Benjamin D Cosgrove

List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/8217158/benjamin-d-cosgrove-publications-by-year.pdf

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

31 1,617 17 38 g-index

38 2,083 10 4.77 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
31	Large-scale integration of single-cell transcriptomic data captures transitional progenitor states in mouse skeletal muscle regeneration. <i>Communications Biology</i> , 2021 , 4, 1280	6.7	9
30	High-resolution single-cell transcriptomics reveals heterogeneity of self-renewing hair follicle stem cells. <i>Experimental Dermatology</i> , 2021 , 30, 457-471	4	4
29	Extracellular serine and glycine are required for mouse and human skeletal muscle stem and progenitor cell function. <i>Molecular Metabolism</i> , 2021 , 43, 101106	8.8	10
28	Single-Cell Analysis of the Muscle Stem Cell Hierarchy Identifies Heterotypic Communication Signals Involved in Skeletal Muscle Regeneration. <i>Cell Reports</i> , 2020 , 30, 3583-3595.e5	10.6	84
27	A reference single-cell transcriptomic atlas of human skeletal muscle tissue reveals bifurcated muscle stem cell populations. <i>Skeletal Muscle</i> , 2020 , 10, 19	5.1	39
26	Cells expressing PAX8 are the main source of homeostatic regeneration of adult mouse endometrial epithelium and give rise to serous endometrial carcinoma. <i>DMM Disease Models and Mechanisms</i> , 2020 , 13,	4.1	11
25	A defined N6-methyladenosine (mA) profile conferred by METTL3 regulates muscle stem cell/myoblast state transitions. <i>Cell Death Discovery</i> , 2020 , 6, 95	6.9	21
24	Single-cell transcriptomic analysis identifies extensive heterogeneity in the cellular composition of mouse Achilles tendons. <i>American Journal of Physiology - Cell Physiology</i> , 2020 , 319, C885-C894	5.4	22
23	Isolation, Culture, Characterization, and Differentiation of Human Muscle Progenitor Cells from the Skeletal Muscle Biopsy Procedure. <i>Journal of Visualized Experiments</i> , 2019 ,	1.6	8
22	How Biophysical Forces Regulate Human B Cell Lymphomas. <i>Cell Reports</i> , 2018 , 23, 499-511	10.6	19
21	High-resolution myogenic lineage mapping by single-cell mass cytometry. <i>Nature Cell Biology</i> , 2017 , 19, 558-567	23.4	79
20	Data-Modeling Identifies Conflicting Signaling Axes Governing Myoblast Proliferation and Differentiation Responses to Diverse Ligand Stimuli. <i>Cellular and Molecular Bioengineering</i> , 2017 , 10, 433-450	3.9	2
19	Microcontact-Printed Hydrogel Microwell Arrays for Clonal Muscle Stem Cell Cultures. <i>Methods in Molecular Biology</i> , 2017 , 1668, 75-92	1.4	4
18	Injectable biomimetic liquid crystalline scaffolds enhance muscle stem cell transplantation. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E7919-E7928	8 ^{11.5}	59
17	The central role of muscle stem cells in regenerative failure with aging. <i>Nature Medicine</i> , 2015 , 21, 854-	63 0.5	247
16	Rejuvenation of the muscle stem cell population restores strength to injured aged muscles. <i>Nature Medicine</i> , 2014 , 20, 255-64	50.5	439
15	Skeletal Muscle Stem Cells 2011 , 347-363		

LIST OF PUBLICATIONS

14	Networks inferred from biochemical data reveal profound differences in toll-like receptor and inflammatory signaling between normal and transformed hepatocytes. <i>Molecular and Cellular Proteomics</i> , 2010 , 9, 1849-65	7.6	92
13	Cytokine-associated drug toxicity in human hepatocytes is associated with signaling network dysregulation. <i>Molecular BioSystems</i> , 2010 , 6, 1195-206		52
12	Model Convolution: A Computational Approach to Digital Image Interpretation. <i>Cellular and Molecular Bioengineering</i> , 2010 , 3, 163-170	3.9	24
11	A multipathway phosphoproteomic signaling network model of idiosyncratic drug- and inflammatory cytokine-induced toxicity in human hepatocytes. <i>Annual International Conference of the IEEE Engineering in Medicine and Biology Society IEEE Engineering in Medicine and Biology Society</i>	0.9	5
10	Three-kinase inhibitor combination recreates multipathway effects of a geldanamycin analogue on hepatocellular carcinoma cell death. <i>Molecular Cancer Therapeutics</i> , 2009 , 8, 2183-92	6.1	15
9	Synergistic drug-cytokine induction of hepatocellular death as an in vitro approach for the study of inflammation-associated idiosyncratic drug hepatotoxicity. <i>Toxicology and Applied Pharmacology</i> , 2009 , 237, 317-30	4.6	115
8	A home away from home: challenges and opportunities in engineering in vitro muscle satellite cell niches. <i>Differentiation</i> , 2009 , 78, 185-94	3.5	98
7	Microfluidic concentration-enhanced cellular kinase activity assay. <i>Journal of the American Chemical Society</i> , 2009 , 131, 10340-1	16.4	62
6	Fusing Tissue Engineering and Systems Biology Toward Fulfilling Their Promise. <i>Cellular and Molecular Bioengineering</i> , 2008 , 1, 33-41	3.9	18
5	An inducible autocrine cascade regulates rat hepatocyte proliferation and apoptosis responses to tumor necrosis factor-alpha. <i>Hepatology</i> , 2008 , 48, 276-88	11.2	63
4	Single-cell analysis of the muscle stem cell hierarchy identifies heterotypic communication signals involved in skeletal muscle regeneration		3
3	A reference single-cell transcriptomic atlas of human skeletal muscle tissue reveals bifurcated muscle stem cell populations		2
2	Strength in numbers: Large-scale integration of single-cell transcriptomic data reveals rare, transient muscle progenitor cell states in muscle regeneration		1
1	Single-cell Transcriptomics Identify Extensive Heterogeneity in the Cellular Composition of Mouse Achilles Tendons		8