Isabelle Janoueix-Lerosey

List of Publications by Year in descending order

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87723 66788 7,592 78 38 citations h-index papers

g-index 82 82 82 10509 docs citations times ranked citing authors all docs

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#	Article	IF	Citations
1	Control-FREEC: a tool for assessing copy number and allelic content using next-generation sequencing data. Bioinformatics, 2012, 28, 423-425.	1.8	847
2	Somatic and germline activating mutations of the ALK kinase receptor in neuroblastoma. Nature, 2008, 455, 967-970.	13.7	787
3	Interaction of a Golgi-Associated Kinesin-Like Protein with Rab6. Science, 1998, 279, 580-585.	6.0	478
4	Relapsed neuroblastomas show frequent RAS-MAPK pathway mutations. Nature Genetics, 2015, 47, 864-871.	9.4	451
5	Heterogeneity of neuroblastoma cell identity defined by transcriptional circuitries. Nature Genetics, 2017, 49, 1408-1413.	9.4	331
6	Germline Mutations of the Paired–Like Homeobox 2B (PHOX2B) Gene in Neuroblastoma. American Journal of Human Genetics, 2004, 74, 761-764.	2.6	288
7	Overall Genomic Pattern Is a Predictor of Outcome in Neuroblastoma. Journal of Clinical Oncology, 2009, 27, 1026-1033.	0.8	288
8	Control-free calling of copy number alterations in deep-sequencing data using GC-content normalization. Bioinformatics, 2011, 27, 268-269.	1.8	249
9	Meta-analysis of Neuroblastomas Reveals a Skewed <i>ALK</i> Mutation Spectrum in Tumors with <i>MYCN</i> Amplification. Clinical Cancer Research, 2010, 16, 4353-4362.	3.2	243
10	SVDetect: a tool to identify genomic structural variations from paired-end and mate-pair sequencing data. Bioinformatics, 2010, 26, 1895-1896.	1.8	178
11	Predicting outcomes for children with neuroblastoma using a multigene-expression signature: a retrospective SIOPEN/COG/GPOH study. Lancet Oncology, The, 2009, 10, 663-671.	5.1	176
12	Emergence of New <i>ALK</i> Mutations at Relapse of Neuroblastoma. Journal of Clinical Oncology, 2014, 32, 2727-2734.	0.8	176
13	Betaâ€catenin status in paediatric medulloblastomas: correlation of immunohistochemical expression with mutational status, genetic profiles, and clinical characteristics. Journal of Pathology, 2009, 218, 86-94.	2.1	171
14	Characterization of Novel Rab6-Interacting Proteins Involved in Endosome-to-TGN Transport. Traffic, 2002, 3, 289-297.	1.3	145
15	Accumulation of Segmental Alterations Determines Progression in Neuroblastoma. Journal of Clinical Oncology, 2010, 28, 3122-3130.	0.8	142
16	Prognostic Impact of Gene Expression–Based Classification for Neuroblastoma. Journal of Clinical Oncology, 2010, 28, 3506-3515.	0.8	129
17	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.	3.2	127
18	VAMP: Visualization and analysis of array-CGH, transcriptome and other molecular profiles. Bioinformatics, 2006, 22, 2066-2073.	1.8	106

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19	Two-hybrid System Screen with the Small GTP-binding Protein Rab6. IDENTIFICATION OF A NOVEL MOUSE GDP DISSOCIATION INHIBITOR ISOFORM AND TWO OTHER POTENTIAL PARTNERS OF Rab6. Journal of Biological Chemistry, 1995, 270, 14801-14808.	1.6	104
20	Transcriptional Programs Define Intratumoral Heterogeneity of Ewing Sarcoma at Single-Cell Resolution. Cell Reports, 2020, 30, 1767-1779.e6.	2.9	96
21	The occurrence of intracranial rhabdoid tumours in mice depends on temporal control of Smarcb1 inactivation. Nature Communications, 2016, 7, 10421.	5. 8	92
22	Recent insights into the biology of neuroblastoma. International Journal of Cancer, 2014, 135, 2249-2261.	2.3	91
23	Activated Alk triggers prolonged neurogenesis and Ret upregulation providing a therapeutic target in ALK-mutated neuroblastoma. Oncotarget, 2014, 5, 2688-2702.	0.8	89
24	Accurate Outcome Prediction in Neuroblastoma across Independent Data Sets Using a Multigene Signature. Clinical Cancer Research, 2010, 16, 1532-1541.	3.2	86
25	Molecular pathogenesis of peripheral neuroblastic tumors. Oncogene, 2010, 29, 1566-1579.	2.6	84
26	Revised Risk Estimation and Treatment Stratification of Low- and Intermediate-Risk Neuroblastoma Patients by Integrating Clinical and Molecular Prognostic Markers. Clinical Cancer Research, 2015, 21, 1904-1915.	3.2	80
27	Upregulation of MAPK Negative Feedback Regulators and RET in Mutant ALK Neuroblastoma: Implications for Targeted Treatment. Clinical Cancer Research, 2015, 21, 3327-3339.	3.2	76
28	Midkine and Alk signaling in sympathetic neuron proliferation and neuroblastoma predisposition. Development (Cambridge), 2011, 138, 4699-4708.	1.2	72
29	Identification and Characterization of Potential Effector Molecules of the Ras-related GTPase Rap2. Journal of Biological Chemistry, 1999, 274, 8737-8745.	1.6	70
30	Germline mutations of the paired-like homeobox 2B (PHOX2B) gene in neuroblastoma. Cancer Letters, 2005, 228, 51-58.	3.2	63
31	Deep Sequencing Reveals Occurrence of Subclonal <i>ALK</i> Mutations in Neuroblastoma at Diagnosis. Clinical Cancer Research, 2015, 21, 4913-4921.	3.2	62
32	Identification of a specific effector of the small GTP-binding protein Rap2. FEBS Journal, 1998, 252, 290-298.	0.2	60
33	Gene expression profiling of 1p35–36 genes in neuroblastoma. Oncogene, 2004, 23, 5912-5922.	2.6	60
34	Characterization of Rearrangements Involving the <i>ALK</i> Gene Reveals a Novel Truncated Form Associated with Tumor Aggressiveness in Neuroblastoma. Cancer Research, 2013, 73, 195-204.	0.4	54
35	Combined 24-color karyotyping and comparative genomic hybridization analysis indicates predominant rearrangements of early replicating chromosome regions in neuroblastoma. Cancer Genetics and Cytogenetics, 2003, 141, 32-42.	1.0	53
36	Repurposing rotavirus vaccines for intratumoral immunotherapy can overcome resistance to immune checkpoint blockade. Science Translational Medicine, 2019, 11 , .	5.8	49

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37	Activated ALK signals through the ERK–ETV5–RET pathway to drive neuroblastoma oncogenesis. Oncogene, 2018, 37, 1417-1429.	2.6	45
38	Breakpoint Features of Genomic Rearrangements in Neuroblastoma with Unbalanced Translocations and Chromothripsis. PLoS ONE, 2013, 8, e72182.	1.1	42
39	Radiogenomics of neuroblastomas: Relationships between imaging phenotypes, tumor genomic profile and survival. PLoS ONE, 2017, 12, e0185190.	1.1	40
40	Characterization of amplicons in neuroblastoma: Highâ€resolution mapping using DNA microarrays, relationship with outcome, and identification of overexpressed genes. Genes Chromosomes and Cancer, 2008, 47, 819-834.	1.5	39
41	Germline gain-of-function mutations of ALK disrupt central nervous system development. Human Mutation, 2011, 32, 272-276.	1.1	38
42	ALK germline mutations in patients with neuroblastoma: a rare and weakly penetrant syndrome. European Journal of Human Genetics, 2012, 20, 291-297.	1.4	38
43	Stepwise occurrence of a complex unbalanced translocation in neuroblastoma leading to insertion of a telomere sequence and late chromosome 17q gain. Oncogene, 2005, 24, 3377-3384.	2.6	36
44	SV-Bay: structural variant detection in cancer genomes using a Bayesian approach with correction for GC-content and read mappability. Bioinformatics, 2016, 32, 984-992.	1.8	36
45	Variety and complexity of chromosome 17 translocations in neuroblastoma. Genes Chromosomes and Cancer, 2004, 39, 143-150.	1.5	35
46	Unraveling Ewing Sarcoma Tumorigenesis Originating from Patient-Derived Mesenchymal Stem Cells. Cancer Research, 2021, 81, 4994-5006.	0.4	35
47	Preferential Occurrence of Chromosome Breakpoints within Early Replicating Regions in Neuroblastoma. Cell Cycle, 2005, 4, 1842-1846.	1.3	33
48	Learning smoothing models of copy number profiles using breakpoint annotations. BMC Bioinformatics, 2013, 14, 164.	1.2	33
49	Lin28B and Let-7 in the Control of Sympathetic Neurogenesis and Neuroblastoma Development. Journal of Neuroscience, 2015, 35, 16531-16544.	1.7	32
50	Wild-type ALK and activating ALK-R1275Q and ALK-F1174L mutations upregulate Myc and initiate tumor formation in murine neural crest progenitor cells. Oncotarget, 2014, 5, 4452-4466.	0.8	32
51	RGS14 is a novel Rap effector that preferentially regulates the GTPase activity of $\widehat{Gl}\pm 0$. Biochemical Journal, 2000, 350, 19.	1.7	31
52	The ALK receptor in sympathetic neuron development and neuroblastoma. Cell and Tissue Research, 2018, 372, 325-337.	1.5	31
53	Plasticity in Neuroblastoma Cell Identity Defines a Noradrenergic-to-Mesenchymal Transition (NMT). Cancers, 2021, 13, 2904.	1.7	29
54	Regulation by miR 181 Family of the Dependence Receptor CDON Tumor Suppressive Activity in Neuroblastoma. Journal of the National Cancer Institute, 2014, 106, .	3.0	27

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55	Internalization and Down-Regulation of the ALK Receptor in Neuroblastoma Cell Lines upon Monoclonal Antibodies Treatment. PLoS ONE, 2012, 7, e33581.	1.1	27
56	Molecular analysis of chromosome arm 17q gain in neuroblastoma. Genes Chromosomes and Cancer, 2000, 28, 276-284.	1.5	26
57	Cholinergic switch associated with morphological differentiation in neuroblastoma. Journal of Pathology, 2009, 219, 463-472.	2.1	26
58	Methylation-associated PHOX2B gene silencing is a rare event in human neuroblastoma. European Journal of Cancer, 2007, 43, 2366-2372.	1.3	20
59	QuantumClone: clonal assessment of functional mutations in cancer based on a genotype-aware method for clonal reconstruction. Bioinformatics, 2018, 34, 1808-1816.	1.8	20
60	Regulation of the GTPase activity of the ras-related rap2 protein. Biochemical and Biophysical Research Communications, 1992, 189, 455-464.	1.0	18
61	Clinical Characteristics and Outcome of Patients with Neuroblastoma Presenting Genomic Amplification of Loci Other than MYCN. PLoS ONE, 2014, 9, e101990.	1.1	17
62	Study of chromatin remodeling genes implicates SMARCA4 as a putative player in oncogenesis in neuroblastoma. International Journal of Cancer, 2019, 145, 2781-2791.	2.3	16
63	Targeting netrinâ€3 in small cell lung cancer and neuroblastoma. EMBO Molecular Medicine, 2021, 13, e12878.	3.3	16
64	Integrative analysis identifies lincRNAs up- and downstream of neuroblastoma driver genes. Scientific Reports, 2019, 9, 5685.	1.6	14
65	High-resolution mapping of amplicons of the short arm of chromosome 1 in two neuroblastoma tumors by microarray-based comparative genomic hybridization. Genes Chromosomes and Cancer, 2004, 40, 266-270.	1.5	13
66	Segmental Chromosomal Aberrations in Localized Neuroblastoma Can be Detected in Formalinâ€Fixed Paraffinâ€Embedded Tissue Samples and Are Associated With Recurrence. Pediatric Blood and Cancer, 2016, 63, 1019-1023.	0.8	13
67	The sympathetic nervous system: malignancy, disease, and novel functions. Cell and Tissue Research, 2018, 372, 163-170.	1.5	12
68	SegAnnDB: interactive Web-based genomic segmentation. Bioinformatics, 2014, 30, 1539-1546.	1.8	10
69	68Ga-DOTATOC and FDG PET Imaging of Preclinical Neuroblastoma Models. Anticancer Research, 2016, 36, 4459-4466.	0.5	10
70	HMCan-diff: a method to detect changes in histone modifications in cells with different genetic characteristics. Nucleic Acids Research, 2017, 45, gkw1319.	6.5	8
71	Use of the Two-Hybrid System to Identify Rab-Interacting Proteins. Methods, 2000, 20, 399-402.	1.9	7
72	Hyperactivation of Alk induces neonatal lethality in knock-in AlkF1178L mice. Oncotarget, 2014, 5, 2703-2713.	0.8	6

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73	ALK mutation dynamics and clonal evolution in a neuroblastoma model exhibiting two ALK mutations. Oncotarget, 2019, 10, 4937-4950.	0.8	5
74	Two cases of localized neuroblastoma with multiple segmental chromosomal alterations and metastatic progression. Pediatric Blood and Cancer, 2013, 60, 332-335.	0.8	3
75	Midkine and Alk signaling in sympathetic neuron proliferation and neuroblastoma predisposition. Journal of Cell Science, 2011, 124, e1-e1.	1.2	1
76	BET and CDK Inhibition Reveal Differences in the Proliferation Control of Sympathetic Ganglion Neuroblasts and Adrenal Chromaffin Cells. Cancers, 2022, 14, 2755.	1.7	1
77	High CD44 expression is not a prognosis marker in patients with high-risk neuroblastoma. EBioMedicine, 2020, 53, 102702.	2.7	O
78	Neuroblastoma Pathogenesis. , 2020, , 29-56.		0