

Junguk Hur

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2747325/publications.pdf>

Version: 2024-02-01

108
papers

4,730
citations

136950

32
h-index

106344

65
g-index

114
all docs

114
docs citations

114
times ranked

8110
citing authors

#	ARTICLE	IF	CITATIONS
1	The impact of methodology on the reproducibility and rigor of DNA methylation data. Scientific Reports, 2022, 12, 380.	3.3	3
2	SMAP is a pipeline for sample matching in proteogenomics. Nature Communications, 2022, 13, 744.	12.8	3
3	Modulation of Inflammatory Signaling Molecules in Bordetella pertussis Antigen-Challenged Human Monocytes in Presence of Adrenergic Agonists. Vaccines, 2022, 10, 321.	4.4	2
4	Interferon- β promotes monocyte-mediated lung injury during influenza infection. Cell Reports, 2022, 38, 110456.	6.4	29
5	Plasma Metabolomics and Lipidomics Differentiate Obese Individuals by Peripheral Neuropathy Status. Journal of Clinical Endocrinology and Metabolism, 2022, 107, 1091-1109.	3.6	17
6	Characterization of Prostanoids Response to Bordetella pertussis Antigen BscF and Tdap in LPS-challenged monocytes. Prostaglandins Leukotrienes and Essential Fatty Acids, 2022, , 102452.	2.2	0
7	COVID-19 vaccine design using reverse and structural vaccinology, ontology-based literature mining and machine learning. Briefings in Bioinformatics, 2022, 23, .	6.5	4
8	Ontological modeling and analysis of experimentally or clinically verified drugs against coronavirus infection. Scientific Data, 2021, 8, 16.	5.3	14
9	Predictability of Macrosomic Birth based on Maternal Factors and Fetal Aneuploidy Screening Biochemical Markers in Hyperglycemic Mothers. International Journal of Medical Sciences, 2021, 18, 2653-2660.	2.5	2
10	Microbial and genetic-based framework identifies drug targets in inflammatory bowel disease. Theranostics, 2021, 11, 7491-7506.	10.0	13
11	Gene expression profiles of diabetic kidney disease and neuropathy in <i>eNOS</i> knockout mice: Predictors of pathology and RAS blockade effects. FASEB Journal, 2021, 35, e21467.	0.5	10
12	Epigenetic Reprogramming Mediated by Maternal Diet Rich in Omega-3 Fatty Acids Protects From Breast Cancer Development in F1 Offspring. Frontiers in Cell and Developmental Biology, 2021, 9, 682593.	3.7	14
13	Anxiety-like behavior and intestinal microbiota changes as strain-and sex-dependent sequelae of mild food allergy in mouse models of cow's milk allergy. Brain, Behavior, and Immunity, 2021, 95, 122-141.	4.1	8
14	Predicting Drug-Induced Liver Injury Using Machine Learning on a Diverse Set of Predictors. Frontiers in Pharmacology, 2021, 12, 648805.	3.5	6
15	Alpha-Synuclein-induced DNA Methylation and Gene Expression in Microglia. Neuroscience, 2021, 468, 186-198.	2.3	8
16	Repurposable drugs for SARS-CoV-2 and influenza sepsis with scRNA-seq data targeting post-transcription modifications. Precision Clinical Medicine, 2021, 4, 215-230.	3.3	3
17	NADPH oxidase 5: A new player in peripheral neuropathy. Journal of the Neurological Sciences, 2021, 429, 119359.	0.6	0
18	Extracellular Vesicles in Serum and Central Nervous System Tissues Contain microRNA Signatures in Sporadic Amyotrophic Lateral Sclerosis. Frontiers in Molecular Neuroscience, 2021, 14, 739016.	2.9	17

#	ARTICLE	IF	CITATIONS
19	Method of Microglial DNA-RNA Purification from a Single Brain of an Adult Mouse. <i>Methods and Protocols</i> , 2021, 4, 86.	2.0	1
20	Integrated lipidomic and transcriptomic analyses identify altered nerve triglycerides in mouse models of prediabetes and type 2 diabetes. <i>DMM Disease Models and Mechanisms</i> , 2020, 13, .	2.4	42
21	oprC Impairs Host Defense by Increasing the Quorum-Sensing-Mediated Virulence of <i>Pseudomonas aeruginosa</i> . <i>Frontiers in Immunology</i> , 2020, 11, 1696.	4.8	11
22	Genome-wide profiling of DNA methylation and gene expression identifies candidate genes for human diabetic neuropathy. <i>Clinical Epigenetics</i> , 2020, 12, 123.	4.1	26
23	Atypical endometriosis is related to a higher recurrence rate. <i>European Journal of Obstetrics, Gynecology and Reproductive Biology</i> , 2020, 254, 44-51.	1.1	5
24	Machine Learning-Based Predictive Modeling of Postpartum Depression. <i>Journal of Clinical Medicine</i> , 2020, 9, 2899.	2.4	48
25	Untargeted metabolomics yields insight into ALS disease mechanisms. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2020, 91, 1329-1338.	1.9	51
26	Cytoplasmic TDP43 Binds microRNAs: New Disease Targets in Amyotrophic Lateral Sclerosis. <i>Frontiers in Cellular Neuroscience</i> , 2020, 14, 117.	3.7	17
27	CIDO, a community-based ontology for coronavirus disease knowledge and data integration, sharing, and analysis. <i>Scientific Data</i> , 2020, 7, 181.	5.3	70
28	2285-PUB: Hippocampal Transcriptomic Changes Due to High-Fat Diet in Prediabetic Mice. <i>Diabetes</i> , 2020, 69, 2285-PUB.	0.6	0
29	537-P: Dietary Reversal Improves Peripheral Neuropathy and Gut Microbiota Profile in a Murine Model of Prediabetes and Obesity. <i>Diabetes</i> , 2020, 69, .	0.6	0
30	Network-Based Assessment of Adverse Drug Reaction Risk in Polypharmacy Using High-Throughput Screening Data. <i>International Journal of Molecular Sciences</i> , 2019, 20, 386.	4.1	12
31	Pathway crosstalk perturbation network modeling for identification of connectivity changes induced by diabetic neuropathy and pioglitazone. <i>BMC Systems Biology</i> , 2019, 13, 1.	3.0	28
32	Toll-like receptors and inflammation in metabolic neuropathy; a role in early versus late disease?. <i>Experimental Neurology</i> , 2019, 320, 112967.	4.1	38
33	Predictive Modeling of Postpartum Depression Using Machine Learning Approaches (P18-130-19). <i>Current Developments in Nutrition</i> , 2019, 3, nzz039.P18-130-19.	0.3	0
34	RBMS1 Methylation and mRNA Expression Are Differentially Regulated in Placenta Tissue from Obese Women (P11-131-19). <i>Current Developments in Nutrition</i> , 2019, 3, nzz048.P11-131-19.	0.3	4
35	Genome-wide DNA methylation profiling of human diabetic peripheral neuropathy in subjects with type 2 diabetes mellitus. <i>Epigenetics</i> , 2019, 14, 766-779.	2.7	28
36	A Computational Platform and Guide for Acceleration of Novel Medicines and Personalized Medicine. <i>Methods in Molecular Biology</i> , 2019, 1939, 181-198.	0.9	1

#	ARTICLE	IF	CITATIONS
37	A Network Pharmacology Approach for the Identification of Common Mechanisms of Drug-Induced Peripheral Neuropathy. CPT: Pharmacometrics and Systems Pharmacology, 2019, 8, 211-219.	2.5	4
38	Temporal evolution of the microbiome, immune system, and epigenome with disease progression in ALS mice. DMM Disease Models and Mechanisms, 2019, 13, .	2.4	50
39	A 2018 workshop: vaccine and drug ontology studies (VDOS 2018). BMC Bioinformatics, 2019, 20, 705.	2.6	1
40	Machine learning-based identification and rule-based normalization of adverse drug reactions in drug labels. BMC Bioinformatics, 2019, 20, 707.	2.6	13
41	Identification of Casiopeina II-gly secondary targets through a systems pharmacology approach. Computational Biology and Chemistry, 2019, 78, 127-132.	2.3	10
42	Post-transcriptional processing at the promoter proximal RNA polymerase II pausing. A possible mechanism for premature termination. FASEB Journal, 2019, 33, 458.13.	0.5	0
43	31-LB: Identification of Repurposable Drug Candidate for Diabetic Peripheral Neuropathy Using High-Throughput Drug-Perturbation Data. Diabetes, 2019, 68, .	0.6	0
44	Tox21 Enricher: Web-based Chemical/Biological Functional Annotation Analysis Tool Based on Tox21 Toxicity Screening Platform. Molecular Informatics, 2018, 37, e1700129.	2.5	7
45	Translational Systems Pharmacology-Based Predictive Assessment of Drug-Induced Cardiomyopathy. CPT: Pharmacometrics and Systems Pharmacology, 2018, 7, 166-174.	2.5	14
46	Acute Biophysical Responses and Psychological Effects of Different Types of Forests in Patients With Metabolic Syndrome. Environment and Behavior, 2018, 50, 298-323.	4.7	28
47	Transcriptional networks of progressive diabetic peripheral neuropathy in the db/db mouse model of type 2 diabetes: An inflammatory story. Experimental Neurology, 2018, 305, 33-43.	4.1	42
48	Advances in omics for informed pharmaceutical research and development in the era of systems medicine. Expert Opinion on Drug Discovery, 2018, 13, 1-4.	5.0	8
49	Association of Sleep Duration and Obesity According to Gender and Age in Korean Adults: Results from the Korea National Health and Nutrition Examination Survey 2007-2015. Journal of Korean Medical Science, 2018, 33, e345.	2.5	16
50	Conserved Transcriptional Signatures in Human and Murine Diabetic Peripheral Neuropathy. Scientific Reports, 2018, 8, 17678.	3.3	40
51	Trends of self-reported sleep duration in Korean Adults: results from the Korea National Health and Nutrition Examination Survey 2007-2015. Sleep Medicine, 2018, 52, 103-106.	1.6	17
52	Ontology-based literature mining and class effect analysis of adverse drug reactions associated with neuropathy-inducing drugs. Journal of Biomedical Semantics, 2018, 9, 17.	1.6	7
53	Exploration of the Anti-Inflammatory Drug Space Through Network Pharmacology: Applications for Drug Repurposing. Frontiers in Physiology, 2018, 9, 151.	2.8	13
54	Abnormal RNA stability in amyotrophic lateral sclerosis. Nature Communications, 2018, 9, 2845.	12.8	113

#	ARTICLE	IF	CITATIONS
55	NOX, NOX, Are You Here? The Emerging Role of NOX5 in Diabetic Neuropathy. Diabetes, 2018, 67, 30-LB.	0.6	4
56	Amelioration of Peripheral Neuropathy in Mouse Models of Diabetes by Dietary Reversal. Diabetes, 2018, 67, .	0.6	6
57	Systems Approach to Assign Expression Based Signatures to Adrenergic Drugs. FASEB Journal, 2018, 32, 690.2.	0.5	0
58	Two-Way Orthogonal Partial Least Squares (O2PLS) Analysis of the Lipidome and Transcriptome in Prediabetic and Diabetic Neuropathy. Diabetes, 2018, 67, 548-P.	0.6	0
59	Large-Scale DNA Methylation Profiling of Human Diabetic Peripheral Neuropathy in Subjects with Type 2 Diabetes Mellitus. Diabetes, 2018, 67, .	0.6	0
60	Comparative RNA-Seq transcriptome analyses reveal distinct metabolic pathways in diabetic nerve and kidney disease. Journal of Cellular and Molecular Medicine, 2017, 21, 2140-2152.	3.6	45
61	Assessment of the DNA damaging potential of environmental chemicals using a quantitative high-throughput screening approach to measure p53 activation. Environmental and Molecular Mutagenesis, 2017, 58, 494-507.	2.2	27
62	Ontology-based literature mining of E. coli vaccine-associated gene interaction networks. Journal of Biomedical Semantics, 2017, 8, 12.	1.6	13
63	Pre-Pregnancy Body Mass Index Is Associated with Dietary Inflammatory Index and C-Reactive Protein Concentrations during Pregnancy. Nutrients, 2017, 9, 351.	4.1	39
64	Prevalences and Management of Diabetes and Pre-diabetes among Korean Teenagers and Young Adults: Results from the Korea National Health and Nutrition Examination Survey 2005-2014. Journal of Korean Medical Science, 2017, 32, 1984.	2.5	14
65	Prediction of Gestational Diabetes Mellitus by Unconjugated Estriol Levels in Maternal Serum. International Journal of Medical Sciences, 2017, 14, 123-127.	2.5	20
66	Genetic Architecture of Group A Streptococcal Necrotizing Soft Tissue Infections in the Mouse. PLoS Pathogens, 2016, 12, e1005732.	4.7	32
67	The Interaction Network Ontology-supported modeling and mining of complex interactions represented with multiple keywords in biomedical literature. BioData Mining, 2016, 9, 41.	4.0	7
68	Gender-specific differences in diabetic neuropathy in BTBR ob/ob mice. Journal of Diabetes and Its Complications, 2016, 30, 30-37.	2.3	40
69	Ontology-based collection, representation and analysis of drug-associated neuropathy adverse events. Journal of Biomedical Semantics, 2016, 7, 29.	1.6	22
70	Computational Approaches to Accelerating Novel Medicine and Better Patient Care from Bedside to Benchtop. Advances in Protein Chemistry and Structural Biology, 2016, 102, 147-179.	2.3	2
71	Transcriptional networks of murine diabetic peripheral neuropathy and nephropathy: common and distinct gene expression patterns. Diabetologia, 2016, 59, 1297-1306.	6.3	34
72	Expression of microRNAs in human post-mortem amyotrophic lateral sclerosis spinal cords provides insight into disease mechanisms. Molecular and Cellular Neurosciences, 2016, 71, 34-45.	2.2	76

#	ARTICLE	IF	CITATIONS
73	Tissue-specific metabolic reprogramming drives nutrient flux in diabetic complications. JCI Insight, 2016, 1, e86976.	5.0	188
74	Literature Mining and Ontology based Analysis of Host-Brucella Geneâ€Gene Interaction Network. Frontiers in Microbiology, 2015, 6, 1386.	3.5	9
75	GLASS: a comprehensive database for experimentally validated GPCR-ligand associations. Bioinformatics, 2015, 31, 3035-3042.	4.1	92
76	The Metabolic Syndrome and Microvascular Complications in a Murine Model of Type 2 Diabetes. Diabetes, 2015, 64, 3294-3304.	0.6	49
77	Systems Pharmacological Analysis of Drugs Inducing Stevensâ€Johnson Syndrome and Toxic Epidermal Necrolysis. Chemical Research in Toxicology, 2015, 28, 927-934.	3.3	18
78	Development and application of an interaction network ontology for literature mining of vaccine-associated gene-gene interactions. Journal of Biomedical Semantics, 2015, 6, 2.	1.6	23
79	BTBR ob/ob mice as a novel diabetic neuropathy model: Neurological characterization and gene expression analyses. Neurobiology of Disease, 2015, 73, 348-355.	4.4	68
80	Early Gestational Weight Gain Rate and Adverse Pregnancy Outcomes in Korean Women. PLoS ONE, 2015, 10, e0140376.	2.5	46
81	Transcriptional changes and developmental abnormalities in a zebrafish model of myotonic dystrophy type 1. DMM Disease Models and Mechanisms, 2014, 7, 143-55.	2.4	25
82	Integrated Systems Pharmacology Analysis of Clinical Drugâ€Induced Peripheral Neuropathy. CPT: Pharmacometrics and Systems Pharmacology, 2014, 3, 1-11.	2.5	31
83	Updates on the web-based VIOLIN vaccine database and analysis system. Nucleic Acids Research, 2014, 42, D1124-D1132.	14.5	66
84	Intraspinal neural stem cell transplantation in amyotrophic lateral sclerosis: Phase 1 trial outcomes. Annals of Neurology, 2014, 75, 363-373.	5.3	184
85	Drug-Induced Rhabdomyolysis: From Systems Pharmacology Analysis to Biochemical Flux. Chemical Research in Toxicology, 2014, 27, 421-432.	3.3	27
86	Fluoxetine prevents dystrophic changes in a zebrafish model of Duchenne muscular dystrophy. Human Molecular Genetics, 2014, 23, 4651-4662.	2.9	55
87	A systems pharmacology approach to model tyrosine kinase inhibitorâ€Induced cardiotoxicity gene interaction networks (844.17). FASEB Journal, 2014, 28, 844.17.	0.5	0
88	Identification of Factors Associated With Sural Nerve Regeneration and Degeneration in Diabetic Neuropathy. Diabetes Care, 2013, 36, 4043-4049.	8.6	31
89	HbA1c is associated with sural nerve regeneration and degeneration in diabetic neuropathy. Journal of the Neurological Sciences, 2013, 333, e697-e698.	0.6	0
90	Epigenetic miRNA dysregulation as a mechanism for sporadic amyotrophic lateral sclerosis. Journal of the Neurological Sciences, 2013, 333, e698.	0.6	0

#	ARTICLE	IF	CITATIONS
91	ER-stress-induced transcriptional regulation increases protein synthesis leading to cell death. <i>Nature Cell Biology</i> , 2013, 15, 481-490.	10.3	1,315
92	The Role of Oxidative Stress in Nervous System Aging. <i>PLoS ONE</i> , 2013, 8, e68011.	2.5	39
93	Diabetic neuropathy. <i>Current Opinion in Neurology</i> , 2012, 25, 536-541.	3.6	131
94	Identification of fever and vaccine-associated gene interaction networks using ontology-based literature mining. <i>Journal of Biomedical Semantics</i> , 2012, 3, 18.	1.6	26
95	Identification of Epigenetically Altered Genes in Sporadic Amyotrophic Lateral Sclerosis. <i>PLoS ONE</i> , 2012, 7, e52672.	2.5	132
96	Oxidative stress and successful antioxidant treatment in models of RYR1-related myopathy. <i>Brain</i> , 2012, 135, 1115-1127.	7.6	114
97	Central nervous system endoplasmic reticulum stress in a murine model of type 2 diabetes. <i>Diabetologia</i> , 2012, 55, 2276-2284.	6.3	43
98	The identification of gene expression profiles associated with progression of human diabetic neuropathy. <i>Brain</i> , 2011, 134, 3222-3235.	7.6	132
99	Ontology-based Brucella vaccine literature indexing and systematic analysis of gene-vaccine association network. <i>BMC Immunology</i> , 2011, 12, 49.	2.2	34
100	Stem cell technology for neurodegenerative diseases. <i>Annals of Neurology</i> , 2011, 70, 353-361.	5.3	219
101	Transcriptional Profiling of Diabetic Neuropathy in the BKS <i>db/db</i> Mouse. <i>Diabetes</i> , 2011, 60, 1981-1989.	0.6	107
102	Pregnancy outcomes and relationship between maternal weight gain and fetal birth weight in Korean pregnant women at risk for gestational diabetes. <i>Journal of Women's Medicine</i> , 2011, 4, 35.	0.1	0
103	Maternal obesity and associated risk of adverse pregnancy outcomes in women with hyperglycemia. <i>Korean Journal of Obstetrics & Gynecology</i> , 2011, 54, 591.	0.1	1
104	Literature-based discovery of diabetes- and ROS-related targets. <i>BMC Medical Genomics</i> , 2010, 3, 49.	1.5	29
105	SciMiner: web-based literature mining tool for target identification and functional enrichment analysis. <i>Bioinformatics</i> , 2009, 25, 838-840.	4.1	78
106	PubChemSR: A search and retrieval tool for PubChem. <i>Chemistry Central Journal</i> , 2008, 2, 11.	2.6	15
107	Twin weight discordance and maternal weight gain in twin pregnancies. <i>International Journal of Gynecology and Obstetrics</i> , 2007, 96, 176-180.	2.3	11
108	A graph-theoretic modeling on GO space for biological interpretation of gene clusters. <i>Bioinformatics</i> , 2004, 20, 381-388.	4.1	65