Michael Taschner

List of Publications by Year in descending order

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MICHAEL TASCHNED

#	Article	IF	CITATIONS
1	The Intraflagellar Transport Machinery. Cold Spring Harbor Perspectives in Biology, 2016, 8, a028092.	2.3	289
2	Architecture and function of IFT complex proteins in ciliogenesis. Differentiation, 2012, 83, S12-S22.	1.0	169
3	Intraflagellar transport proteins 172, 80, 57, 54, 38,Âand 20 form a stable tubulinâ€binding <scp>IFT</scp> â€82 complex. EMBO Journal, 2016, 35, 773-790.	3.5	162
4	Crystal structures of IFT70/52 and IFT52/46 provide insight into intraflagellar transport B core complex assembly. Journal of Cell Biology, 2014, 207, 269-282.	2.3	115
5	Self-organization of <i>parS</i> centromeres by the ParB CTP hydrolase. Science, 2019, 366, 1129-1133.	6.0	110
6	Biochemical Mapping of Interactions within the Intraflagellar Transport (IFT) B Core Complex. Journal of Biological Chemistry, 2011, 286, 26344-26352.	1.6	71
7	Structural Studies of Ciliary Components. Journal of Molecular Biology, 2012, 422, 163-180.	2.0	69
8	CTCF loss has limited effects on global genome architecture in Drosophila despite critical regulatory functions. Nature Communications, 2021, 12, 1011.	5.8	60
9	Structural basis of outer dynein arm intraflagellar transport by the transport adaptor protein ODA16 and the intraflagellar transport protein IFT46. Journal of Biological Chemistry, 2017, 292, 7462-7473.	1.6	48
10	Direct induction of microtubule branching by microtubule nucleation factor SSNA1. Nature Cell Biology, 2018, 20, 1172-1180.	4.6	48
11	Atomic resolution structure of human α-tubulin acetyltransferase bound to acetyl-CoA. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 19649-19654.	3.3	42
12	Crystal structure of intraflagellar transport protein 80 reveals a homo-dimer required for ciliogenesis. ELife, 2018, 7, .	2.8	36
13	Intraflagellar transport protein IFT52 recruits IFT46 to the basal body and flagella. Journal of Cell Science, 2017, 130, 1662-1674.	1.2	35
14	<i>IFT81</i> , encoding an IFT-B core protein, as a very rare cause of a ciliopathy phenotype. Journal of Medical Genetics, 2015, 52, 657-665.	1.5	32
15	Nse5/6 inhibits the Smc5/6 ATPase and modulates DNA substrate binding. EMBO Journal, 2021, 40, e107807.	3.5	30
16	Membrane association and remodeling by intraflagellar transport protein IFT172. Nature Communications, 2018, 9, 4684.	5.8	28
17	Essential role of Cp190 in physical and regulatory boundary formation. Science Advances, 2022, 8, eabl8834.	4.7	27
18	IFT proteins spatially control the geometry of cleavage furrow ingression and lumen positioning. Nature Communications, 2017, 8, 1928.	5.8	20

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19	<scp>IFT</scp> proteins interact with <scp>HSET</scp> to promote supernumerary centrosome clustering in mitosis. EMBO Reports, 2020, 21, e49234.	2.0	19
20	Purification and crystal structure of human ODA16 : Implications for ciliary import of outer dynein arms by the intraflagellar transport machinery. Protein Science, 2020, 29, 1502-1510.	3.1	12
21	Recombinant Reconstitution and Purification of the IFT-B Core Complex from Chlamydomonas reinhardtii. Methods in Molecular Biology, 2016, 1454, 69-82.	0.4	7
22	Crystal structure of the invertebrate bifunctional purine biosynthesis enzyme PAICS at 2.8 Ã resolution. Proteins: Structure, Function and Bioinformatics, 2013, 81, 1473-1478.	1.5	6
23	Complex Reconstitution from Individual Protein Modules. Advances in Experimental Medicine and Biology, 2016, 896, 305-314.	0.8	3