Xinjie Xu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1616004/publications.pdf

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		1684188	1372567
19	104	5	10
papers	citations	h-index	g-index
19	19	19	203
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Guiding the global evolution of cytogenetic testing for hematologic malignancies. Blood, 2022, 139, 2273-2284.	1.4	29
2	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. Cancer Genetics, 2018, 228-229, 197-217.	0.4	25
3	Detection of cryptic CCND1 rearrangements in mantle cell lymphoma by next generation sequencing. Annals of Diagnostic Pathology, 2020, 46, 151533.	1.3	8
4	Myeloid malignancies with 5q and 7q deletions are associated with extreme genomic complexity, biallelic TP53 variants, and very poor prognosis. Blood Cancer Journal, 2021, 11, 18.	6.2	8
5	Clinical utility of next generation sequencing to detect IGH/IL3 rearrangements [t(5;14)(q31.1;q32.1)] in B-lymphoblastic leukemia/lymphoma. Annals of Diagnostic Pathology, 2021, 53, 151761.	1.3	8
6	Cryptic and atypical <scp>KMT2Aâ€USP2</scp> and <scp>KMT2Aâ€USP8</scp> rearrangements identified by mate pair sequencing in infant and childhood leukemia. Genes Chromosomes and Cancer, 2020, 59, 422-427.	2.8	7
7	MYC break-apart FISH probe set reveals frequent unbalanced patterns of uncertain significance when evaluating aggressive B-cell lymphoma. Blood Cancer Journal, 2021, 11, 184.	6.2	6
8	Identification of a novel <i><scp>KMT2A</scp>/<scp>GIMAP8</scp></i> gene fusion in a pediatric patient with acute undifferentiated leukemia. Genes Chromosomes and Cancer, 2021, 60, 108-111.	2.8	5
9	A community approach to the cancer-variant-interpretation bottleneck. Nature Cancer, 2022, 3, 522-525.	13.2	3
10	Siblings with ETV6/RUNX1-positive B-lymphoblastic leukemia: A single site experience and review of the literature. Annals of Diagnostic Pathology, 2020, 48, 151588.	1.3	1
11	OUP accepted manuscript. Laboratory Medicine, 2021, , .	1.2	1
12	A Novel <i>USP25::PDGFRA</i> Gene Fusion in a 78 Year Old Patient with a Myeloid Neoplasm. Laboratory Medicine, 2022, 53, e134-e138.	1.2	1
13	Identification of EWSR1 rearrangements in patients with immature hematopoietic neoplasms: A case series and review of literature. Annals of Diagnostic Pathology, 2022, 58, 151942.	1.3	1
14	Apparent coexistence of <i>ETV6::RUNX1</i> and <i>KMT2A::MLLT3</i> fusions due to a nonproductive <i>KMT2A</i> rearrangement in B-ALL. Leukemia and Lymphoma, 2022, , 1-4.	1.3	1
15	Utilizing next-generation sequencing to characterize a case of acute myeloid leukemia with $t(4;12)(q12;p13)$ in the absence of ETV6/CHIC2 and ETV6/PDGFRA gene fusions. Cancer Genetics, 2021, 260-261, 1-5.	0.4	O
16	Expert Curation of Somatic FLT3 Variants By the Clingen Somatic Hematologic Cancer Taskforce (ClinGen HCT). Blood, 2021, 138, 4387-4387.	1.4	0
17	Expert Variant Curation Combined with in-Silico analysis for Clinical Interpretation of BCL2 variants in Resistance to BCL2 Inhibitors in Chronic Lymphocytic Leukemia/ Small Lymphocytic Lymphoma. Blood, 2020, 136, 42-43.	1.4	0
18	Clinical Value of Next Generation Sequencing in the Detection of Recurring Structural Rearrangements and Copy Number Abnormalities in Acute Myeloid Leukemia. Blood, 2020, 136, 21-22.	1.4	0

#	Article	IF	CITATIONS
19	Expert Curation of Somatic Variants in Hematological Malignancies By the Clingen Somatic Hematological Cancer Taskforce (ClinGen HCT). Blood, 2020, 136, 23-23.	1.4	O