

David Wu

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

52
papers

1,914
citations

18
h-index

43
g-index

55
ext. papers

2,441
ext. citations

4.4
avg, IF

4.41
L-index

| # | Paper | IF | Citations |
|----|---|------|-----------|
| 52 | Acquisition of a CD19-negative myeloid phenotype allows immune escape of MLL-rearranged B-ALL from CD19 CAR-T-cell therapy. <i>Blood</i> , 2016 , 127, 2406-10 | 2.2 | 436 |
| 51 | International, evidence-based consensus diagnostic criteria for HHV-8-negative/idiopathic multicentric Castleman disease. <i>Blood</i> , 2017 , 129, 1646-1657 | 2.2 | 234 |
| 50 | Validation and implementation of targeted capture and sequencing for the detection of actionable mutation, copy number variation, and gene rearrangement in clinical cancer specimens. <i>Journal of Molecular Diagnostics</i> , 2014 , 16, 56-67 | 5.1 | 203 |
| 49 | High-throughput sequencing detects minimal residual disease in acute T lymphoblastic leukemia. <i>Science Translational Medicine</i> , 2012 , 4, 134ra63 | 17.5 | 175 |
| 48 | Detection of minimal residual disease in B lymphoblastic leukemia by high-throughput sequencing of IGH. <i>Clinical Cancer Research</i> , 2014 , 20, 4540-8 | 12.9 | 110 |
| 47 | Measurable residual disease detection by high-throughput sequencing improves risk stratification for pediatric B-ALL. <i>Blood</i> , 2018 , 131, 1350-1359 | 2.2 | 108 |
| 46 | Mass cytometry of Hodgkin lymphoma reveals a CD4 regulatory T-cell-rich and exhausted T-effector microenvironment. <i>Blood</i> , 2018 , 132, 825-836 | 2.2 | 85 |
| 45 | ClinGen Myeloid Malignancy Variant Curation Expert Panel recommendations for germline RUNX1 variants. <i>Blood Advances</i> , 2019 , 3, 2962-2979 | 7.8 | 55 |
| 44 | Clinicopathologic Features and Prognostic Impact of Lymph Node Involvement in Patients With Breast Implant-associated Anaplastic Large Cell Lymphoma. <i>American Journal of Surgical Pathology</i> , 2018 , 42, 293-305 | 6.7 | 52 |
| 43 | Detection of minimal residual disease in NPM1-mutated acute myeloid leukemia by next-generation sequencing. <i>Modern Pathology</i> , 2014 , 27, 1438-46 | 9.8 | 45 |
| 42 | Pathology of Castleman Disease. <i>Hematology/Oncology Clinics of North America</i> , 2018 , 32, 37-52 | 3.1 | 45 |
| 41 | Genomic analyses of flow-sorted Hodgkin Reed-Sternberg cells reveal complementary mechanisms of immune evasion. <i>Blood Advances</i> , 2019 , 3, 4065-4080 | 7.8 | 38 |
| 40 | CADD score has limited clinical validity for the identification of pathogenic variants in noncoding regions in a hereditary cancer panel. <i>Genetics in Medicine</i> , 2016 , 18, 1269-1275 | 8.1 | 30 |
| 39 | Pretransplantation Minimal Residual Disease Predicts Survival in Patients with Mantle Cell Lymphoma Undergoing Autologous Stem Cell Transplantation in Complete Remission. <i>Biology of Blood and Marrow Transplantation</i> , 2016 , 22, 380-385 | 4.7 | 28 |
| 38 | International evidence-based consensus diagnostic and treatment guidelines for unicentric Castleman disease. <i>Blood Advances</i> , 2020 , 4, 6039-6050 | 7.8 | 24 |
| 37 | Recurrent somatic loss of TNFRSF14 in classical Hodgkin lymphoma. <i>Genes Chromosomes and Cancer</i> , 2016 , 55, 278-87 | 5 | 22 |
| 36 | Deep NPM1 Sequencing Following Allogeneic Hematopoietic Cell Transplantation Improves Risk Assessment in Adults with NPM1-Mutated AML. <i>Biology of Blood and Marrow Transplantation</i> , 2018 , 24, 1615-1620 | 4.7 | 20 |

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| 35 | Ultrasensitive detection of acute myeloid leukemia minimal residual disease using single molecule molecular inversion probes. <i>Haematologica</i> , 2017 , 102, 1549-1557 | 6.6 | 19 |
| 34 | A novel disease-causing synonymous exonic mutation in affecting RNA splicing. <i>Blood</i> , 2018 , 132, 1211-1215 | | 17 |
| 33 | Flow cytometry for non-Hodgkin and classical Hodgkin lymphoma. <i>Methods in Molecular Biology</i> , 2013 , 971, 27-47 | 1.4 | 17 |
| 32 | How I curate: applying American Society of Hematology-Clinical Genome Resource Myeloid Malignancy Variant Curation Expert Panel rules for variant curation for germline predisposition to myeloid malignancies. <i>Haematologica</i> , 2020 , 105, 870-887 | 6.6 | 14 |
| 31 | Flow cytometry of ALK-negative anaplastic large cell lymphoma of breast implant-associated effusion and capsular tissue. <i>Cytometry Part B - Clinical Cytometry</i> , 2014 , | 3.4 | 14 |
| 30 | Pattern associated leukemia immunophenotypes and measurable disease detection in acute myeloid leukemia or myelodysplastic syndrome with mutated NPM1. <i>Cytometry Part B - Clinical Cytometry</i> , 2019 , 96, 67-72 | 3.4 | 14 |
| 29 | Flow cytometry of ALK-negative anaplastic large cell lymphoma of breast implant-associated effusion and capsular tissue. <i>Cytometry Part B - Clinical Cytometry</i> , 2015 , 88, 58-63 | 3.4 | 12 |
| 28 | Reactive T cells by flow cytometry distinguish Hodgkin lymphomas from T cell/histiocyte-rich large B cell lymphoma. <i>Cytometry Part B - Clinical Cytometry</i> , 2016 , 90, 424-32 | 3.4 | 12 |
| 27 | Insufficient evidence exists to use histopathologic subtype to guide treatment of idiopathic multicentric Castleman disease. <i>American Journal of Hematology</i> , 2020 , 95, 1553-1561 | 7.1 | 9 |
| 26 | Clinical Experience With Modified, Single-Tube T-Cell Receptor V β Flow Cytometry Analysis for T-Cell Clonality. <i>American Journal of Clinical Pathology</i> , 2016 , 145, 467-85 | 1.9 | 9 |
| 25 | Cross-Platform DNA Encoding for Single-Cell Imaging of Gene Expression. <i>Angewandte Chemie - International Edition</i> , 2016 , 55, 8975-8 | 16.4 | 8 |
| 24 | Full-Length Isoforms of Kaposi's Sarcoma-Associated Herpesvirus Latency-Associated Nuclear Antigen Accumulate in the Cytoplasm of Cells Undergoing the Lytic Cycle of Replication. <i>Journal of Virology</i> , 2017 , 91, | 6.6 | 7 |
| 23 | Jumping translocations in myelodysplastic syndromes. <i>Cancer Genetics</i> , 2016 , 209, 395-402 | 2.3 | 7 |
| 22 | Cutaneous T-cell lymphoma in sub-Saharan Africa. <i>Journal of the National Comprehensive Cancer Network: JNCCN</i> , 2013 , 11, 275-280 | 7.3 | 6 |
| 21 | Bendamustine with rituximab, etoposide and carboplatin (T(R)EC) in relapsed or refractory aggressive lymphoma: a prospective multicentre phase 1/2 clinical trial. <i>British Journal of Haematology</i> , 2018 , 183, 601-607 | 4.5 | 6 |
| 20 | Flow cytometric features of incidental indolent T lymphoblastic proliferations. <i>Cytometry Part B - Clinical Cytometry</i> , 2020 , 98, 282-287 | 3.4 | 5 |
| 19 | Myeloid/lymphoid neoplasms with FLT3 rearrangement. <i>Modern Pathology</i> , 2021 , 34, 1673-1685 | 9.8 | 5 |
| 18 | Early T-Cell Precursor Acute Lymphoblastic Leukemia in an Infant With an NRAS Q61R Mutation and Clinical Features of Juvenile Myelomonocytic Leukemia. <i>Pediatric Blood and Cancer</i> , 2016 , 63, 1667-70 | 3 | 5 |

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| 17 | Computer-aided detection of rare tumor populations in flow cytometry: an example with classic Hodgkin lymphoma. <i>American Journal of Clinical Pathology</i> , 2015 , 144, 517-24 | 1.9 | 3 |
| 16 | Targeted Next-Generation Sequencing of Acute Leukemia. <i>Methods in Molecular Biology</i> , 2017 , 1633, 163-184 | 1.4 | 3 |
| 15 | Flow Cytometry for Non-Hodgkin and Hodgkin Lymphomas. <i>Methods in Molecular Biology</i> , 2019 , 1956, 35-60 | 1.4 | 3 |
| 14 | Ultrasensitive Detection of Chimerism by Single-Molecule Molecular Inversion Probe Capture and High-Throughput Sequencing of Copy Number Deletion Polymorphisms. <i>Clinical Chemistry</i> , 2018 , 64, 938-949 | 5.5 | 2 |
| 13 | Residual Disease Monitoring By High Throughput Sequencing Provides Risk Stratification in Childhood B-ALL and Identifies a Novel Subset of Patients Having Poor Outcome. <i>Blood</i> , 2016 , 128, 1086-1086 ^{2,2} | 2.2 | 2 |
| 12 | Robust Detection Of Minimal Residual Disease In Unselected Patients With B-Cell Precursor Acute Lymphoblastic Leukemia By High-Throughput Sequencing Of IGH. <i>Blood</i> , 2013 , 122, 2550-2550 | 2.2 | 1 |
| 11 | Comprehensive Evaluation and Validation of a Next-Generation Sequencing Assay for Minimal Residual Disease Detection in T-Lymphoblastic Leukemia/Lymphoma. <i>Blood</i> , 2019 , 134, 1475-1475 | 2.2 | 1 |
| 10 | De Novo Identification and Visualization of Important Cell Populations for Classic Hodgkin Lymphoma Using Flow Cytometry and Machine Learning. <i>American Journal of Clinical Pathology</i> , 2021 , 156, 1092-1102 | 1.9 | 1 |
| 9 | A Curriculum for Genomic Education of Molecular Genetic Pathology Fellows: A Report of the Association for Molecular Pathology Training and Education Committee. <i>Journal of Molecular Diagnostics</i> , 2021 , 23, 1218-1240 | 5.1 | 1 |
| 8 | Commentary on A Case of Rapid Deterioration with Marked Hypergammaglobulinemia. <i>Clinical Chemistry</i> , 2020 , 66, 1379-1380 | 5.5 | |
| 7 | Expert Curation of Somatic Variants in Hematological Malignancies By the ClinGen Somatic Hematological Cancer Taskforce (ClinGen HCT). <i>Blood</i> , 2020 , 136, 23-23 | 2.2 | |
| 6 | Comparative Genomic Analyses Defines Shared and Unique Features of cHL and PMBL and New Mechanisms of Sensitivity to PD-1 Blockade. <i>Blood</i> , 2019 , 134, 1493-1493 | 2.2 | |
| 5 | Bone Marrow Involvement Detected By Multi-Parameter Flow Cytometry Predicts Poor Outcome after Autologous Stem Cell Transplantation for Peripheral T-Cell Lymphoma. <i>Blood</i> , 2015 , 126, 1972-1972 ^{2,2} | 2.2 | |
| 4 | Detection of Mutations in Inherited Bone Marrow Failure and Myelodysplastic Syndrome Genes Using Genomic Capture and Massively Parallel Sequencing in Clinical Diagnostics. <i>Blood</i> , 2016 , 128, 1507-1507 ^{2,2} | 2.2 | |
| 3 | On-Going Evolution Of IGH In B-Cell Precursor Acute Lymphoblastic Leukemia Does Not Substantially Affect Day 29, Post-Treatment MRD Quantification By High-Throughput Sequencing. <i>Blood</i> , 2013 , 122, 1341-1341 | 2.2 | |
| 2 | Detection Of Recurrent/Persistent Disease By T-Cell Receptor Repertoire Profiling In Patients With Mature T-Cell Neoplasm. <i>Blood</i> , 2013 , 122, 2614-2614 | 2.2 | |
| 1 | Cross-Platform DNA Encoding for Single-Cell Imaging of Gene Expression. <i>Angewandte Chemie</i> , 2016 , 128, 9121-9124 | 3.6 | |