Scott T R Walsh

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Surface-fill hydrogel attenuates the oncogenic signature of complex anatomical surface cancer in a single application. Nature Nanotechnology, 2021, 16, 1251-1259.	31.5	41
2	New anti-IL-7Rα monoclonal antibodies show efficacy against T cell acute lymphoblastic leukemia in pre-clinical models. Leukemia, 2020, 34, 35-49.	7.2	26
3	The Cytokine Receptor IL-7Rα Impairs IL-2 Receptor Signaling and Constrains the InÂVitro Differentiation of Foxp3+ Treg Cells. IScience, 2020, 23, 101421.	4.1	15
4	SOCS3 is a suppressor of γc cytokine signaling and constrains generation of murine Foxp3 ⁺ regulatory T cells. European Journal of Immunology, 2020, 50, 986-999.	2.9	6
5	Design of a Multicompartment Hydrogel that Facilitates Timeâ€Resolved Delivery of Combination Therapy and Synergized Killing of Clioblastoma. Angewandte Chemie - International Edition, 2018, 57, 15040-15044.	13.8	87
6	Design of a Multicompartment Hydrogel that Facilitates Timeâ€Resolved Delivery of Combination Therapy and Synergized Killing of Glioblastoma. Angewandte Chemie, 2018, 130, 15260-15264.	2.0	17
7	Triggered Formation of Anionic Hydrogels from Self-Assembling Acidic Peptide Amphiphiles. Macromolecules, 2017, 50, 5643-5651.	4.8	17
8	Activated T Cells Secrete an Alternatively Spliced Form of Common Î ³ -Chain that Inhibits Cytokine Signaling and Exacerbates Inflammation. Immunity, 2014, 40, 910-923.	14.3	53
9	Soluble IL7Rα potentiates IL-7 bioactivity and promotes autoimmunity. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, E1761-70.	7.1	129
10	The human histone chaperone sNASP interacts with linker and core histones through distinct mechanisms. Nucleic Acids Research, 2012, 40, 660-669.	14.5	44
11	Structural reorganization of the interleukin-7 signaling complex. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 2503-2508.	7.1	54
12	Biophysical characterization of glycosaminoglycan-IL-7 interactions using SPR. Biochimie, 2012, 94, 242-249.	2.6	21
13	Structural insights into the common γâ€chain family of cytokines and receptors from the interleukinâ€7 pathway. Immunological Reviews, 2012, 250, 303-316.	6.0	50
14	A Biosensor Study Indicating That Entropy, Electrostatics, and Receptor Glycosylation Drive the Binding Interaction between Interleukin-7 and Its Receptor. Biochemistry, 2010, 49, 8766-8778.	2.5	17
15	Structural and Biophysical Studies of the Human IL-7/IL-7Rα Complex. Structure, 2009, 17, 54-65.	3.3	77
16	Crystal structure of a 3B3 variant—A broadly neutralizing HIVâ€1 scFv antibody. Protein Science, 2009, 18, 2429-2441.	7.6	16
17	Expanded binding specificity of the human histone chaperone NASP. Nucleic Acids Research, 2008, 36, 5763-5772.	14.5	56
18	Crystallization and preliminary X-ray diffraction of human interleukin-7 bound to unglycosylated and glycosylated forms of its α-receptor. Acta Crystallographica Section F: Structural Biology Communications, 2007, 63, 865-869.	0.7	12

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19	Crystal Structure and Site 1 Binding Energetics of Human Placental Lactogen. Journal of Molecular Biology, 2006, 358, 773-784.	4.2	21
20	The high- and low-affinity receptor binding sites of growth hormone are allosterically coupled. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 17078-17083.	7.1	30
21	The hydration of amides in helices; a comprehensive picture from molecular dynamics, IR, and NMR. Protein Science, 2003, 12, 520-531.	7.6	150
22	Site2 binding energetics of the regulatory step of growth hormone–induced receptor homodimerization. Protein Science, 2003, 12, 1960-1970.	7.6	39
23	Structure of a phage display-derived variant of human growth hormone complexed to two copies of the extracellular domain of its receptor: evidence for strong structural coupling between receptor binding sites. Journal of Molecular Biology, 2002, 316, 277-289.	4.2	31
24	Hydrophobic Core Malleability of a De Novo Designed Three-helix Bundle Protein. Journal of Molecular Biology, 2001, 305, 361-373.	4.2	35
25	Dynamics of a De Novo Designed Three-Helix Bundle Protein Studied by15N,13C, and2H NMR Relaxation Methodsâ€. Biochemistry, 2001, 40, 9560-9569.	2.5	44
26	Role of an α-helical bulge in the yeast heat shock transcription factor 1 1Edited by F. E. Cohen. Journal of Molecular Biology, 2000, 295, 393-409.	4.2	29