

Stefan Kurtenbach

List of Publications by Year in descending order

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Version: 2024-02-01

43
papers

1,725
citations

430874

18
h-index

414414

32
g-index

49
all docs

49
docs citations

49
times ranked

2802
citing authors

#	ARTICLE	IF	CITATIONS
1	Single-cell analysis reveals new evolutionary complexity in uveal melanoma. <i>Nature Communications</i> , 2020, 11, 496.	12.8	268
2	PRAME as an Independent Biomarker for Metastasis in Uveal Melanoma. <i>Clinical Cancer Research</i> , 2016, 22, 1234-1242.	7.0	205
3	Single-cell analysis of olfactory neurogenesis and differentiation in adult humans. <i>Nature Neuroscience</i> , 2020, 23, 323-326.	14.8	165
4	Punctuated evolution of canonical genomic aberrations in uveal melanoma. <i>Nature Communications</i> , 2018, 9, 116.	12.8	144
5	Gain of function of ASXL1 truncating protein in the pathogenesis of myeloid malignancies. <i>Blood</i> , 2018, 131, 328-341.	1.4	133
6	Pannexin1 Stabilizes Synaptic Plasticity and Is Needed for Learning. <i>PLoS ONE</i> , 2012, 7, e51767.	2.5	121
7	Tmem16b is Specifically Expressed in the Cilia of Olfactory Sensory Neurons. <i>Chemical Senses</i> , 2010, 35, 239-245.	2.0	94
8	Epigenetic reprogramming and aberrant expression of PRAME are associated with increased metastatic risk in Class 1 and Class 2 uveal melanomas. <i>Oncotarget</i> , 2016, 7, 59209-59219.	1.8	94
9	BAP1 regulates epigenetic switch from pluripotency to differentiation in developmental lineages giving rise to BAP1-mutant cancers. <i>Science Advances</i> , 2019, 5, eaax1738.	10.3	57
10	Gap junction modulation and its implications for heart function. <i>Frontiers in Physiology</i> , 2014, 5, 82.	2.8	44
11	BAP1 Loss Is Associated with DNA Methylomic Repatterning in Highly Aggressive Class 2 Uveal Melanomas. <i>Clinical Cancer Research</i> , 2019, 25, 5663-5673.	7.0	41
12	Pannexin1 Channel Proteins in the Zebrafish Retina Have Shared and Unique Properties. <i>PLoS ONE</i> , 2013, 8, e77722.	2.5	41
13	Sleep-wakefulness cycle and behavior in pannexin1 knockout mice. <i>Behavioural Brain Research</i> , 2017, 318, 24-27.	2.2	35
14	Olfaction in Three Genetic and Two MPTP-Induced Parkinson's Disease Mouse Models. <i>PLoS ONE</i> , 2013, 8, e77509.	2.5	32
15	The smelling of Hedione results in sex-differentiated human brain activity. <i>NeuroImage</i> , 2015, 113, 365-373.	4.2	27
16	Investigation of olfactory function in a Panx1 knock out mouse model. <i>Frontiers in Cellular Neuroscience</i> , 2014, 8, 266.	3.7	23
17	Dual Screen for Efficacy and Toxicity Identifies HDAC Inhibitor with Distinctive Activity Spectrum for BAP1-Mutant Uveal Melanoma. <i>Molecular Cancer Research</i> , 2021, 19, 215-222.	3.4	21
18	Odorant-Dependent Generation of Nitric Oxide in Mammalian Olfactory Sensory Neurons. <i>PLoS ONE</i> , 2009, 4, e5499.	2.5	21

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19	Impact of the Usher syndrome on olfaction. <i>Human Molecular Genetics</i> , 2016, 25, 524-533.	2.9	19
20	Emerging functions of pannexin 1 in the eye. <i>Frontiers in Cellular Neuroscience</i> , 2014, 8, 263.	3.7	17
21	Uphylotplot2: visualizing phylogenetic trees from single-cell RNA-seq data. <i>BMC Genomics</i> , 2021, 22, 419.	2.8	17
22	Molecular evolution of a chordate specific family of G protein-coupled receptors. <i>BMC Evolutionary Biology</i> , 2011, 11, 234.	3.2	16
23	A Potential Compensatory Role of Panx3 in the VNO of a Panx1 Knock Out Mouse Model. <i>Frontiers in Molecular Neuroscience</i> , 2018, 11, 135.	2.9	15
24	The BEACH Protein LRBA Promotes the Localization of the Heterotrimeric G-protein Golf to Olfactory Cilia. <i>Scientific Reports</i> , 2017, 7, 8409.	3.3	10
25	ChIPprimersDB: a public repository of verified qPCR primers for chromatin immunoprecipitation (ChIP). <i>Nucleic Acids Research</i> , 2019, 47, D46-D49.	14.5	9
26	Isolation, culture optimization and functional characterization of stem cell neurospheres from mouse neonatal olfactory bulb and epithelium. <i>European Archives of Oto-Rhino-Laryngology</i> , 2017, 274, 3071-3085.	1.6	7
27	Establishment and Characterization of a Novel Human Ocular Adnexal Sebaceous Carcinoma Cell Line. <i>Translational Vision Science and Technology</i> , 2021, 10, 34.	2.2	4
28	PieParty: visualizing cells from scRNA-seq data as pie charts. <i>Life Science Alliance</i> , 2021, 4, e202000986.	2.8	2
29	Abstract 4861: The role of PRAME in promoting uveal melanoma metastasis. <i>Cancer Research</i> , 2017, 77, 4861-4861.	0.9	2
30	Array data extractor (ADE): a LabVIEW program to extract and merge gene array data. <i>BMC Research Notes</i> , 2013, 6, 496.	1.4	1
31	Abstract 1541: The tumor suppressor BAP1 promotes a developmental switch from pluripotency to differentiation. <i>Cancer Research</i> , 2017, 77, 1541-1541.	0.9	1
32	Abstract 4025: New candidate therapy for BAP1-mutant cancer identified using novel screen. , 2020, , .		1
33	Abstract 3027: Role of BCOR in retinoblastoma. , 2021, , .		0
34	Abstract 2764: Mechanisms of genomic-microenvironmental interactions in uveal melanoma. , 2021, , .		0
35	Abstract 4348: Methylation analysis of uveal melanoma reveals definitive patterns in tumors harboring BAP1 mutations. , 2017, , .		0
36	Abstract 794: Potential role of DLL4 in uveal melanoma vascular mimicry. , 2017, , .		0

#	ARTICLE	IF	CITATIONS
37	Abstract 5369: Epigenetic, transcriptomic and ubiquitomic changes associated with BAP1 loss in uveal melanoma. , 2017, , .		0
38	Abstract 3390: Clonal evolution in uveal melanoma. , 2017, , .		0
39	Abstract 5199: A novel role for BAP1 in development and tumor suppression. , 2019, , .		0
40	Abstract 4244: Novel expressed long non-coding RNAs in uveal melanoma. , 2019, , .		0
41	Abstract 1591: Single cell analysis of uveal melanoma reveals new evolutionary complexity. , 2020, , .		0
42	Abstract 2737: PRAME misexpression alters protein ubiquitination and leads to increased invasion and metastasis in uveal melanoma. , 2019, , .		0
43	Abstract 855: Analysis of canonical uveal melanoma mutations reveals novel signaling effects. Cancer Research, 2022, 82, 855-855.	0.9	0