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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.

28 papers	877 citations	13 h-index	29 g-index
32 ext. papers	1,248 ext. citations	7.4 avg, IF	3.68 L-index

#	Paper	IF	Citations
28	Novel molecular subgroups for clinical classification and outcome prediction in childhood medulloblastoma: a cohort study. <i>Lancet Oncology, The</i> , 2017 , 18, 958-971	21.7	248
27	DNA methylation profiling of medulloblastoma allows robust subclassification and improved outcome prediction using formalin-fixed biopsies. <i>Acta Neuropathologica</i> , 2013 , 125, 359-71	14.3	95
26	Second-generation molecular subgrouping of medulloblastoma: an international meta-analysis of Group 3 and Group 4 subtypes. <i>Acta Neuropathologica</i> , 2019 , 138, 309-326	14.3	90
25	MBCL-31. A WHOLE CHROMOSOME ABERRATION PHENOTYPE IN NON-WNT/NON-SHH TUMORS PREDICTS OUTCOME WITHIN STANDARD-RISK MEDULLOBLASTOMAS FROM THE HIT-SIOP-PNET4 CLINICAL TRIAL. <i>Neuro-Oncology</i> , 2018 , 20, i123-i123	1	78
24	ATRT-20. INTRA- AND EXTRA-CRANIAL MALIGNANT RHABDOID TUMOURS SHARE COMMON LOCATION-INDEPENDENT CLINICAL AND MOLECULAR DISEASE CHARACTERISTICS. <i>Neuro-Oncology</i> , 2018 , 20, i31-i32	1	78
23	Next-generation systematics: An innovative approach to resolve the structure of complex prokaryotic taxa. <i>Scientific Reports</i> , 2016 , 6, 38392	4.9	72
22	Prognostic effect of whole chromosomal aberration signatures in standard-risk, non-WNT/non-SHH medulloblastoma: a retrospective, molecular analysis of the HIT-SIOP PNET 4 trial. <i>Lancet Oncology, The</i> , 2018 , 19, 1602-1616	21.7	33
21	Pediatric pan-central nervous system tumor analysis of immune-cell infiltration identifies correlates of antitumor immunity. <i>Nature Communications</i> , 2020 , 11, 4324	17.4	32
20	Reducing Viability Bias in Analysis of Gut Microbiota in Preterm Infants at Risk of NEC and Sepsis. <i>Frontiers in Cellular and Infection Microbiology</i> , 2017 , 7, 237	5.9	29
19	Genomic analysis of endemic clones of toxigenic and non-toxigenic <i>Corynebacterium diphtheriae</i> in Belarus during and after the major epidemic in 1990s. <i>BMC Genomics</i> , 2017 , 18, 873	4.5	19
18	Analysis of <i>Listeria</i> using exogenous volatile organic compound metabolites and their detection by static headspace-multi-capillary column-gas chromatography-ion mobility spectrometry (SHS-MCC-GC-IMS). <i>Analytical and Bioanalytical Chemistry</i> , 2017 , 409, 4247-4256	4.4	18
17	Epigenetic landscape correlates with genetic subtype but does not predict outcome in childhood acute lymphoblastic leukemia. <i>Epigenetics</i> , 2015 , 10, 717-26	5.7	17
16	SH2B3 inactivation through CN-LOH 12q is uniquely associated with B-cell precursor ALL with iAMP21 or other chromosome 21 gain. <i>Leukemia</i> , 2019 , 33, 1881-1894	10.7	14
15	A chromatographic approach to distinguish Gram-positive from Gram-negative bacteria using exogenous volatile organic compound metabolites. <i>Journal of Chromatography A</i> , 2017 , 1501, 79-88	4.5	13
14	Time, pattern, and outcome of medulloblastoma relapse and their association with tumour biology at diagnosis and therapy: a multicentre cohort study. <i>The Lancet Child and Adolescent Health</i> , 2020 , 4, 865-874	14.5	12
13	The Treatment of Possible Severe Infection in Infants: An Open Randomized Safety Trial of Parenteral Benzylpenicillin and Gentamicin Versus Ceftriaxone in Infants . <i>Pediatric Infectious Disease Journal</i> , 2017 , 36, e328-e333	3.4	7
12	Imaging Characteristics of Wingless Pathway Subgroup Medulloblastomas: Results from the German HIT/SIOP-Trial Cohort. <i>American Journal of Neuroradiology</i> , 2019 , 40, 1811-1817	4.4	5

11	Human Bone Proteomes before and after Decomposition: Investigating the Effects of Biological Variation and Taphonomic Alteration on Bone Protein Profiles and the Implications for Forensic Proteomics. <i>Journal of Proteome Research</i> , 2021 , 20, 2533-2546	5.6	4
10	Advanced molecular pathology for rare tumours: A national feasibility study and model for centralised medulloblastoma diagnostics. <i>Neuropathology and Applied Neurobiology</i> , 2021 , 47, 736-747	5.2	3
9	Emergence and maintenance of actionable genetic drivers at medulloblastoma relapse. <i>Neuro-Oncology</i> , 2021 ,	1	3
8	The Effects of Inter-Individual Biological Differences and Taphonomic Alteration on Human Bone Protein Profiles: Implications for the Development of PMI/AAD Estimation Methods		2
7	Inter and intra-tumoral heterogeneity as a platform for personalized therapies in medulloblastoma. <i>Pharmacology & Therapeutics</i> , 2021 , 228, 107828	13.9	2
6	Challenges of starting treatment protocols for acute lymphoblastic leukaemia in a low-income setting - the Blantyre experience. <i>British Journal of Haematology</i> , 2020 , 191, e87-e90	4.5	1
5	Response: Commentary: Reducing Viability Bias in Analysis of Gut Microbiota in Preterm Infants at Risk of NEC and Sepsis. <i>Frontiers in Cellular and Infection Microbiology</i> , 2018 , 8, 374	5.9	1
4	Integration of genome-level data to allow identification of subtype-specific vulnerability genes as novel therapeutic targets. <i>Oncogene</i> , 2021 , 40, 5213-5223	9.2	1
3	MEDB-71. Molecular characterisation of group 4 medulloblastoma improves risk-stratification and its biological understanding. <i>Neuro-Oncology</i> , 2022 , 24, i123-i123	1	
2	MEDB-43. Development of a bioinformatics pipeline for identification of differential DNA methylation events associated with medulloblastoma relapse. <i>Neuro-Oncology</i> , 2022 , 24, i115-i115	1	
1	MEDB-65. Molecular subclassification of a national cohort of pediatric medulloblastoma based on methylation profile. <i>Neuro-Oncology</i> , 2022 , 24, i121-i121	1	