

# Patricia T Greipp

## List of Publications by Year in descending order

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Version: 2024-02-01

168  
papers

3,042  
citations

159585

30  
h-index

197818

49  
g-index

169  
all docs

169  
docs citations

169  
times ranked

4637  
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparative study of therapy-related and de novo adult B-cell acute lymphoblastic leukaemia. <i>British Journal of Haematology</i> , 2022, 196, 963-968.	2.5	6
2	<i>BRAF</i> Rearrangements and <i>BRAF</i> V600E Mutations Are Seen in a Subset of Pancreatic Carcinomas With Acinar Differentiation. <i>Archives of Pathology and Laboratory Medicine</i> , 2022, 146, 840-845.	2.5	5
3	Detection of a Cryptic <i>KMT2A/AFDN</i> Gene Fusion [ins(6;11)(q27;q23q23)] in a Pediatric Patient with Newly Diagnosed Acute Myeloid Leukemia. <i>Laboratory Medicine</i> , 2022, 53, e95-e99.	1.2	1
4	Myeloid malignancies in cancer patients treated with poly(ADP-ribose) polymerase (PARP) inhibitors: a case series. <i>Blood Cancer Journal</i> , 2022, 12, 11.	6.2	9
5	A Novel <i>USP25::PDGFRA</i> Gene Fusion in a 78 Year Old Patient with a Myeloid Neoplasm. <i>Laboratory Medicine</i> , 2022, 53, e134-e138.	1.2	1
6	A simple additive staging system for newly diagnosed multiple myeloma. <i>Blood Cancer Journal</i> , 2022, 12, 21.	6.2	30
7	Gene fusions in gastrointestinal tract cancers. <i>Genes Chromosomes and Cancer</i> , 2022, 61, 285-297.	2.8	4
8	Identification of <i>EWSR1</i> rearrangements in patients with immature hematopoietic neoplasms: A case series and review of literature. <i>Annals of Diagnostic Pathology</i> , 2022, 58, 151942.	1.3	1
9	BAP1 Immunostain Status in Intraocular Biopsy Specimens for Uveal Melanoma Highly Correlates with Other Prognostic Markers. <i>Ocular Oncology and Pathology</i> , 2022, 8, 22-29.	1.0	0
10	Apparent coexistence of <i>ETV6::RUNX1</i> and <i>KMT2A::MLLT3</i> fusions due to a nonproductive <i>KMT2A</i> rearrangement in B-ALL. <i>Leukemia and Lymphoma</i> , 2022, , 1-4.	1.3	1
11	CD117, BAP1, MTAP, and TdT Is a Useful Immunohistochemical Panel to Distinguish Thymoma from Thymic Carcinoma. <i>Cancers</i> , 2022, 14, 2299.	3.7	6
12	Outcomes following venetoclax-based treatment in therapy-related myeloid neoplasms. <i>American Journal of Hematology</i> , 2022, 97, 1013-1022.	4.1	7
13	Unique characteristics and outcomes of therapy-related acute lymphoblastic leukemia following treatment for multiple myeloma. <i>Blood Cancer Journal</i> , 2022, 12, .	6.2	6
14	Spurious <i>CD34</i> expression in B-cell lymphoma due to nonspecific binding to <i>PerCP-Cy5.5</i> fluorochrome conjugates: A rare phenomenon and a diagnostic pitfall. <i>Cytometry Part B - Clinical Cytometry</i> , 2022, 102, 326-328.	1.5	3
15	Typical, atypical and cryptic t(15;17)(q24;q21) ( <i>PML::RARA</i> ) observed in acute promyelocytic leukemia: A retrospective review of 831 patients with concurrent chromosome and <i>PML::RARA</i> dual-color dual-fusion FISH studies. <i>Genes Chromosomes and Cancer</i> , 2022, 61, 629-634.	2.8	7
16	Mediastinal B-cell lymphoma with <i>MYC</i> , <i>BCL2</i> , and <i>BCL6</i> rearrangements. <i>Journal of Hematopathology</i> , 2022, 15, 151-155.	0.4	2
17	Novel t(1;8)(p31.3;q21.3) <i>NFIA</i> - <i>RUNX1T1</i> Translocation in an Infant Erythroblastic Sarcoma. <i>American Journal of Clinical Pathology</i> , 2021, 156, 129-138.	0.7	10
18	Detection of a Cryptic <i>EP300/ZNF384</i> Gene Fusion by Chromosomal Microarray and Next-Generation Sequencing Studies in a Pediatric Patient with B-Lymphoblastic Leukemia. <i>Laboratory Medicine</i> , 2021, 52, 297-302.	1.2	0

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19	Identification of a novel <i>KMT2A</i> / <i>GIMAP8</i> gene fusion in a pediatric patient with acute undifferentiated leukemia. <i>Genes Chromosomes and Cancer</i> , 2021, 60, 108-111.	2.8	5
20	Impact of Novel Targeted Therapies and Cytogenetic Risk Groups on Outcome After Allogeneic Transplantation for Adult ALL. <i>Transplantation and Cellular Therapy</i> , 2021, 27, 165.e1-165.e11.	1.2	11
21	Erythroblastic sarcoma transformation from a chronic myeloid neoplasm with <i>FGFR1</i> rearrangement presenting as a pleural effusion: a case report. <i>Journal of Hematopathology</i> , 2021, 14, 157-162.	0.4	0
22	Myeloid malignancies with 5q and 7q deletions are associated with extreme genomic complexity, biallelic <i>TP53</i> variants, and very poor prognosis. <i>Blood Cancer Journal</i> , 2021, 11, 18.	6.2	8
23	Inhibition of ATM Induces Hypersensitivity to Proton Irradiation by Upregulating Toxic End Joining. <i>Cancer Research</i> , 2021, 81, 3333-3346.	0.9	16
24	Treatment outcome of clonal cytopenias of undetermined significance: a single-institution retrospective study. <i>Blood Cancer Journal</i> , 2021, 11, 43.	6.2	11
25	Adult Philadelphia-like B-cell acute lymphoblastic leukemia: Characteristics, outcomes, and role of allogeneic hematopoietic cell transplantation in comparison to Philadelphia-positive and Philadelphia-negative acute lymphoblastic leukemia.. <i>Journal of Clinical Oncology</i> , 2021, 39, 7022-7022.	1.6	2
26	Assessment of fixed-duration therapies for treatment-naïve <i>Waldenström</i> macroglobulinemia. <i>American Journal of Hematology</i> , 2021, 96, 945-953.	4.1	12
27	Clinical and biological characteristics and prognostic impact of somatic <i>GATA2</i> mutations in myeloid malignancies: a single institution experience. <i>Blood Cancer Journal</i> , 2021, 11, 122.	6.2	7
28	Increased complexity of t(11;14) rearrangements in plasma cell neoplasms compared with mantle cell lymphoma. <i>Genes Chromosomes and Cancer</i> , 2021, 60, 678-686.	2.8	2
29	Assessment of isochromosome 12p and 12p abnormalities in germ cell tumors using fluorescence in situ hybridization, single-nucleotide polymorphism arrays, and next-generation sequencing/mate-pair sequencing. <i>Human Pathology</i> , 2021, 112, 20-34.	2.0	19
30	The Prognostic Role of <i>MYC</i> Structural Variants Identified by NGS and FISH in Multiple Myeloma. <i>Clinical Cancer Research</i> , 2021, 27, 5430-5439.	7.0	14
31	Clinical utility of next generation sequencing to detect <i>IGH/IL3</i> rearrangements [t(5;14)(q31.1;q32.1)] in B-lymphoblastic leukemia/lymphoma. <i>Annals of Diagnostic Pathology</i> , 2021, 53, 151761.	1.3	8
32	Disease outcomes and biomarkers of progression in smouldering <i>Waldenström</i> macroglobulinaemia. <i>British Journal of Haematology</i> , 2021, 195, 210-216.	2.5	12
33	Identification of adult Philadelphia-like acute lymphoblastic leukemia using a FISH-based algorithm distinguishes prognostic groups and outcomes. <i>Blood Cancer Journal</i> , 2021, 11, 156.	6.2	4
34	Dual Primary <i>IGH</i> Translocations in Multiple Myeloma: A Novel Finding. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2021, 21, e710-e713.	0.4	0
35	OUP accepted manuscript. <i>Laboratory Medicine</i> , 2021, , .	1.2	1
36	Prognostic significance of acquired 1q22 gain in multiple myeloma. <i>American Journal of Hematology</i> , 2021, , .	4.1	6

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37	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/PDGFR $\alpha$ gene fusions. <i>Cancer Genetics</i> , 2021, 260-261, 1-5.	0.4	0
38	MYC break-apart FISH probe set reveals frequent unbalanced patterns of uncertain significance when evaluating aggressive B-cell lymphoma. <i>Blood Cancer Journal</i> , 2021, 11, 184.	6.2	6
39	Myelodysplastic/myeloproliferative neoplasms-unclassifiable with isolated isochromosome 17q represents a distinct clinico-biologic subset: a multi-institutional collaborative study from the Bone Marrow Pathology Group. <i>Modern Pathology</i> , 2021, , .	5.5	9
40	Characterization of Atypical t(11;14) CCND1/IGH Translocations in Multiple Myeloma. <i>Blood</i> , 2021, 138, 3771-3771.	1.4	1
41	Unique Characteristics and Outcomes of Therapy-Related Acute Lymphoblastic Leukemia (trALL) Following Therapy for Multiple Myeloma (MM). <i>Blood</i> , 2021, 138, 2285-2285.	1.4	0
42	False-Negative Centromere 15 Probe Results in Association with African Ancestry in Plasma Cell Dyscrasias. <i>Blood</i> , 2021, 138, 4101-4101.	1.4	0
43	Clinical Characteristics and Prognosis of Thirty-Three Patients with Myeloid Neoplasms and DDX41 Mutation: Mayo Clinic Experience. <i>Blood</i> , 2021, 138, 3691-3691.	1.4	1
44	Fluorescence <i>in situ</i> hybridisation for TP63 rearrangements in T cell lymphomas: single site experience of 470 patients and implications for clinical testing. <i>Histopathology</i> , 2020, 76, 481-485.	2.9	8
45	Ibrutinib monotherapy outside of clinical trial setting in Waldenström macroglobulinaemia: practice patterns, toxicities and outcomes. <i>British Journal of Haematology</i> , 2020, 188, 394-403.	2.5	41
46	Clinical utility of fluorescence <i>in situ</i> hybridization-based diagnosis of BCR-ABL1 like (<math>P</math>-hiladelphia chromosome like) B-acute lymphoblastic leukemia. <i>American Journal of Hematology</i> , 2020, 95, E68-E72.	4.1	4
47	Secondary acquisition of BCR-ABL1 fusion in de novo GATA2-MECOM positive acute myeloid leukemia with subsequent emergence of a rare KMT2A-ASXL2 fusion. <i>Cancer Genetics</i> , 2020, 241, 67-71.	0.4	3
48	Characterization of a cryptic PML-RARA fusion by mate-pair sequencing in a case of acute promyelocytic leukemia with a normal karyotype and negative RARA FISH studies. <i>Leukemia and Lymphoma</i> , 2020, 61, 975-978.	1.3	7
49	Impact of MYD88 <sup>L265P</sup> mutation status on histological transformation of Waldenström Macroglobulinemia. <i>American Journal of Hematology</i> , 2020, 95, 274-281.	4.1	33
50	IgM AL amyloidosis: delineating disease biology and outcomes with clinical, genomic and bone marrow morphological features. <i>Leukemia</i> , 2020, 34, 1373-1382.	7.2	40
51	Implications of MYC Rearrangements in Newly Diagnosed Multiple Myeloma. <i>Clinical Cancer Research</i> , 2020, 26, 6581-6588.	7.0	32
52	Siblings with ETV6/RUNX1-positive B-lymphoblastic leukemia: A single site experience and review of the literature. <i>Annals of Diagnostic Pathology</i> , 2020, 48, 151588.	1.3	1
53	Clinical characteristics and treatment outcomes of newly diagnosed multiple myeloma with chromosome 1q abnormalities. <i>Blood Advances</i> , 2020, 4, 3509-3519.	5.2	58
54	Cytogenetic abnormalities in multiple myeloma: association with disease characteristics and treatment response. <i>Blood Cancer Journal</i> , 2020, 10, 82.	6.2	59

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55	Detection of cryptic CCND1 rearrangements in mantle cell lymphoma by next generation sequencing. <i>Annals of Diagnostic Pathology</i> , 2020, 46, 151533.	1.3	8
56	Twin-to-twin transmission of transient abnormal myelopoiesis without constitutional trisomy 21: A case report. <i>Cancer Genetics</i> , 2020, 244, 62-64.	0.4	1
57	Rare MDM2 amplification in a fat-predominant angiomyolipoma. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2020, 477, 661-666.	2.8	1
58	Characteristics of patients with myelodysplastic syndrome with balanced translocations. <i>British Journal of Haematology</i> , 2020, 190, 244-248.	2.5	1
59	Systematic use of fluorescence <i>in situ</i> hybridisation and clinicopathological features in the screening of PDGFRB rearrangements of patients with myeloid/lymphoid neoplasms. <i>Histopathology</i> , 2020, 76, 1042-1054.	2.9	13
60	Impact of marrow blasts percentage on high-grade myelodysplastic syndrome assessed using revised international prognostic scoring system. <i>Annals of Hematology</i> , 2020, 99, 513-518.	1.8	1
61	Case Report with Review of the Literature: Uveal Melanoma in a Patient with Carney Complex – Another Rare Component of the Syndrome?. <i>Ocular Oncology and Pathology</i> , 2020, 6, 311-317.	1.0	2
62	Increased ERBB2 Gene Copy Numbers Reveal a Subset of Salivary Duct Carcinomas with High Densities of Tumor Infiltrating Lymphocytes and PD-L1 Expression. <i>Head and Neck Pathology</i> , 2020, 14, 951-965.	2.6	11
63	Characterizing false-positive fluorescence in situ hybridization results by mate-pair sequencing in a patient with chronic myeloid leukemia and progression to myeloid blast crisis. <i>Cancer Genetics</i> , 2020, 243, 48-51.	0.4	6
64	Cryptic and atypical KMT2A- <i>USP2</i> and KMT2A- <i>USP8</i> rearrangements identified by mate pair sequencing in infant and childhood leukemia. <i>Genes Chromosomes and Cancer</i> , 2020, 59, 422-427.	2.8	7
65	Metaphase cytogenetics and plasma cell proliferation index for risk stratification in newly diagnosed multiple myeloma. <i>Blood Advances</i> , 2020, 4, 2236-2244.	5.2	20
66	The Prognostic Significance of Acquired 1q22 Gain in Multiple Myeloma. <i>Blood</i> , 2020, 136, 9-10.	1.4	0
67	Predictors of Survival and Time to Progression to Myeloid Neoplasm in Patients with Clonal Cytopenias. <i>Blood</i> , 2020, 136, 26-27.	1.4	1
68	Treatment Outcome for Symptomatic Patients with Clonal Cytopenia of Undetermined Significance: A Single-Institution Retrospective Study. <i>Blood</i> , 2020, 136, 44-44.	1.4	0
69	Heterogeneity of MYC Abnormalities in Multiple Myeloma. <i>Blood</i> , 2020, 136, 2-3.	1.4	0
70	Clinical Value of Next Generation Sequencing in the Detection of Recurring Structural Rearrangements and Copy Number Abnormalities in Acute Myeloid Leukemia. <i>Blood</i> , 2020, 136, 21-22.	1.4	0
71	Hepatic YAP1-TFE3 Rearranged Epithelioid Hemangioendothelioma. <i>Case Reports in Gastrointestinal Medicine</i> , 2019, 2019, 1-5.	0.3	5
72	Identification of a Novel ZBTB20-JAK2 Fusion by Mate-Pair Sequencing in a Young Adult With B-Lymphoblastic Leukemia/Lymphoma. <i>Mayo Clinic Proceedings</i> , 2019, 94, 1381-1384.	3.0	7

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73	Targeting MYC activity in double-hit lymphoma with MYC and BCL2 and/or BCL6 rearrangements with epigenetic bromodomain inhibitors. <i>Journal of Hematology and Oncology</i> , 2019, 12, 73.	17.0	71
74	Whole Genome Mate-pair Sequencing of Plasma Cell Neoplasm as a Novel Diagnostic Strategy: A Case of Unrecognized t(2;11) Structural Variation. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2019, 19, 598-602.	0.4	2
75	Characterization of a rarely reported STAT5B/RARA gene fusion in a young adult with newly diagnosed acute promyelocytic leukemia with resistance to ATRA therapy. <i>Cancer Genetics</i> , 2019, 237, 51-54.	0.4	5
76	Characterization of TCF3 rearrangements in pediatric B-lymphoblastic leukemia/lymphoma by mate-pair sequencing (MPseq) identifies complex genomic rearrangements and a novel TCF3/TEF gene fusion. <i>Blood Cancer Journal</i> , 2019, 9, 81.	6.2	14
77	Atypical lipomatous tumour/well-differentiated liposarcoma and dedifferentiated liposarcoma in patients aged 40 years: a study of 116 patients. <i>Histopathology</i> , 2019, 75, 833-842.	2.9	16
78	Amplification of 9p24.1 in diffuse large B-cell lymphoma identifies a unique subset of cases that resemble primary mediastinal large B-cell lymphoma. <i>Blood Cancer Journal</i> , 2019, 9, 73.	6.2	37
79	Tetraploidy is associated with poor prognosis at diagnosis in multiple myeloma. <i>American Journal of Hematology</i> , 2019, 94, E117-E120.	4.1	13
80	Aurora kinase B-phosphorylated HP1± functions in chromosomal instability. <i>Cell Cycle</i> , 2019, 18, 1407-1421.	2.6	7
81	Characterization of a t(1;2)(p36;p21) involving the PRDM16 gene region by mate-pair sequencing (MPseq) in a patient with newly diagnosed acute myeloid leukemia with myelodysplasia-related changes. <i>Journal of Hematopathology</i> , 2019, 12, 85-90.	0.4	0
82	Characterization of a cryptic KMT2A/AFF1 gene fusion by mate-pair sequencing (MPseq) in a young adult with newly diagnosed B-lymphoblastic leukemia. <i>Journal of Hematopathology</i> , 2019, 12, 99-104.	0.4	1
83	Elucidating a false-negative MYC break-apart fluorescence in situ hybridization probe study by next-generation sequencing in a patient with high-grade B-cell lymphoma with IGH/MYC and IGH/BCL2 rearrangements. <i>Journal of Physical Education and Sports Management</i> , 2019, 5, a004077.	1.2	14
84	Immunohistochemistry for TFE3 lacks specificity and sensitivity in the diagnosis of TFE3-rearranged neoplasms: a comparative, 2-laboratory study. <i>Human Pathology</i> , 2019, 87, 65-74.	2.0	41
85	Natural history of multiple myeloma with de novo del(17p). <i>Blood Cancer Journal</i> , 2019, 9, 32.	6.2	38
86	Hyperhaploid plasma cell myeloma characterized by poor outcome and monosomy 17 with frequently co-occurring TP53 mutations. <i>Blood Cancer Journal</i> , 2019, 9, 20.	6.2	10
87	Detection of a cryptic NUP214/ABL1 gene fusion by mate-pair sequencing (MPseq) in a newly diagnosed case of pediatric T-lymphoblastic leukemia. <i>Journal of Physical Education and Sports Management</i> , 2019, 5, a003533.	1.2	8
88	A near-chaploid clone harboring a BCR/ABL1 gene fusion in an adult patient with newly diagnosed B-lymphoblastic leukemia. <i>Genes Chromosomes and Cancer</i> , 2019, 58, 665-668.	2.8	3
89	Acute leukemias harboring KMT2A/MLLT10 fusion: a 10-year experience from a single genomics laboratory. <i>Genes Chromosomes and Cancer</i> , 2019, 58, 567-577.	2.8	19
90	Impact of acquired del(17p) in multiple myeloma. <i>Blood Advances</i> , 2019, 3, 1930-1938.	5.2	41

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91	Characterization of a cryptic IGH/CCND1 rearrangement in a case of mantle cell lymphoma with negative CCND1 FISH studies. <i>Blood Advances</i> , 2019, 3, 1298-1302.	5.2	16
92	Mate pair sequencing outperforms fluorescence in situ hybridization and improves diagnostic yield in the genomic characterization of multiple myeloma. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2019, 19, e58.	0.4	0
93	Mate pair sequencing outperforms fluorescence in situ hybridization in the genomic characterization of multiple myeloma. <i>Blood Cancer Journal</i> , 2019, 9, 103.	6.2	27
94	Rapid assessment of hyperdiploidy in plasma cell disorders using a novel multiparametric flow cytometry method. <i>American Journal of Hematology</i> , 2019, 94, 424-430.	4.1	11
95	Large Chromosomal Rearrangements Yield Biomarkers to Distinguish Low-Risk From Intermediate- and High-Risk Prostate Cancer. <i>Mayo Clinic Proceedings</i> , 2019, 94, 27-36.	3.0	7
96	Multiple isodicentric Y chromosomes in myeloid malignancies: a unique cytogenetic entity and potential therapeutic target. <i>Leukemia and Lymphoma</i> , 2019, 60, 821-824.	1.3	2
97	False-negative rates for <i>MYC</i> fluorescence in situ hybridization probes in B-cell neoplasms. <i>Haematologica</i> , 2019, 104, e248-e251.	3.5	43
98	Constitutional chromosome rearrangements that mimic the 2017 world health organization acute myeloid leukemia with recurrent genetic abnormalities: A study of three cases and review of the literature. <i>Cancer Genetics</i> , 2019, 230, 37-46.	0.4	8
99	Use of mate-pair sequencing to characterize a complex cryptic BCR/ABL1 rearrangement observed in a newly diagnosed case of chronic myeloid leukemia. <i>Human Pathology</i> , 2019, 89, 109-114.	2.0	7
100	Elderly acute lymphoblastic leukemia: a Mayo Clinic study of 124 patients. <i>Leukemia and Lymphoma</i> , 2019, 60, 990-999.	1.3	9
101	Mate pair sequencing improves detection of genomic abnormalities in acute myeloid leukemia. <i>European Journal of Haematology</i> , 2019, 102, 87-96.	2.2	35
102	Reassignment of HER2 status for subgroups of breast cancer according to the 2018 updated American Society of Clinical Oncology and College of American Pathologists guidelines: The impact of combined immunohistochemistry (IHC) and fluorescence in situ hybridization (FISH) reflex testing in a large national reference laboratory. <i>Journal of Clinical Oncology</i> , 2019, 37, 3144-3144.	1.6	0
103	Identification of Targetable Tumor Associated Proteins in Adult T-Acute Lymphoblastic Leukemia/Lymphoma (T-ALL/LBL) Including a Novel CC-Chemokine 4 (CCR4)-Positive T-ALL/LBL with Unique Immunophenotype. <i>Blood</i> , 2019, 134, 5210-5210.	1.4	0
104	Impact of Targeted Immunotherapies and Novel Cytogenetic and Clinical Risk Groups on Outcome after Allogeneic Hematopoietic Stem Cell Transplant (AlloHCT) for Acute Lymphoblastic Leukemia (ALL): The Mayo Clinic Cohort. <i>Blood</i> , 2019, 134, 2588-2588.	1.4	0
105	A Novel Approach to Risk Stratification in Multiple Myeloma Using ISS Stage and FISH. <i>Blood</i> , 2019, 134, 1800-1800.	1.4	1
106	Prognostic significance of interphase FISH in monoclonal gammopathy of undetermined significance. <i>Leukemia</i> , 2018, 32, 1811-1815.	7.2	28
107	De novo pure erythroid leukemia: refining the clinicopathologic and cytogenetic characteristics of a rare entity. <i>Modern Pathology</i> , 2018, 31, 705-717.	5.5	25
108	Recurrent STAT3-JAK2 fusions in indolent T-cell lymphoproliferative disorder of the gastrointestinal tract. <i>Blood</i> , 2018, 131, 2262-2266.	1.4	77

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109	Polypoid fibroadipose tumors of the esophagus: "giant fibrovascular polyp" or liposarcoma? A clinicopathological and molecular cytogenetic study of 13 cases. <i>Modern Pathology</i> , 2018, 31, 337-342.	5.5	37
110	Molecular testing for the clinical diagnosis of fibrolamellar carcinoma. <i>Modern Pathology</i> , 2018, 31, 141-149.	5.5	47
111	Fibrolamellar carcinoma in the Carney complex: PRKAR1A loss instead of the classic DNAJB1-PRKACA fusion. <i>Hepatology</i> , 2018, 68, 1441-1447.	7.3	48
112	Loss of BAP1 Expression in Atypical Mesothelial Proliferations Helps to Predict Malignant Mesothelioma. <i>American Journal of Surgical Pathology</i> , 2018, 42, 256-263.	3.7	34
113	PDGFB Rearrangements in Dermatofibrosarcoma Protuberans of the Vulva: A Study of 11 Cases Including Myxoid and Fibrosarcomatous Variants. <i>International Journal of Gynecological Pathology</i> , 2018, 37, 537-546.	1.4	9
114	A 10-Year Review of KMT2A Gene Fusion Partners Observed in Pediatric T-Lymphoblastic Leukemia/Lymphoma: The Mayo Clinic Experience. <i>American Journal of Clinical Pathology</i> , 2018, 150, S132-S132.	0.7	0
115	Use of Mate-Pair Sequencing (MPseq) to Elucidate a Complex BCR-ABL1 Rearrangement Observed in a Newly Diagnosed Case of Chronic Myeloid Leukemia. <i>American Journal of Clinical Pathology</i> , 2018, 150, S131-S132.	0.7	0
116	Loss of TNFAIP3 enhances MYD88L265P-driven signaling in non-Hodgkin lymphoma. <i>Blood Cancer Journal</i> , 2018, 8, 97.	6.2	36
117	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
118	Impact of clone size with a single cytogenetic abnormality on the revised International Prognostic Scoring System in myelodysplastic syndromes. <i>American Journal of Hematology</i> , 2018, 93, E398-E401.	4.1	1
119	KMT2A (MLL) rearrangements observed in pediatric/young adult T-lymphoblastic leukemia/lymphoma: A 10-year review from a single cytogenetic laboratory. <i>Genes Chromosomes and Cancer</i> , 2018, 57, 541-546.	2.8	21
120	14. Clinical mate-pair sequencing reveals complex genomic rearrangements in B-lymphoblastic leukemia/lymphoma (B-ALL). <i>Cancer Genetics</i> , 2018, 224-225, 55-56.	0.4	0
121	Defining Lymphoplasmacytic Lymphoma. <i>American Journal of Clinical Pathology</i> , 2018, 150, 168-176.	0.7	5
122	12. Mate pair sequencing: Unveiling underappreciated complexity and providing clarity to the previously unanswered questions of cytogenetics. <i>Cancer Genetics</i> , 2018, 224-225, 54-55.	0.4	0
123	A Test Utilization Approach to the Diagnostic Workup of Isolated Eosinophilia in Otherwise Morphologically Unremarkable Bone Marrow. <i>American Journal of Clinical Pathology</i> , 2018, 150, 421-431.	0.7	12
124	The Utilization of Chromosomal Microarray Technologies for Hematologic Neoplasms. <i>American Journal of Clinical Pathology</i> , 2018, 150, 375-384.	0.7	13
125	The clinical outcomes of reclassified erythroleukemia (erythroid/myeloid) as myelodysplastic syndrome (MDS) per 2017 WHO guideline compared to MDS. <i>American Journal of Hematology</i> , 2018, 93, E355-E357.	4.1	2
126	13. Clinical utility of mate pair sequencing to detect diagnostic and prognostic chromosomal rearrangements and copy number changes in patients with acute myeloid leukemia. <i>Cancer Genetics</i> , 2018, 224-225, 55.	0.4	0



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127	IgM Associated Light Chain (AL) Amyloidosis: Delineating Disease Biology with Clinical, Genomic and Bone Marrow Morphological Features. <i>Blood</i> , 2018, 132, 4460-4460.	1.4	1
128	The impact of clonal size on the revised international prognostic scoring system (R-IPSS) in myelodysplastic syndromes (MDS) with a single cytogenetic abnormality.. <i>Journal of Clinical Oncology</i> , 2018, 36, 7068-7068.	1.6	0
129	The clinical outcomes of reclassified erythroleukemia (erythroid/myeloid) as myelodysplastic syndrome per 2017 WHO guideline compared to de novo MDS.. <i>Journal of Clinical Oncology</i> , 2018, 36, e19026-e19026.	1.6	0
130	Abstract 655: Hyperdiploidy in plasma cell disorders using multi-parametric flow cytometry (MFC) vs. FISH. , 2018, , .		0
131	Marrow Blast Percentage Impact on High-Grade Myelodysplastic Syndrome By the Revised International Prognostic Scoring System. <i>Blood</i> , 2018, 132, 5510-5510.	1.4	0
132	Impact of Acquired Del(17p) in Patients with Multiple Myeloma. <i>Blood</i> , 2018, 132, 4449-4449.	1.4	0
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