2 in neurodegeneration: evidence for association of the p.R47H variant with frontotemporal dementia and Parkinson's disease. <i>Molecular Neurodegeneration</i> , 2013, 8, 19.	4.4	323
8	Genetic variation in PCDH11X is associated with susceptibility to late-onset Alzheimer's disease. <i>Nature Genetics</i> , 2009, 41, 192-198.	9.4	279
9	A novel Alzheimer disease locus located near the gene encoding tau protein. <i>Molecular Psychiatry</i> , 2016, 21, 108-117.	4.1	260
10	Brain Expression Genome-Wide Association Study (eGWAS) Identifies Human Disease-Associated Variants. <i>PLoS Genetics</i> , 2012, 8, e1002707.	1.5	225
11	Genetics of Alzheimer's Disease: A Centennial Review. <i>Neurologic Clinics</i> , 2007, 25, 611-667.	0.8	206
12	Meta-Analysis of the Alzheimer's Disease Human Brain Transcriptome and Functional Dissection in Mouse Models. <i>Cell Reports</i> , 2020, 32, 107908.	2.9	199
13	Assessment of the genetic variance of late-onset Alzheimer's disease. <i>Neurobiology of Aging</i> , 2016, 41, 200.e13-200.e20.	1.5	174
14	APOE4 exacerbates synapse loss and neurodegeneration in Alzheimer's disease patient iPSC-derived cerebral organoids. <i>Nature Communications</i> , 2020, 11, 5540.	5.8	172
15	Genome-wide association study of corticobasal degeneration identifies risk variants shared with progressive supranuclear palsy. <i>Nature Communications</i> , 2015, 6, 7247.	5.8	170
16	Effects of Multiple Genetic Loci on Age at Onset in Late-Onset Alzheimer Disease. <i>JAMA Neurology</i> , 2014, 71, 1394.	4.5	166
17	Transethnic genome-wide scan identifies novel Alzheimer's disease loci. <i>Alzheimer's and Dementia</i> , 2017, 13, 727-738.	0.4	166
18	Alzheimer's Risk Factors Age, APOE Genotype, and Sex Drive Distinct Molecular Pathways. <i>Neuron</i> , 2020, 106, 727-742.e6.	3.8	152

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19	Novel late-onset Alzheimer disease loci variants associate with brain gene expression. <i>Neurology</i> , 2012, 79, 221-228.	1.5	144
20	Novel Alzheimer Disease Risk Loci and Pathways in African American Individuals Using the African Genome Resources Panel. <i>JAMA Neurology</i> , 2021, 78, 102.	4.5	144
21	TREM2 is associated with increased risk for Alzheimer's disease in African Americans. <i>Molecular Neurodegeneration</i> , 2015, 10, 19.	4.4	130
22	<i>APOE</i> ϵ 4 is associated with severity of Lewy body pathology independent of Alzheimer pathology. <i>Neurology</i> , 2018, 91, e1182-e1195.	1.5	122
23	Systematic analysis of dark and camouflaged genes reveals disease-relevant genes hiding in plain sight. <i>Genome Biology</i> , 2019, 20, 97.	3.8	122
24	Conserved brain myelination networks are altered in Alzheimer's and other neurodegenerative diseases. <i>Alzheimer's and Dementia</i> , 2018, 14, 352-366.	0.4	116
25	Association of MAPT haplotypes with Alzheimer's disease risk and MAPT brain gene expression levels. <i>Alzheimer's Research and Therapy</i> , 2014, 6, 39.	3.0	106
26	Potential genetic modifiers of disease risk and age at onset in patients with frontotemporal lobar degeneration and GRN mutations: a genome-wide association study. <i>Lancet Neurology</i> , The, 2018, 17, 548-558.	4.9	97
27	Two rare <i>AKAP9</i> variants are associated with Alzheimer's disease in African Americans. <i>Alzheimer's and Dementia</i> , 2014, 10, 609.	0.4	94
28	Genetic risk factors for the posterior cortical atrophy variant of Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2016, 12, 862-871.	0.4	93
29	Genetic variants in a haplotype block spanning <i>IDE</i> are significantly associated with plasma A β 42 levels and risk for Alzheimer disease. <i>Human Mutation</i> , 2004, 23, 334-342.	1.1	91
30	Late-onset Alzheimer's risk variants in memory decline, incident mild cognitive impairment, and Alzheimer's disease. <i>Neurobiology of Aging</i> , 2015, 36, 60-67.	1.5	90
31	Genome-wide analyses as part of the international FTLD-TDP whole-genome sequencing consortium reveals novel disease risk factors and increases support for immune dysfunction in FTLD. <i>Acta Neuropathologica</i> , 2019, 137, 879-899.	3.9	90
32	Two novel loci, <i>COBL</i> and <i>SLC10A2</i> , for Alzheimer's disease in African Americans. <i>Alzheimer's and Dementia</i> , 2017, 13, 119-129.	0.4	87
33	A nonsynonymous mutation in <i>PLCG2</i> reduces the risk of Alzheimer's disease, dementia with Lewy bodies and frontotemporal dementia, and increases the likelihood of longevity. <i>Acta Neuropathologica</i> , 2019, 138, 237-250.	3.9	87
34	Genetics of Alzheimer disease in the pre- and post-GWAS era. <i>Alzheimer's Research and Therapy</i> , 2010, 2, 3.	3.0	85
35	Imaging correlations of tau, amyloid, metabolism, and atrophy in typical and atypical Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2018, 14, 1005-1014.	0.4	80
36	Glutathione S-transferase omega genes in Alzheimer and Parkinson disease risk, age-at-diagnosis and brain gene expression: an association study with mechanistic implications. <i>Molecular Neurodegeneration</i> , 2012, 7, 13.	4.4	75

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37	ABI3 and PLCG2 missense variants as risk factors for neurodegenerative diseases in Caucasians and African Americans. <i>Molecular Neurodegeneration</i> , 2018, 13, 53.	4.4	75
38	Large eQTL meta-analysis reveals differing patterns between cerebral cortical and cerebellar brain regions. <i>Scientific Data</i> , 2020, 7, 340.	2.4	75
39	Fine mapping of the \hat{A} -T catenin gene to a quantitative trait locus on chromosome 10 in late-onset Alzheimer's disease pedigrees. <i>Human Molecular Genetics</i> , 2003, 12, 3133-3143.	1.4	72
40	TYROBP genetic variants in early-onset Alzheimer's disease. <i>Neurobiology of Aging</i> , 2016, 48, 222.e9-222.e15.	1.5	69
41	Sex and age interact to determine clinicopathologic differences in Alzheimer's disease. <i>Acta Neuropathologica</i> , 2018, 136, 873-885.	3.9	69
42	Linkage, whole genome sequence, and biological data implicate variants in RAB10 in Alzheimer's disease resilience. <i>Genome Medicine</i> , 2017, 9, 100.	3.6	67
43	[¹⁸ F]AV-1451 clustering of entorhinal and cortical uptake in Alzheimer's disease. <i>Annals of Neurology</i> , 2018, 83, 248-257.	2.8	67
44	Elevated amyloid β protein ($A\beta_{42}$) and late onset Alzheimer's disease are associated with single nucleotide polymorphisms in the urokinase-type plasminogen activator gene. <i>Human Molecular Genetics</i> , 2005, 14, 447-460.	1.4	64
45	Late-onset Alzheimer disease risk variants mark brain regulatory loci. <i>Neurology: Genetics</i> , 2015, 1, e15.	0.9	64
46	Genome-wide association interaction analysis for Alzheimer's disease. <i>Neurobiology of Aging</i> , 2014, 35, 2436-2443.	1.5	61
47	Expression and processing analyses of wild type and p.R47H TREM2 variant in Alzheimer's disease brains. <i>Molecular Neurodegeneration</i> , 2016, 11, 72.	4.4	55
48	Late-onset Alzheimer disease genetic variants in posterior cortical atrophy and posterior AD. <i>Neurology</i> , 2014, 82, 1455-1462.	1.5	51
49	TLR5 decoy receptor as a novel anti-amyloid therapeutic for Alzheimer's disease. <i>Journal of Experimental Medicine</i> , 2018, 215, 2247-2264.	4.2	50
50	Selective Vulnerability of the Nucleus Basalis of Meynert Among Neuropathologic Subtypes of Alzheimer Disease. <i>JAMA Neurology</i> , 2020, 77, 225.	4.5	50
51	Gene expression, methylation and neuropathology correlations at progressive supranuclear palsy risk loci. <i>Acta Neuropathologica</i> , 2016, 132, 197-211.	3.9	49
52	Heritability of plasma amyloid β in typical late-onset Alzheimer's disease pedigrees. <i>Genetic Epidemiology</i> , 2001, 21, 19-30.	0.6	48
53	Concordant Association of Insulin Degrading Enzyme Gene (IDE) Variants with IDE mRNA, $A\beta$, and Alzheimer's Disease. <i>PLoS ONE</i> , 2010, 5, e8764.	1.1	48
54	A candidate regulatory variant at the <i>TREM</i> gene cluster associates with decreased Alzheimer's disease risk and increased <i>TREML1</i> and <i>TREM2</i> brain gene expression. <i>Alzheimer's and Dementia</i> , 2017, 13, 663-673.	0.4	48

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55	Divergent brain gene expression patterns associate with distinct cell-specific tau neuropathology traits in progressive supranuclear palsy. <i>Acta Neuropathologica</i> , 2018, 136, 709-727.	3.9	47
56	Transcriptomic analysis to identify genes associated with selective hippocampal vulnerability in Alzheimer's disease. <i>Nature Communications</i> , 2021, 12, 2311.	5.8	44
57	Identifying drug targets for neurological and psychiatric disease via genetics and the brain transcriptome. <i>PLoS Genetics</i> , 2021, 17, e1009224.	1.5	43
58	Global and local ancestry in African-Americans: Implications for Alzheimer's disease risk. <i>Alzheimer's and Dementia</i> , 2016, 12, 233-243.	0.4	42
59	Deciphering cellular transcriptional alterations in Alzheimer's disease brains. <i>Molecular Neurodegeneration</i> , 2020, 15, 38.	4.4	42
60	Evaluation of memory endophenotypes for association with <i>CLU</i> , <i>CR1</i> , and <i>PICALM</i> variants in black and white subjects. , 2014, 10, 205-213.		40
61	Systems biology approach to late-onset Alzheimer's disease genome-wide association study identifies novel candidate genes validated using brain expression data and <i>Caenorhabditis elegans</i> experiments. , 2017, 13, 1133-1142.		40
62	Association of Long Runs of Homozygosity With Alzheimer Disease Among African American Individuals. <i>JAMA Neurology</i> , 2015, 72, 1313.	4.5	39
63	Tau and apolipoprotein E modulate cerebrovascular tight junction integrity independent of cerebral amyloid angiopathy in Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2020, 16, 1372-1383.	0.4	34
64	Mitophagy alterations in Alzheimer's disease are associated with granulovacuolar degeneration and early tau pathology. <i>Alzheimer's and Dementia</i> , 2021, 17, 417-430.	0.4	34
65	<i>MAPT</i> haplotype H1G is associated with increased risk of dementia with Lewy bodies. <i>Alzheimer's and Dementia</i> , 2016, 12, 1297-1304.	0.4	32
66	Gene expression endophenotypes: a novel approach for gene discovery in Alzheimer's disease. <i>Molecular Neurodegeneration</i> , 2011, 6, 31.	4.4	31
67	Role for the microtubule-associated protein tau variant p.A152T in risk of α -synucleinopathies. <i>Neurology</i> , 2015, 85, 1680-1686.	1.5	31
68	<i>LRRK2</i> variation and dementia with Lewy bodies. <i>Parkinsonism and Related Disorders</i> , 2016, 31, 98-103.	1.1	30
69	<i>TMEM106B</i> haplotypes have distinct gene expression patterns in aged brain. <i>Molecular Neurodegeneration</i> , 2018, 13, 35.	4.4	30
70	Ethnoracial differences in Alzheimer's disease from the Florida Autopsied Multi-Ethnic (FLAME) cohort. <i>Alzheimer's and Dementia</i> , 2019, 15, 635-643.	0.4	29
71	Genetics of Gene Expression in the Aging Human Brain Reveal TDP-43 Proteinopathy Pathophysiology. <i>Neuron</i> , 2020, 107, 496-508.e6.	3.8	29
72	Atlas of Transcription Factor Binding Sites from ENCODE DNase Hypersensitivity Data across 27 Tissue Types. <i>Cell Reports</i> , 2020, 32, 108029.	2.9	28

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73	Genome-wide pleiotropy analysis of neuropathological traits related to Alzheimer's disease. <i>Alzheimer's Research and Therapy</i> , 2018, 10, 22.	3.0	27
74	<i>ABCA7</i> loss-of-function variants, expression, and neurologic disease risk. <i>Neurology: Genetics</i> , 2017, 3, e126.	0.9	26
75	Molecular estimation of neurodegeneration pseudotime in older brains. <i>Nature Communications</i> , 2020, 11, 5781.	5.8	26
76	African American exome sequencing identifies potential risk variants at Alzheimer disease loci. <i>Neurology: Genetics</i> , 2017, 3, e141.	0.9	25
77	Integrative functional genomic analysis of intron retention in human and mouse brain with Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2021, 17, 984-1004.	0.4	25
78	Apolipoprotein E regulates lipid metabolism and β -synuclein pathology in human iPSC-derived cerebral organoids. <i>Acta Neuropathologica</i> , 2021, 142, 807-825.	3.9	25
79	Male-specific epistasis between <i>WWC1</i> and <i>TLN2</i> genes is associated with Alzheimer's disease. <i>Neurobiology of Aging</i> , 2018, 72, 188.e3-188.e12.	1.5	24
80	Evaluating pathogenic dementia variants in posterior cortical atrophy. <i>Neurobiology of Aging</i> , 2016, 37, 38-44.	1.5	23
81	Genome-wide transcriptome analysis identifies novel dysregulated genes implicated in Alzheimer's pathology. <i>Alzheimer's and Dementia</i> , 2020, 16, 1213-1223.	0.4	23
82	Transcriptional landscape of human microglia implicates age, sex, and <i>APOE</i> -related immunometabolic pathway perturbations. <i>Aging Cell</i> , 2022, 21, e13606.	3.0	23
83	Risk factors for severe COVID-19 differ by age for hospitalized adults. <i>Scientific Reports</i> , 2022, 12, 6568.	1.6	23
84	Genetic architecture of resilience of executive functioning. <i>Brain Imaging and Behavior</i> , 2012, 6, 621-633.	1.1	22
85	<i>RAB39B</i> gene mutations are not a common cause of Parkinson's disease or dementia with Lewy bodies. <i>Neurobiology of Aging</i> , 2016, 45, 107-108.	1.5	21
86	<i>MAPT</i> haplotype-stratified GWAS reveals differential association for AD risk variants. <i>Alzheimer's and Dementia</i> , 2020, 16, 983-1002.	0.4	21
87	Atlas of RNA editing events affecting protein expression in aged and Alzheimer's disease human brain tissue. <i>Nature Communications</i> , 2021, 12, 7035.	5.8	19
88	Association and heterogeneity at the <i>GAPDH</i> locus in Alzheimer's disease. <i>Neurobiology of Aging</i> , 2012, 33, 203.e25-203.e33.	1.5	17
89	MRI and flortaucipir relationships in Alzheimer's phenotypes are heterogeneous. <i>Annals of Clinical and Translational Neurology</i> , 2020, 7, 707-721.	1.7	17
90	Microglia show differential transcriptomic response to $A\beta$ peptide aggregates ex vivo and in vivo. <i>Life Science Alliance</i> , 2021, 4, e202101108.	1.3	17

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91	TREM2 p.R47H substitution is not associated with dementia with Lewy bodies. <i>Neurology: Genetics</i> , 2016, 2, e85.	0.9	16
92	Utility of Plasma Neurofilament Light in the 1Florida Alzheimerâ€™s Disease Research Center (ADRC). <i>Journal of Alzheimer's Disease</i> , 2021, 79, 59-70.	1.2	16
93	Prominent amyloid plaque pathology and cerebral amyloid angiopathy in APP V717I (London) carrier â€™ phenotypic variability in autosomal dominant Alzheimerâ€™s disease. <i>Acta Neuropathologica Communications</i> , 2020, 8, 31.	2.4	14
94	Alternative Approaches in Gene Discovery and Characterization in Alzheimerâ€™s Disease. <i>Current Genetic Medicine Reports</i> , 2013, 1, 39-51.	1.9	13
95	Alzheimerâ€™s disease and progressive supranuclear palsy share similar transcriptomic changes in distinct brain regions. <i>Journal of Clinical Investigation</i> , 2022, 132, .	3.9	13
96	Whole genome sequencingâ€™based copy number variations reveal novel pathways and targets in Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2022, 18, 1846-1867.	0.4	13
97	LRRTM3 Interacts with APP and BACE1 and Has Variants Associating with Late-Onset Alzheimerâ€™s Disease (LOAD). <i>PLoS ONE</i> , 2013, 8, e64164.	1.1	12
98	An alternative transcript of the Alzheimer's disease risk gene SORL1 encodes a truncated receptor. <i>Neurobiology of Aging</i> , 2018, 71, 266.e11-266.e24.	1.5	12
99	Comparative evaluation for the globin gene depletion methods for mRNA sequencing using the whole blood-derived total RNAs. <i>BMC Genomics</i> , 2020, 21, 890.	1.2	12
100	Manifestations of Alzheimerâ€™s disease genetic risk in the blood are evident in a multiomic analysis in healthy adults aged 18 to 90. <i>Scientific Reports</i> , 2022, 12, 6117.	1.6	12
101	Investigating Heterogeneity and Neuroanatomic Correlates of Longitudinal Clinical Decline in Atypical Alzheimer Disease. <i>Neurology</i> , 2022, 98, .	1.5	12
102	Plasma Biomarkers of Alzheimerâ€™s Disease in African Americans. <i>Journal of Alzheimer's Disease</i> , 2021, 79, 323-334.	1.2	11
103	Relationship of APOE, age at onset, amyloid and clinical phenotype in Alzheimer disease. <i>Neurobiology of Aging</i> , 2021, 108, 90-98.	1.5	11
104	Identification of missing variants by combining multiple analytic pipelines. <i>BMC Bioinformatics</i> , 2018, 19, 139.	1.2	10
105	The influence of Î²-amyloid on [¹⁸ F]AV-1451 in semantic variant of primary progressive aphasia. <i>Neurology</i> , 2019, 92, e710-e722.	1.5	10
106	Genome-wide analysis identifies a novel LINC-PINT splice variant associated with vascular amyloid pathology in Alzheimerâ€™s disease. <i>Acta Neuropathologica Communications</i> , 2021, 9, 93.	2.4	9
107	Epigenomic features related to microglia are associated with attenuated effect of <i>APOE</i> Î¼4 on Alzheimer's disease risk in humans. <i>Alzheimer's and Dementia</i> , 2022, 18, 688-699.	0.4	9
108	Association of ABI3 and PLCG2 missense variants with disease risk and neuropathology in Lewy body disease and progressive supranuclear palsy. <i>Acta Neuropathologica Communications</i> , 2020, 8, 172.	2.4	8

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109	Longitudinal Amyloid- β^2 PET in Atypical Alzheimer's Disease and Frontotemporal Lobar Degeneration. <i>Journal of Alzheimer's Disease</i> , 2020, 74, 377-389.	1.2	7
110	Cilostazol Versus Aspirin for Secondary Stroke Prevention: Systematic Review and Meta-Analysis. <i>Journal of Stroke and Cerebrovascular Diseases</i> , 2021, 30, 105581.	0.7	7
111	Expression of an alternatively spliced variant of SORL1 in neuronal dendrites is decreased in patients with Alzheimer's disease. <i>Acta Neuropathologica Communications</i> , 2021, 9, 43.	2.4	7
112	Analysis of intraoperative human brain tissue transcriptome reveals putative risk genes and altered molecular pathways in glioma-related seizures. <i>Epilepsy Research</i> , 2021, 173, 106618.	0.8	7
113	Evaluation of Associations of Alzheimer's Disease Risk Variants that Are Highly Expressed in Microglia with Neuropathological Outcome Measures. <i>Journal of Alzheimer's Disease</i> , 2019, 70, 659-666.	1.2	6
114	Effects of sex and APOE on Parkinson's Disease-related cognitive decline. <i>Neurologia I Neurochirurgia Polska</i> , 2021, 55, 559-566.	0.6	6
115	The quest for Alzheimer disease genes' focus on CSF tau. <i>Nature Reviews Neurology</i> , 2013, 9, 368-370.	4.9	5
116	Latent trait modeling of tau neuropathology in progressive supranuclear palsy. <i>Acta Neuropathologica</i> , 2021, 141, 667-680.	3.9	5
117	Impact of variant-level batch effects on identification of genetic risk factors in large sequencing studies. <i>PLoS ONE</i> , 2021, 16, e0249305.	1.1	5
118	Comprehensive Screening for Disease Risk Variants in Early-Onset Alzheimer's Disease Genes in African Americans Identifies Novel PSEN Variants. <i>Journal of Alzheimer's Disease</i> , 2017, 56, 1215-1222.	1.2	4
119	Modulating innate immune activation states impacts the efficacy of specific $\text{A}\beta^2$ immunotherapy. <i>Molecular Neurodegeneration</i> , 2021, 16, 32.	4.4	4
120	Clinical, Imaging, and Pathologic Characteristics of Patients With Right vs Left Hemisphere-Predominant Logopenic Progressive Aphasia. <i>Neurology</i> , 2021, 97, e523-e534.	1.5	4
121	Clinical Deep Phenotyping of <i>ABCA7</i> Mutation Carriers. <i>Neurology: Genetics</i> , 2022, 8, e655.	0.9	4
122	Epigenomic features related to microglia are associated with attenuated effect of APOE ϵ^4 on Alzheimer's disease risk in humans. <i>Alzheimer's and Dementia</i> , 2020, 16, e043533.	0.4	2
123	Transcript levels in plasma contribute substantial predictive value as potential Alzheimer's disease biomarkers in African Americans. <i>EBioMedicine</i> , 2022, , 103929.	2.7	2
124	Target-enriched sequencing of chromosome 17q21.31 in sporadic tauopathies reveals no candidate variants. <i>Neurobiology of Aging</i> , 2018, 66, 177.e7-177.e10.	1.5	1
125	Other Genes Implicated in Alzheimer's Disease. , 2013, , 209-230.		0
126	O3-04-01: NEXT-GENERATION RNA SEQUENCING IN ALZHEIMER'S DISEASE AND PROGRESSIVE SUPRANUCLEAR PALSY. , 2014, 10, P214-P215.		0

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127	S4â€²â€²04: Accelerating Medicines Partnership: Identifying Therapeutic Targets for Alzheimer's Disease with Comparative Transcriptomics. <i>Alzheimer's and Dementia</i> , 2016, 12, P322.	0.4	0