## Sylvain Baulande

List of Publications by Year in descending order

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147801 149698 3,700 59 31 56 citations h-index g-index papers 67 67 67 6940 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Single-Cell Analysis Reveals Fibroblast Clusters Linked to Immunotherapy Resistance in Cancer. Cancer Discovery, 2020, 10, 1330-1351.	9.4	424
2	Heterogeneity of neuroblastoma cell identity defined by transcriptional circuitries. Nature Genetics, 2017, 49, 1408-1413.	21.4	331
3	Aryl Hydrocarbon Receptor Controls Monocyte Differentiation into Dendritic Cells versus Macrophages. Immunity, 2017, 47, 582-596.e6.	14.3	282
4	Adiponutrin, a Transmembrane Protein Corresponding to a Novel Dietary- and Obesity-linked mRNA Specifically Expressed in the Adipose Lineage. Journal of Biological Chemistry, 2001, 276, 33336-33344.	3.4	206
5	Tissue-resident FOLR2+ macrophages associate with CD8+ TÂcell infiltration in human breast cancer. Cell, 2022, 185, 1189-1207.e25.	28.9	166
6	Human in vivo-generated monocyte-derived dendritic cells and macrophages cross-present antigens through a vacuolar pathway. Nature Communications, 2018, 9, 2570.	12.8	157
7	FLASH Irradiation Spares Lung Progenitor Cells and Limits the Incidence of Radio-induced Senescence. Clinical Cancer Research, 2020, 26, 1497-1506.	7.0	148
8	CD44 regulates epigenetic plasticity by mediating iron endocytosis. Nature Chemistry, 2020, 12, 929-938.	13.6	132
9	Whole-Exome Sequencing of Cell-Free DNA Reveals Temporo-spatial Heterogeneity and Identifies Treatment-Resistant Clones in Neuroblastoma. Clinical Cancer Research, 2018, 24, 939-949.	7.0	127
10	Clonally Expanded T Cells Reveal Immunogenicity of Rhabdoid Tumors. Cancer Cell, 2019, 36, 597-612.e8.	16.8	100
11	Transcriptional Programs Define Intratumoral Heterogeneity of Ewing Sarcoma at Single-Cell Resolution. Cell Reports, 2020, 30, 1767-1779.e6.	6.4	96
12	Whole mitochondrial genome screening in maternally inherited non-syndromic hearing impairment using a microarray resequencing mitochondrial DNA chip. European Journal of Human Genetics, 2007, 15, 1145-1155.	2.8	85
13	Contribution of resident and circulating precursors to tumor-infiltrating CD8 <sup>+</sup> T cell populations in lung cancer. Science Immunology, 2021, 6, .	11.9	82
14	PLK1 inhibition exhibits strong anti-tumoral activity in CCND1-driven breast cancer metastases with acquired palbociclib resistance. Nature Communications, 2020, 11, 4053.	12.8	77
15	Analysis of the Molecular Dialogue Between Gray Mold ( <i>Botrytis cinerea</i> ) and Grapevine ( <i>Vitis vinifera</i> ) Reveals a Clear Shift in Defense Mechanisms During Berry Ripening. Molecular Plant-Microbe Interactions, 2015, 28, 1167-1180.	2.6	73
16	Combined transcript, proteome, and metabolite analysis of transgenic maize seeds engineered for enhanced carotenoid synthesis reveals pleotropic effects in core metabolism. Journal of Experimental Botany, 2015, 66, 3141-3150.	4.8	65
17	Human lymphoid organ cDC2 and macrophages play complementary roles in T follicular helper responses. Journal of Experimental Medicine, 2019, 216, 1561-1581.	8.5	63
18	Differential Endothelial Transcriptomics Identifies Semaphorin 3G as a Vascular Class 3 Semaphorin. Arteriosclerosis, Thrombosis, and Vascular Biology, 2011, 31, 151-159.	2.4	60

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19	The bipartite TAD organization of the X-inactivation center ensures opposing developmental regulation of Tsix and Xist. Nature Genetics, 2019, 51, 1024-1034.	21.4	60
20	Cacnb4 directly couples electrical activity to gene expression, a process defective in juvenile epilepsy. EMBO Journal, 2012, 31, 3730-3744.	7.8	57
21	An element within the 5′ untranslated region of humanHsp70mRNA which acts as a general enhancer of mRNA translation. FEBS Journal, 2001, 268, 1908-1917.	0.2	55
22	A Conserved Noncoding Locus Regulates Random Monoallelic Xist Expression across a Topological Boundary. Molecular Cell, 2020, 77, 352-367.e8.	9.7	48
23	STAG2 mutations alter CTCF-anchored loop extrusion, reduce cis-regulatory interactions and EWSR1-FLI1 activity in Ewing sarcoma. Cancer Cell, 2021, 39, 810-826.e9.	16.8	48
24	Evolutionary Routes in Metastatic Uveal Melanomas Depend on <i>MBD4</i> Alterations. Clinical Cancer Research, 2019, 25, 5513-5524.	7.0	46
25	Activated ALK signals through the ERK–ETV5–RET pathway to drive neuroblastoma oncogenesis. Oncogene, 2018, 37, 1417-1429.	5.9	45
26	A high-risk retinoblastoma subtype with stemness features, dedifferentiated cone states and neuronal/ganglion cell gene expression. Nature Communications, 2021, 12, 5578.	12.8	45
27	Human papilloma virus (HPV) integration signature in Cervical Cancer: identification of MACROD2 gene as HPV hot spot integration site. British Journal of Cancer, 2021, 124, 777-785.	6.4	44
28	H3K27me3 conditions chemotolerance in triple-negative breast cancer. Nature Genetics, 2022, 54, 459-468.	21.4	44
29	Follicular lymphoma triggers phenotypic and functional remodeling of the human lymphoid stromal cell landscape. Immunity, 2021, 54, 1788-1806.e7.	14.3	43
30	Circulating tumor DNA analysis enables molecular characterization of pediatric renal tumors at diagnosis. International Journal of Cancer, 2019, 144, 68-79.	5.1	37
31	A large collection of integrated genomically characterized patientâ€derived xenografts highlighting the heterogeneity of tripleâ€negative breast cancer. International Journal of Cancer, 2019, 145, 1902-1912.	5.1	37
32	Frequency and Prognostic Impact of <i>ALK</i> Amplifications and Mutations in the European Neuroblastoma Study Group (SIOPEN) High-Risk Neuroblastoma Trial (HR-NBL1). Journal of Clinical Oncology, 2021, 39, 3377-3390.	1.6	30
33	Sulfiredoxin Protects Mice from Lipopolysaccharide-Induced Endotoxic Shock. Antioxidants and Redox Signaling, 2011, 14, 2071-2080.	5.4	29
34	Trajectory and uniqueness of mutational signatures in yeast mutators. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 24947-24956.	7.1	29
35	Gene Expression Signature for Spontaneous Cancer Regression in Melanoma Pigs. Neoplasia, 2008, 10, 714-IN5.	5.3	27
36	Detection of TRIM32 deletions in LGMD patients analyzed by a combined strategy of CGH array and massively parallel sequencing. European Journal of Human Genetics, 2015, 23, 929-934.	2.8	21

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37	Deciphering the Mammary Stem Cell Niche: A Role for Laminin-Binding Integrins. Stem Cell Reports, 2019, 12, 831-844.	4.8	20
38	Mechanistic Signatures of Human Papillomavirus Insertions in Anal Squamous Cell Carcinomas. Cancers, 2019, 11, 1846.	3.7	19
39	A novel 3D culture model recapitulates primary FL B-cell features and promotes their survival. Blood Advances, 2021, 5, 5372-5386.	5.2	18
40	TNFSF10/TRAIL regulates human T4 effector memory lymphocyte radiosensitivity and predicts radiation-induced acute and subacute dermatitis. Oncotarget, 2016, 7, 21416-21427.	1.8	16
41	Study of chromatin remodeling genes implicates SMARCA4 as a putative player in oncogenesis in neuroblastoma. International Journal of Cancer, 2019, 145, 2781-2791.	5.1	16
42	Identification of Tissue of Origin and Guided Therapeutic Applications in Cancers of Unknown Primary Using Deep Learning and RNA Sequencing (TransCUPtomics). Journal of Molecular Diagnostics, 2021, 23, 1380-1392.	2.8	15
43	Oncogenic chimeric transcription factors drive tumor-specific transcription, processing, and translation of silent genomic regions. Molecular Cell, 2022, 82, 2458-2471.e9.	9.7	14
44	Biological effect of human serum collected before and after oral intake of Pygeum africanum on various benign prostate cell cultures. Asian Journal of Andrology, 2012, 14, 499-504.	1.6	13
45	PD-L1 and ICOSL discriminate human Secretory and Helper dendritic cells in cancer, allergy and autoimmunity. Nature Communications, 2022, 13, 1983.	12.8	12
46	ERG transcription factors have a splicing regulatory function involving RBFOX2 that is altered in the EWS-FLI1 oncogenic fusion. Nucleic Acids Research, 2021, 49, 5038-5056.	14.5	11
47	CD8+T cell responsiveness to anti-PD-1 is epigenetically regulated by Suv39h1 in melanomas. Nature Communications, 2022, 13, .	12.8	11
48	Tetratricopeptide repeat domain 7A is a nuclear factor that modulates transcription and chromatin structure. Cell Discovery, 2018, 4, 61.	6.7	10
49	Definition of Biologically Distinct Groups of Conjunctival Melanomas According to Etiological Factors and Implications for Precision Medicine. Cancers, 2021, 13, 3836.	3.7	10
50	Detection and quantification of extracellular microRNAs in medulloblastoma. Journal of Cancer Metastasis and Treatment, 2015, 1, 67.	0.8	10
51	Kronos scRT: a uniform framework for single-cell replication timing analysis. Nature Communications, 2022, 13, 2329.	12.8	9
52	Molecular diagnosis of retinoblastoma by circulating tumor DNA analysis. European Journal of Cancer, 2021, 154, 277-287.	2.8	7
53	Human papilloma virus integration sites and genomic signatures in head and neck squamous cell carcinoma. Molecular Oncology, 2022, 16, 3001-3016.	4.6	7
54	Decentralization of Next-Generation RNA Sequencing-Based MammaPrint® and BluePrint® Kit at University Hospitals Leuven and Curie Institute Paris. Translational Oncology, 2019, 12, 1557-1565.	3.7	6

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55	ALK mutation dynamics and clonal evolution in a neuroblastoma model exhibiting two ALK mutations. Oncotarget, 2019, 10, 4937-4950.	1.8	5
56	Recombination in a sterile polyploid hybrid yeast upon meiotic Return-To-Growth. Microbiological Research, 2021, 250, 126789.	5.3	5
57	Lack of evidence for CDK12 as an ovarian cancer predisposing gene. Familial Cancer, 2020, 19, 203-209.	1.9	1
58	Clinical and Molecular Update on the Fourth Reported Family with Hamamy Syndrome. Molecular Syndromology, 2021, 12, 342-350.	0.8	1
59	BET and CDK Inhibition Reveal Differences in the Proliferation Control of Sympathetic Ganglion Neuroblasts and Adrenal Chromaffin Cells. Cancers, 2022, 14, 2755.	3.7	1