Laurentijn Tilleman

List of Publications by Year in descending order

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Version: 2024-02-01

933447 713466 26 536 10 21 citations g-index h-index papers 30 30 30 963 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	The transcriptome of Mycobacterium tuberculosis in a lipid-rich dormancy model through RNAseq analysis. Scientific Reports, 2017, 7, 17665.	3.3	88
2	Direct comparison of distinct naive pluripotent states in human embryonic stem cells. Nature Communications, 2017, 8, 15055.	12.8	60
3	Performance of four modern whole genome amplification methods for copy number variant detection in single cells. Scientific Reports, 2017, 7, 3422.	3.3	54
4	A multi-omics analysis of the grapevine pathogen Lasiodiplodia theobromae reveals that temperature affects the expression of virulence- and pathogenicity-related genes. Scientific Reports, 2019, 9, 13144.	3.3	47
5	Gain of 20q11.21 in Human Pluripotent Stem Cells Impairs TGF-Î ² -Dependent Neuroectodermal Commitment. Stem Cell Reports, 2019, 13, 163-176.	4.8	39
6	The transcription factor ETS1 is an important regulator of human NK cell development and terminal differentiation. Blood, 2020, 136, 288-298.	1.4	33
7	Dual RNA Sequencing of Vitis vinifera during Lasiodiplodia theobromae Infection Unveils Host–Pathogen Interactions. International Journal of Molecular Sciences, 2019, 20, 6083.	4.1	28
8	HES1 and HES4 have non-redundant roles downstream of Notch during early human T-cell development. Haematologica, 2020, 106, 130-141.	3.5	20
9	Myeloid-specific IRE1alpha deletion reduces tumour development in a diabetic, non-alcoholic steatohepatitis-induced hepatocellular carcinoma mouse model. Metabolism: Clinical and Experimental, 2020, 107, 154220.	3.4	19
10	RRM2 enhances MYCN-driven neuroblastoma formation and acts as a synergistic target with CHK1 inhibition. Science Advances, 2022, 8, .	10.3	15
11	Contemporary pharmacogenetic assays in view of the PharmGKB database. Pharmacogenomics, 2019, 20, 261-272.	1.3	13
12	Uncovering low-level mosaicism in human embryonic stem cells using high throughput single cell shallow sequencing. Scientific Reports, 2019, 9, 14844.	3.3	12
13	Marine biogenics in sea spray aerosols interact with the mTOR signaling pathway. Scientific Reports, 2019, 9, 675.	3.3	12
14	Comparative study of preimplantation development following distinct assisted oocyte activation protocols in a PLC-zeta knockout mouse model. Molecular Human Reproduction, 2020, 26, 801-815.	2.8	11
15	Activin A-derived human embryonic stem cells show increased competence to differentiate into primordial germ cell-like cells. Stem Cells, 2021, 39, 551-563.	3.2	11
16	Maternal Recognition of Pregnancy in the Horse: Are MicroRNAs the Secret Messengers?. International Journal of Molecular Sciences, 2020, 21, 419.	4.1	10
17	YtrASa, a GntR-Family Transcription Factor, Represses Two Genetic Loci Encoding Membrane Proteins in Sulfolobus acidocaldarius. Frontiers in Microbiology, 2019, 10, 2084.	3.5	9
18	WNT Inhibition and Increased FGF Signaling Promotes Derivation of Less Heterogeneous Primed Human Embryonic Stem Cells, Compatible with Differentiation. Stem Cells and Development, 2019, 28, 579-592.	2.1	9

#	Article	IF	CITATIONS
19	STR profiling and Copy Number Variation analysis on single, preserved cells using current Whole Genome Amplification methods. Scientific Reports, 2017, 7, 17189.	3.3	8
20	Metallothioneins alter macrophage phenotype and represent novel therapeutic targets for acetaminophen-induced liver injury. Journal of Leukocyte Biology, 2021, 111, 123-133.	3.3	8
21	Kinship analysis on single cells after whole genome amplification. Scientific Reports, 2020, 10, 14647.	3.3	6
22	Endogenous suppression of WNT signalling in human embryonic stem cells leads to low differentiation propensity towards definitive endoderm. Scientific Reports, 2021, 11, 6137.	3.3	6
23	Transcriptional landscape changes during human embryonic stem cell derivation. Molecular Human Reproduction, 2018, 24, 543-555.	2.8	5
24	Comparative genomics of Flavobacterium columnare unveils novel insights in virulence and antimicrobial resistance mechanisms. Veterinary Research, 2021, 52, 18.	3.0	5
25	Pan-cancer pharmacogenetics: targeted sequencing panels or exome sequencing?. Pharmacogenomics, 2020, 21, 1073-1084.	1.3	3
26	GATA-3 Suppresses IRF8 to promote human T-cell lineage commitment. Experimental Hematology, 2017, 53, S83.	0.4	0