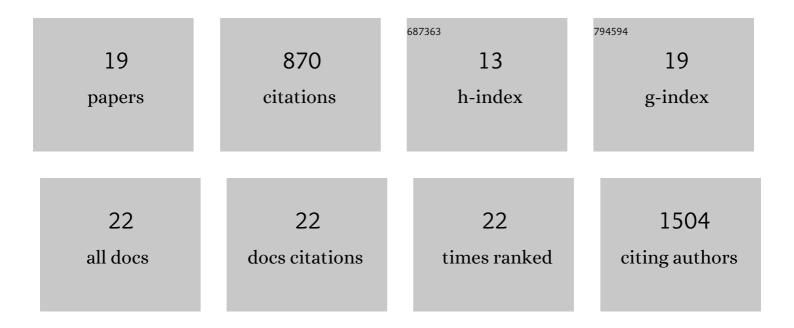
## Kenneth B Hoehn

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7377664/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Single-cell multi-omics reveals dyssynchrony of the innate and adaptive immune system in progressive COVID-19. Nature Communications, 2022, 13, 440.	12.8	100
2	Phylogenetic analysis of migration, differentiation, and class switching in B cells. PLoS Computational Biology, 2022, 18, e1009885.	3.2	40
3	Immune dysregulation and autoreactivity correlate with disease severity in SARS-CoV-2-associated multisystem inflammatory syndrome in children. Immunity, 2021, 54, 1083-1095.e7.	14.3	164
4	Cutting Edge: Distinct B Cell Repertoires Characterize Patients with Mild and Severe COVID-19. Journal of Immunology, 2021, 206, 2785-2790.	0.8	31
5	Single-cell immunophenotyping of the skin lesion erythema migrans identifies IgM memory B cells. JCI Insight, 2021, 6, .	5.0	10
6	Elevated N-Linked Glycosylation of IgG V Regions in Myasthenia Gravis Disease Subtypes. Journal of Immunology, 2021, 207, 2005-2014.	0.8	14
7	B Cell Mobilization, Dissemination, Fine Tuning of Local Antigen Specificity and Isotype Selection in Asthma. Frontiers in Immunology, 2021, 12, 702074.	4.8	4
8	Human B cell lineages associated with germinal centers following influenza vaccination are measurably evolving. ELife, 2021, 10, .	6.0	28
9	BTG1 Mutation Promotes Aggressive Lymphoma Development By Lowering the Threshold to MYC Activation and Generating "Super-Competitor" B Cells. Blood, 2021, 138, 359-359.	1.4	2
10	Thymus-derived B cell clones persist in the circulation after thymectomy in myasthenia gravis. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 30649-30660.	7.1	33
11	Mutant EZH2 Induces a Pre-malignant Lymphoma Niche by Reprogramming the Immune Response. Cancer Cell, 2020, 37, 655-673.e11.	16.8	93
12	Single-cell repertoire tracing identifies rituximab-resistant B cells during myasthenia gravis relapses. JCI Insight, 2020, 5, .	5.0	37
13	Overexpression of T-bet in HIV infection is associated with accumulation of B cells outside germinal centers and poor affinity maturation. Science Translational Medicine, 2019, 11, .	12.4	65
14	Repertoire-wide phylogenetic models of B cell molecular evolution reveal evolutionary signatures of aging and vaccination. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 22664-22672.	7.1	71
15	A Phylogenetic Codon Substitution Model for Antibody Lineages. Genetics, 2017, 206, 417-427.	2.9	56
16	A framework for detecting natural selection on traits above the species level. Methods in Ecology and Evolution, 2016, 7, 331-339.	5.2	3
17	The Diversity and Molecular Evolution of B-Cell Receptors during Infection. Molecular Biology and Evolution, 2016, 33, 1147-1157.	8.9	72
18	Dynamics of immunoglobulin sequence diversity in HIV-1 infected individuals. Philosophical Transactions of the Royal Society B: Biological Sciences, 2015, 370, 20140241.	4.0	33

#	Article	IF	CITATIONS
19	How Big Is Your Y? A Genome Sequence-Based Estimate of the Size of the Male-Specific Region in <i>Megaselia scalaris</i> . G3: Genes, Genomes, Genetics, 2015, 5, 45-48.	1.8	3