Andrea Grioni

List of Publications by Year
in descending order

15 papers	420 citations	1163117 8 h-index	1199594 12 g-index
17 all docs	17 docs citations	17 times ranked	728 citing authors

#	Article	IF	Citations
1	Transcriptome analysis reveals rice MADS13 as an important repressor of the carpel development pathway in ovules. Journal of Experimental Botany, 2021, 72, 398-414.	4.8	7
2	Recurrent genetic fusions redefine *MLL* germ line acute lymphoblastic leukemia in infants. Blood, 2021, 137, 1980-1984.	1.4	12
3	Multi-branch Convolutional Neural Network for Identification of Small Non-coding RNA genomic loci. Scientific Reports, 2020, 10, 9486.	3.3	21
4	High *EVI1* Expression due to *NRIP1/EVI1* Fusion in Therapy–related Acute Myeloid Leukemia: Description of the First Pediatric Case. HemaSphere, 2020, 4, e471.	2.7	3
5	Standardized next-generation sequencing of immunoglobulin and T-cell receptor gene recombinations for MRD marker identification in acute lymphoblastic leukaemia; a EuroClonality-NGS validation study. Leukemia, 2019, 33, 2241-2253.	7.2	177
6	Quality control and quantification in IG/TR next-generation sequencing marker identification: protocols and bioinformatic functionalities by EuroClonality-NGS. Leukemia, 2019, 33, 2254-2265.	7.2	70
7	First evidence of a paediatric patient with Cornelia de Lange syndrome with acute lymphoblastic leukaemia. Journal of Clinical Pathology, 2019, 72, 558-561.	2.0	10
8	A Simple RNA Target Capture NGS Strategy for Fusion Genes Assessment in the Diagnostics of Pediatric B–cell Acute Lymphoblastic Leukemia. HemaSphere, 2019, 3, e250.	2.7	13
9	Pre-Clinical Efficacy of the Novel Kinase Inhibitor Nintedanib on PAX5 Fusion Genes in Pediatric Ph-like B-Cell Precursor Acute Lymphoblastic Leukemia. Blood, 2019, 134, 745-745.	1.4	0
10	A novel *EP300* mutation associated with Rubinstein–Taybi syndrome type 2 presenting as combined immunodeficiency. Pediatric Allergy and Immunology, 2018, 29, 776-781.	2.6	4
11	ARResT/Interrogate: an interactive immunoprofiler for IG/TR NGS data. Bioinformatics, 2017, 33, 435-437.	4.1	85
12	High resolution IgH repertoire analysis reveals fetal liver as the likely origin of life-long, innate B lymphopoiesis in humans. Clinical Immunology, 2017, 183, 8-16.	3.2	15
13	A Versatile DNA/RNA NGS Targeted Capture Strategy for Detection of Fusion Genes in Pediatric ALL. Blood, 2016, 128, 2913-2913.	1.4	0
14	High Resolution Igh Repertoire Analysis Reveals the Human Fetal Liver As the Origin of Life-Long, Innate B Lymphopoiesis. Blood, 2016, 128, 127-127.	1.4	0
15	Library Preparation Is the Major Factor Affecting Differences in Results of Immunoglobulin Gene Rearrangements Detection on Two Major Next-Generation Sequencing Platforms. Blood, 2015, 126, 1411-1411.	1.4	1