Magali Olivier

List of Publications by Year in descending order

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80 papers 11,086 citations

39 h-index 91884 69 g-index

90 all docs

90 docs citations

90 times ranked 16721 citing authors

#	Article	IF	CITATIONS
1	Molecular profiles and urinary biomarkers of upper tract urothelial carcinomas associated with aristolochic acid exposure. International Journal of Cancer, 2022, 150, 374-386.	5.1	4
2	The TP53 Database: transition from the International Agency for Research on Cancer to the US National Cancer Institute. Cell Death and Differentiation, 2022, 29, 1071-1073.	11.2	53
3	Specifications of the ACMG/AMP variant interpretation guidelines for germline <i>TP53</i> variants. Human Mutation, 2021, 42, 223-236.	2.5	81
4	The International Collaboration for Cancer Classification and Research. International Journal of Cancer, 2021, 148, 560-571.	5.1	32
5	PVAmpliconFinder: a workflow for the identification of human papillomaviruses from high-throughput amplicon sequencing. BMC Bioinformatics, 2020, 21, 233.	2.6	2
6	Anthropometry, body shape in early-life and risk of premenopausal breast cancer among Latin American women: results from the PRECAMA study. Scientific Reports, 2020, 10, 2294.	3.3	10
7	A quantitative model to predict pathogenicity of missense variants in the <i>TP53</i> gene. Human Mutation, 2019, 40, 788-800.	2.5	21
8	Experimental and pan-cancer genome analyses reveal widespread contribution of acrylamide exposure to carcinogenesis in humans. Genome Research, 2019, 29, 521-531.	5.5	57
9	Molecular features of premenopausal breast cancers in Latin American women: Pilot results from the PRECAMA study. PLoS ONE, 2019, 14, e0210372.	2.5	12
10	Project profile: a multicenter study on breast cancer in young women in Latin America (PRECAMA) Tj ETQq0 0 0	rgBT /Ove	rlock 10 Tf 50
11	Improved, ACMG-compliant, in silico prediction of pathogenicity for missense substitutions encoded by <i>TP53</i> variants. Human Mutation, 2018, 39, 1061-1069.	2.5	29
12	MA11.05 A Case-Control Study to Test the Use of ctDNA in the Early Detection of SCLC Reveals TP53 Mutations in Non-Cancer Controls. Journal of Thoracic Oncology, 2017, 12, S405-S406.	1.1	0
13	Genome-scale mutational signatures of aflatoxin in cells, mice, and human tumors. Genome Research, 2017, 27, 1475-1486.	5.5	90
14	IARC TP53 Database. , 2017, , 2193-2198.		O
15	<i>TP53</i> Variations in Human Cancers: New Lessons from the IARC TP53 Database and Genomics Data. Human Mutation, 2016, 37, 865-876.	2.5	589
16	Identification of Circulating Tumor DNA for the Early Detection of Small-cell Lung Cancer. EBioMedicine, 2016, 10, 117-123.	6.1	153
17	MutSpec: a Galaxy toolbox for streamlined analyses of somatic mutation spectra in human and mouse cancer genomes. BMC Bioinformatics, 2016, 17, 170.	2.6	44
18	Revealing the Molecular Portrait of Triple Negative Breast Tumors in an Understudied Population through Omics Analysis of Formalin-Fixed and Paraffin-Embedded Tissues. PLoS ONE, 2015, 10, e0126762.	2.5	18

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19	Low-Coverage Exome Sequencing Screen in Formalin-Fixed Paraffin-Embedded Tumors Reveals Evidence of Exposure to Carcinogenic Aristolochic Acid. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1873-1881.	2.5	21
20	Circulating free DNA concentration is an independent prognostic biomarker in lung cancer. European Respiratory Journal, 2015, 46, 1773-1780.	6.7	114
21	Abstract 4748: Revealing the molecular portrait of triple negative breast tumors from an understudied population through omics analysis of formalin-fixed and paraffin-embedded tissues., 2015,,.		0
22	Noninvasive Diagnosis of Actionable Mutations by Deep Sequencing of Circulating Free DNA in Lung Cancer from Never-Smokers: A Proof-of-Concept Study from BioCAST/IFCT-1002. Clinical Cancer Research, 2014, 20, 4613-4624.	7.0	195
23	Age at cancer onset in germline TP53 mutation carriers: association with polymorphisms in predicted G-quadruplex structures. Carcinogenesis, 2014, 35, 807-815.	2.8	29
24	Modelling mutational landscapes of human cancers in vitro. Scientific Reports, 2014, 4, 4482.	3.3	83
25	Abstract 305: Ultra-low coverage exome sequencing of FFPE tumor specimens identifies exposure to carcinogenic aristolochic acid., 2014, , .		0
26	Analysis of TP53 mutation spectra reveals the fingerprint of the potent environmental carcinogen, aristolochic acid. Mutation Research - Reviews in Mutation Research, 2013, 753, 41-49.	5.5	55
27	TP53 Somatic Mutations: Prognostic and Predictive Value in Human Cancers. , 2013, , 127-146.		2
28	A meta-analysis of cancer risk associated with the TP53 intron 3 duplication polymorphism (rs17878362): geographic and tumor-specific effects. Cell Death and Disease, 2013, 4, e492-e492.	6.3	43
29	Assessing TP53 Status in Human Tumors: Lessons from Breast Cancer. , 2013, , 147-165.		1
30	Low Prevalence of <i>TP53 </i> Mutations and <i>MDM2 </i> Rhabdomyosarcoma. Sarcoma, 2012, 2012, 1-6.	1.3	23
31	Prognostic value of <i>TP53</i> , <i>KRAS</i> and <i>EGFR</i> mutations in nonsmall cell lung cancer: the EUELC cohort. European Respiratory Journal, 2012, 40, 177-184.	6.7	92
32	Upper urinary tract urothelial cancers: where it is A:T. Nature Reviews Cancer, 2012, 12, 503-504.	28.4	22
33	Prognostic and predictive value of TP53mutations in node-positive breast cancer patients treated with anthracycline- or anthracycline/taxane-based adjuvant therapy: results from the BIG 02-98 phase III trial. Breast Cancer Research, 2012, 14, R70.	5.0	52
34	Sarcomas in <i>TP53</i> germline mutation carriers. Cancer, 2012, 118, 1387-1396.	4.1	189
35	p53-Dependent repression of focal adhesion kinase in response to estradiol in breast cancer cell-lines. Cancer Letters, 2011, 300, 215-224.	7.2	25
36	Biological functions of p53 isoforms through evolution: lessons from animal and cellular models. Cell Death and Differentiation, 2011, 18, 1815-1824.	11.2	173

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37	Somatic mutations in cancer prognosis and prediction: lessons from TP53 and EGFR genes. Current Opinion in Oncology, 2011, 23, 88-92.	2.4	41
38	Understanding wild-type and mutant p53 activities in human cancer: new landmarks on the way to targeted therapies. Cancer Gene Therapy, 2011, 18, 2-11.	4.6	151
39	Estrogen levels act as a rheostat on p53 levels and modulate p53-dependent responses in breast cancer cell lines. Breast Cancer Research and Treatment, 2011, 125, 35-42.	2.5	27
40	TP53 mutation p.R337H in gastric cancer tissues of a 12-year-old male child - evidence for chimerism involving a common mutant founder haplotype: case report. BMC Cancer, 2011, 11, 449.	2.6	34
41	p53 status influences response to tamoxifen but not to fulvestrant in breast cancer cell lines. International Journal of Cancer, 2011, 128, 1813-1821.	5.1	29
42	IARC TP53 Database., 2011,, 1799-1802.		1
43	Abstract 5538: Sarcomas in TP53 germline mutation carriers. , 2011, , .		0
44	Tumor protein 53 mutations and inherited cancer: beyond Li-Fraumeni syndrome. Current Opinion in Oncology, 2010, 22, 64-69.	2.4	91
45	Detailed haplotype analysis at the <i>TP53 < /i>locus in p.R337H mutation carriers in the population of Southern Brazil: evidence for a founder effect. Human Mutation, 2010, 31, 143-150.</i>	2.5	116
46	p53 regulates the transcription of its î"133p53 isoform through specific response elements contained within the TP53 P2 internal promoter. Oncogene, 2010, 29, 2691-2700.	5.9	60
47	TP53 Mutations in Human Cancers: Origins, Consequences, and Clinical Use. Cold Spring Harbor Perspectives in Biology, 2010, 2, a001008-a001008.	5.5	1,494
48	TP53 Mutations in Human Cancers: Selection versus Mutagenesis. Molecular Biology Intelligence Unit, 2010, , 1-18.	0.2	1
49	Biomarkers Predict <i>p53</i> Gene Therapy Efficacy in Recurrent Squamous Cell Carcinoma of the Head and Neck. Clinical Cancer Research, 2009, 15, 7719-7725.	7.0	87
50	Somatic mutation databases as tools for molecular epidemiology and molecular pathology of cancer: Proposed guidelines for improving data collection, distribution, and integration. Human Mutation, 2009, 30, 275-282.	2.5	14
51	Recent advances in p53 research: an interdisciplinary perspective. Cancer Gene Therapy, 2009, 16, 1-12.	4.6	140
52	TP53 PIN3 and MDM2 SNP309 polymorphisms as genetic modifiers in the Li-Fraumeni syndrome: impact on age at first diagnosis. Journal of Medical Genetics, 2009, 46, 766-772.	3.2	64
53	Locus-specific databases and recommendations to strengthen their contribution to the classification of variants in cancer susceptibility genes. Human Mutation, 2008, 29, 1273-1281.	2.5	41
54	Influence of TP53 gene status on treatment response in breast cancer cells. European Journal of Cancer, Supplement, 2008, 6, 104.	2.2	0

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55	Detection of R337H, a germline TP53 mutation predisposing to multiple cancers, in asymptomatic women participating in a breast cancer screening program in Southern Brazil. Cancer Letters, 2008, 261, 21-25.	7.2	94
56	Prognostic and Predictive Value of TP53 Mutations in Human Cancer., 2007,, 321-338.		13
57	The TP53 mutation, R337H, is associated with Li-Fraumeni and Li-Fraumeni-like syndromes in Brazilian families. Cancer Letters, 2007, 245, 96-102.	7.2	170
58	Response to "Germline TP53 R337H mutation is not sufficient to establish Li-Fraumeni or Li-Fraumeni-like syndromeâ€, by Ribeiro et al Cancer Letters, 2007, 247, 356-358.	7.2	1
59	Impact of mutant p53 functional properties on <i>TP53</i> mutation patterns and tumor phenotype: lessons from recent developments in the IARC TP53 database. Human Mutation, 2007, 28, 622-629.	2.5	1,441
60	TP53 mutations in human cancers: functional selection and impact on cancer prognosis and outcomes. Oncogene, 2007, 26, 2157-2165.	5.9	796
61	Patterns of TP53 Mutations in Human Cancer: Interplay Between Mutagenesis, DNA Repair and Selection., 2007,, 293-319.		4
62	Patterns of TP53 Mutations in Human Cancer: Interplay between Mutagenesis, DNA Repair and Selection., 2007,, 293-319.		0
63	Prognostic and Predictive Value of TP53 Mutations in Human Cancer. , 2007, , 321-338.		0
64	Predicting the transactivation activity of p53 missense mutants using a four-body potential score derived from Delaunay tessellations. Human Mutation, 2006, 27, 163-172.	2.5	21
65	Computational approaches for predicting the biological effect of p53 missense mutations: a comparison of three sequence analysis based methods. Nucleic Acids Research, 2006, 34, 1317-1325.	14.5	295
66	The clinical value of somatic TP53 gene mutations in 1,794 patients with breast cancer Clinical Cancer Research, 2006, 12, 1157-1167.	7.0	495
67	TP53 mutations as biomarkers for cancer epidemiology in Latin America: Current knowledge and perspectives. Mutation Research - Reviews in Mutation Research, 2005, 589, 192-207.	5.5	16
68	Independent prognostic value of somatic TP53gene mutations in 1794 breast cancer patients. Breast Cancer Research, 2005, 7, 1.	5.0	0
69	Effect of the codon 72 polymorphism (c.215G>C, p.Arg72Pro) in combination with somatic sequence variants in theTP53gene on survival in patients with advanced ovarian carcinoma. Human Mutation, 2004, 24, 21-34.	2.5	46
70	The Tumor Suppressor Gene TP53: Implications for Cancer Management and Therapy. Critical Reviews in Clinical Laboratory Sciences, 2004, 41, 551-583.	6.1	31
71	TP53 mutation spectra and load: a tool for generating hypotheses on the etiology of cancer. larc (international Agency for Research on Cancer) Scientific Publications, 2004, , 247-70.	0.4	83
72	Li-Fraumeni and related syndromes: correlation between tumor type, family structure, and TP53 genotype. Cancer Research, 2003, 63, 6643-50.	0.9	350

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73	Differential Activation of p53 by the Various Adducts of Mitomycin C. Journal of Biological Chemistry, 2002, 277, 40513-40519.	3.4	50
74	Integrating mutation data and structural analysis of the TP53 tumor-suppressor protein. Human Mutation, 2002, 19, 149-164.	2.5	122
75	The IARC TP53 database: New online mutation analysis and recommendations to users. Human Mutation, 2002, 19, 607-614.	2.5	1,107
76	Tobacco smoke carcinogens, DNA damage and p53 mutations in smoking-associated cancers. Oncogene, 2002, 21, 7435-7451.	5.9	961
77	TP53 mutation patterns in breast cancers: searching for clues of environmental carcinogenesis. Seminars in Cancer Biology, 2001, 11, 353-360.	9.6	56
78	TP53 mutation spectrum in lung cancers and mutagenic signature of components of tobacco smoke: lessons from the IARC TP53 mutation database. Mutagenesis, 2001, 16, 551-553.	2.6	47
79	Relaxed cell-cycle arrests and propagation of unrepaired chromosomal damage in cancer cell lines with wild-typep53. Molecular Carcinogenesis, 1998, 23, 1-12.	2.7	23
80	Modulation of DNA Topoisomerase I Activity byp53. Biochemistry, 1996, 35, 5778-5786.	2.5	99