Ruben Boers

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4012793/publications.pdf

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		1684188	1474206
9	157	5	9
papers	citations	h-index	g-index
10 all docs	10 docs citations	10 times ranked	201 citing authors

#	Article	lF	CITATIONS
1	Genome-wide DNA methylation profiling using the methylation-dependent restriction enzyme LpnPl. Genome Research, 2018, 28, 88-99.	5.5	54
2	Rapid in vitro generation of bona fide exhausted CD8+ T cells is accompanied by Tcf7Âpromotor methylation. PLoS Pathogens, 2020, 16, e1008555.	4.7	36
3	CTCF chromatin residence time controls three-dimensional genome organization, gene expression and DNA methylation in pluripotent cells. Nature Cell Biology, 2021, 23, 881-893.	10.3	30
4	Genome-wide aberrant methylation in primary metastatic UM and their matched metastases. Scientific Reports, 2022, 12, 42.	3.3	11
5	Differentially Methylated Regions in Desmoid-Type Fibromatosis: A Comparison Between CTNNB1 S45F and T41A Tumors. Frontiers in Oncology, 2020, 10, 565031.	2.8	7
6	Characterization of Histone Modifications Associated with Inactive X-Chromosome in Trophoblast Stem Cells, eXtra-Embryonic Endoderm Cells and in In Vitro Derived Undifferentiated and Differentiated Epiblast Like Stem Cells. PLoS ONE, 2016, 11, e0167154.	2.5	7
7	Genome wide DNA methylation analysis of alveolar capillary dysplasia lung tissue reveals aberrant methylation of genes involved in development including the FOXF1 locus. Clinical Epigenetics, 2021, 13, 148.	4.1	6
8	MicroRNA expression and DNA methylation profiles do not distinguish between primary and recurrent well-differentiated liposarcoma. PLoS ONE, 2020, 15, e0228014.	2.5	3
9	Kidney Organoids Are Capable of Forming Tumors, but Not Teratomas. Stem Cells, 2022, 40, 577-591.	3.2	3