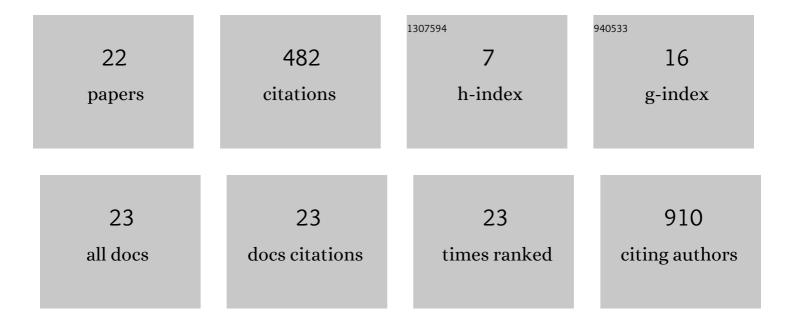
Midhat S Farooqi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4903330/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Trial of Transplantation of HCV-Infected Kidneys into Uninfected Recipients. New England Journal of Medicine, 2017, 376, 2394-2395.	27.0	315
2	Outcomes of Hematopoietic Cell Transplantation in Patients with Germline SAMD9/SAMD9L Mutations. Biology of Blood and Marrow Transplantation, 2019, 25, 2186-2196.	2.0	30
3	Decitabine and Vorinostat with Chemotherapy in Relapsed Pediatric Acute Lymphoblastic Leukemia: A TACL Pilot Study. Clinical Cancer Research, 2020, 26, 2297-2307.	7.0	28
4	Role of high-throughput sequencing in the diagnosis of cutaneous T-cell lymphoma. Journal of Clinical Pathology, 2018, 71, 814-820.	2.0	26
5	Precision Medicine in Pediatric Cancer: Current Applications and Future Prospects. High-Throughput, 2018, 7, 39.	4.4	18
6	Factors Affecting Migration to GRCh38 in Laboratories Performing Clinical Next-Generation Sequencing. Journal of Molecular Diagnostics, 2021, 23, 651-657.	2.8	13
7	Hepatitis C virus genotyping of organ donor samples to aid in transplantation of <scp>HCV</scp> â€positive organs. Clinical Transplantation, 2018, 32, e13172.	1.6	9
8	NanoString Digital Molecular Profiling of Protein and microRNA in Rhabdomyosarcoma. Cancers, 2022, 14, 522.	3.7	7
9	Validation of a next-generation sequencing oncology panel optimized for low input DNA. Cancer Genetics, 2018, 228-229, 55-63.	0.4	6
10	Cuplike nuclear morphology is highly associated with IKZF1 deletion in pediatric precursor B-cell ALL. Blood, 2019, 134, 324-329.	1.4	5
11	MYOD1 as a prognostic indicator in rhabdomyosarcoma. Pediatric Blood and Cancer, 2021, 68, e29085.	1.5	5
12	Clinical and molecular characterization of novel deletions causing epsilon gamma delta beta thalassemia: Report of two cases. Pathology Research and Practice, 2019, 215, 152578.	2.3	4
13	Next-Generation Sequencing in the Diagnosis of Rare Pediatric Sinonasal Tumors. Ear, Nose and Throat Journal, 2021, 100, NP263-NP268.	0.8	4
14	Reinterpretation of Chromosomal Microarrays with Detailed Medical History. Journal of Pediatrics, 2020, 222, 180-185.e1.	1.8	4
15	Clinical Validation of Somatic Mutation Detection by the OncoScan CNV Plus Assay. Journal of Molecular Diagnostics, 2021, 23, 29-37.	2.8	3
16	T-lymphoblastic leukemia/lymphoma with interfollicular growth pattern and Castleman-like morphologic features. Journal of Hematopathology, 2021, 14, 163-169.	0.4	3
17	Occurrence and characterization of medulloblastoma in a patient with Curryâ€Jones syndrome. Clinical Genetics, 2020, 97, 670-671.	2.0	2
18	Whole Genome Bisulfite Sequencing (WGBS) Robustly Measures the Pharmacodynamic Effect of Decitabine/Vorinostat Epigenetic Treatment in Relapsed Pediatric ALL Demonstrating Potent Hypomethylation Associated with Upregulation of PRC2 and TP53 Targets. Blood, 2018, 132, 918-918.	1.4	0

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#	Article	IF	CITATIONS
19	Single Cell Sequencing Reveals Heterogeneity of Gene Expression in KMT2A Rearranged Infant ALL at Relapse Compared to Diagnosis. Blood, 2019, 134, 2756-2756.	1.4	0
20	Germline Variants Associated with Cancer Predisposition and Bone Marrow Failure Are Common in KMT2A-r Infant Acute Lymphoblastic Leukemia Patients. Blood, 2020, 136, 41-41.	1.4	0
21	Chromosomal Microarray Reinterpretation: Applications to Pediatric Practice. Journal of Pediatrics, 2022, 243, 219-223.	1.8	0
22	Abstract 2503: Identification of clinically relevant gene fusions in archived pediatric solid and liquid tumor samples using Arima-HiC sequencing. Cancer Research, 2022, 82, 2503-2503.	0.9	0