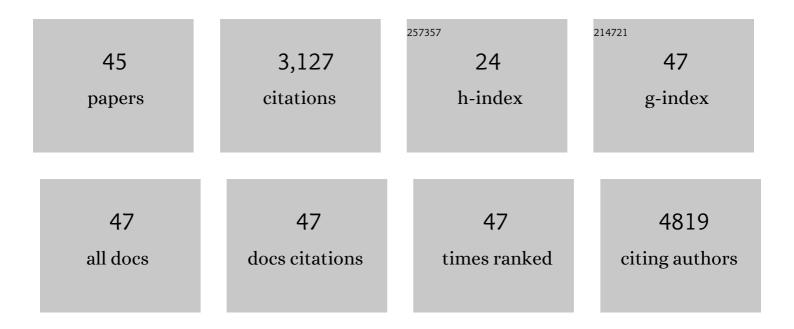
Arne Schon

List of Publications by Year in descending order

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ADNE SCHON

#	Article	IF	CITATIONS
1	The light chain of the L9 antibody is critical for binding circumsporozoite protein minor repeats and preventing malaria. Cell Reports, 2022, 38, 110367.	2.9	11
2	Development of high-affinity nanobodies specific for NaV1.4 and NaV1.5 voltage-gated sodium channel isoforms. Journal of Biological Chemistry, 2022, 298, 101763.	1.6	7
3	Highly protective antimalarial antibodies via precision library generation and yeast display screening. Journal of Experimental Medicine, 2022, 219, .	4.2	9
4	Enhancing durability of CIS43 monoclonal antibody by Fc mutation or AAV delivery for malaria prevention. JCI Insight, 2021, 6, .	2.3	25
5	Mutational fitness landscapes reveal genetic and structural improvement pathways for a vaccine-elicited HIV-1 broadly neutralizing antibody. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	21
6	Functional human IgA targets a conserved site on malaria sporozoites. Science Translational Medicine, 2021, 13, .	5.8	21
7	Reversibility and irreversibility in the temperature denaturation of monoclonal antibodies. Analytical Biochemistry, 2021, 626, 114240.	1.1	12
8	Impact of temperature on the affinity of SARS-CoV-2 Spike glycoprotein for host ACE2. Journal of Biological Chemistry, 2021, 297, 101151.	1.6	42
9	Vaccination in a humanized mouse model elicits highly protective PfCSP-targeting anti-malarial antibodies. Immunity, 2021, 54, 2859-2876.e7.	6.6	19
10	Protective effects of combining monoclonal antibodies and vaccines against the Plasmodium falciparum circumsporozoite protein. PLoS Pathogens, 2021, 17, e1010133.	2.1	20
11	Optimization of Small Molecules That Sensitize HIV-1 Infected Cells to Antibody-Dependent Cellular Cytotoxicity. ACS Medicinal Chemistry Letters, 2020, 11, 371-378.	1.3	8
12	A Potent Anti-Malarial Human Monoclonal Antibody Targets Circumsporozoite Protein Minor Repeats and Neutralizes Sporozoites in the Liver. Immunity, 2020, 53, 733-744.e8.	6.6	99
13	Cryo-EM Structures of SARS-CoV-2 Spike without and with ACE2 Reveal a pH-Dependent Switch to Mediate Endosomal Positioning of Receptor-Binding Domains. Cell Host and Microbe, 2020, 28, 867-879.e5.	5.1	316
14	Disulfide stabilization of human norovirus GI.1 virus-like particles focuses immune response toward blockade epitopes. Npj Vaccines, 2020, 5, 110.	2.9	6
15	Garcinoic Acid Is a Natural and Selective Agonist of Pregnane X Receptor. Journal of Medicinal Chemistry, 2020, 63, 3701-3712.	2.9	27
16	Preclinical Development of a Fusion Peptide Conjugate as an HIV Vaccine Immunogen. Scientific Reports, 2020, 10, 3032.	1.6	36
17	Targeting the pregnane X receptor using microbial metabolite mimicry. EMBO Molecular Medicine, 2020, 12, e11621.	3.3	53
18	Binding Thermodynamics to Intrinsically Disordered Protein Domains. Methods in Molecular Biology, 2020, 2141, 449-462.	0.4	4

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19	Antibody Lineages with Vaccine-Induced Antigen-Binding Hotspots Develop Broad HIV Neutralization. Cell, 2019, 178, 567-584.e19.	13.5	106
20	Lattice engineering enables definition of molecular features allowing for potent small-molecule inhibition of HIV-1 entry. Nature Communications, 2019, 10, 47.	5.8	50
21	Bioinspired supramolecular engineering of self-assembling immunofibers for high affinity binding of immunoglobulin G. Biomaterials, 2018, 178, 448-457.	5.7	14
22	A human monoclonal antibody prevents malaria infection by targeting a new site of vulnerability on the parasite. Nature Medicine, 2018, 24, 408-416.	15.2	235
23	SOSIP Changes Affect Human Immunodeficiency Virus Type 1 Envelope Glycoprotein Conformation and CD4 Engagement. Journal of Virology, 2018, 92, .	1.5	24
24	Long term stability of a HIV-1 neutralizing monoclonal antibody using isothermal calorimetry. Analytical Biochemistry, 2018, 554, 61-69.	1.1	8
25	The β20–β21 of gp120 is a regulatory switch for HIV-1 Env conformational transitions. Nature Communications, 2017, 8, 1049.	5.8	88
26	Crystal structures of trimeric HIV envelope with entry inhibitors BMS-378806 and BMS-626529. Nature Chemical Biology, 2017, 13, 1115-1122.	3.9	110
27	A novel lipoate attachment enzyme is shared by Plasmodium and Chlamydia species. Molecular Microbiology, 2017, 106, 439-451.	1.2	11
28	Temperature stability of proteins: Analysis of irreversible denaturation using isothermal calorimetry. Proteins: Structure, Function and Bioinformatics, 2017, 85, 2009-2016.	1.5	57
29	Enthalpy screen of drug candidates. Analytical Biochemistry, 2016, 513, 1-6.	1.1	20
30	Small-Molecule CD4-Mimics: Structure-Based Optimization of HIV-1 Entry Inhibition. ACS Medicinal Chemistry Letters, 2016, 7, 330-334.	1.3	86
31	Three easy pieces. Biochimica Et Biophysica Acta - General Subjects, 2016, 1860, 975-980.	1.1	6
32	Conformational stability and self-association equilibrium in biologics. Drug Discovery Today, 2016, 21, 342-347.	3.2	20
33	Single-Chain Soluble BG505.SOSIP gp140 Trimers as Structural and Antigenic Mimics of Mature Closed HIV-1 Env. Journal of Virology, 2015, 89, 5318-5329.	1.5	125
34	Crystal structure, conformational fixation and entry-related interactions of mature ligand-free HIV-1 Env. Nature Structural and Molecular Biology, 2015, 22, 522-531.	3.6	333
35	CD4 mimetics sensitize HIV-1-infected cells to ADCC. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, E2687-94.	3.3	118
36	Impact of Surface Polyethylene Glycol (PEG) Density on Biodegradable Nanoparticle Transport in Mucus <i>ex Vivo</i> and Distribution <i>in Vivo</i> . ACS Nano, 2015, 9, 9217-9227.	7.3	425

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37	Denatured state aggregation parameters derived from concentration dependence of protein stability. Analytical Biochemistry, 2015, 488, 45-50.	1.1	24
38	CD4-Mimetic Small Molecules Sensitize Human Immunodeficiency Virus to Vaccine-Elicited Antibodies. Journal of Virology, 2014, 88, 6542-6555.	1.5	55
39	Ligand binding analysis and screening by chemical denaturation shift. Analytical Biochemistry, 2013, 443, 52-57.	1.1	35
40	Some Binding-Related Drug Properties are Dependent on Thermodynamic Signature. Chemical Biology and Drug Design, 2011, 77, 161-165.	1.5	35
41	Thermodynamics-based drug design: strategies for inhibiting protein–protein interactions. Future Medicinal Chemistry, 2011, 3, 1129-1137.	1.1	51
42	Chapter 5 Isothermal Titration Calorimetry. Methods in Enzymology, 2009, 455, 127-155.	0.4	142
43	Strategies for targeting HIV-1 envelope glycoprotein gp120 in the development of new antivirals. Future HIV Therapy, 2007, 1, 223-229.	0.5	2
44	Thermodynamics of Binding of a Low-Molecular-Weight CD4 Mimetic to HIV-1 gp120â€. Biochemistry, 2006, 45, 10973-10980.	1.2	151
45	The binding of HIV-1 protease inhibitors to human serum proteins. Biophysical Chemistry, 2003, 105, 221-230.	1.5	55