Silvia Regina Caminada Toledo

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Genome-wide association study identifies two susceptibility loci for osteosarcoma. Nature Genetics, 2013, 45, 799-803.	21.4	181
2	Frequency of Pathogenic Germline Variants in Cancer-Susceptibility Genes in Patients With Osteosarcoma. JAMA Oncology, 2020, 6, 724.	7.1	139
3	Maternal embryonic leucine zipper kinase transcript abundance correlates with malignancy grade in human astrocytomas. International Journal of Cancer, 2008, 122, 807-815.	5.1	128
4	A Genome-Wide Scan Identifies Variants in <i>NFIB</i> Associated with Metastasis in Patients with Osteosarcoma. Cancer Discovery, 2015, 5, 920-931.	9.4	88
5	Comparative genomic hybridization analysis identifies gains of 1p35â^¼p36 and chromosome 19 in osteosarcoma. Cancer Genetics and Cytogenetics, 2001, 130, 14-21.	1.0	69
6	Tropism of mesenchymal stem cell toward CD133+ stem cell of glioblastoma in vitro and promote tumor proliferation in vivo. Stem Cell Research and Therapy, 2018, 9, 310.	5.5	52
7	GNAS mutations are not detected in parosteal and low-grade central osteosarcomas. Modern Pathology, 2015, 28, 1336-1342.	5.5	47
8	Expression analysis of stem cell-related genes reveal OCT4 as a predictor of poor clinical outcome in medulloblastoma. Journal of Neuro-Oncology, 2012, 106, 71-79.	2.9	40
9	DNA methylation landscape of hepatoblastomas reveals arrest at early stages of liver differentiation and cancer-related alterations. Oncotarget, 2017, 8, 97871-97889.	1.8	32
10	Analysis of KIAA1549–BRAF fusion gene expression and IDH1/IDH2 mutations in low grade pediatric astrocytomas. Journal of Neuro-Oncology, 2014, 117, 235-242.	2.9	31
11	Genomeâ€wide association study identifies the <i>GLDC</i> / <i>IL33</i> locus associated with survival of osteosarcoma patients. International Journal of Cancer, 2018, 142, 1594-1601.	5.1	31
12	Genetic and Immunohistochemical Expression of Integrins ITGAV, ITGA6, and ITGA3 As Prognostic Factor for Colorectal Cancer: Models for Global and Disease-Free Survival. PLoS ONE, 2015, 10, e0144333.	2.5	30
13	MYCN Gene Amplification. American Journal of Pathology, 1999, 155, 1439-1443.	3.8	29
14	Bone deposition, bone resorption, and osteosarcoma. Journal of Orthopaedic Research, 2010, 28, 1142-1148.	2.3	27
15	TNF-alpha, TNF-beta, IL-6, IL-10, PECAM-1 and the MPO Inflammatory Gene Polymorphisms in Osteosarcoma. Journal of Pediatric Hematology/Oncology, 2007, 29, 293-297.	0.6	25
16	MAPK7 and MAP2K4 as prognostic markers in osteosarcoma. Human Pathology, 2012, 43, 994-1002.	2.0	25
17	SHH, WNT, and NOTCH pathways in medulloblastoma: when cancer stem cells maintain self-renewal and differentiation properties. Child's Nervous System, 2014, 30, 1165-72.	1.1	25
18	<i>MAPK7</i> gene controls proliferation, migration and cell invasion in osteosarcoma. Molecular Carcinogenesis, 2016, 55, 1700-1713.	2.7	25

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19	MAPK pathways regulation by DUSP1 in the development of osteosarcoma: Potential markers and therapeutic targets. Molecular Carcinogenesis, 2017, 56, 1630-1641.	2.7	24
20	Tumor Dynamics in Response to Antiangiogenic Therapy with Oral Metronomic Topotecan and Pazopanib in Neuroblastoma Xenografts. Translational Oncology, 2013, 6, 493-503.	3.7	23
21	mRNA expression of matrix metalloproteinases (MMPs) 2 and 9 and tissue inhibitor of matrix metalloproteinases (TIMPs) 1 and 2 in childhood acute lymphoblastic leukemia: Potential role of TIMP1 as an adverse prognostic factor. Leukemia Research, 2010, 34, 32-37.	0.8	22
22	The metastatic behavior of osteosarcoma by gene expression and cytogenetic analyses. Human Pathology, 2013, 44, 2188-2198.	2.0	18
23	ASPM gene expression in medulloblastoma. Child's Nervous System, 2011, 27, 71-74.	1.1	17
24	TET Upregulation Leads to 5-Hydroxymethylation Enrichment in Hepatoblastoma. Frontiers in Genetics, 2019, 10, 553.	2.3	17
25	Aberrant DNA methylation of ESR1 and p14ARF genes could be useful as prognostic indicators in osteosarcoma. OncoTargets and Therapy, 2013, 6, 713.	2.0	14
26	Expression Profiling Using a cDNA Array and Immunohistochemistry for the Extracellular Matrix Genes FN-1, ITGA-3, ITGB-5, MMP-2, and MMP-9 in Colorectal Carcinoma Progression and Dissemination. Scientific World Journal, The, 2014, 2014, 1-27.	2.1	14
27	PRAME gene expression profile in medulloblastoma. Arquivos De Neuro-Psiquiatria, 2011, 69, 9-12.	0.8	13
28	Expression of major vault protein gene in osteosarcoma patients. Journal of Orthopaedic Research, 2007, 25, 958-963.	2.3	12
29	Insights on PRAME and osteosarcoma by means of gene expression profiling. Journal of Orthopaedic Science, 2011, 16, 458-466.	1.1	12
30	Insights Into the Somatic Mutation Burden of Hepatoblastomas From Brazilian Patients. Frontiers in Oncology, 2020, 10, 556.	2.8	12
31	CYP genes in osteosarcoma: Their role in tumorigenesis, pulmonary metastatic microenvironment and treatment response. Oncotarget, 2017, 8, 38530-38540.	1.8	12
32	Aberrant signaling pathways in medulloblastomas: a stem cell connection. Arquivos De Neuro-Psiquiatria, 2010, 68, 947-952.	0.8	11
33	Hepatoblastomas exhibit marked <i>NNMT</i> downregulation driven by promoter DNA hypermethylation. Tumor Biology, 2020, 42, 101042832097712.	1.8	11
34	Comparative genomic hybridization analysis of pediatric adamantinomatous craniopharyngiomas and a review of the literature. Journal of Neurosurgery: Pediatrics, 2004, 101, 85-90.	1.3	9
35	Investigation of PAX3/7-FKHR fusion genes and IGF2 gene expression in rhabdomyosarcoma tumors. Growth Hormone and IGF Research, 2012, 22, 245-249.	1.1	9
36	Myelodysplastic syndrome in childhood: report of two cases with deletion of chromosome 4 and the Philadelphia chromosome. Leukemia Research, 2002, 26, 533-538.	0.8	8

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37	Investigation of IGF2, Hedgehog and fusion gene expression profiles in pediatric sarcomas. Growth Hormone and IGF Research, 2014, 24, 130-136.	1.1	8
38	Establishment of primary cell culture and an intracranial xenograft model of pediatric ependymoma: a prospect for therapy development and understanding of tumor biology. Oncotarget, 2018, 9, 21731-21743.	1.8	8
39	Expression Profile Analysis of Genes Related to Resistance/Sensibility to Prednisolone, Daunorubicin, L-Asparaginase and Vincristine in Childhood Acute Lymphoblastic Leukemia Blood, 2007, 110, 3463-3463.	1.4	8
40	Intravenous Grafts of Human Amniotic Fluid-Derived Stem Cells Reduce Behavioral Deficits in Experimental Ischemic Stroke. Cell Transplantation, 2019, 28, 1306-1320.	2.5	7
41	New therapeutic target for pediatric anaplastic ependymoma control: study of anti-tumor activity by a Kunitz-type molecule, Amblyomin-X. Scientific Reports, 2019, 9, 9973.	3.3	6
42	Unraveling the Genetic Architecture of Hepatoblastoma Risk: Birth Defects and Increased Burden of Germline Damaging Variants in Gastrointestinal/Renal Cancer Predisposition and DNA Repair Genes. Frontiers in Genetics, 2022, 13, 858396.	2.3	6
43	Molecular profiling of osteosarcoma in children and adolescents from different age groups using a next-generation sequencing panel. Cancer Genetics, 2021, 258-259, 85-92.	0.4	5
44	Copy Number Alterations in Hepatoblastoma: Literature Review and a Brazilian Cohort Analysis Highlight New Biological Pathways. Frontiers in Oncology, 2021, 11, 741526.	2.8	5
45	Establishment and cytogenetic characterization of a cell line from a pulmonary metastasis of osteosarcoma. Cytotechnology, 2013, 65, 347-353.	1.6	4
46	Valproic acid treatment response in vitro is determined by TP53 status in medulloblastoma. Child's Nervous System, 2018, 34, 1497-1509.	1.1	4
47	MAPK7 variants related to prognosis and chemotherapy response in osteosarcoma. Annals of Diagnostic Pathology, 2020, 46, 151482.	1.3	3
48	Gliomas in children and adolescents: investigation of molecular alterations with a potential prognostic and therapeutic impact. Journal of Cancer Research and Clinical Oncology, 2022, 148, 107-119.	2.5	2
49	Abnormal spindle-like microcephaly-associated (ASPM) gene expression in posterior fossa brain tumors of childhood and adolescence. Child's Nervous System, 2021, 37, 137-145.	1.1	1
50	Molecular profiling of pediatric and adolescent ependymomas: identification of genetic variants using a next-generation sequencing panel. Journal of Neuro-Oncology, 2021, 155, 13-23.	2.9	1