

Qiang Pan-hammarstrom

List of Publications by Year in descending order

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132
papers

9,007
citations

36303

51
h-index

48315

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all docs

163
docs citations

163
times ranked

13257
citing authors

#	ARTICLE	IF	CITATIONS
1	Deleterious Mutations in LRBA Are Associated with a Syndrome of Immune Deficiency and Autoimmunity. <i>American Journal of Human Genetics</i> , 2012, 90, 986-1001.	6.2	452
2	Autoantibodies neutralizing type I IFNs are present in ~4% of uninfected individuals over 70 years old and account for ~20% of COVID-19 deaths. <i>Science Immunology</i> , 2021, 6, .	11.9	357
3	B-cell activating factor receptor deficiency is associated with an adult-onset antibody deficiency syndrome in humans. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 13945-13950.	7.1	332
4	Genomic basis for RNA alterations in cancer. <i>Nature</i> , 2020, 578, 129-136.	27.8	280
5	X-linked recessive TLR7 deficiency in ~1% of men under 60 years old with life-threatening COVID-19. <i>Science Immunology</i> , 2021, 6, .	11.9	267
6	Relevance of biallelic versus monoallelic TNFRSF13B mutations in distinguishing disease-causing from risk-increasing TNFRSF13B variants in antibody deficiency syndromes. <i>Blood</i> , 2009, 113, 1967-1976.	1.4	254
7	B Cell Super-Enhancers and Regulatory Clusters Recruit AID Tumorigenic Activity. <i>Cell</i> , 2014, 159, 1524-1537.	28.9	234
8	Mapping of multiple susceptibility variants within the MHC region for 7 immune-mediated diseases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 18680-18685.	7.1	231
9	Human genetic and immunological determinants of critical COVID-19 pneumonia. <i>Nature</i> , 2022, 603, 587-598.	27.8	216
10	Reexamining the role of TACI coding variants in common variable immunodeficiency and selective IgA deficiency. <i>Nature Genetics</i> , 2007, 39, 429-430.	21.4	210
11	Genetic Landscapes of Relapsed and Refractory Diffuse Large B-Cell Lymphomas. <i>Clinical Cancer Research</i> , 2016, 22, 2290-2300.	7.0	186
12	Neonatal screening for severe primary immunodeficiency diseases using high-throughput triplex real-time PCR. <i>Blood</i> , 2012, 119, 2552-2555.	1.4	183
13	Genetic basis of PD-L1 overexpression in diffuse large B-cell lymphomas. <i>Blood</i> , 2016, 127, 3026-3034.	1.4	168
14	Persistence of SARS-CoV-2-specific B and T cell responses in convalescent COVID-19 patients 6-8 months after the infection. <i>Med</i> , 2021, 2, 281-295.e4.	4.4	153
15	Selective IgA deficiency in early life: Association to infections and allergic diseases during childhood. <i>Clinical Immunology</i> , 2009, 133, 78-85.	3.2	147
16	Role for Msh5 in the regulation of Ig class switch recombination. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 7193-7198.	7.1	142
17	Risk for myasthenia gravis maps to a ¹⁵¹ Pro→Ala change in TNIP1 and to human leukocyte antigen*08. <i>Annals of Neurology</i> , 2012, 72, 927-935.	5.3	137
18	Association of IFIH1 and other autoimmunity risk alleles with selective IgA deficiency. <i>Nature Genetics</i> , 2010, 42, 777-780.	21.4	134

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19	Combined immunodeficiency and Epstein-Barr virus-induced B cell malignancy in humans with inherited CD70 deficiency. <i>Journal of Experimental Medicine</i> , 2017, 214, 91-106.	8.5	134
20	Impact of DNA ligase IV on nonhomologous end joining pathways during class switch recombination in human cells. <i>Journal of Experimental Medicine</i> , 2005, 201, 189-194.	8.5	131
21	Genetic heterogeneity in primary and relapsed mantle cell lymphomas: Impact of recurrent CARD11 mutations. <i>Oncotarget</i> , 2016, 7, 38180-38190.	1.8	130
22	The risk of COVID-19 death is much greater and age dependent with type I IFN autoantibodies. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2200413119.	7.1	110
23	Bispecific IgG neutralizes SARS-CoV-2 variants and prevents escape in mice. <i>Nature</i> , 2021, 593, 424-428.	27.8	108
24	Heterologous immunization with inactivated vaccine followed by mRNA-booster elicits strong immunity against SARS-CoV-2 Omicron variant. <i>Nature Communications</i> , 2022, 13, 2670.	12.8	108
25	Identification of IgF, a hinge-region-containing Ig class, and IgD in <i>Xenopus tropicalis</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 12087-12092.	7.1	102
26	Exome sequencing reveals novel mutation targets in diffuse large B-cell lymphomas derived from Chinese patients. <i>Blood</i> , 2014, 124, 2544-2553.	1.4	102
27	AID from bony fish catalyzes class switch recombination. <i>Journal of Experimental Medicine</i> , 2005, 202, 733-738.	8.5	100
28	Interleukin-21 restores immunoglobulin production ex vivo in patients with common variable immunodeficiency and selective IgA deficiency. <i>Blood</i> , 2009, 114, 4089-4098.	1.4	100
29	SARS-CoV-2-related MIS-C: A key to the viral and genetic causes of Kawasaki disease?. <i>Journal of Experimental Medicine</i> , 2021, 218, .	8.5	100
30	Frequent alterations in cytoskeleton remodelling genes in primary and metastatic lung adenocarcinomas. <i>Nature Communications</i> , 2015, 6, 10131.	12.8	93
31	Quantification of λ -deleting recombination excision circles in Guthrie cards for the identification of early B-cell maturation defects. <i>Journal of Allergy and Clinical Immunology</i> , 2011, 128, 223-225.e2.	2.9	91
32	A hypomorphic recombination-activating gene 1 (RAG1) mutation resulting in a phenotype resembling common variable immunodeficiency. <i>Journal of Allergy and Clinical Immunology</i> , 2014, 134, 1375-1380.	2.9	91
33	RAC2 loss-of-function mutation in 2 siblings with characteristics of common variable immunodeficiency. <i>Journal of Allergy and Clinical Immunology</i> , 2015, 135, 1380-1384.e5.	2.9	89
34	Class Switch Recombination: A Comparison Between Mouse and Human. <i>Advances in Immunology</i> , 2007, 93, 1-61.	2.2	87
35	DNA repair genes are selectively mutated in diffuse large B cell lymphomas. <i>Journal of Experimental Medicine</i> , 2013, 210, 1729-1742.	8.5	87
36	ATM Is Not Required in Somatic Hypermutation of VH, but Is Involved in the Introduction of Mutations in the Switch μ Region. <i>Journal of Immunology</i> , 2003, 170, 3707-3716.	0.8	86

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37	Clinical implications of systematic phenotyping and exome sequencing in patients with primary antibody deficiency. <i>Genetics in Medicine</i> , 2019, 21, 243-251.	2.4	86
38	Human T-bet Governs Innate and Innate-like Adaptive IFN- β Immunity against Mycobacteria. <i>Cell</i> , 2020, 183, 1826-1847.e31.	28.9	83
39	Inflammatory Gene Profiles in Gastric Mucosa during <i>Helicobacter pylori</i> Infection in Humans. <i>Journal of Immunology</i> , 2004, 172, 2595-2606.	0.8	77
40	Genetic landscape of hepatitis B virus-associated diffuse large B-cell lymphoma. <i>Blood</i> , 2018, 131, 2670-2681.	1.4	77
41	Serum Microarrays for Large Scale Screening of Protein Levels. <i>Molecular and Cellular Proteomics</i> , 2005, 4, 1942-1947.	3.8	76
42	DNA repair: the link between primary immunodeficiency and cancer. <i>Annals of the New York Academy of Sciences</i> , 2011, 1246, 50-63.	3.8	75
43	Delineation of the Role of the Mre11 Complex in Class Switch Recombination. <i>Journal of Biological Chemistry</i> , 2004, 279, 16479-16487.	3.4	73
44	<i>DCLRE1C</i> (ARTEMIS) mutations causing phenotypes ranging from atypical severe combined immunodeficiency to mere antibody deficiency. <i>Human Molecular Genetics</i> , 2015, 24, 7361-7372.	2.9	72
45	Mapping of Switch Recombination Junctions, a Tool for Studying DNA Repair Pathways during Immunoglobulin Class Switching. <i>Advances in Immunology</i> , 2010, 108, 45-109.	2.2	67
46	Common variants at PVT1, ATG13-AMBRA1, AHI1 and CLEC16A are associated with selective IgA deficiency. <i>Nature Genetics</i> , 2016, 48, 1425-1429.	21.4	67
47	Human serum from SARS-CoV-2-vaccinated and COVID-19 patients shows reduced binding to the RBD of SARS-CoV-2 Omicron variant. <i>BMC Medicine</i> , 2022, 20, 102.	5.5	67
48	Extended clinical and immunological phenotype and transplant outcome in CD27 and CD70 deficiency. <i>Blood</i> , 2020, 136, 2638-2655.	1.4	64
49	Antibody deficiency diseases. <i>European Journal of Immunology</i> , 2008, 38, 327-333.	2.9	63
50	Recessive inborn errors of type I IFN immunity in children with COVID-19 pneumonia. <i>Journal of Experimental Medicine</i> , 2022, 219, .	8.5	59
51	Impact of a 3-Months Vegetarian Diet on the Gut Microbiota and Immune Repertoire. <i>Frontiers in Immunology</i> , 2018, 9, 908.	4.8	56
52	Co-expression of PD-L1 and p-AKT is associated with poor prognosis in diffuse large B-cell lymphoma via PD-1/PD-L1 axis activating intracellular AKT/mTOR pathway in tumor cells. <i>Oncotarget</i> , 2016, 7, 33350-33362.	1.8	56
53	Immunity to SARS-CoV-2 up to 15 months after infection. <i>iScience</i> , 2022, 25, 103743.	4.1	56
54	Non-homologous end joining in class switch recombination: the beginning of the end. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2009, 364, 653-665.	4.0	55

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55	High-Density SNP Mapping of the HLA Region Identifies Multiple Independent Susceptibility Loci Associated with Selective IgA Deficiency. <i>PLoS Genetics</i> , 2012, 8, e1002476.	3.5	53
56	Common variable immunodeficiency classification by quantifying T-cell receptor and immunoglobulin \hat{P} -deleting recombination excision circles. <i>Journal of Allergy and Clinical Immunology</i> , 2013, 131, 1437-1440.e5.	2.9	52
57	CD27 \hat{a} B-Cells Produce Class Switched and Somatic Hyper-Mutated Antibodies during Chronic HIV-1 Infection. <i>PLoS ONE</i> , 2009, 4, e5427.	2.5	51
58	Novel Mutations in TAC1 (TNFRSF13B) Causing Common Variable Immunodeficiency. <i>Journal of Clinical Immunology</i> , 2009, 29, 777-785.	3.8	48
59	Current Understanding and Future Research Priorities in Malignancy Associated With Inborn Errors of Immunity and DNA Repair Disorders: The Perspective of an Interdisciplinary Working Group. <i>Frontiers in Immunology</i> , 2018, 9, 2912.	4.8	48
60	Identification of the activation-induced cytidine deaminase gene from zebrafish: an evolutionary analysis. <i>Developmental and Comparative Immunology</i> , 2005, 29, 61-71.	2.3	47
61	A primary immunodeficiency characterized by defective immunoglobulin class switch recombination and impaired DNA repair. <i>Journal of Experimental Medicine</i> , 2007, 204, 1207-1216.	8.5	47
62	The H2B deubiquitinase Usp22 promotes antibody class switch recombination by facilitating non-homologous end joining. <i>Nature Communications</i> , 2018, 9, 1006.	12.8	47
63	Unique DNA Repair Gene Variations and Potential Associations with the Primary Antibody Deficiency Syndromes IgAD and CVID. <i>PLoS ONE</i> , 2010, 5, e12260.	2.5	45
64	The Porcine Ig \hat{I} Gene: Unique Chimeric Splicing of the First Constant Region Domain in its Heavy Chain Transcripts. <i>Journal of Immunology</i> , 2003, 171, 1312-1318.	0.8	44
65	Disparate roles of ATR and ATM in immunoglobulin class switch recombination and somatic hypermutation. <i>Journal of Experimental Medicine</i> , 2006, 203, 99-110.	8.5	44
66	Cernunnos influences human immunoglobulin class switch recombination and may be associated with B cell lymphomagenesis. <i>Journal of Experimental Medicine</i> , 2012, 209, 291-305.	8.5	44
67	XRCC4 deficiency in human subjects causes a marked neurological phenotype but no overt immunodeficiency. <i>Journal of Allergy and Clinical Immunology</i> , 2015, 136, 1007-1017.	2.9	44
68	Inherited IFNAR1 Deficiency in a Child with Both Critical COVID-19 Pneumonia and Multisystem Inflammatory Syndrome. <i>Journal of Clinical Immunology</i> , 2022, 42, 471-483.	3.8	44
69	Involvement of Artemis in nonhomologous end-joining during immunoglobulin class switch recombination. <i>Journal of Experimental Medicine</i> , 2008, 205, 3031-3040.	8.5	41
70	A global effort to dissect the human genetic basis of resistance to SARS-CoV-2 infection. <i>Nature Immunology</i> , 2022, 23, 159-164.	14.5	41
71	M7 \hat{a} FLIPI is not prognostic in follicular lymphoma patients with first \hat{a} line rituximab chemo \hat{a} free therapy. <i>British Journal of Haematology</i> , 2020, 188, 259-267.	2.5	40
72	Genetic mechanisms of HLA-I loss and immune escape in diffuse large B cell lymphoma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	38

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73	Immunodeficiency in Bloom's Syndrome. <i>Journal of Clinical Immunology</i> , 2018, 38, 35-44.	3.8	36
74	Extensive diversification of IgH subclass-encoding genes and IgM subclass switching in crocodylians. <i>Nature Communications</i> , 2013, 4, 1337.	12.8	35
75	Differential role of nonhomologous end joining factors in the generation, DNA damage response, and myeloid differentiation of human induced pluripotent stem cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 8889-8894.	7.1	34
76	Gut Microbiota Perturbation in IgA Deficiency Is Influenced by IgA-Autoantibody Status. <i>Gastroenterology</i> , 2021, 160, 2423-2434.e5.	1.3	34
77	X-Linked TLR7 Deficiency Underlies Critical COVID-19 Pneumonia in a Male Patient with Ataxia-Telangiectasia. <i>Journal of Clinical Immunology</i> , 2022, 42, 1-9.	3.8	34
78	A regulatory role for the cohesin loader NIPBL in nonhomologous end joining during immunoglobulin class switch recombination. <i>Journal of Experimental Medicine</i> , 2013, 210, 2503-2513.	8.5	33
79	APRIL-dependent lifelong plasmacyte maintenance and immunoglobulin production in humans. <i>Journal of Allergy and Clinical Immunology</i> , 2020, 146, 1109-1120.e4.	2.9	33
80	A single-cell atlas of diffuse large B cell lymphoma. <i>Cell Reports</i> , 2022, 39, 110713.	6.4	33
81	DNA-PKcs Is Involved in Ig Class Switch Recombination in Human B Cells. <i>Journal of Immunology</i> , 2015, 195, 5608-5615.	0.8	30
82	Synergistic antitumor effect of histone deacetylase inhibitor and Doxorubicin in peripheral T-cell lymphoma. <i>Leukemia Research</i> , 2017, 56, 29-35.	0.8	30
83	Study of an extended family with CTLA-4 deficiency suggests a CD28/CTLA-4 independent mechanism responsible for differences in disease manifestations and severity. <i>Clinical Immunology</i> , 2018, 188, 94-102.	3.2	30
84	Genome-wide mutational signatures revealed distinct developmental paths for human B cell lymphomas. <i>Journal of Experimental Medicine</i> , 2021, 218, .	8.5	29
85	Expression of <i>Helicobacter pylori</i> Virulence Factors and Associated Expression Profiles of Inflammatory Genes in the Human Gastric Mucosa. <i>Infection and Immunity</i> , 2007, 75, 5118-5126.	2.2	27
86	Novel mutation of the activation-induced cytidine deaminase gene in a Tajik family: special review on hyper-immunoglobulin M syndrome. <i>Expert Review of Clinical Immunology</i> , 2012, 8, 539-546.	3.0	27
87	Loss of ZBTB24 impairs nonhomologous end-joining and class-switch recombination in patients with ICF syndrome. <i>Journal of Experimental Medicine</i> , 2020, 217, .	8.5	27
88	SARS-CoV-2-specific B- and T-cell immunity in a population-based study of young Swedish adults. <i>Journal of Allergy and Clinical Immunology</i> , 2022, 149, 65-75.e8.	2.9	27
89	Multiple IgH Isotypes Including IgD, Subclasses of IgM, and IgY Are Expressed in the Common Ancestors of Modern Birds. <i>Journal of Immunology</i> , 2016, 196, 5138-5147.	0.8	25
90	Development of passive immunity against SARS-CoV-2 for management of immunodeficient patients—a perspective. <i>Journal of Allergy and Clinical Immunology</i> , 2020, 146, 58-60.	2.9	24

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91	Analysis of B Cell Receptor Repertoires Reveals Key Signatures of the Systemic B Cell Response after SARS-CoV-2 Infection. <i>Journal of Virology</i> , 2022, 96, JVI0160021.	3.4	24
92	Placental Transfer of Maternally-Derived IgA Precludes the Use of Guthrie Card Eluates as a Screening Tool for Primary Immunodeficiency Diseases. <i>PLoS ONE</i> , 2012, 7, e43419.	2.5	23
93	Genome-wide CRISPR screens reveal synthetic lethal interaction between CREBBP and EP300 in diffuse large B-cell lymphoma. <i>Cell Death and Disease</i> , 2021, 12, 419.	6.3	21
94	From Your Nose to Your Toes: A Review of Severe Acute Respiratory Syndrome Coronavirus 2 Pandemic-Associated Pernio. <i>Journal of Investigative Dermatology</i> , 2021, 141, 2791-2796.	0.7	21
95	Respiratory viral infections in otherwise healthy humans with inherited IRF7 deficiency. <i>Journal of Experimental Medicine</i> , 2022, 219, .	8.5	21
96	New facets of antibody deficiencies. <i>Current Opinion in Immunology</i> , 2013, 25, 629-638.	5.5	20
97	Severe Congenital Neutropenia or Hyper-IgM Syndrome? A Novel Mutation of CD40 Ligand in a Patient with Severe Neutropenia. <i>International Archives of Allergy and Immunology</i> , 2008, 147, 255-259.	2.1	18
98	Altered spectrum of somatic hypermutation in common variable immunodeficiency disease characteristic of defective repair of mutations. <i>Immunogenetics</i> , 2011, 63, 1-11.	2.4	18
99	Defective formation of IgA memory B cells, Th1 and Th17 cells in symptomatic patients with selective IgA deficiency. <i>Clinical and Translational Immunology</i> , 2020, 9, e1130.	3.8	17
100	Antibody therapy for COVID-19. <i>Current Opinion in Allergy and Clinical Immunology</i> , 2021, 21, 553-558.	2.3	17
101	Lack of correlation between the reduction of serum immunoglobulin concentration and the CTG repeat expansion in patients with type 1 Dystrofia Myotonica. <i>Journal of Neuroimmunology</i> , 2003, 144, 100-104.	2.3	15
102	Physical mapping of the giant panda immunoglobulin heavy chain constant region genes. <i>Developmental and Comparative Immunology</i> , 2007, 31, 1034-1049.	2.3	15
103	Relation of activation-induced deaminase (AID) expression with antibody response to A(H1N1)pdm09 vaccination in HIV-1 infected patients. <i>Vaccine</i> , 2013, 31, 2231-2237.	3.8	15
104	Identification of a functional variant of estrogen receptor beta in an African population. <i>Carcinogenesis</i> , 2004, 25, 2067-2073.	2.8	14
105	Cytokine Gene Expression Profiles in Human Lymphocytes Induced by a Formula of Traditional Chinese Medicine, Vigconic VI-28. <i>Journal of Interferon and Cytokine Research</i> , 2006, 26, 628-636.	1.2	14
106	A Regulatory Role for NBS1 in Strand-Specific Mutagenesis during Somatic Hypermutation. <i>PLoS ONE</i> , 2008, 3, e2482.	2.5	14
107	Hallmarks of Cancers: Primary Antibody Deficiency Versus Other Inborn Errors of Immunity. <i>Frontiers in Immunology</i> , 2021, 12, 720025.	4.8	14
108	Aberrant recombination and repair during immunoglobulin class switching in BRCA1-deficient human B cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 2157-2162.	7.1	13

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109	Association of Short-term Air Pollution Exposure With SARS-CoV-2 Infection Among Young Adults in Sweden. <i>JAMA Network Open</i> , 2022, 5, e228109.	5.9	12
110	Genomic characterization of lymphomas in patients with inborn errors of immunity. <i>Blood Advances</i> , 2022, 6, 5403-5414.	5.2	12
111	Clinical and functional impact of recurrent S1PR1 mutations in mantle cell lymphoma. <i>Blood Advances</i> , 2018, 2, 621-625.	5.2	10
112	Distinct subtype distribution and somatic mutation spectrum of lymphomas in East Asia. <i>Current Opinion in Hematology</i> , 2017, 24, 367-376.	2.5	9
113	Compound Heterozygous Mutations of IL2-Inducible T cell Kinase in a Swedish Patient: the Importance of Early Genetic Diagnosis. <i>Journal of Clinical Immunology</i> , 2019, 39, 131-134.	3.8	9
114	Towards precision medicine in lymphoid malignancies. <i>Journal of Internal Medicine</i> , 2022, 292, 221-242.	6.0	9
115	Distinct clinical and genetic features of hepatitis B virus-associated follicular lymphoma in Chinese patients. <i>Blood Advances</i> , 2022, 6, 2731-2744.	5.2	8
116	Mutation of TNFRSF13B in a child with 22q11 deletion syndrome associated with granulomatous lymphoproliferation. <i>Journal of Allergy and Clinical Immunology</i> , 2015, 135, 559-561.	2.9	7
117	Reduced immunoglobulin gene diversity in patients with Cornelia de Lange syndrome. <i>Journal of Allergy and Clinical Immunology</i> , 2018, 141, 408-411.e8.	2.9	6
118	Lung function before and after COVID-19 in young adults: A population-based study. , 2022, 1, 37-42.		6
119	Selective IgG2 deficiency due to a point mutation causing abnormal splicing of the C λ 2 gene. <i>International Immunology</i> , 2004, 17, 95-101.	4.0	5
120	NEIL1 is a candidate gene associated with common variable immunodeficiency in a patient with a chromosome 15q24 deletion. <i>Clinical Immunology</i> , 2017, 176, 71-76.	3.2	5
121	Lack of Association between Human Switch Recombination Breakpoints and the Secondary Structure of Targeted DNA Regions. <i>Journal of Immunology</i> , 2004, 172, 2727.1-2727.	0.8	4
122	Defects in plasma cell differentiation are associated with primary immunodeficiency in human subjects. <i>Journal of Allergy and Clinical Immunology</i> , 2018, 141, 1217-1219.	2.9	4
123	Comment on "Reassessment of the Role of Mut S Homolog 5 in Ig Class Switch Recombination Shows Lack of Involvement in cis- and trans-Switching". <i>Journal of Immunology</i> , 2009, 182, 4495-4496.	0.8	3
124	SARS-CoV-2 infection in patients with inborn errors of immunity due to DNA repair defects. <i>Acta Biochimica Et Biophysica Sinica</i> , 2022, , .	2.0	3
125	F.112. Human BAFF-R Deficiency is Associated with Primary Antibody Deficiency Syndrome. <i>Clinical Immunology</i> , 2009, 131, S123.	3.2	2
126	Generation of a human induced pluripotent stem cell line (PHAi003) from a primary immunodeficient patient with CD70 mutation. <i>Stem Cell Research</i> , 2019, 41, 101612.	0.7	1

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127	A Novel SPINK5 Gene Mutation Associated with Netherton Syndrome in an Omani Patient. Sultan Qaboos University Medical Journal, 2021, 21, 652-656.	1.0	1
128	M7-FLIPI Not Valid in Follicular Lymphoma Patients with First-Line Rituximab Chemo-Free Therapy. Blood, 2018, 132, 4154-4154.	1.4	1
129	Nurture your scientific curiosity early in your research career. Nature Genetics, 2013, 45, 116-118.	21.4	0
130	Generation of human induced pluripotent stem cell lines from patients with selective IgA deficiency. Stem Cell Research, 2019, 41, 101613.	0.7	0
131	Class-Switch Recombination Defects. Rare Diseases of the Immune System, 2019, , 179-199.	0.1	0
132	Covid-19 and its Impact on Medical Research and Society. International Business and Management, 2021, , 73-84.	0.1	0