Rex K H Au-Yeung

List of Publications by Year
in descending order

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47 papers	3,088 citations	394421 19 h-index	276875 41 g-index
49 all docs	49 docs citations	49 times ranked	5971 citing authors

#	Article	IF	Citations
1	Genetic and Functional Drivers of Diffuse Large BÂCell Lymphoma. Cell, 2017, 171, 481-494.e15.	28.9	804
2	PD1 blockade with pembrolizumab is highly effective in relapsed or refractory NK/T-cell lymphoma failing l-asparaginase. Blood, 2017, 129, 2437-2442.	1.4	399
3	Human intestinal tract serves as an alternative infection route for Middle East respiratory syndrome coronavirus. Science Advances, 2017, 3, eaao4966.	10.3	317
4	Clinical and Molecular Epidemiology of Human Bocavirus in Respiratory and Fecal Samples from Children in Hong Kong. Journal of Infectious Diseases, 2007, 196, 986-993.	4.0	172
5	Rat Hepatitis E Virus as Cause of Persistent Hepatitis after Liver Transplant. Emerging Infectious Diseases, 2018, 24, 2241-2250.	4.3	167
6	The Genetic Basis of Hepatosplenic T-cell Lymphoma. Cancer Discovery, 2017, 7, 369-379.	9.4	163
7	Middle East respiratory syndrome coronavirus and bat coronavirus HKU9 both can utilize GRP78 for attachment onto host cells. Journal of Biological Chemistry, 2018, 293, 11709-11726.	3.4	153
8	Enteropathy-associated T cell lymphoma subtypes are characterized by loss of function of SETD2. Journal of Experimental Medicine, 2017, 214, 1371-1386.	8.5	144
9	The whole-genome landscape of Burkitt lymphoma subtypes. Blood, 2019, 134, 1598-1607.	1.4	113
10	Receptor-type tyrosine-protein phosphatase Îº directly targets STAT3 activation for tumor suppression in nasal NK/T-cell lymphoma. Blood, 2015, 125, 1589-1600.	1.4	108
11	Zika Virus Infection in Dexamethasone-immunosuppressed Mice Demonstrating Disseminated Infection with Multi-organ Involvement Including Orchitis Effectively Treated by Recombinant Type I Interferons. EBioMedicine, 2016, 14, 112-122.	6.1	77
12	The mutational landscape of Burkitt-like lymphoma with 11q aberration is distinct from that of Burkitt lymphoma. Blood, 2019, 133, 962-966.	1.4	69
13	Discovery and Sequence Analysis of Four Deltacoronaviruses from Birds in the Middle East Reveal Interspecies Jumping with Recombination as a Potential Mechanism for Avian-to-Avian and Avian-to-Mammalian Transmission. Journal of Virology, 2018, 92, .	3.4	62
14	Targeting highly pathogenic coronavirus-induced apoptosis reduces viral pathogenesis and disease severity. Science Advances, 2021, 7, .	10.3	48
15	Experience with provisional WHOâ€entities large Bâ€cell lymphoma with <i>IRF4</i>â€rearrangement and Burkittâ€like lymphoma with 11q aberration in paediatric patients of the NHLâ€BFM group. British Journal of Haematology, 2020, 190, 753-763.	2.5	46
16	Whole-genome sequencing identifies responders to Pembrolizumab in relapse/refractory natural-killer/T cell lymphoma. Leukemia, 2020, 34, 3413-3419.	7.2	42
17	Disseminated fusarium infection after ibrutinib therapy in chronic lymphocytic leukaemia. Annals of Hematology, 2017, 96, 871-872.	1.8	29
18	Mp1p Is a Virulence Factor in Talaromyces (Penicillium) marneffei. PLoS Neglected Tropical Diseases, 2016, 10, e0004907.	3.0	29

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19	Recent advances in the diagnosis and treatment of natural killer/T-cell lymphomas. Expert Review of Hematology, 2019, 12, 927-935.	2.2	21
20	Isolation of MERS-related coronavirus from lesser bamboo bats that uses DPP4 and infects human-DPP4-transgenic mice. Nature Communications, 2021, 12, 216.	12.8	20
21	SPARC-positive macrophages are the superior prognostic factor in the microenvironment of diffuse large B-cell lymphoma and independent of MYC rearrangement and double-/triple-hit status. Annals of Oncology, 2021, 32, 1400-1409.	1.2	18
22	In silico analysis of 16S rRNA gene sequencing based methods for identification of medically important aerobic Gram-negative bacteria. Journal of Medical Microbiology, 2011, 60, 1281-1286.	1.8	11
23	Low-dose nivolumab induced durable complete response in relapsed primary central nervous system diffuse large B cell lymphoma. Annals of Hematology, 2019, 98, 2227-2230.	1.8	8
24	Hepatic phaeohyphomycosis due to a novel dematiaceous fungus, <i>Pleurostoma hongkongense</i> sp. nov., and importance of antifungal susceptibility testing. Emerging Microbes and Infections, 2021, 10, 81-96.	6.5	7
25	T-Cell Clustering in Neoplastic Follicles of Follicular Lymphoma. Cancer Microenvironment, 2018, 11, 135-140.	3.1	6
26	Molecular features of nonâ€anaplastic peripheral Tâ€cell lymphoma in children and adolescents. Pediatric Blood and Cancer, 2021, 68, e29285.	1.5	6
27	Mp1p homologues as virulence factors in Aspergillus fumigatus. Medical Mycology, 2018, 56, 350-360.	0.7	5
28	Low-dose pembrolizumabÂinduced complete radiologic and molecular response of posttransplant lymphoproliferative disorder presenting as classical Hodgkin lymphoma. Annals of Hematology, 2020, 99, 385-388.	1.8	5
29	Addictive response of primary cutaneous diffuse large B cell lymphoma leg type to low-dose ibrutinib. Annals of Hematology, 2019, 98, 2433-2436.	1.8	4
30	Integrative Genetic and Clinical Analysis through Whole Exome Sequencing in 1001 Diffuse Large B Cell Lymphoma (DLBCL) Patients Reveals Novel Disease Drivers and Risk Groups. Blood, 2016, 128, 1087-1087.	1.4	4
31	Whole Exome and Transcriptome Sequencing in 1042 Cases Reveals Distinct Clinically Relevant Genetic Subgroups of Follicular Lymphoma. Blood, 2019, 134, 19-19.	1.4	4
32	Epigenetic Silencing of Tumor Suppressor lncRNA NKILA: Implication on NF-Î°B Signaling in Non-Hodgkinâ€™s Lymphoma. Genes, 2022, 13, 128.	2.4	4
33	A rare disease in an atypical locationâ€"Kimuraâ€™s Disease of the upper extremity. Skeletal Radiology, 2015, 44, 1833-1837.	2.0	3
34	T-Cell Lymphoma Clonality by Copy Number Variation Analysis of T-Cell Receptor Genes. Cancers, 2021, 13, 340.	3.7	3
35	Whole Exome Sequencing of Type 1 and Type 2 Enteropathy-Associated T Cell Lymphoma Reveals Genetic Basis of Eatl Oncogenesis. Blood, 2015, 126, 575-575.	1.4	3
36	A soft-computing based approach to overlapped cells analysis in histopathology images with genetic algorithm. Applied Soft Computing Journal, 2022, 126, 109279.	7.2	3

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37	Spectrum of B-cell neoplasms associated with immunoglobulin G4-related disease. Annals of Hematology, 2022, 101, 99-108.	1.8	2
38	Epstein-Barr virusâ€“positive diffuse large B-cell lymphoma after frontline brentuximab vedotin treatment of classical Hodgkin lymphoma. Annals of Hematology, 2021, , 1.	1.8	2
39	Complete response of age-related Epstein-Barr virus-associated polymorphic nodal lymphoproliferative disease of plasmacytic type to low-dose lenalidomide. Annals of Hematology, 2018, 97, 363-366.	1.8	1
40	CD38 is not expressed in pediatric ALKâ€positive anaplastic large cell lymphoma. Pediatric Blood and Cancer, 2019, 66, e27541.	1.5	1
41	CD30 expression: a diagnostic pitfall for primary cutaneous gamma delta T cell lymphoma. Annals of Hematology, 2020, 100, 2653-2657.	1.8	1
42	Clonal heterogeneity of polymorphic B-cell lymphoproliferative disease, EBV-positive, iatrogenic/immune senescence: implications on pathogenesis and treatment. Hematology, 2022, 27, 684-690.	1.5	1
43	Unique evolution of angioimmunoblastic T cell lymphoma to Epstein-Barr virusâ€“positive plasma cell myeloma. Annals of Hematology, 2020, 99, 2949-2952.	1.8	0
44	CD3+/CD56+ EBV+ neoplasms in the nose and upper aerodigestive tract: potential misdiagnosis of plasma cell malignancies as NK/T cell lymphoma. Annals of Hematology, 2021, 100, 1101-1104.	1.8	0
45	CD3+CD4+ lymphoid neoplasm: diagnostic clues for plasmablastic lymphoma instead of peripheral T-cell lymphoma. Annals of Hematology, 2021, , 1.	1.8	0
46	Whole-Genome Genomics Correlates of Response to Anti-PD1 Therapy in Relapsed/Refractory Natural Killer/T Cell Lymphoma. Blood, 2018, 132, 2915-2915.	1.4	0
47	Heterogeneous Genetic Alterations and Novel Pathogenic Pathways in Relapsed DLBCL Revealed By Whole Exome Sequencing. Blood, 2019, 134, 2770-2770.	1.4	0