Hagit Baris-Feldman

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/9938391/publications.pdf

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26 papers	5,050 citations	686830 13 h-index	580395 25 g-index
27 all docs	27 docs citations	27 times ranked	8787 citing authors

#	Article	IF	Citations
1	Autoantibodies against type I IFNs in patients with life-threatening COVID-19. Science, 2020, 370, .	6.0	1,983
2	Inborn errors of type I IFN immunity in patients with life-threatening COVID-19. Science, 2020, 370, .	6.0	1,749
3	Autoantibodies neutralizing type I IFNs are present in ~4% of uninfected individuals over 70 years old and account for ~20% of COVID-19 deaths. Science Immunology, 2021, 6, .	5.6	357
4	X-linked recessive TLR7 deficiency in ~1% of men under 60 years old with life-threatening COVID-19. Science Immunology, 2021, 6, .	5.6	267
5	Human genetic and immunological determinants of critical COVID-19 pneumonia. Nature, 2022, 603, 587-598.	13.7	216
6	The risk of COVID-19 death is much greater and age dependent with type I IFN autoantibodies. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e2200413119.	3.3	110
7	Recessive inborn errors of type I IFN immunity in children with COVID-19 pneumonia. Journal of Experimental Medicine, 2022, 219, .	4.2	59
8	Diagnostic criteria for constitutional mismatch repair deficiency (CMMRD): recommendations from the international consensus working group. Journal of Medical Genetics, 2022, 59, 318-327.	1.5	57
9	A global effort to dissect the human genetic basis of resistance to SARS-CoV-2 infection. Nature Immunology, 2022, 23, 159-164.	7.0	41
10	PI(3,4)P2-mediated cytokinetic abscission prevents early senescence and cataract formation. Science, 2021, 374, eabk0410.	6.0	37
11	Vaccine breakthrough hypoxemic COVID-19 pneumonia in patients with auto-Abs neutralizing type I IFNs. Science Immunology, 2023, 8, .	5.6	35
12	The National Autism Database of Israel: a Resource for Studying Autism Risk Factors, Biomarkers, Outcome Measures, and Treatment Efficacy. Journal of Molecular Neuroscience, 2020, 70, 1303-1312.	1.1	22
13	Clinical outcomes after 4.5â€‰years of eliglustat therapy for <scp>Gaucher</scp> disease type 1: Phase 3 <scp>ENGAGE</scp> trial final results. American Journal of Hematology, 2021, 96, 1156-1165.	2.0	22
14	Pathogenic variants in SMARCA5 , a chromatin remodeler, cause a range of syndromic neurodevelopmental features. Science Advances, 2021, 7, .	4.7	17
15	Spectrum of genes for inherited hearing loss in the Israeli Jewish population, including the novel human deafness gene <scp>ATOH1</scp>. Clinical Genetics, 2020, 98, 353-364.	1.0	15
16	Heterozygous loss of WBP11 function causes multiple congenital defects in humans and mice. Human Molecular Genetics, 2021, 29, 3662-3678.	1.4	14
17	Prenatal clubfoot increases the risk for clinically significant chromosomal microarray results â€“ Analysis of 269 singleton pregnancies. Early Human Development, 2020, 145, 105047.	0.8	10
18	Bi-allelic variants in neuronal cell adhesion molecule cause a neurodevelopmental disorder characterized by developmental delay, hypotonia, neuropathy/spasticity. American Journal of Human Genetics, 2022, 109, 518-532.	2.6	8

#	Article	IF	Citations
19	Diagnostic yield of multigene panel testing in an Israeli cohort: enrichment of low-penetrance variants. Breast Cancer Research and Treatment, 2020, 181, 445-453.	1.1	7
20	Eculizumab-Responsive Adult Onset Protein Losing Enteropathy, Caused by Germline CD55-Deficiency and Complicated by Aggressive Angiosarcoma. Journal of Clinical Immunology, 2021, 41, 477-481.	2.0	6
21	Non-immune Hemolysis in Gaucher Disease and Review of the Literature. Rambam Maimonides Medical Journal, 2021, 12, e0025.	0.4	4
22	Expertsâ€™ views on COVIDâ€19 vaccination and the impact of the pandemic on patients with Gaucher disease. British Journal of Haematology, 2021, 195, e135-e137.	1.2	3
23	CD55-deficiency in Jews of Bukharan descent is caused by the Cromer blood type Dr(aâˆ’) variant. Human Genetics, 2023, 142, 683-690.	1.8	3
24	Nonâ€immune hydrops fetalis caused by PIEZO1 compound heterozygous deletions detected only by exome sequencing. Prenatal Diagnosis, 2022, , .	1.1	3
25	A recurrent pathogenic BRCA2 exon 5â€“11 duplication in the Christian Arab population in Israel. Familial Cancer, 2022, 21, 289-294.	0.9	2
26	A novel truncating variant in the FGD1 gene associated with Aarskogâ€“Scott syndrome in a family previously diagnosed with Tel Hashomer camptodactyly. American Journal of Medical Genetics, Part A, 2021, 185, 3161-3166.	0.7	0