Shannon Leigh Risacher

List of Publications by Year
in descending order

369	13,249	31976	29157
papers	citations	53	104
		h-index	g-index

462	462	462	17297
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Common genetic variants influence human subcortical brain structures. Nature, 2015, 520, 224-229.	27.8	772
2	The ENIGMA Consortium: large-scale collaborative analyses of neuroimaging and genetic data. Brain Imaging and Behavior, 2014, 8, 153-182.	2.1	696
3	Identification of common variants associated with human hippocampal and intracranial volumes. Nature Genetics, 2012, 44, 552-561.	21.4	594
4	Baseline MRI Predictors of Conversion from MCI to Probable AD in the ADNI Cohort. Current Alzheimer Research, 2009, 6, 347-361.	1.4	484
5	The genetic architecture of the human cerebral cortex. Science, 2020, 367, .	12.6	450
6	Altered bile acid profile associates with cognitive impairment in Alzheimer's disease—An emerging role for gut microbiome. Alzheimer's and Dementia, 2019, 15, 76-92.	0.8	396
7	Alzheimer's Disease Neuroimaging Initiative biomarkers as quantitative phenotypes: Genetics core aims, progress, and plans. Alzheimer's and Dementia, 2010, 6, 265-273.	0.8	378
8	Metabolic network failures in Alzheimer's disease: A biochemical roadÂmap. Alzheimer's and Dementia, 2017, 13, 965-984.	0.8	362
9	Whole genome association study of brain-wide imaging phenotypes for identifying quantitative trait loci in MCI and AD: A study of the ADNI cohort. NeuroImage, 2010, 53, 1051-1063.	4.2	340
10	Subjective Cognitive Decline in Older Adults: An Overview of Self-Report Measures Used Across 19 International Research Studies. Journal of Alzheimer's Disease, 2015, 48, S63-S86.	2.6	317
11	Novel genetic loci associated with hippocampal volume. Nature Communications, 2017, 8, 13624.	12.8	250
12	Genetic studies of quantitative MCI and AD phenotypes in ADNI: Progress, opportunities, and plans. Alzheimer's and Dementia, 2015, 11, 792-814.	0.8	241
13	Association Between Anticholinergic Medication Use and Cognition, Brain Metabolism, and Brain Atrophy in Cognitively Normal Older Adults. JAMA Neurology, 2016, 73, 721.	9.0	235
14	Subjective cognitive decline and rates of incident Alzheimer's disease and non—Alzheimer's disease dementia. Alzheimer's and Dementia, 2019, 15, 465-476.	0.8	232
15	Longitudinal MRI atrophy biomarkers: Relationship to conversion in the ADNI cohort. Neurobiology of Aging, 2010, 31, 1401-1418.	3.1	230
16	Novel genetic loci underlying human intracranial volume identified through genome-wide association. Nature Neuroscience, 2016, 19, 1569-1582.	14.8	213
17	Altered bile acid profile in mild cognitive impairment and Alzheimer's disease: Relationship to neuroimaging and CSF biomarkers. Alzheimer's and Dementia, 2019, 15, 232-244.	0.8	198
18	Genetic architecture of subcortical brain structures in 38,851 individuals. Nature Genetics, 2019, 51, 1624-1636.	21.4	192

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19	Network approaches to systems biology analysis of complex disease: integrative methods for multi-omics data. Briefings in Bioinformatics, 2018, 19, 1370-1381.	6.5	185
20	APOE and BCHE as modulators of cerebral amyloid deposition: a florbetapir PET genome-wide association study. Molecular Psychiatry, 2014, 19, 351-357.	7.9	181
21	Genetic analysis of quantitative phenotypes in AD and MCI: imaging, cognition and biomarkers. Brain Imaging and Behavior, 2014, 8, 183-207.	2.1	161
22	*APOE* effect on Alzheimer's disease biomarkers in older adults with significant memory concern. Alzheimer's and Dementia, 2015, 11, 1417-1429.	0.8	157
23	Identifying quantitative trait loci via group-sparse multitask regression and feature selection: an imaging genetics study of the ADNI cohort. Bioinformatics, 2012, 28, 229-237.	4.1	149
24	Association of Altered Liver Enzymes With Alzheimer Disease Diagnosis, Cognition, Neuroimaging Measures, and Cerebrospinal Fluid Biomarkers. JAMA Network Open, 2019, 2, e197978.	5.9	142
25	Altered Default Mode Network Connectivity in Older Adults with Cognitive Complaints and Amnestic Mild Cognitive Impairment. Journal of Alzheimer's Disease, 2013, 35, 751-760.	2.6	135
26	The role of apolipoprotein E (APOE) genotype in early mild cognitive impairment (E-MCI). Frontiers in Aging Neuroscience, 2013, 5, 11.	3.4	126
27	Visual contrast sensitivity in Alzheimer's disease, mild cognitive impairment, and older adults with cognitive complaints. Neurobiology of Aging, 2013, 34, 1133-1144.	3.1	123
28	Neuropathological correlates and genetic architecture of microglial activation in elderly human brain. Nature Communications, 2019, 10, 409.	12.8	121
29	APOE genotype and neuroimaging markers of Alzheimer's disease: systematic review and meta-analysis. Journal of Neurology, Neurosurgery and Psychiatry, 2015, 86, 127-134.	1.9	118
30	GWAS of longitudinal amyloid accumulation on ^{18}F-florbetapir PET in Alzheimer's disease implicates microglial activation gene *IL1RAP*. Brain, 2015, 138, 3076-3088.	7.6	117
31	Alzheimer disease brain atrophy subtypes are associated with cognition and rate of decline. Neurology, 2017, 89, 2176-2186.	1.1	115
32	Identifying disease sensitive and quantitative trait-relevant biomarkers from multidimensional heterogeneous imaging genetics data via sparse multimodal multitask learning. Bioinformatics, 2012, 28, i127-i136.	4.1	114
33	A large scale multivariate parallel ICA method reveals novel imaging–genetic relationships for Alzheimer's disease in the ADNI cohort. NeuroImage, 2012, 60, 1608-1621.	4.2	111
34	Neuroimaging and Other Biomarkers for Alzheimer's Disease: The Changing Landscape of Early Detection. Annual Review of Clinical Psychology, 2013, 9, 621-648.	12.3	110
35	Neuroimaging Biomarkers of Neurodegenerative Diseases and Dementia. Seminars in Neurology, 2013, 33, 386-416.	1.4	110
36	Associations of the Top 20 Alzheimer Disease Risk Variants With Brain Amyloidosis. JAMA Neurology, 2018, 75, 328.	9.0	101

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37	Multi-modal neuroimaging feature selection with consistent metric constraint for diagnosis of Alzheimer's disease. Medical Image Analysis, 2020, 60, 101625.	11.6	99
38	Structured sparse canonical correlation analysis for brain imaging genetics: an improved GraphNet method. Bioinformatics, 2016, 32, 1544-1551.	4.1	96
39	The Cognitive Change Index as a Measure of Self and Informant Perception of Cognitive Decline: Relation to Neuropsychological Tests. Journal of Alzheimer's Disease, 2016, 51, 1145-1155.	2.6	93
40	Selfâ€rated and informantâ€rated everyday function in comparison to objective markers of Alzheimer's disease. Alzheimer's and Dementia, 2015, 11, 1080-1089.	0.8	85
41	Whole-exome sequencing and imaging genetics identify functional variants for rate of change in hippocampal volume in mild cognitive impairment. Molecular Psychiatry, 2013, 18, 781-787.	7.9	81
42	Type 2 diabetes mellitus is associated with brain atrophy and hypometabolism in the ADNI cohort. Neurology, 2016, 87, 595-600.	1.1	81
43	Plasma amyloid beta levels are associated with cerebral amyloid and tau deposition. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2019, 11, 510-519.	2.4	77
44	Comparison of Manual and Automated Determination of Hippocampal Volumes in MCI and Early AD. Brain Imaging and Behavior, 2010, 4, 86-95.	2.1	74
45	Sparse multi-task regression and feature selection to identify brain imaging predictors for memory performance. , 2011, , 557-562.		72
46	Targeted neurogenesis pathway-based gene analysis identifies ADORA2A associated with hippocampal volume in mild cognitive impairment and Alzheimer's disease. Neurobiology of Aging, 2017, 60, 92-103.	3.1	70
47	Voxel and surface-based topography of memory and executive deficits in mild cognitive impairment and Alzheimerâ€™s disease. Brain Imaging and Behavior, 2012, 6, 551-567.	2.1	66
48	Serum triglycerides in Alzheimer disease. Neurology, 2020, 94, e2088-e2098.	1.1	63
49	From phenotype to genotype: an association study of longitudinal phenotypic markers to Alzheimer's disease relevant SNPs. Bioinformatics, 2012, 28, i619-i625.	4.1	62
50	Genetic correlations and genome-wide associations of cortical structure in general population samples of 22,824 adults. Nature Communications, 2020, 11, 4796.	12.8	61
51	Multiple loci influencing hippocampal degeneration identified by genome scan. Annals of Neurology, 2012, 72, 65-75.	5.3	59
52	Traumatic brain injury and age at onset of cognitive impairment in older adults. Journal of Neurology, 2016, 263, 1280-1285.	3.6	59
53	Genome-wide pathway analysis of memory impairment in the Alzheimerâ€™s Disease Neuroimaging Initiative (ADNI) cohort implicates gene candidates, canonical pathways, and networks. Brain Imaging and Behavior, 2012, 6, 634-648.	2.1	58
54	Transcriptome-guided amyloid imaging genetic analysis via a novel structured sparse learning algorithm. Bioinformatics, 2014, 30, i564-i571.	4.1	57

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55	Bundle analytics, a computational framework for investigating the shapes and profiles of brain pathways across populations. Scientific Reports, 2020, 10, 17149.	3.3	57
56	Identifying AD-Sensitive and Cognition-Relevant Imaging Biomarkers via Joint Classification and Regression. Lecture Notes in Computer Science, 2011, 14, 115-123.	1.3	57
57	Influence of *TSPO* Genotype on ^{11}C-PBR28 Standardized Uptake Values. Journal of Nuclear Medicine, 2013, 54, 1320-1322.	5.0	56
58	Plasma Tau Association with Brain Atrophy in Mild Cognitive Impairment and Alzheimer's Disease. Journal of Alzheimer's Disease, 2017, 58, 1245-1254.	2.6	54
59	Detecting genetic associations with brain imaging phenotypes in Alzheimer's disease via a novel structured SCCA approach. Medical Image Analysis, 2020, 61, 101656.	11.6	53
60	Identifying Neuroimaging and Proteomic Biomarkers for MCI and AD via the Elastic Net. Lecture Notes in Computer Science, 2011, 7012, 27-34.	1.3	53
61	Cognitive complaints in older adults at risk for Alzheimer's disease are associated with altered resting-state networks. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2017, 6, 40-49.	2.4	52
62	Genomic Copy Number Analysis in Alzheimer's Disease and Mild Cognitive Impairment: An ADNI Study. International Journal of Alzheimer's Disease, 2011, 2011, 1-10.	2.0	51
63	Resting state network modularity along the prodromal late onset Alzheimer's disease continuum. NeuroImage: Clinical, 2019, 22, 101687.	2.7	51
64	White matter alterations in early-stage Alzheimer's disease: A tract-specific study. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2019, 11, 576-587.	2.4	50
65	Influence of Genetic Variation on Plasma Protein Levels in Older Adults Using a Multi-Analyte Panel. PLoS ONE, 2013, 8, e70269.	2.5	50
66	Targeted metabolomics and medication classification data from participants in the ADNI1 cohort. Scientific Data, 2017, 4, 170140.	5.3	49
67	Protective variant for hippocampal atrophy identified by whole exome sequencing. Annals of Neurology, 2015, 77, 547-552.	5.3	48
68	Relationship between baseline brain metabolism measured using [18F]FDG PET and memory and executive function in prodromal and early Alzheimer's disease. Brain Imaging and Behavior, 2012, 6, 568-583.	2.1	47
69	Amyloid pathway-based candidate gene analysis of [11C]PiB-PET in the Alzheimer's Disease Neuroimaging Initiative (ADNI) cohort. Brain Imaging and Behavior, 2012, 6, 1-15.	2.1	47
70	Genome-wide association study of brain amyloid deposition as measured by Pittsburgh Compound-B (PiB)-PET imaging. Molecular Psychiatry, 2021, 26, 309-321.	7.9	47
71	Mining Outcome-relevant Brain Imaging Genetic Associations via Three-way Sparse Canonical Correlation Analysis in Alzheimer's Disease. Scientific Reports, 2017, 7, 44272.	3.3	44
72	Olfactory identification in subjective cognitive decline and mild cognitive impairment: Association with tau but not amyloid positron emission tomography. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2017, 9, 57-66.	2.4	44

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73	Association of cancer history with Alzheimer's disease onset and structural brain changes. Frontiers in Physiology, 2014, 5, 423.	2.8	43
74	Association of plasma and cortical amyloid beta is modulated by <i>APOE</i> Îµ4 status. Alzheimer's and Dementia, 2014, 10, e9-e18.	0.8	43
75	Cortical surface biomarkers for predicting cognitive outcomes using group l2,1 norm. Neurobiology of Aging, 2015, 36, S185-S193.	3.1	43
76	Identification of associations between genotypes and longitudinal phenotypes via temporally-constrained group sparse canonical correlation analysis. Bioinformatics, 2017, 33, i341-i349.	4.1	42
77	The Structural and Functional Connectome and Prediction of Risk for Cognitive Impairment in Older Adults. Current Behavioral Neuroscience Reports, 2015, 2, 234-245.	1.3	41
78	Topographic staging of tau positron emission tomography images. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2018, 10, 221-231.	2.4	41
79	Exercise prevents obesity-induced cognitive decline and white matter damage in mice. Neurobiology of Aging, 2019, 80, 154-172.	3.1	40
80	Deep Fusion of Brain Structure-Function in Mild Cognitive Impairment. Medical Image Analysis, 2021, 72, 102082.	11.6	37
81	Deep learning detection of informative features in tau PET for Alzheimerâ€™s disease classification. BMC Bioinformatics, 2020, 21, 496.	2.6	37
82	Identifying progressive imaging genetic patterns via multi-task sparse canonical correlation analysis: a longitudinal study of the ADNI cohort. Bioinformatics, 2019, 35, i474-i483.	4.1	36
83	A Novel Structure-Aware Sparse Learning Algorithm for Brain Imaging Genetics. Lecture Notes in Computer Science, 2014, 17, 329-336.	1.3	36
84	The effect of the top 20 Alzheimer disease risk genes on grayâ€matter density and FDG PET brain metabolism. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2016, 5, 53-66.	2.4	35
85	Comprehensive Gene- and Pathway-Based Analysis of Depressive Symptoms in Older Adults. Journal of Alzheimer's Disease, 2015, 45, 1197-1206.	2.6	33
86	FASTKD2 is associated with memory and hippocampal structure in older adults. Molecular Psychiatry, 2015, 20, 1197-1204.	7.9	33
87	Cognitive Dysfunction and Greater Visitâ€toâ€Visit Systolic Blood Pressure Variability. Journal of the American Geriatrics Society, 2013, 61, 2168-2173.	2.6	32
88	A novel SCCA approach via truncated <i> â„" </i> 1-norm and truncated group lasso for brain imaging genetics. Bioinformatics, 2018, 34, 278-285.	4.1	31
89	Plasma phosphorylated-tau181 as a predictive biomarker for Alzheimerâ€™s amyloid, tau and FDG PET status. Translational Psychiatry, 2021, 11, 585.	4.8	31
90	Targeted genetic analysis of cerebral blood flow imaging phenotypes implicates the INPP5D gene. Neurobiology of Aging, 2019, 81, 213-221.	3.1	30

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91	Tau-related white-matter alterations along spatially selective pathways. NeuroImage, 2021, 226, 117560.	4.2	30
92	Development and validation of language and visuospatial composite scores in ADNI. Alzheimer's and Dementia: Translational Research and Clinical Interventions, 2020, 6, e12072.	3.7	29
93	Network-based analysis of genetic variants associated with hippocampal volume in Alzheimer's disease: a study of ADNI cohorts. BioData Mining, 2016, 9, 3.	4.0	28
94	Association analysis of rare variants near the APOE region with CSF and neuroimaging biomarkers of Alzheimer's disease. BMC Medical Genomics, 2017, 10, 29.	1.5	28
95	Type 2 diabetes mellitus and cerebrospinal fluid Alzheimer's disease biomarker amyloid Î²1â€42 in Alzheimer's Disease Neuroimaging Initiative participants. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2018, 10, 94-98.	2.4	28
96	Dysregulated Fc gamma receptorâ€"mediated phagocytosis pathway in Alzheimer's disease: network-based gene expression analysis. Neurobiology of Aging, 2020, 88, 24-32.	3.1	28
97	Associating Multi-Modal Brain Imaging Phenotypes and Genetic Risk Factors via a Dirty Multi-Task Learning Method. IEEE Transactions on Medical Imaging, 2020, 39, 3416-3428.	8.9	27
98	A rat head holder for simultaneous scanning of two rats in small animal PET scanners: Design, construction, feasibility testing and kinetic validation. Journal of Neuroscience Methods, 2009, 176, 24-33.	2.5	26
99	Identifying Multimodal Intermediate Phenotypes Between Genetic Risk Factors and Disease Status in Alzheimer's Disease. Neuroinformatics, 2016, 14, 439-452.	2.8	26
100	Tau Imaging in Alzheimer's Disease Diagnosis and Clinical Trials. Neurotherapeutics, 2017, 14, 62-68.	4.4	26
101	Visual contrast sensitivity is associated with the presence of cerebral amyloid and tau deposition. Brain Communications, 2020, 2, fcaa019.	3.3	26
102	Neuroimaging in aging and neurologic diseases. Handbook of Clinical Neurology / Edited By P J Vinken and G W Bruyn, 2019, 167, 191-227.	1.8	25
103	Multi-Task Sparse Canonical Correlation Analysis with Application to Multi-Modal Brain Imaging Genetics. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2021, 18, 227-239.	3.0	25
104	Cholinergic Enhancement of Brain Activation in Mild Cognitive Impairment during Episodic Memory Encoding. Frontiers in Psychiatry, 2013, 4, 105.	2.6	23
105	Hippocampal transcriptome-guided genetic analysis of correlated episodic memory phenotypes in Alzheimer's disease. Frontiers in Genetics, 2015, 6, 117.	2.3	23
106	Tissue-specific network-based genome wide study of amygdala imaging phenotypes to identify functional interaction modules. Bioinformatics, 2017, 33, 3250-3257.	4.1	23
107	Genomeâ€wide transcriptome analysis identifies novel dysregulated genes implicated in Alzheimer's pathology. Alzheimer's and Dementia, 2020, 16, 1213-1223.	0.8	23
108	Staging tau pathology with tau PET in Alzheimer's disease: a longitudinal study. Translational Psychiatry, 2021, 11, 483.	4.8	23

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109	Genetic architecture of resilience of executive functioning. Brain Imaging and Behavior, 2012, 6, 621-633.	2.1	22
110	Memory concerns in the early Alzheimer's disease prodrome: Regional association with tau deposition. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2018, 10, 322-331.	2.4	22
111	Genome-wide association analysis of hippocampal volume identifies enrichment of neurogenesis-related pathways. Scientific Reports, 2019, 9, 14498.	3.3	22
112	Serum metabolites associated with brain amyloid beta deposition, cognition and dementia progression. Brain Communications, 2021, 3, fcab139.	3.3	21
113	Sparse Bayesian Learning for Identifying Imaging Biomarkers in AD Prediction. Lecture Notes in Computer Science, 2010, 13, 611-618.	1.3	21
114	Integration of bioinformatics and imaging informatics for identifying rare PSEN1 variants in Alzheimer's disease. BMC Medical Genomics, 2016, 9, 30.	1.5	20
115	Identifying diagnosis-specific genotype–phenotype associations via joint multitask sparse canonical correlation analysis and classification. Bioinformatics, 2020, 36, i371-i379.	4.1	20
116	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
117	Genome-wide association study of language performance in Alzheimer's disease. Brain and Language, 2017, 172, 22-29.	1.6	20
118	Hippocampal-subfield microstructures and their relation to plasma biomarkers in Alzheimer's disease. Brain, 2022, 145, 2149-2160.	7.6	20
119	Differences in Medication Use in the Alzheimer¼s Disease Neuroimaging Initiative. Drugs and Aging, 2010, 27, 677-686.	2.7	19
120	Optimizing differential identifiability improves connectome predictive modeling of cognitive deficits from functional connectivity in Alzheimer's disease. Human Brain Mapping, 2021, 42, 3500-3516.	3.6	18
121	Age at injury is associated with the long-term cognitive outcome of traumatic brain injuries. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2017, 6, 196-200.	2.4	17
122	Exome Chip Analysis Identifies Low-Frequency and Rare Variants in *MRPL38* for White Matter Hyperintensities on Brain Magnetic Resonance Imaging. Stroke, 2018, 49, 1812-1819.	2.0	17
123	Identification of exon skipping events associated with Alzheimer's disease in the human hippocampus. BMC Medical Genomics, 2019, 12, 13.	1.5	17
124	Neurodegenerative changes in early- and late-onset cognitive impairment with and without brain amyloidosis. Alzheimer's Research and Therapy, 2020, 12, 93.	6.2	17
125	Regional imaging genetic enrichment analysis. Bioinformatics, 2020, 36, 2554-2560.	4.1	16
126	Sparse Bayesian multi-task learning for predicting cognitive outcomes from neuroimaging measures in Alzheimer's disease. , 2012, , .		15

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127	Bile acids targeted metabolomics and medication classification data in the ADNI1 and ADNIGO/2 cohorts. Scientific Data, 2019, 6, 212.	5.3	15
128	IDENTIFICATION OF DISCRIMINATIVE IMAGING PROTEOMICS ASSOCIATIONS IN ALZHEIMER'S DISEASE VIA A NOVEL SPARSE CORRELATION MODEL. , 2017, 22, 94-104.		14
129	Telomere Shortening in the Alzheimer's Disease Neuroimaging Initiative Cohort. Journal of Alzheimer's Disease, 2019, 71, 33-43.	2.6	14
130	GN-SCCA: GraphNet Based Sparse Canonical Correlation Analysis for Brain Imaging Genetics. Lecture Notes in Computer Science, 2015, 9250, 275-284.	1.3	14
131	Two-dimensional enrichment analysis for mining high-level imaging genetic associations. Brain Informatics, 2017, 4, 27-37.	3.0	13
132	Volumetric comparison of hippocampal subfields extracted from 4-minute accelerated vs. 8-minute high-resolution T2-weighted 3T MRI scans. Brain Imaging and Behavior, 2018, 12, 1583-1595.	2.1	13
133	Fast Multi-Task SCCA Learning with Feature Selection for Multi-Modal Brain Imaging Genetics. , 2018, 2018, 356-361.		13
134	Joint High-Order Multi-Task Feature Learning to Predict the Progression of Alzheimer's Disease. Lecture Notes in Computer Science, 2018, 11070, 555-562.	1.3	13
135	Differential patterns of gray matter volumes and associated gene expression profiles in cognitively-defined Alzheimer's disease subgroups. NeuroImage: Clinical, 2021, 30, 102660.	2.7	13
136	Rare variants in the splicing regulatory elements of EXOC3L4 are associated with brain glucose metabolism in Alzheimer's disease. BMC Medical Genomics, 2018, 11, 76.	1.5	12
137	Novel Markers of Angiogenesis in the Setting of Cognitive Impairment and Dementia. Journal of Alzheimer's Disease, 2020, 75, 959-969.	2.6	12
138	Identifying Associations Between Brain Imaging Phenotypes and Genetic Factors via a Novel Structured SCCA Approach. Lecture Notes in Computer Science, 2017, 10265, 543-555.	1.3	12
139	Cerebral hypometabolism and grey matter density in MAPT intron 10 +3 mutation carriers. American Journal of Neurodegenerative Disease, 2014, 3, 103-14.	0.1	12
140	Quantitative trait loci identification for brain endophenotypes via new additive model with random networks. Bioinformatics, 2018, 34, i866-i874.	4.1	11
141	Genome-wide Network-assisted Association and Enrichment Study of Amyloid Imaging Phenotype in Alzheimer's Disease. Current Alzheimer Research, 2020, 16, 1163-1174.	1.4	11
142	Multivariate genome wide association and network analysis of subcortical imaging phenotypes in Alzheimer's disease. BMC Genomics, 2020, 21, 896.	2.8	11
143	Building a surface atlas of hippocampal subfields from MRI scans using FreeSurfer, FIRST and SPHARM. , 2014, 2014, 813-816.		10
144	Detection of tau in Gerstmann-Sträussler-Scheinker disease (PRNP F198S) by [18F]Flortaucipir PET. Acta Neuropathologica Communications, 2018, 6, 114.	5.2	10

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145	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
146	Social Networks and Cognitive Reserve: Network Structure Moderates the Association Between Amygdalar Volume and Cognitive Outcomes. Journals of Gerontology - Series B Psychological Sciences and Social Sciences, 2022, 77, 1490-1500.	3.9	10
147	Structured sparse CCA for brain imaging genetics via graph OSCAR. BMC Systems Biology, 2016, 10, 68.	3.0	9
148	Pattern Discovery in Brain Imaging Genetics via SCCA Modeling with a Generic Non-convex Penalty. Scientific Reports, 2017, 7, 14052.	3.3	9
149	Diagnosis Status Guided Brain Imaging Genetics Via Integrated Regression And Sparse Canonical Correlation Analysis. , 2019, 2019, 356-359.		9
150	Neurodegenerative Patterns of Cognitive Clusters of Early-Onset Alzheimer's Disease Subjects: Evidence for Disease Heterogeneity. Dementia and Geriatric Cognitive Disorders, 2019, 48, 131-142.	1.5	9
151	Differential trajectories of hypometabolism across cognitively-defined Alzheimer's disease subgroups. NeuroImage: Clinical, 2021, 31, 102725.	2.7	9
152	Identification of functional variants from whole-exome sequencing, combined with neuroimaging genetics. Molecular Psychiatry, 2013, 18, 739-739.	7.9	8
153	Identifying Candidate Genetic Associations with MRI-Derived AD-Related ROI via Tree-Guided Sparse Learning. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2019, 16, 1986-1996.	3.0	8
154	Integrative-omics for discovery of network-level disease biomarkers: a case study in Alzheimer's disease. Briefings in Bioinformatics, 2021, 22, .	6.5	8
155	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
156	Longitudinal Genotype-Phenotype Association Study via Temporal Structure Auto-learning Predictive Model. Lecture Notes in Computer Science, 2017, 10229, 287-302.	1.3	8
157	The effect of reference panels and software tools on genotype imputation. AMIA ... Annual Symposium proceedings, 2011, 2011, 1013-8.	0.2	8
158	[(11)C]PiB PET in Gerstmann-Sträussler-Scheinker disease. American Journal of Nuclear Medicine and Molecular Imaging, 2016, 6, 84-93.	1.0	8
159	Imaging of alcohol-induced dopamine release in rats:Preliminary findings with [11C]raclopride PET. Synapse, 2011, 65, 929-937.	1.2	7
160	Volumetric GWAS of medial temporal lobe structures identifies an ERC1 locus using ADNI high-resolution T2-weighted MRI data. Neurobiology of Aging, 2020, 95, 81-93.	3.1	7
161	Temporal stability of the ventral attention network and general cognition along the Alzheimer's disease spectrum. NeuroImage: Clinical, 2021, 31, 102726.	2.7	7
162	Association of the top 20 Alzheimer's disease risk genes with [18F]flortaucipir PET. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2022, 14, e12308.	2.4	7

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163	Sparse Canonical Correlation Analysis via truncated ℓ<inf>1</inf>-norm with application to brain imaging genetics. , 2016, 2016, 707-711.		6
164	Codon bias among synonymous rare variants is associated with Alzheimer's disease imaging biomarker. , 2018, , .		6
165	Predicting progressions of cognitive outcomes via high-order multi-modal multi-task feature learning. , 2018, , .		6
166	Longitudinal Genotype–Phenotype Association Study through Temporal Structure Auto-Learning Predictive Model. Journal of Computational Biology, 2018, 25, 809-824.	1.6	6
167	Codon bias among synonymous rare variants is associated with Alzheimer's disease imaging biomarker. Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing, 2018, 23, 365-376.	0.7	6
168	Genetic variation affecting exon skipping contributes to brain structural atrophy in Alzheimer's disease. AMIA Summits on Translational Science Proceedings, 2018, 2017, 124-131.	0.4	6
169	Surface-based morphometric analysis of hippocampal subfields in mild cognitive impairment and Alzheimer's disease. , 2015, 2015, .		5
170	Multiple incomplete views clustering via non-negative matrix factorization with its application in Alzheimer's disease analysis. , 2018, , .		5
171	Neuroimaging Advances in Neurologic and Neurodegenerative Diseases. Neurotherapeutics, 2021, 18, 659-660.	4.4	5
172	Tau deposition and structural connectivity demonstrate differential association patterns with neurocognitive tests. Brain Imaging and Behavior, 2022, 16, 702-714.	2.1	5
173	Head injury is associated with tau deposition on PET in MCI and AD patients. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2021, 13, e12230.	2.4	5
174	Network-Guided Sparse Learning for Predicting Cognitive Outcomes from MRI Measures. Lecture Notes in Computer Science, 2013, 8159, 202-210.	1.3	5
175	Transcriptome-Guided Imaging Genetic Analysis via a Novel Sparse CCA Algorithm. Lecture Notes in Computer Science, 2017, 10551, 220-229.	1.3	5
176	Multimodal Neuroimaging Predictors for Cognitive Performance Using Structured Sparse Learning. Lecture Notes in Computer Science, 2012, , 1-17.	1.3	4
177	DIAGNOSIS-GUIDED METHOD FOR IDENTIFYING MULTI-MODALITY NEUROIMAGING BIOMARKERS ASSOCIATED WITH GENETIC RISK FACTORS IN ALZHEIMER'S DISEASE. , 2016, , .		4
178	Brain explorer for connectomic analysis. Brain Informatics, 2017, 4, 253-269.	3.0	4
179	Joint exploration and mining of memory-relevant brain anatomic and connectomic patterns via a three-way association model. , 2018, 2018, 6-9.		4
180	MIND food and speed of processing training in older adults with low education, the MINDSpeed Alzheimer's disease prevention pilot trial. Contemporary Clinical Trials, 2019, 84, 105814.	1.8	4

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