

Jane Synnergren

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9429918/publications.pdf>

Version: 2024-02-01

44
papers

1,514
citations

331670

21
h-index

315739

38
g-index

47
all docs

47
docs citations

47
times ranked

2616
citing authors

#	ARTICLE	IF	CITATIONS
1	Multimodal deep learning for biomedical data fusion: a review. <i>Briefings in Bioinformatics</i> , 2022, 23, .	6.5	118
2	Multi-Omics Characterization of a Human Stem Cell-Based Model of Cardiac Hypertrophy. <i>Life</i> , 2022, 12, 293.	2.4	5
3	Data Mining Identifies CCN2 and THBS1 as Biomarker Candidates for Cardiac Hypertrophy. <i>Life</i> , 2022, 12, 726.	2.4	2
4	Multi-assignment clustering: Machine learning from a biological perspective. <i>Journal of Biotechnology</i> , 2021, 326, 1-10.	3.8	2
5	Unraveling the Metabolic Derangements Occurring in Non-infarcted Areas of Pig Hearts With Chronic Heart Failure. <i>Frontiers in Cardiovascular Medicine</i> , 2021, 8, 753470.	2.4	0
6	Unraveling the Metabolic Derangements Occurring in Non-infarcted Areas of Pig Hearts With Chronic Heart Failure. <i>Frontiers in Cardiovascular Medicine</i> , 2021, 8, 753470.	2.4	1
7	Transcriptional sex and regional differences in paired human atrial and ventricular cardiac biopsies collected in vivo. <i>Physiological Genomics</i> , 2020, 52, 110-120.	2.3	7
8	Models of the blood-brain barrier using iPSC-derived cells. <i>Molecular and Cellular Neurosciences</i> , 2020, 107, 103533.	2.2	44
9	Cardiac hypertrophy in a dish: A human stem cell based model. <i>Biology Open</i> , 2020, 9, .	1.2	10
10	Human Pluripotent Stem Cell-Derived Hepatocytes Show Higher Transcriptional Correlation with Adult Liver Tissue than with Fetal Liver Tissue. <i>ACS Omega</i> , 2020, 5, 4816-4827.	3.5	4
11	Characterization of Human Induced Pluripotent Stem Cell-Derived Hepatocytes with Mature Features and Potential for Modeling Metabolic Diseases. <i>International Journal of Molecular Sciences</i> , 2020, 21, 469.	4.1	14
12	Enhanced xeno-free differentiation of hiPSC-derived astroglia applied in a blood-brain barrier model. <i>Fluids and Barriers of the CNS</i> , 2019, 16, 27.	5.0	8
13	Diabetic Cardiomyopathy Modelling Using Induced Pluripotent Stem Cell Derived Cardiomyocytes: Recent Advances and Emerging Models. <i>Stem Cell Reviews and Reports</i> , 2019, 15, 13-22.	5.6	25
14	Human iPSC-Derived Astroglia from a Stable Neural Precursor State Show Improved Functionality Compared with Conventional Astrocytic Models. <i>Stem Cell Reports</i> , 2018, 10, 1030-1045.	4.8	81
15	Expression Profiling of Human Pluripotent Stem Cell-Derived Cardiomyocytes Exposed to Doxorubicin-Integration and Visualization of Multi-Omics Data. <i>Toxicological Sciences</i> , 2018, 163, 182-195.	3.1	30
16	Barrier Properties and Transcriptome Expression in Human iPSC-Derived Models of the Blood-Brain Barrier. <i>Stem Cells</i> , 2018, 36, 1816-1827.	3.2	81
17	CSF/serum albumin ratio in dementias: a cross-sectional study on 1861 patients. <i>Neurobiology of Aging</i> , 2017, 59, 1-9.	3.1	84
18	Comparative transcriptomics of hepatic differentiation of human pluripotent stem cells and adult human liver tissue. <i>Physiological Genomics</i> , 2017, 49, 430-446.	2.3	11

#	ARTICLE	IF	CITATIONS
19	Maintenance of drug metabolism and transport functions in human precision-cut liver slices during prolonged incubation for 5 days. <i>Archives of Toxicology</i> , 2017, 91, 2079-2092.	4.2	33
20	A data analysis framework for biomedical big data: Application on mesoderm differentiation of human pluripotent stem cells. <i>PLoS ONE</i> , 2017, 12, e0179613.	2.5	8
21	Highly Synchronized Expression of Lineage-Specific Genes during <i>In Vitro</i> Hepatic Differentiation of Human Pluripotent Stem Cell Lines. <i>Stem Cells International</i> , 2016, 2016, 1-22.	2.5	11
22	Towards Creating the Perfect <i>In Vitro</i> Cell Model. <i>Stem Cells International</i> , 2016, 2016, 1-2.	2.5	7
23	Comparative transcriptomic analysis identifies genes differentially expressed in human epicardial progenitors and hiPSC-derived cardiac progenitors. <i>Physiological Genomics</i> , 2016, 48, 771-784.	2.3	2
24	MicroRNAs as potential biomarkers for doxorubicin-induced cardiotoxicity. <i>Toxicology in Vitro</i> , 2016, 34, 26-34.	2.4	51
25	Interactive Visualization of Large-Scale Gene Expression Data. , 2016, , .		1
26	Identification of novel biomarkers for doxorubicin-induced toxicity in human cardiomyocytes derived from pluripotent stem cells. <i>Toxicology</i> , 2015, 328, 102-111.	4.2	71
27	Identification of stable reference genes in differentiating human pluripotent stem cells. <i>Physiological Genomics</i> , 2015, 47, 232-239.	2.3	18
28	Gene networks and transcription factor motifs defining the differentiation of stem cells into hepatocyte-like cells. <i>Journal of Hepatology</i> , 2015, 63, 934-942.	3.7	165
29	Long-Term Chronic Toxicity Testing Using Human Pluripotent Stem Cell-Derived Hepatocytes. <i>Drug Metabolism and Disposition</i> , 2014, 42, 1401-1406.	3.3	87
30	Clinical Outcome 3 Years After Autologous Chondrocyte Implantation Does Not Correlate With the Expression of a Predefined Gene Marker Set in Chondrocytes Prior to Implantation but Is Associated With Critical Signaling Pathways. <i>Orthopaedic Journal of Sports Medicine</i> , 2014, 2, 232596711455078.	1.7	13
31	Hepatic Differentiation and Maturation of Human Embryonic Stem Cells Cultured in a Perfused Three-Dimensional Bioreactor. <i>Stem Cells and Development</i> , 2013, 22, 581-594.	2.1	56
32	Comparison of human cardiac gene expression profiles in paired samples of right atrium and left ventricle collected in vivo. <i>Physiological Genomics</i> , 2012, 44, 89-98.	2.3	43
33	Global transcriptional profiling reveals similarities and differences between human stem cell-derived cardiomyocyte clusters and heart tissue. <i>Physiological Genomics</i> , 2012, 44, 245-258.	2.3	65
34	High expression of arachidonate 15-lipoxygenase and proinflammatory markers in human ischemic heart tissue. <i>Biochemical and Biophysical Research Communications</i> , 2012, 424, 327-330.	2.1	13
35	Expression of microRNAs and their target mRNAs in human stem cell-derived cardiomyocyte clusters and in heart tissue. <i>Physiological Genomics</i> , 2011, 43, 581-594.	2.3	24
36	Human Embryonic Stem Cell Derived Hepatocyte-Like Cells as a Tool for In Vitro Hazard Assessment of Chemical Carcinogenicity. <i>Toxicological Sciences</i> , 2011, 124, 278-290.	3.1	66

#	ARTICLE	IF	CITATIONS
37	Human Embryonic Mesodermal Progenitors Highly Resemble Human Mesenchymal Stem Cells and Display High Potential for Tissue Engineering Applications. <i>Tissue Engineering - Part A</i> , 2010, 16, 2161-2182.	3.1	64
38	Transcriptional Profiling of Human Embryonic Stem Cells Differentiating to Definitive and Primitive Endoderm and Further Toward the Hepatic Lineage. <i>Stem Cells and Development</i> , 2010, 19, 961-978.	2.1	17
39	Classification of information fusion methods in systems biology. <i>In Silico Biology</i> , 2009, 9, 65-76.	0.9	3
40	Molecular Signature of Cardiomyocyte Clusters Derived from Human Embryonic Stem Cells. <i>Stem Cells</i> , 2008, 26, 1831-1840.	3.2	78
41	Cardiomyogenic gene expression profiling of differentiating human embryonic stem cells. <i>Journal of Biotechnology</i> , 2008, 134, 162-170.	3.8	26
42	A data integration method for exploring gene regulatory mechanisms. , 2008, , .		0
43	Mapping of the JDL data fusion model to bioinformatics. , 2007, , .		5
44	Differentiating Human Embryonic Stem Cells Express a Unique Housekeeping Gene Signature. <i>Stem Cells</i> , 2007, 25, 473-480.	3.2	58