

# Hiroko Tanaka

## List of Publications by Citations

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**Version:** 2024-04-25

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141  
papers

6,917  
citations

40  
h-index

82  
g-index

147  
ext. papers

8,912  
ext. citations

8.8  
avg, IF

4.7  
L-index

#	Paper	IF	Citations
141	Mutational landscape and clonal architecture in grade II and III gliomas. <i>Nature Genetics</i> , <b>2015</b> , 47, 458-68	36.3	543
140	Integrated molecular analysis of adult T cell leukemia/lymphoma. <i>Nature Genetics</i> , <b>2015</b> , 47, 1304-15	36.3	469
139	Whole-genome mutational landscape and characterization of noncoding and structural mutations in liver cancer. <i>Nature Genetics</i> , <b>2016</b> , 48, 500-9	36.3	423
138	Somatic RHOA mutation in angioimmunoblastic T cell lymphoma. <i>Nature Genetics</i> , <b>2014</b> , 46, 171-5	36.3	411
137	Aberrant PD-L1 expression through 3'UTR disruption in multiple cancers. <i>Nature</i> , <b>2016</b> , 534, 402-6	50.4	403
136	Age-related remodelling of oesophageal epithelia by mutated cancer drivers. <i>Nature</i> , <b>2019</b> , 565, 312-317	50.4	270
135	The landscape of somatic mutations in Down syndrome-related myeloid disorders. <i>Nature Genetics</i> , <b>2013</b> , 45, 1293-9	36.3	244
134	Dynamics of clonal evolution in myelodysplastic syndromes. <i>Nature Genetics</i> , <b>2017</b> , 49, 204-212	36.3	228
133	Inherited and Somatic Defects in DDX41 in Myeloid Neoplasms. <i>Cancer Cell</i> , <b>2015</b> , 27, 658-70	24.3	228
132	Genetic abnormalities in myelodysplasia and secondary acute myeloid leukemia: impact on outcome of stem cell transplantation. <i>Blood</i> , <b>2017</b> , 129, 2347-2358	2.2	184
131	Somatic SETBP1 mutations in myeloid malignancies. <i>Nature Genetics</i> , <b>2013</b> , 45, 942-6	36.3	178
130	Exome sequencing identifies secondary mutations of SETBP1 and JAK3 in juvenile myelomonocytic leukemia. <i>Nature Genetics</i> , <b>2013</b> , 45, 937-41	36.3	175
129	Acquired initiating mutations in early hematopoietic cells of CLL patients. <i>Cancer Discovery</i> , <b>2014</b> , 4, 1088-1091	10.1	172
128	ACTN1 mutations cause congenital macrothrombocytopenia. <i>American Journal of Human Genetics</i> , <b>2013</b> , 92, 431-8	11	154
127	Genomic characterization of biliary tract cancers identifies driver genes and predisposing mutations. <i>Journal of Hepatology</i> , <b>2018</b> , 68, 959-969	13.4	149
126	Recurrent somatic mutations underlie corticotropin-independent Cushing's syndrome. <i>Science</i> , <b>2014</b> , 344, 917-20	33.3	142
125	Whole-genome mutational landscape of liver cancers displaying biliary phenotype reveals hepatitis impact and molecular diversity. <i>Nature Communications</i> , <b>2015</b> , 6, 6120	17.4	139

124	Profiling of somatic mutations in acute myeloid leukemia with FLT3-ITD at diagnosis and relapse. <i>Blood</i> , <b>2015</b> , 126, 2491-501	2.2	134
123	Integrated genetic and epigenetic analysis defines novel molecular subgroups in rhabdomyosarcoma. <i>Nature Communications</i> , <b>2015</b> , 6, 7557	17.4	110
122	Mutations in the gene encoding the E2 conjugating enzyme UBE2T cause Fanconi anemia. <i>American Journal of Human Genetics</i> , <b>2015</b> , 96, 1001-7	11	90
121	Clonal evolution in myelodysplastic syndromes. <i>Nature Communications</i> , <b>2017</b> , 8, 15099	17.4	86
120	Defective Epstein-Barr virus in chronic active infection and haematological malignancy. <i>Nature Microbiology</i> , <b>2019</b> , 4, 404-413	26.6	80
119	Variegated RHOA mutations in adult T-cell leukemia/lymphoma. <i>Blood</i> , <b>2016</b> , 127, 596-604	2.2	77
118	Frequent mutations that converge on the NFKBIZ pathway in ulcerative colitis. <i>Nature</i> , <b>2020</b> , 577, 260-265	50.4	77
117	Prognostic relevance of integrated genetic profiling in adult T-cell leukemia/lymphoma. <i>Blood</i> , <b>2018</b> , 131, 215-225	2.2	76
116	Genomic landscape of liposarcoma. <i>Oncotarget</i> , <b>2015</b> , 6, 42429-44	3.3	75
115	Recurrent SPI1 (PU.1) fusions in high-risk pediatric T cell acute lymphoblastic leukemia. <i>Nature Genetics</i> , <b>2017</b> , 49, 1274-1281	36.3	74
114	Molecular heterogeneity in peripheral T-cell lymphoma, not otherwise specified revealed by comprehensive genetic profiling. <i>Leukemia</i> , <b>2019</b> , 33, 2867-2883	10.7	71
113	Haploinsufficiency of TNFAIP3 (A20) by germline mutation is involved in autoimmune lymphoproliferative syndrome. <i>Journal of Allergy and Clinical Immunology</i> , <b>2017</b> , 139, 1914-1922	11.5	69
112	Loss of function mutations in RPL27 and RPS27 identified by whole-exome sequencing in Diamond-Blackfan anaemia. <i>British Journal of Haematology</i> , <b>2015</b> , 168, 854-64	4.5	67
111	Abnormal hematopoiesis and autoimmunity in human subjects with germline IKZF1 mutations. <i>Journal of Allergy and Clinical Immunology</i> , <b>2017</b> , 140, 223-231	11.5	63
110	Whole genome sequencing discriminates hepatocellular carcinoma with intrahepatic metastasis from multi-centric tumors. <i>Journal of Hepatology</i> , <b>2017</b> , 66, 363-373	13.4	62
109	Phosphatase and tensin homolog (PTEN) mutation can cause activated phosphatidylinositol 3-kinase $\beta$ syndrome-like immunodeficiency. <i>Journal of Allergy and Clinical Immunology</i> , <b>2016</b> , 138, 1672-1680.e10	11.5	61
108	Integrated analysis of whole genome and transcriptome sequencing reveals diverse transcriptomic aberrations driven by somatic genomic changes in liver cancers. <i>PLoS ONE</i> , <b>2014</b> , 9, e114263	3.7	58
107	Frequent structural variations involving programmed death ligands in Epstein-Barr virus-associated lymphomas. <i>Leukemia</i> , <b>2019</b> , 33, 1687-1699	10.7	57

106	Biallelic DICER1 mutations in sporadic pleuropulmonary blastoma. <i>Cancer Research</i> , <b>2014</b> , 74, 2742-9	10.1	56
105	Integrated molecular profiling of juvenile myelomonocytic leukemia. <i>Blood</i> , <b>2018</b> , 131, 1576-1586	2.2	51
104	Clinical utility of next-generation sequencing for inherited bone marrow failure syndromes. <i>Genetics in Medicine</i> , <b>2017</b> , 19, 796-802	8.1	45
103	Whole-exome sequencing reveals the spectrum of gene mutations and the clonal evolution patterns in paediatric acute myeloid leukaemia. <i>British Journal of Haematology</i> , <b>2016</b> , 175, 476-489	4.5	42
102	Landscape and function of multiple mutations within individual oncogenes. <i>Nature</i> , <b>2020</b> , 582, 95-99	50.4	41
101	GATA2 and secondary mutations in familial myelodysplastic syndromes and pediatric myeloid malignancies. <i>Haematologica</i> , <b>2015</b> , 100, e398-401	6.6	38
100	A comprehensive characterization of -acting splicing-associated variants in human cancer. <i>Genome Research</i> , <b>2018</b> , 28, 1111-1125	9.7	32
99	Frequent germline mutations of in sporadic subcutaneous panniculitis-like T-cell lymphoma. <i>Blood Advances</i> , <b>2019</b> , 3, 588-595	7.8	31
98	Characterization of HBV integration patterns and timing in liver cancer and HBV-infected livers. <i>Oncotarget</i> , <b>2018</b> , 9, 25075-25088	3.3	30
97	Invariant patterns of clonal succession determine specific clinical features of myelodysplastic syndromes. <i>Nature Communications</i> , <b>2019</b> , 10, 5386	17.4	29
96	TERT promoter mutations and chromosome 8p loss are characteristic of nonalcoholic fatty liver disease-related hepatocellular carcinoma. <i>International Journal of Cancer</i> , <b>2016</b> , 139, 2512-8	7.5	28
95	Classification of primary liver cancer with immunosuppression mechanisms and correlation with genomic alterations. <i>EBioMedicine</i> , <b>2020</b> , 53, 102659	8.8	26
94	Exome sequencing identified as a novel causative gene for Diamond-Blackfan anemia. <i>Haematologica</i> , <b>2017</b> , 102, e93-e96	6.6	25
93	Recurrent genetic defects on chromosome 5q in myeloid neoplasms. <i>Oncotarget</i> , <b>2017</b> , 8, 6483-6495	3.3	24
92	Loss of DNA Damage Response in Neuroblastoma and Utility of a PARP Inhibitor. <i>Journal of the National Cancer Institute</i> , <b>2017</b> , 109,	9.7	23
91	Genomic landscape of colitis-associated cancer indicates the impact of chronic inflammation and its stratification by mutations in the Wnt signaling. <i>Oncotarget</i> , <b>2018</b> , 9, 969-981	3.3	23
90	Transcriptome analysis offers a comprehensive illustration of the genetic background of pediatric acute myeloid leukemia. <i>Blood Advances</i> , <b>2019</b> , 3, 3157-3169	7.8	21
89	Frequent genetic alterations in immune checkpoint-related genes in intravascular large B-cell lymphoma. <i>Blood</i> , <b>2021</b> , 137, 1491-1502	2.2	19

88	Late-Onset Combined Immunodeficiency with a Novel IL2RG Mutation and Probable Revertant Somatic Mosaicism. <i>Journal of Clinical Immunology</i> , <b>2015</b> , 35, 610-4	5.7	18
87	PIEZO1 gene mutation in a Japanese family with hereditary high phosphatidylcholine hemolytic anemia and hemochromatosis-induced diabetes mellitus. <i>International Journal of Hematology</i> , <b>2016</b> , 104, 125-9	2.3	18
86	De Novo Mutations Activating Germline TP53 in an Inherited Bone-Marrow-Failure Syndrome. <i>American Journal of Human Genetics</i> , <b>2018</b> , 103, 440-447	11	18
85	Genetic and transcriptional landscape of plasma cells in POEMS syndrome. <i>Leukemia</i> , <b>2019</b> , 33, 1723-1735.	5.7	18
84	BRCC3 mutations in myeloid neoplasms. <i>Haematologica</i> , <b>2015</b> , 100, 1051-7	6.6	17
83	Identification of the genetic and clinical characteristics of neuroblastomas using genome-wide analysis. <i>Oncotarget</i> , <b>2017</b> , 8, 107513-107529	3.3	17
82	Common Variable Immunodeficiency Caused by FANC Mutations. <i>Journal of Clinical Immunology</i> , <b>2017</b> , 37, 434-444	5.7	15
81	Integrated Molecular Characterization of the Lethal Pediatric Cancer Pancreatoblastoma. <i>Cancer Research</i> , <b>2018</b> , 78, 865-876	10.1	15
80	Clinical and genetic features of dyskeratosis congenita, cryptic dyskeratosis congenita, and Hoyeraal-Hreidarsson syndrome in Japan. <i>International Journal of Hematology</i> , <b>2015</b> , 102, 544-52	2.3	14
79	Diagnostic challenge of Diamond-Blackfan anemia in mothers and children by whole-exome sequencing. <i>International Journal of Hematology</i> , <b>2017</b> , 105, 515-520	2.3	14
78	Novel neuroblastoma amplified sequence () mutations in a Japanese boy with fever-triggered recurrent acute liver failure. <i>Human Genome Variation</i> , <b>2019</b> , 6, 2	1.8	14
77	Molecular classification and diagnostics of upper urinary tract urothelial carcinoma. <i>Cancer Cell</i> , <b>2021</b> , 39, 793-809.e8	24.3	13
76	Genetic heterogeneity of uncharacterized childhood autoimmune diseases with lymphoproliferation. <i>Pediatric Blood and Cancer</i> , <b>2018</b> , 65, e26831	3	12
75	Genomic Heterogeneity Within Individual Prostate Cancer Foci Impacts Predictive Biomarkers of Targeted Therapy. <i>European Urology Focus</i> , <b>2019</b> , 5, 416-424	5.1	12
74	Recurrent mutations in -rearranged acute myeloid leukemia. <i>Blood Advances</i> , <b>2018</b> , 2, 2879-2889	7.8	12
73	ASXL2 mutations are frequently found in pediatric AML patients with t(8;21)/ RUNX1-RUNX1T1 and associated with a better prognosis. <i>Genes Chromosomes and Cancer</i> , <b>2017</b> , 56, 382-393	5	11
72	Whole-exome sequence analysis of ataxia telangiectasia-like phenotype. <i>Journal of the Neurological Sciences</i> , <b>2014</b> , 340, 86-90	3.2	11
71	Integrated multiomics analysis of hepatoblastoma unravels its heterogeneity and provides novel druggable targets. <i>Npj Precision Oncology</i> , <b>2020</b> , 4, 20	9.8	11

70	Somatic mosaicism in chronic myeloid leukemia in remission. <i>Blood</i> , <b>2016</b> , 128, 2863-2866	2.2	11
69	Genomic analysis of pancreatic juice DNA assesses malignant risk of intraductal papillary mucinous neoplasm of pancreas. <i>Cancer Medicine</i> , <b>2019</b> , 8, 4565-4573	4.8	10
68	Dysregulation of Epstein-Barr Virus Infection in Hypomorphic ZAP70 Mutation. <i>Journal of Infectious Diseases</i> , <b>2018</b> , 218, 825-834	7	10
67	Unbiased Detection of Driver Mutations in Extramammary Paget Disease. <i>Clinical Cancer Research</i> , <b>2021</b> , 27, 1756-1765	12.9	10
66	Landscape of driver mutations and their clinical impacts in pediatric B-cell precursor acute lymphoblastic leukemia. <i>Blood Advances</i> , <b>2020</b> , 4, 5165-5173	7.8	10
65	Combined landscape of single-nucleotide variants and copy number alterations in clonal hematopoiesis. <i>Nature Medicine</i> , <b>2021</b> , 27, 1239-1249	50.5	10
64	Clonally related diffuse large B-cell lymphoma and interdigitating dendritic cell sarcoma sharing translocation. <i>Haematologica</i> , <b>2018</b> , 103, e553-e556	6.6	9
63	Integrated genetic and epigenetic analysis revealed heterogeneity of acute lymphoblastic leukemia in Down syndrome. <i>Cancer Science</i> , <b>2019</b> , 110, 3358-3367	6.9	8
62	Genetic and clinical landscape of breast cancers with germline BRCA1/2 variants. <i>Communications Biology</i> , <b>2020</b> , 3, 578	6.7	7
61	Clonal evolution and clinical implications of genetic abnormalities in blastic transformation of chronic myeloid leukaemia. <i>Nature Communications</i> , <b>2021</b> , 12, 2833	17.4	7
60	Genomic analysis of clonal origin of Langerhans cell histiocytosis following acute lymphoblastic leukaemia. <i>British Journal of Haematology</i> , <b>2016</b> , 175, 169-72	4.5	7
59	Constitutional abnormalities of IDH1 combined with secondary mutations predispose a patient with Maffucci syndrome to acute lymphoblastic leukemia. <i>Pediatric Blood and Cancer</i> , <b>2017</b> , 64, e26647	3	6
58	Whole genome sequencing analysis identifies recurrent structural alterations in esophageal squamous cell carcinoma. <i>PeerJ</i> , <b>2020</b> , 8, e9294	3.1	6
57	Genetic Predispositions to Myeloid Neoplasms Caused By Germline DDX41 Mutations. <i>Blood</i> , <b>2015</b> , 126, 2843-2843	2.2	5
56	Fusion partner-specific mutation profiles and KRAS mutations as adverse prognostic factors in MLL-rearranged AML. <i>Blood Advances</i> , <b>2020</b> , 4, 4623-4631	7.8	5
55	Modification of cellular and humoral immunity by somatically reverted T cells in X-linked lymphoproliferative syndrome type 1. <i>Journal of Allergy and Clinical Immunology</i> , <b>2019</b> , 143, 421-424.e11	11.5	5
54	Molecular studies reveal and gene fusions displaced in a case of infantile acute lymphoblastic leukemia with complex karyotype. <i>Oncology Letters</i> , <b>2017</b> , 14, 2295-2299	2.6	4
53	In Analogy to AML, MDS Can be Sub-Classified By Ancestral Mutations. <i>Blood</i> , <b>2014</b> , 124, 823-823	2.2	4

52	Landscape Of Genetic Lesions In 944 Patients With Myelodysplastic Syndromes. <i>Blood</i> , <b>2013</b> , 122, 521-5212		3
51	Impact of Somatic Mutations on Outcome in Patients with MDS after Stem-Cell Transplantation. <i>Blood</i> , <b>2015</b> , 126, 711-711	2.2	3
50	Comprehensive genetic analysis of pediatric germ cell tumors identifies potential drug targets. <i>Communications Biology</i> , <b>2020</b> , 3, 544	6.7	3
49	Frequent mutations in HLA and related genes in extranodal NK/T cell lymphomas. <i>Leukemia and Lymphoma</i> , <b>2021</b> , 62, 95-103	1.9	3
48	Molecular Classification and Tumor Microenvironment Characterization of Gallbladder Cancer by Comprehensive Genomic and Transcriptomic Analysis. <i>Cancers</i> , <b>2021</b> , 13,	6.6	3
47	Clinical Impacts of Germline DDX41 Mutations on Myeloid Neoplasms. <i>Blood</i> , <b>2020</b> , 136, 38-40	2.2	2
46	Somatic G17V Rhoa Mutation Specifies Angioimmunoblastic T-Cell Lymphoma. <i>Blood</i> , <b>2013</b> , 122, 815-815	2.2	2
45	Chronological Analysis of Clonal Evolution in Acquired Aplastic Anemia. <i>Blood</i> , <b>2014</b> , 124, 253-253	2.2	2
44	Frequent Activating Somatic Alterations in T-Cell Receptor / NF- $\kappa$ B Signaling in Adult T-Cell Leukemia/Lymphoma. <i>Blood</i> , <b>2015</b> , 126, 113-113	2.2	2
43	Serial Sequencing in Myelodysplastic Syndromes Reveals Dynamic Changes in Clonal Architecture and Allows for a New Prognostic Assessment of Mutations Detected in Cross-Sectional Testing. <i>Blood</i> , <b>2015</b> , 126, 709-709	2.2	2
42	Distinct gene alterations with a high percentage of myeloperoxidase-positive leukemic blasts in de novo acute myeloid leukemia. <i>Leukemia Research</i> , <b>2018</b> , 65, 34-41	2.7	1
41	Genotype-Phenotype Relationships and Therapeutic Targets in Acute Erythroid Leukemia. <i>Blood</i> , <b>2020</b> , 136, 17-18	2.2	1
40	Prognostic Relevance of Genetic Abnormalities in Blastic Transformation of Chronic Myeloid Leukemia. <i>Blood</i> , <b>2020</b> , 136, 3-4	2.2	1
39	Landscape of Genetic Alterations in Adult T-Cell Leukemia/Lymphoma. <i>Blood</i> , <b>2014</b> , 124, 75-75	2.2	1
38	Prognostic Relevance of Integrated Genetic Profiling in Adult T-Cell Leukemia/Lymphoma. <i>Blood</i> , <b>2015</b> , 126, 2643-2643	2.2	1
37	Two Novel Distinct Subtypes of Myeloid Neoplasms Molecularly Associated with Histone H3K36 Methylations. <i>Blood</i> , <b>2015</b> , 126, 2841-2841	2.2	1
36	Genetic Background of Idiopathic Bone Marrow Failure Syndromes in Children. <i>Blood</i> , <b>2015</b> , 126, 3610-3610		1
35	EPOR/JAK/STAT Signaling Pathway As Therapeutic Target of Acute Erythroid Leukemia. <i>Blood</i> , <b>2021</b> , 138, 610-610	2.2	1

34	Alteration of the immune environment in bone marrow from children with recurrent B cell precursor acute lymphoblastic leukemia. <i>Cancer Science</i> , <b>2021</b> ,	6.9	1
33	Genetic Basis of Primary Central Nervous System Lymphoma. <i>Blood</i> , <b>2015</b> , 126, 2687-2687	2.2	1
32	DDX41 Is a Tumor Suppressor Gene Associated with Inherited and Acquired Mutations. <i>Blood</i> , <b>2014</b> , 124, 125-125	2.2	0
31	Whole-Genome Sequencing of Primary Central Nervous System Lymphoma and Diffuse Large B-Cell Lymphoma. <i>Blood</i> , <b>2016</b> , 128, 4112-4112	2.2	0
30	Development of Philadelphia chromosome-negative acute myeloid leukemia with IDH2 and NPM1 mutations in a patient with chronic myeloid leukemia who showed a major molecular response to tyrosine kinase inhibitor therapy. <i>International Journal of Hematology</i> , <b>2021</b> , 113, 936-940	2.3	0
29	Genomic analysis of multiple myeloma using targeted capture sequencing in the Japanese cohort. <i>British Journal of Haematology</i> , <b>2020</b> , 191, 755-763	4.5	
28	Post-Treatment Clone Size Predicts Survival Independently of IPSS-R and Response after Azacitidine Therapy for MDS.. <i>Blood</i> , <b>2020</b> , 136, 12-13	2.2	
27	ASXL1 Mutations Predict a Poor Response to Darbepoetin Alfa in Anemic Patients with Low-Risk MDS: A Multicenter, Phase II Study. <i>Blood</i> , <b>2020</b> , 136, 28-29	2.2	
26	KRAS mutations Frequently Coexist with High-Risk MLL Fusions and Are Independent Adverse Prognostic Factors in MLL-Rearranged Acute Myeloid Leukemia. <i>Blood</i> , <b>2020</b> , 136, 28-29	2.2	
25	Analysis of Genomic Predispositions to Sporadic Myeloid Neoplasms Mediated By DDX41 in Japan. <i>Blood</i> , <b>2018</b> , 132, 4371-4371	2.2	
24	Molecular Characteristics That Predict Response to Azacitidine Therapy. <i>Blood</i> , <b>2019</b> , 134, 4246-4246	2.2	
23	Integrated Analysis of Copy-Number Alterations and Gene Mutations in 2,000 Patients with Myeloid Neoplasms. <i>Blood</i> , <b>2019</b> , 134, 4216-4216	2.2	
22	Whole Exome and Transcriptome Analyses in Pediatric T-Cell Acute Lymphoblastic Leukemia. <i>Blood</i> , <b>2014</b> , 124, 3527-3527	2.2	
21	Novel Biological Effects and Distinct Patterns of Rhoa Mutations in Adult T-Cell Leukemia/Lymphoma and Angioimmunoblastic T Cell Lymphoma. <i>Blood</i> , <b>2014</b> , 124, 2215-2215	2.2	
20	Clinical and Molecular Significance of Peripheral Blood Cell-Free DNA in B-Cell Lymphomas for Detection of Genetic Mutations and Correlation with Disease Status. <i>Blood</i> , <b>2014</b> , 124, 1658-1658	2.2	
19	Diagnostic Efficacy of Whole-Exome Sequencing in 250 Patients with Congenital Bone Marrow Failure. <i>Blood</i> , <b>2014</b> , 124, 4385-4385	2.2	
18	Whole-Exome Sequencing Reveals a Paucity of Somatic Gene Mutations in Aplastic Anemia and Refractory Cytopenia of Childhood. <i>Blood</i> , <b>2014</b> , 124, 4388-4388	2.2	
17	Next-Generation Sequencing Reveal Proviral Genome and Transcriptome in Adult T-Cell Leukemia/Lymphoma. <i>Blood</i> , <b>2015</b> , 126, 3882-3882	2.2	



- 16 Landscape of DNA Methylation and Genetic Profiles in 291 Patients with Myelodysplastic Syndromes. *Blood*, **2015**, 126, 5205-5205 2.2
- 15 Clonal Evolution in Relapsed Pediatric Acute Lymphoblastic Leukemia. *Blood*, **2015**, 126, 1425-1425 2.2
- 14 TAL1 and MYB Abnormalities in Childhood T-Cell Acute Lymphoblastic Leukemia. *Blood*, **2015**, 126, 2628-2628 2.2
- 13 Whole-Exome Analysis of Autoimmune Lymphoproliferative Syndrome-like Diseases. *Blood*, **2015**, 126, 1022-1022 2.2
- 12 Genome-Wide Mutational Landscape of Infant Acute Lymphoblastic Leukemia. *Blood*, **2016**, 128, 4070-4070 2.2
- 11 Transcriptome Analysis Revealed the Entire Genetic Understanding of Pediatric Acute Myeloid Leukemia with a Normal Karyotype. *Blood*, **2016**, 128, 2850-2850 2.2
- 10 Identifications of Highly Aggressive Phenotype with SPI1 Overexpression in Pediatric T Cell Acute Lymphoblastic Leukemia/Lymphoma. *Blood*, **2016**, 128, 909-909 2.2
- 9 Comprehensive Genetic Analysis in Cases of Juvenile Myelomonocytic Leukemia for Prognostic Estimation. *Blood*, **2016**, 128, 3159-3159 2.2
- 8 Gene Expression Profiles and Methylation Analysis in Down Syndrome Related Acute Lymphoblastic Leukemia. *Blood*, **2016**, 128, 4084-4084 2.2
- 7 TAL1 Super Enhancer Aberration and Stil-TAL1 Fusion in Pediatric T Cell Acute Lymphoblastic Leukemia. *Blood*, **2016**, 128, 1734-1734 2.2
- 6 Structural Variations Involving Programmed Death Ligands in B-Cell and T-Cell Lymphomas. *Blood*, **2016**, 128, 4105-4105 2.2
- 5 Distinctive Genetic Features of Plasma Cells in POEMS Syndrome. *Blood*, **2016**, 128, 4404-4404 2.2
- 4 Genetic Profile of Acute Erythroid Leukemia. *Blood*, **2016**, 128, 40-40 2.2
- 3 Landscape of Driver Mutations and Their Clinical Impacts in Pediatric Acute Lymphoblastic Leukemia. *Blood*, **2016**, 128, 912-912 2.2
- 2 Whole Exome Analysis Reveals Spectrum of Gene Mutations in Juvenile Myelomonocytic Leukemia. *Blood*, **2012**, 120, 170-170 2.2
- 1 Genetic Landscapes Of Childhood T-Cell Acute Lymphoblastic Leukemia. *Blood*, **2013**, 122, 3786-3786 2.2