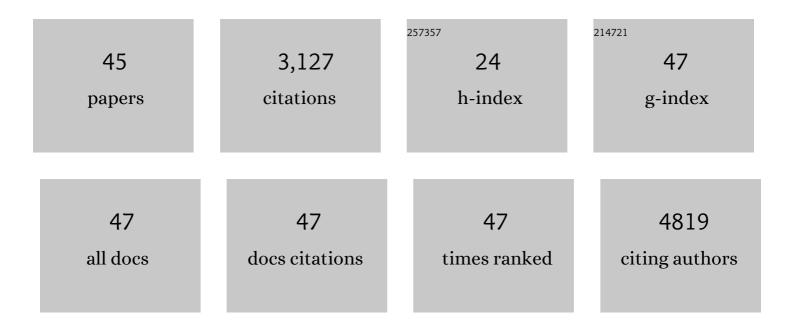
Arne Schon

List of Publications by Year in descending order

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ADNE SCHON

| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 1 | 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 |
| 2 | 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 |
| 3 | 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 |
| 4 | 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 |
| 5 | Thermodynamics of Binding of a Low-Molecular-Weight CD4 Mimetic to HIV-1 gp120â€. Biochemistry, 2006, 45, 10973-10980. | 1.2 | 151 |
| 6 | Chapter 5 Isothermal Titration Calorimetry. Methods in Enzymology, 2009, 455, 127-155. | 0.4 | 142 |
| 7 | 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 |
| 8 | 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 |
| 9 | Crystal structures of trimeric HIV envelope with entry inhibitors BMS-378806 and BMS-626529. Nature Chemical Biology, 2017, 13, 1115-1122. | 3.9 | 110 |
| 10 | Antibody Lineages with Vaccine-Induced Antigen-Binding Hotspots Develop Broad HIV Neutralization. Cell, 2019, 178, 567-584.e19. | 13.5 | 106 |
| 11 | 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 |
| 12 | The β20–β21 of gp120 is a regulatory switch for HIV-1 Env conformational transitions. Nature Communications, 2017, 8, 1049. | 5.8 | 88 |
| 13 | Small-Molecule CD4-Mimics: Structure-Based Optimization of HIV-1 Entry Inhibition. ACS Medicinal Chemistry Letters, 2016, 7, 330-334. | 1.3 | 86 |
| 14 | Temperature stability of proteins: Analysis of irreversible denaturation using isothermal calorimetry. Proteins: Structure, Function and Bioinformatics, 2017, 85, 2009-2016. | 1.5 | 57 |
| 15 | The binding of HIV-1 protease inhibitors to human serum proteins. Biophysical Chemistry, 2003, 105, 221-230. | 1.5 | 55 |
| 16 | CD4-Mimetic Small Molecules Sensitize Human Immunodeficiency Virus to Vaccine-Elicited Antibodies. Journal of Virology, 2014, 88, 6542-6555. | 1.5 | 55 |
| 17 | Targeting the pregnane X receptor using microbial metabolite mimicry. EMBO Molecular Medicine, 2020, 12, e11621. | 3.3 | 53 |
| 18 | Thermodynamics-based drug design: strategies for inhibiting protein–protein interactions. Future Medicinal Chemistry, 2011, 3, 1129-1137. | 1.1 | 51 |

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| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 19 | 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 |
| 20 | 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 |
| 21 | Preclinical Development of a Fusion Peptide Conjugate as an HIV Vaccine Immunogen. Scientific Reports, 2020, 10, 3032. | 1.6 | 36 |
| 22 | Some Binding-Related Drug Properties are Dependent on Thermodynamic Signature. Chemical Biology and Drug Design, 2011, 77, 161-165. | 1.5 | 35 |
| 23 | Ligand binding analysis and screening by chemical denaturation shift. Analytical Biochemistry, 2013, 443, 52-57. | 1.1 | 35 |
| 24 | Garcinoic Acid Is a Natural and Selective Agonist of Pregnane X Receptor. Journal of Medicinal Chemistry, 2020, 63, 3701-3712. | 2.9 | 27 |
| 25 | Enhancing durability of CIS43 monoclonal antibody by Fc mutation or AAV delivery for malaria prevention. JCI Insight, 2021, 6, . | 2.3 | 25 |
| 26 | Denatured state aggregation parameters derived from concentration dependence of protein stability. Analytical Biochemistry, 2015, 488, 45-50. | 1.1 | 24 |
| 27 | SOSIP Changes Affect Human Immunodeficiency Virus Type 1 Envelope Glycoprotein Conformation and CD4 Engagement. Journal of Virology, 2018, 92, . | 1.5 | 24 |
| 28 | 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 |
| 29 | Functional human IgA targets a conserved site on malaria sporozoites. Science Translational Medicine, 2021, 13, . | 5.8 | 21 |
| 30 | Enthalpy screen of drug candidates. Analytical Biochemistry, 2016, 513, 1-6. | 1.1 | 20 |
| 31 | Conformational stability and self-association equilibrium in biologics. Drug Discovery Today, 2016, 21, 342-347. | 3.2 | 20 |
| 32 | Protective effects of combining monoclonal antibodies and vaccines against the Plasmodium falciparum circumsporozoite protein. PLoS Pathogens, 2021, 17, e1010133. | 2.1 | 20 |
| 33 | Vaccination in a humanized mouse model elicits highly protective PfCSP-targeting anti-malarial antibodies. Immunity, 2021, 54, 2859-2876.e7. | 6.6 | 19 |
| 34 | Bioinspired supramolecular engineering of self-assembling immunofibers for high affinity binding of immunoglobulin G. Biomaterials, 2018, 178, 448-457. | 5.7 | 14 |
| 35 | Reversibility and irreversibility in the temperature denaturation of monoclonal antibodies. Analytical Biochemistry, 2021, 626, 114240. | 1.1 | 12 |
| 36 | A novel lipoate attachment enzyme is shared by Plasmodium and Chlamydia species. Molecular Microbiology, 2017, 106, 439-451. | 1.2 | 11 |

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|----|---|-----|-----------|
| 37 | 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 |
| 38 | Highly protective antimalarial antibodies via precision library generation and yeast display screening. Journal of Experimental Medicine, 2022, 219, . | 4.2 | 9 |
| 39 | Long term stability of a HIV-1 neutralizing monoclonal antibody using isothermal calorimetry. Analytical Biochemistry, 2018, 554, 61-69. | 1.1 | 8 |
| 40 | 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 |
| 41 | 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 |
| 42 | Three easy pieces. Biochimica Et Biophysica Acta - General Subjects, 2016, 1860, 975-980. | 1.1 | 6 |
| 43 | Disulfide stabilization of human norovirus Gl.1 virus-like particles focuses immune response toward blockade epitopes. Npj Vaccines, 2020, 5, 110. | 2.9 | 6 |
| 44 | Binding Thermodynamics to Intrinsically Disordered Protein Domains. Methods in Molecular Biology, 2020, 2141, 449-462. | 0.4 | 4 |
| 45 | Strategies for targeting HIV-1 envelope glycoprotein gp120 in the development of new antivirals. Future HIV Therapy, 2007, 1, 223-229. | 0.5 | 2 |