Yoonsung Lee

List of Publications by Year in descending order

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759233 794594 19 896 12 19 h-index citations g-index papers 20 20 20 1178 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Fgf signaling instructs position-dependent growth rate during zebrafish fin regeneration. Development (Cambridge), 2005, 132, 5173-5183.	2.5	300
2	Reiterative roles for FGF signaling in the establishment of size and proportion of the zebrafish heart. Developmental Biology, 2008, 321, 397-406.	2.0	113
3	Maintenance of blastemal proliferation by functionally diverse epidermis in regenerating zebrafish fins. Developmental Biology, 2009, 331, 270-280.	2.0	90
4	ADictyosteliumHomologue of WASP Is Required for Polarized F-Actin Assembly during Chemotaxis. Molecular Biology of the Cell, 2005, 16, 2191-2206.	2.1	75
5	Neuronal Regulation of the Spatial Patterning of Neurogenesis. Developmental Cell, 2010, 18, 136-147.	7.0	75
6	FGF signalling restricts haematopoietic stem cell specification via modulation of the BMP pathway. Nature Communications, 2014, 5, 5588.	12.8	45
7	Restriction of hepatic competence by Fgf signaling. Development (Cambridge), 2011, 138, 1339-1348.	2.5	38
8	FGF signalling specifies haematopoietic stem cells through its regulation of somitic Notch signalling. Nature Communications, 2014, 5, 5583.	12.8	37
9	Cobll1 is linked to drug resistance and blastic transformation in chronic myeloid leukemia. Leukemia, 2017, 31, 1532-1539.	7.2	22
10	Heterozygous variants in <i>MYBPC1</i> are associated with an expanded neuromuscular phenotype beyond arthrogryposis. Human Mutation, 2019, 40, 1115-1126.	2.5	19
11	Labelâ€free optical projection tomography for quantitative threeâ€dimensional anatomy of mouse embryo. Journal of Biophotonics, 2019, 12, e201800481.	2.3	16
12	Ras controls melanocyte expansion during zebrafish fin stripe regeneration. DMM Disease Models and Mechanisms, 2010, 3, 496-503.	2.4	14
13	Large-scale generation and phenotypic characterization of zebrafish CRISPR mutants of DNA repair genes. DNA Repair, 2021, 107, 103173.	2.8	13
14	<scp><i>FAM213A</i></scp> is linked to prognostic significance in acute myeloid leukemia through regulation of oxidative stress and myelopoiesis. Hematological Oncology, 2020, 38, 381-389.	1.7	10
15	Haematopoietic stem cell-dependent Notch transcription is mediated by p53 through the Histone chaperone Supt16h. Nature Cell Biology, 2020, 22, 1411-1422.	10.3	9
16	FRZB as a key molecule in abdominal aortic aneurysm progression affecting vascular integrity. Bioscience Reports, 2021, 41, .	2.4	8
17	AML poor prognosis factor, TPD52, is associated with the maintenance of haematopoietic stem cells through regulation of cell proliferation. Journal of Cellular Biochemistry, 2021, 122, 403-412.	2.6	6
18	Role of kif2c, A Gene Related to ALL Relapse, in Embryonic Hematopoiesis in Zebrafish. International Journal of Molecular Sciences, 2020, 21, 3127.	4.1	4

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#	Article	IF	CITATIONS
19	Reciprocal interactions among Cobll1, PACSIN2, and SH3BP1 regulate drug resistance in chronic myeloid leukemia. Cancer Medicine, 2022, , .	2.8	2