

Sungeun Kim

List of Publications by Year in descending order

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Version: 2024-02-01

130
papers

7,187
citations

87888

38
h-index

66911

78
g-index

167
all docs

167
docs citations

167
times ranked

11026
citing authors

#	ARTICLE	IF	CITATIONS
1	Common genetic variants influence human subcortical brain structures. <i>Nature</i> , 2015, 520, 224-229.	27.8	772
2	The ENIGMA Consortium: large-scale collaborative analyses of neuroimaging and genetic data. <i>Brain Imaging and Behavior</i> , 2014, 8, 153-182.	2.1	696
3	Identification of common variants associated with human hippocampal and intracranial volumes. <i>Nature Genetics</i> , 2012, 44, 552-561.	21.4	594
4	Alzheimer's Disease Neuroimaging Initiative biomarkers as quantitative phenotypes: Genetics core aims, progress, and plans. <i>Alzheimer's and Dementia</i> , 2010, 6, 265-273.	0.8	378
5	Metabolic network failures in Alzheimer's disease: A biochemical road map. <i>Alzheimer's and Dementia</i> , 2017, 13, 965-984.	0.8	362
6	Whole genome association study of brain-wide imaging phenotypes for identifying quantitative trait loci in MCI and AD: A study of the ADNI cohort. <i>NeuroImage</i> , 2010, 53, 1051-1063.	4.2	340
7	Novel genetic loci associated with hippocampal volume. <i>Nature Communications</i> , 2017, 8, 13624.	12.8	250
8	Genetic studies of quantitative MCI and AD phenotypes in ADNI: Progress, opportunities, and plans. <i>Alzheimer's and Dementia</i> , 2015, 11, 792-814.	0.8	241
9	Longitudinal MRI atrophy biomarkers: Relationship to conversion in the ADNI cohort. <i>Neurobiology of Aging</i> , 2010, 31, 1401-1418.	3.1	230
10	Novel genetic loci underlying human intracranial volume identified through genome-wide association. <i>Nature Neuroscience</i> , 2016, 19, 1569-1582.	14.8	213
11	Genome-wide association study identifies four novel loci associated with Alzheimer's endophenotypes and disease modifiers. <i>Acta Neuropathologica</i> , 2017, 133, 839-856.	7.7	199
12	Genetic architecture of subcortical brain structures in 38,851 individuals. <i>Nature Genetics</i> , 2019, 51, 1624-1636.	21.4	192
13	Genetic analysis of quantitative phenotypes in AD and MCI: imaging, cognition and biomarkers. <i>Brain Imaging and Behavior</i> , 2014, 8, 183-207.	2.1	161
14	APOE effect on Alzheimer's disease biomarkers in older adults with significant memory concern. <i>Alzheimer's and Dementia</i> , 2015, 11, 1417-1429.	0.8	157
15	Identifying quantitative trait loci via group-sparse multitask regression and feature selection: an imaging genetics study of the ADNI cohort. <i>Bioinformatics</i> , 2012, 28, 229-237.	4.1	149
16	The role of apolipoprotein E (APOE) genotype in early mild cognitive impairment (E-MCI). <i>Frontiers in Aging Neuroscience</i> , 2013, 5, 11.	3.4	126
17	CWAS of longitudinal amyloid accumulation on ¹⁸ F-florbetapir PET in Alzheimer's disease implicates microglial activation gene IL1RAP. <i>Brain</i> , 2015, 138, 3076-3088.	7.6	117
18	Voxelwise gene-wide association study (vGeneWAS): Multivariate gene-based association testing in 731 elderly subjects. <i>NeuroImage</i> , 2011, 56, 1875-1891.	4.2	116

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19	Structured sparse canonical correlation analysis for brain imaging genetics: an improved GraphNet method. <i>Bioinformatics</i> , 2016, 32, 1544-1551.	4.1	96
20	Comparison of Manual and Automated Determination of Hippocampal Volumes in MCI and Early AD. <i>Brain Imaging and Behavior</i> , 2010, 4, 86-95.	2.1	74
21	Targeted neurogenesis pathway-based gene analysis identifies ADORA2A associated with hippocampal volume in mild cognitive impairment and Alzheimer's disease. <i>Neurobiology of Aging</i> , 2017, 60, 92-103.	3.1	70
22	Voxel and surface-based topography of memory and executive deficits in mild cognitive impairment and Alzheimer's disease. <i>Brain Imaging and Behavior</i> , 2012, 6, 551-567.	2.1	66
23	Analysis of Copy Number Variation in Alzheimer's Disease: The NIALOAD/ NCRAD Family Study. <i>Current Alzheimer Research</i> , 2012, 9, 801-814.	1.4	64
24	From phenotype to genotype: an association study of longitudinal phenotypic markers to Alzheimer's disease relevant SNPs. <i>Bioinformatics</i> , 2012, 28, i619-i625.	4.1	62
25	Multiple loci influencing hippocampal degeneration identified by genome scan. <i>Annals of Neurology</i> , 2012, 72, 65-75.	5.3	59
26	Genome-wide pathway analysis of memory impairment in the Alzheimer's Disease Neuroimaging Initiative (ADNI) cohort implicates gene candidates, canonical pathways, and networks. <i>Brain Imaging and Behavior</i> , 2012, 6, 634-648.	2.1	58
27	Transcriptome-guided amyloid imaging genetic analysis via a novel structured sparse learning algorithm. <i>Bioinformatics</i> , 2014, 30, i564-i571.	4.1	57
28	Harnessing peripheral DNA methylation differences in the Alzheimer's Disease Neuroimaging Initiative (ADNI) to reveal novel biomarkers of disease. <i>Clinical Epigenetics</i> , 2020, 12, 84.	4.1	57
29	Influence of <i>TSP0</i> Genotype on ¹¹ C-PBR28 Standardized Uptake Values. <i>Journal of Nuclear Medicine</i> , 2013, 54, 1320-1322.	5.0	56
30	Plasma Tau Association with Brain Atrophy in Mild Cognitive Impairment and Alzheimer's Disease. <i>Journal of Alzheimer's Disease</i> , 2017, 58, 1245-1254.	2.6	54
31	Identifying Neuroimaging and Proteomic Biomarkers for MCI and AD via the Elastic Net. <i>Lecture Notes in Computer Science</i> , 2011, 7012, 27-34.	1.3	53
32	Genomic Copy Number Analysis in Alzheimer's Disease and Mild Cognitive Impairment: An ADNI Study. <i>International Journal of Alzheimer's Disease</i> , 2011, 2011, 1-10.	2.0	51
33	Influence of Genetic Variation on Plasma Protein Levels in Older Adults Using a Multi-Analyte Panel. <i>PLoS ONE</i> , 2013, 8, e70269.	2.5	50
34	Analysis of Copy Number Variation in Alzheimer's Disease in a Cohort of Clinically Characterized and Neuropathologically Verified Individuals. <i>PLoS ONE</i> , 2012, 7, e50640.	2.5	49
35	Targeted metabolomics and medication classification data from participants in the ADNI1 cohort. <i>Scientific Data</i> , 2017, 4, 170140.	5.3	49
36	Protective variant for hippocampal atrophy identified by whole exome sequencing. <i>Annals of Neurology</i> , 2015, 77, 547-552.	5.3	48

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37	Relationship between baseline brain metabolism measured using [18F]FDG PET and memory and executive function in prodromal and early Alzheimer's disease. <i>Brain Imaging and Behavior</i> , 2012, 6, 568-583.	2.1	47
38	Amyloid pathway-based candidate gene analysis of [11C]PiB-PET in the Alzheimer's Disease Neuroimaging Initiative (ADNI) cohort. <i>Brain Imaging and Behavior</i> , 2012, 6, 1-15.	2.1	47
39	Association of plasma and cortical amyloid beta is modulated by APOE ϵ 4 status. <i>Alzheimer's and Dementia</i> , 2014, 10, e9-e18.	0.8	43
40	Cortical surface biomarkers for predicting cognitive outcomes using group l2,1 norm. <i>Neurobiology of Aging</i> , 2015, 36, S185-S193.	3.1	43
41	A deep learning framework identifies dimensional representations of Alzheimer's Disease from brain structure. <i>Nature Communications</i> , 2021, 12, 7065.	12.8	38
42	A Novel Structure-Aware Sparse Learning Algorithm for Brain Imaging Genetics. <i>Lecture Notes in Computer Science</i> , 2014, 17, 329-336.	1.3	36
43	Genome-wide association and interaction studies of CSF T-tau/A β 242 ratio in ADNI cohort. <i>Neurobiology of Aging</i> , 2017, 57, 247.e1-247.e8.	3.1	34
44	Gene-based GWAS and biological pathway analysis of the resilience of executive functioning. <i>Brain Imaging and Behavior</i> , 2014, 8, 110-118.	2.1	33
45	Comprehensive Gene- and Pathway-Based Analysis of Depressive Symptoms in Older Adults. <i>Journal of Alzheimer's Disease</i> , 2015, 45, 1197-1206.	2.6	33
46	Network-based analysis of genetic variants associated with hippocampal volume in Alzheimer's disease: a study of ADNI cohorts. <i>BioData Mining</i> , 2016, 9, 3.	4.0	28
47	Association analysis of rare variants near the APOE region with CSF and neuroimaging biomarkers of Alzheimer's disease. <i>BMC Medical Genomics</i> , 2017, 10, 29.	1.5	28
48	Genetic Interactions Explain Variance in Cingulate Amyloid Burden: An AV-45 PET Genome-Wide Association and Interaction Study in the ADNI Cohort. <i>BioMed Research International</i> , 2015, 2015, 1-11.	1.9	24
49	Fourier method for large-scale surface modeling and registration. <i>Computers and Graphics</i> , 2009, 33, 299-311.	2.5	23
50	Hippocampal transcriptome-guided genetic analysis of correlated episodic memory phenotypes in Alzheimer's disease. <i>Frontiers in Genetics</i> , 2015, 6, 117.	2.3	23
51	Tissue-specific network-based genome wide study of amygdala imaging phenotypes to identify functional interaction modules. <i>Bioinformatics</i> , 2017, 33, 3250-3257.	4.1	23
52	Genome-wide transcriptome analysis identifies novel dysregulated genes implicated in Alzheimer's pathology. <i>Alzheimer's and Dementia</i> , 2020, 16, 1213-1223.	0.8	23
53	Genetic architecture of resilience of executive functioning. <i>Brain Imaging and Behavior</i> , 2012, 6, 621-633.	2.1	22
54	Sparse Bayesian Learning for Identifying Imaging Biomarkers in AD Prediction. <i>Lecture Notes in Computer Science</i> , 2010, 13, 611-618.	1.3	21

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55	Integration of bioinformatics and imaging informatics for identifying rare PSEN1 variants in Alzheimer's disease. BMC Medical Genomics, 2016, 9, 30.	1.5	20
56	Hippocampal Surface Mapping of Genetic Risk Factors in AD via Sparse Learning Models. Lecture Notes in Computer Science, 2011, 14, 376-383.	1.3	20
57	Genome-wide association study of language performance in Alzheimer's disease. Brain and Language, 2017, 172, 22-29.	1.6	20
58	Telomere Shortening in the Alzheimer's Disease Neuroimaging Initiative Cohort. Journal of Alzheimer's Disease, 2019, 71, 33-43.	2.6	14
59	GN-SCCA: GraphNet Based Sparse Canonical Correlation Analysis for Brain Imaging Genetics. Lecture Notes in Computer Science, 2015, 9250, 275-284.	1.3	14
60	Genome-wide network-based pathway analysis of CSF t-tau/Á ² 1-42 ratio in the ADNI cohort. BMC Genomics, 2017, 18, 421.	2.8	13
61	Two-dimensional enrichment analysis for mining high-level imaging genetic associations. Brain Informatics, 2017, 4, 27-37.	3.0	13
62	Genetic Influences on Plasma Homocysteine Levels in African Americans and Yoruba Nigerians. Journal of Alzheimer's Disease, 2016, 49, 991-1003.	2.6	12
63	A telescope GWAS analysis strategy, based on SNPs-genes-pathways ensemble and on multivariate algorithms, to characterize late onset Alzheimer's disease. Scientific Reports, 2020, 10, 12063.	3.3	11
64	A Graph-Based Integration of Multimodal Brain Imaging Data for the Detection of Early Mild Cognitive Impairment (E-MCI). Lecture Notes in Computer Science, 2013, 8159, 159-169.	1.3	10
65	Structured sparse CCA for brain imaging genetics via graph OSCAR. BMC Systems Biology, 2016, 10, 68.	3.0	9
66	PARP1 Gene Variation and Microglial Activity on [11C]PBR28 PET in Older Adults at Risk for Alzheimer's Disease. Lecture Notes in Computer Science, 2013, 8159, 150-158.	1.3	8
67	Longitudinal Genotype-Phenotype Association Study via Temporal Structure Auto-learning Predictive Model. Lecture Notes in Computer Science, 2017, 10229, 287-302.	1.3	8
68	Automatic Prediction of Conversion from Mild Cognitive Impairment to Probable Alzheimer's Disease using Structural Magnetic Resonance Imaging. AMIA ... Annual Symposium proceedings, 2010, 2010, 542-6.	0.2	8
69	The effect of reference panels and software tools on genotype imputation. AMIA ... Annual Symposium proceedings, 2011, 2011, 1013-8.	0.2	8
70	Glucose metabolism patterns: A potential index to characterize brain ageing and predict high conversion risk into cognitive impairment. GeroScience, 2022, 44, 2319-2336.	4.6	8
71	Genetic Clustering on the Hippocampal Surface for Genome-Wide Association Studies. Lecture Notes in Computer Science, 2013, 16, 690-697.	1.3	7
72	Data synthesis and method evaluation for brain imaging genetics. , 2014, 2014, 1202-1205.		6

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73	Longitudinal Genotype-Phenotype Association Study through Temporal Structure Auto-Learning Predictive Model. <i>Journal of Computational Biology</i> , 2018, 25, 809-824.	1.6	6
74	Multimodal Neuroimaging Predictors for Cognitive Performance Using Structured Sparse Learning. <i>Lecture Notes in Computer Science</i> , 2012, , 1-17.	1.3	4
75	P4-008: Mapre2 as a novel Alzheimer's disease target gene from gwas of CSF amyloid beta 1-42, tau and hyperphosphorylated tau in the ADNI cohort. , 2015, 11, P767-P768.		3
76	[P1-142]: DNA METHYLATION DYNAMICS IN ALZHEIMER'S DISEASE DIAGNOSIS AND PROGRESSION. <i>Alzheimer's and Dementia</i> , 2017, 13, P297.	0.8	3
77	FOURIER METHODS FOR 3D SURFACE MODELING AND ANALYSIS. <i>Series in Computer Vision</i> , 2011, , 175-196.	0.1	2
78	P3-002: GWAS of the joint ADGC data set identifies novel common variants associated with late-onset Alzheimer's disease. , 2013, 9, P550-P550.		2
79	P2-098: Whole Brain Surface-Based Analysis Identified Brain Atrophy Associated with SNPs in <i>FRMD6</i> Linked to Alzheimer's Disease. <i>Alzheimer's and Dementia</i> , 2016, 12, P648.	0.8	2
80	O3-03-01: Genome-wide association study of CSF biomarkers amyloid beta 1-42, tau and tau phosphorylated at threonine 181 in the ADNI cohort. , 2010, 6, S129-S129.		1
81	O3-06-01: Association analysis of candidate SNPs on hippocampal volume and shape in mild cognitive impairment and older adults with cognitive complaints. , 2010, 6, S137-S138.		1
82	Joint identification of imaging and proteomics biomarkers of Alzheimer's disease using network-guided sparse learning. , 2014, 2014, 665-668.		1
83	O3-03-02: TWO-YEAR LONGITUDINAL CHANGE IN AMYLOID DEPOSITION, GLUCOSE METABOLISM, AND HIPPOCAMPAL ATROPHY IN ADNI-2 PARTICIPANTS: RELATION TO GENETIC RISK. , 2014, 10, P211-P212.		1
84	IC-P-042: Influence of rare reelin variants on quantitative PET imaging and CSF phenotypes in late-onset Alzheimer's disease. , 2015, 11, P36-P36.		1
85	P1-201: Genetic findings using ADNI multimodal quantitative phenotypes: A 2014 update. , 2015, 11, P426-P426.		1
86	O3-13-04: Genome-wide rare variant analysis identifies candidate genes significantly associated with composite scores for memory. , 2015, 11, P251-P252.		1
87	Two-Dimensional Enrichment Analysis for Mining High-Level Imaging Genetic Associations. <i>Lecture Notes in Computer Science</i> , 2015, 9250, 115-124.	1.3	1
88	[IC-P-063]: <i>KLK8</i> AS A MODULATOR OF ALZHEIMER'S DISEASE PATHOLOGY: NEUROIMAGING GENETICS. <i>Alzheimer's and Dementia</i> , 2017, 13, P51.	0.8	1
89	IC-O1-03: Hippocampal transcriptome-guided gene-gene interaction of memory phenotype in MCI and Alzheimer's disease. , 2013, 9, P4-P4.		0
90	P3-018: INFLUENCE OF RARE PSEN1 VARIANTS ON QUANTITATIVE STRUCTURAL IMAGING AND CSF PHENOTYPES IN LATE ONSET ALZHEIMER'S DISEASE. , 2014, 10, P633-P633.		0

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91	IC-P-172: GENOME-WIDE PROTEIN INTERACTION GUIDED EPISTATIC ANALYSIS ON MEMORY PERFORMANCE: AN ADNI STUDY. , 2014, 10, P95-P96.		0
92	IC-P-173: EFFECTS OF NEWLY IDENTIFIED TOP AD CANDIDATE GENES ON MEMORY PERFORMANCE: SNP, GENE, AND EPISTASIS ANALYSES IN ADNI. , 2014, 10, P96-P97.		0
93	P1-230: EFFECTS OF NEWLY IDENTIFIED TOP AD CANDIDATE GENES ON MEMORY PERFORMANCE: SNP, GENE, AND EPISTASIS ANALYSES IN ADNI. , 2014, 10, P388-P388.		0
94	IC-P-095: TWO-YEAR LONGITUDINAL CHANGE IN AMYLOID DEPOSITION, GLUCOSE METABOLISM, AND HIPPOCAMPAL ATROPHY IN ADNI-2 PARTICIPANTS: RELATION TO GENETIC RISK. , 2014, 10, P53-P54.		0
95	IC-P-174: RARE VARIANT IN PLD3 IS ASSOCIATED WITH ALZHEIMER'S PATTERN OF NEURODEGENERATIVE CHANGES. , 2014, 10, P97-P97.		0
96	P3-024: NEXT-GENERATION SEQUENCING OF THE BCHE LOCUS IDENTIFIES A FUNCTIONAL SNP ASSOCIATED WITH ALZHEIMER'S DISEASE BIOMARKERS AND AGE OF ONSET. , 2014, 10, P636-P636.		0
97	O2-03-01: INCREASED AMYLOID DEPOSITION IN OLDER ADULTS AT RISK FOR PROGRESSION TO ALZHEIMER'S DISEASE DUE TO GENETIC BACKGROUND AND/OR THE PRESENCE OF SIGNIFICANT MEMORY CONCERNS. , 2014, 10, P167-P167.		0
98	IC-P-096: INCREASED AMYLOID DEPOSITION IN OLDER ADULTS AT RISK FOR PROGRESSION TO ALZHEIMER'S DISEASE DUE TO GENETIC BACKGROUND AND/OR THE PRESENCE OF SIGNIFICANT MEMORY CONCERNS. , 2014, 10, P54-P54.		0
99	P1-213: GENOME-WIDE PROTEIN INTERACTION-GUIDED EPISTATIC ANALYSIS ON MEMORY PERFORMANCE: AN ADNI STUDY. , 2014, 10, P381-P382.		0
100	P3-017: ASSOCIATION ANALYSIS OF RARE VARIANTS NEAR THE APOE REGION WITH CEREBROSPINAL FLUID (CSF) BIOMARKERS OF ALZHEIMER'S DISEASE. , 2014, 10, P632-P633.		0
101	P3-019: RARE VARIANT IN PLD3 IS ASSOCIATED WITH ALZHEIMER'S PATTERN OF NEURODEGENERATIVE CHANGES. , 2014, 10, P634-P634.		0
102	P2-132: Association of cerebral microhemorrhages with amyloid deposition and hyperlipidemia. , 2015, 11, P534-P535.		0
103	P3-011: Genome-wide association of plasma homocysteine in the indianapolis-ibadan dementia study cohort. , 2015, 11, P623-P624.		0
104	P3-014: Influence of rare RELN variants on quantitative PET imaging and CSF phenotypes in late-onset Alzheimer's disease. , 2015, 11, P624-P625.		0
105	P4-191: Gwas identifies gli3 as a novel gene for language deficits and cortical changes in older adults at-risk for Alzheimer's disease. , 2015, 11, P853-P853.		0
106	P1-193: Anticholinergic medication use in older adults is associated with memory and hippocampal volume. , 2015, 11, P422-P422.		0
107	IC-P-035: Effect of hypertension and antihypertensive medication on executive function, brain atrophy, and white matter hyperintensities. , 2015, 11, P32-P33.		0
108	P4-002: Genome-wide network-based pathway analysis of CSF biomarker t-tau in the ADNI cohort. , 2015, 11, P765-P765.		0

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109	IC-P-034: Anticholinergic medication use in older adults is associated with memory and hippocampal volume. , 2015, 11, P32-P32.		0
110	P4-197: Gene expression profiling identifies altered networks in late-onset Alzheimer's disease: Immune response and mitochondrial process. , 2015, 11, P855-P856.		0
111	O4-05-01: Gwas of longitudinal amyloid PET identifies IL1RAP as a new potential Alzheimer's disease target. , 2015, 11, P277-P278.		0
112	IC-P-037: Association of cerebral microhemorrhages with amyloid deposition and hyperlipidemia. , 2015, 11, P33-P34.		0
113	O1-04-04: Effect of hypertension and antihypertensive medication on executive function, brain atrophy, and white matter hyperintensities. , 2015, 11, P133-P134.		0
114	P1-009: The nav2 (neuron navigator 2) gene as a common genetic influence across correlated episodic memory performances. , 2015, 11, P339-P340.		0
115	O4-12-06: The Alzheimer's metabolome: Identification of novel markers and treatment targets. , 2015, 11, P301-P302.		0
116	ICâ€Pâ€072: Gene Expression Of ABCA7 Dysregulated in Peripheral Blood is Associated With Decreased Metabolic Activity in Hippocampus. Alzheimer's and Dementia, 2016, 12, P56.	0.8	0
117	ICâ€Pâ€074: Genomeâ€Wide Metaâ€Analysis of Transcriptome Profiling Identifies Novel Dysregulated Genes Implicated in Alzheimerâ€™s Disease. Alzheimer's and Dementia, 2016, 12, P58.	0.8	0
118	ICâ€Pâ€075: The Growth and Impact of ADNI Genetics Publications as Measured by Science Mapping. Alzheimer's and Dementia, 2016, 12, P60.	0.8	0
119	IC-P-109: Plasma TAU Levels in Mild Cognitive Impairment and Alzheimerâ€™s Disease. , 2016, 12, P82-P83.		0
120	P2â€258: The Growth and Impact of ADNI Genetics Publications as Measured by Science Mapping. Alzheimer's and Dementia, 2016, 12, P725.	0.8	0
121	P3â€087: Gene Expression of <i>ABCA7</i> Dysregulated in Peripheral Blood is Associated With Decreased Metabolic Activity in Hippocampus. Alzheimer's and Dementia, 2016, 12, P851.	0.8	0
122	P3-089: Influence of Parkinsonâ€™s Disease Candidate Genes On Lewy Body Pathology in Autopsy-Confirmed Alzheimer's Disease Cases. , 2016, 12, P854-P854.		0
123	F1â€02â€01: The Alzheimerâ€™s Metabolome: Relationship to Pathological Markers and Cognitive Decline in the Alzheimerâ€™s Disease Neuroimaging Initiative (ADNI). Alzheimer's and Dementia, 2016, 12, P164.	0.8	0
124	F1-02-02: Genetic Influence on Levels of Targeted Metabolites Associated with Alzheimerâ€™s Disease. , 2016, 12, P164-P165.		0
125	O2-06-02: Genome-Wide Meta-Analysis of Transcriptome Profiling Identifies Novel Dysregulated Genes Implicated in Alzheimer's Disease. , 2016, 12, P238-P239.		0
126	O4â€10â€04: Plasma TAU Levels in Mild Cognitive Impairment and Alzheimer's Disease. Alzheimer's and Dementia, 2016, 12, P358.	0.8	0

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127	02â€“10â€“05: Cerebrospinal Fluid Levels of Amyloid Beta and Tau as Endophenotypes Reveal Novel Variants Potentially Informative for Alzheimer's Disease. Alzheimer's and Dementia, 2016, 12, P252.	0.8	0
128	[F1â€“02â€“04]: INTEGRATING MULTIâ€“MODALITY IMAGING AND MULTIâ€“LAYER â€“OMICS TO ADVANCE THE SYSTEMS BIOLOGY OF ALZHEIMER'S DISEASE. Alzheimer's and Dementia, 2017, 13, P175.	0.8	0
129	Bootstrapped Sparse Canonical Correlation Analysis. , 2018, , 101-117.		0
130	A Network-Based Framework for Mining High-Level Imaging Genetic Associations. , 2018, , 119-134.		0