

Gordana Raca

List of Publications by Year in descending order

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Version: 2024-02-01

35
papers

1,409
citations

687363

13
h-index

454955

30
g-index

38
all docs

38
docs citations

38
times ranked

2906
citing authors

#	ARTICLE	IF	CITATIONS
1	A harmonized meta-knowledgebase of clinical interpretations of somatic genomic variants in cancer. <i>Nature Genetics</i> , 2020, 52, 448-457.	21.4	104
2	Diagnostic testing for uniparental disomy: a points to consider statement from the American College of Medical Genetics and Genomics (ACMG). <i>Genetics in Medicine</i> , 2020, 22, 1133-1141.	2.4	89
3	Myeloid lineage switch following chimeric antigen receptor T cell therapy in a patient with TCF3&ZNF384 fusionεpositive B lymphoblastic leukemia. <i>Pediatric Blood and Cancer</i> , 2018, 65, e27265.	1.5	67
4	OncoKids. <i>Journal of Molecular Diagnostics</i> , 2018, 20, 765-776.	2.8	58
5	A semiautomated whole-exome sequencing workflow leads to increased diagnostic yield and identification of novel candidate variants. <i>Journal of Physical Education and Sports Management</i> , 2019, 5, a003756.	1.2	41
6	Technical laboratory standards for interpretation and reporting of acquired copy-number abnormalities and copy-neutral loss of heterozygosity in neoplastic disorders: a joint consensus recommendation from the American College of Medical Genetics and Genomics (ACMG) and the Cancer Genomics Consortium (CGC). <i>Genetics in Medicine</i> , 2019, 21, 1903-1916.	2.4	39
7	Assessing copy number aberrations and copy neutral loss of heterozygosity across the genome as best practice: An evidence based review of clinical utility from the cancer genomics consortium (CGC) working group for myelodysplastic syndrome, myelodysplastic/myeloproliferative and myeloproliferative neoplasms. <i>Cancer Genetics</i> , 2018, 228-229, 197-217.	0.4	25
8	Assessing copy number abnormalities and copy-neutral loss-of-heterozygosity across the genome as best practice in diagnostic evaluation of acute myeloid leukemia: An evidence-based review from the cancer genomics consortium (CGC) myeloid neoplasms working group. <i>Cancer Genetics</i> , 2018, 228-229, 218-235.	0.4	21
9	Increased Incidence of IKZF1 deletions and IGH-CRLF2 translocations in B-ALL of Hispanic/Latino childrenεa novel health disparity. <i>Leukemia</i> , 2021, 35, 2399-2402.	7.2	19
10	Standard operating procedure for curation and clinical interpretation of variants in cancer. <i>Genome Medicine</i> , 2019, 11, 76.	8.2	16
11	Adapting crowdsourced clinical cancer curation in CIViC to the ClinGen minimum variant level data communityεdriven standards. <i>Human Mutation</i> , 2018, 39, 1721-1732.	2.5	15
12	Evidence-based review of genomic aberrations in B-lymphoblastic leukemia/lymphoma: Report from the cancer genomics consortium working group for lymphoblastic leukemia. <i>Cancer Genetics</i> , 2020, 243, 52-72.	0.4	14
13	Increased prevalence of CRLF2 rearrangements in obesity-associated acute lymphoblastic leukemia. <i>Blood</i> , 2021, 138, 199-202.	1.4	8
14	A neoplasm with FIP1L1-PDGFRα fusion presenting as pediatric T-cell lymphoblastic leukemia/lymphoma without eosinophilia. <i>Cancer Genetics</i> , 2017, 216-217, 91-99.	0.4	6
15	A multimodal genomics approach to diagnostic evaluation of pediatric hematologic malignancies. <i>Cancer Genetics</i> , 2021, 254-255, 25-33.	0.4	6
16	Dilated cardiomyopathy in a patient with autosomal dominant EEF1A2-related neurodevelopmental disorder. <i>European Journal of Medical Genetics</i> , 2021, 64, 104121.	1.3	5
17	Standardized evidence-based approach for assessment of oncogenic and clinical significance of NTRK fusions. <i>Cancer Genetics</i> , 2022, 264-265, 50-59.	0.4	5
18	Response to Maya et al.. <i>Genetics in Medicine</i> , 2020, 22, 1278-1279.	2.4	4

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19	Primary Knee Intra-articular Synovial Sarcoma in Pediatric and Adolescent Patients. <i>Pediatric and Developmental Pathology</i> , 2021, 24, 159-163.	1.0	3
20	Conventional Cytogenetic Analysis of Hematologic Neoplasms: A 20-Year Review of Proficiency Test Results From the College of American Pathologists/American College of Medical Genetics and Genomics Cytogenetics Committee. <i>Archives of Pathology and Laboratory Medicine</i> , 2021, 145, 176-190.	2.5	3
21	Characterization of <i>PAX5</i> Intragenic Tandem Multiplication in Pediatric B-Lymphoblastic Leukemia by Optical Genome Mapping. <i>Blood Advances</i> , 2022, , .	5.2	3
22	A community approach to the cancer-variant-interpretation bottleneck. <i>Nature Cancer</i> , 2022, 3, 522-525.	13.2	3
23	Pediatric Metastatic Hepatoblastoma With an <i>ARID1A</i> Mutation and Rhabdoid Cells. <i>International Journal of Surgical Pathology</i> , 2022, 30, 307-312.	0.8	2
24	Congenital B-lymphoblastic leukemia with a cryptic MLL rearrangement and post-treatment evolution to mixed phenotype acute leukemia. <i>Leukemia Research Reports</i> , 2016, 6, 29-32.	0.4	1
25	Response to Mounts and Besser. <i>Genetics in Medicine</i> , 2021, 23, 240-242.	2.4	1
26	PLAG1 Immunohistochemical Staining Is a Surrogate Marker for PLAG1 Fusions in Lipoblastomas. <i>Pediatric and Developmental Pathology</i> , 2021, , 109352662110433.	1.0	1
27	Recommendations for future extensions to the HGNC gene fusion nomenclature. <i>Leukemia</i> , 2021, 35, 3611-3612.	7.2	1
28	insufficient for pathogenesis. <i>Cancer Genetics</i> , 2022, 260-261, 37-40.	0.4	1
29	Primary Adrenal Malignant Rhabdoid Tumor in a 14-Year-Old Female: A Case Report and Literature Review. <i>International Journal of Surgical Pathology</i> , 2022, 30, 172-176.	0.8	0
30	Abstract 210: Advancing knowledgebase representation of pediatric cancer variants through ClinGen/CIViC collaboration. , 2021, , .		0
31	Abstract 449: A standard operating procedure for the curation of gene fusions. , 2021, , .		0
32	Whole genome SNP arrays for best practice for detection of diagnostic, prognostic and therapy related copy number changes and copy neutral-loss of heterozygosity across solid tumors and hematologic malignancies.. <i>Journal of Clinical Oncology</i> , 2020, 38, e15575-e15575.	1.6	0
33	Efficacy and Safety of FLAG-Ida As Frontline Therapy in a Pediatric AML Population: A Single Institution Experience. <i>Blood</i> , 2021, 138, 2342-2342.	1.4	0
34	Expert Curation of Somatic FLT3 Variants By the Clingen Somatic Hematologic Cancer Taskforce (ClinGen HCT). <i>Blood</i> , 2021, 138, 4387-4387.	1.4	0
35	Expert Curation of Somatic Variants in Hematological Malignancies By the Clingen Somatic Hematological Cancer Taskforce (ClinGen HCT). <i>Blood</i> , 2020, 136, 23-23.	1.4	0