

Davide Pellacani

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2644897/publications.pdf>

Version: 2024-02-01

20
papers

948
citations

567281

15
h-index

794594

19
g-index

20
all docs

20
docs citations

20
times ranked

2968
citing authors

#	ARTICLE	IF	CITATIONS
1	Mammary epithelial cells have lineage-rooted metabolic identities. <i>Nature Metabolism</i> , 2021, 3, 665-681.	11.9	24
2	Initiation of human mammary cell tumorigenesis by mutant KRAS requires YAP inactivation. <i>Oncogene</i> , 2020, 39, 1957-1968.	5.9	18
3	MYC-induced human acute myeloid leukemia requires a continuing IL-3/GM-CSF costimulus. <i>Blood</i> , 2020, 136, 2764-2773.	1.4	15
4	A topological view of human CD34+ cell state trajectories from integrated single-cell output and proteomic data. <i>Blood</i> , 2019, 133, 927-939.	1.4	17
5	Transcriptional regulation of normal human mammary cell heterogeneity and its perturbation in breast cancer. <i>EMBO Journal</i> , 2019, 38, e100330.	7.8	35
6	Growth Factor-Dependent Activation of a MYC-Induced Latent AML Program in Human Hematopoietic Cells. <i>Blood</i> , 2019, 134, 2533-2533.	1.4	0
7	Phenotype-independent DNA methylation changes in prostate cancer. <i>British Journal of Cancer</i> , 2018, 119, 1133-1143.	6.4	14
8	Single-cell analysis identifies a CD33+ subset of human cord blood cells with high regenerative potential. <i>Nature Cell Biology</i> , 2018, 20, 710-720.	10.3	36
9	Dissociation of Survival, Proliferation, and State Control in Human Hematopoietic Stem Cells. <i>Stem Cell Reports</i> , 2017, 8, 152-162.	4.8	22
10	Distinct signaling programs control human hematopoietic stem cell survival and proliferation. <i>Blood</i> , 2017, 129, 307-318.	1.4	35
11	Mass Cytometric Analysis Reveals Viable Activated Caspase-3+ Luminal Progenitors in the Normal Adult Human Mammary Gland. <i>Cell Reports</i> , 2017, 21, 1116-1126.	6.4	20
12	Fate mapping of human glioblastoma reveals an invariant stem cell hierarchy. <i>Nature</i> , 2017, 549, 227-232.	27.8	321
13	Analysis of Normal Human Mammary Epigenomes Reveals Cell-Specific Active Enhancer States and Associated Transcription Factor Networks. <i>Cell Reports</i> , 2016, 17, 2060-2074.	6.4	90
14	Barcoding reveals complex clonal dynamics of de novo transformed human mammary cells. <i>Nature</i> , 2015, 528, 267-271.	27.8	101
15	MicroRNA Expression Profile of Primary Prostate Cancer Stem Cells as a Source of Biomarkers and Therapeutic Targets. <i>European Urology</i> , 2015, 67, 7-10.	1.9	61
16	Conserved Two-Step Regulatory Mechanism of Human Epithelial Differentiation. <i>Stem Cell Reports</i> , 2014, 2, 180-188.	4.8	18
17	Differential Cytotoxic Activity of a Novel Palladium-Based Compound on Prostate Cell Lines, Primary Prostate Epithelial Cells and Prostate Stem Cells. <i>PLoS ONE</i> , 2013, 8, e64278.	2.5	35
18	Retinoic acid and androgen receptors combine to achieve tissue specific control of human prostatic transglutaminase expression: a novel regulatory network with broader significance. <i>Nucleic Acids Research</i> , 2012, 40, 4825-4840.	14.5	26

#	ARTICLE	IF	CITATIONS
19	Prostate cancer stem cells: Are they androgen-responsive?. <i>Molecular and Cellular Endocrinology</i> , 2012, 360, 14-24.	3.2	37
20	Development and limitations of lentivirus vectors as tools for tracking differentiation in prostate epithelial cells. <i>Experimental Cell Research</i> , 2010, 316, 3161-3171.	2.6	23