

Anne Houdusse-JuillÃ©©

List of Publications by Year in descending order

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92
papers

7,183
citations

50276

46
h-index

60623

81
g-index

105
all docs

105
docs citations

105
times ranked

6561
citing authors

#	ARTICLE	IF	CITATIONS
1	Myosin VIIa, harmonin and cadherin 23, three Usher I gene products that cooperate to shape the sensory hair cell bundle. <i>EMBO Journal</i> , 2002, 21, 6689-6699.	7.8	392
2	Structural and Functional Insights into the Myosin Motor Mechanism. <i>Annual Review of Biophysics</i> , 2010, 39, 539-557.	10.0	352
3	Atomic Structure of Scallop Myosin Subfragment S1 Complexed with MgADP. <i>Cell</i> , 1999, 97, 459-470.	28.9	345
4	A structural state of the myosin V motor without bound nucleotide. <i>Nature</i> , 2003, 425, 419-423.	27.8	288
5	Three myosin V structures delineate essential features of chemo-mechanical transduction. <i>EMBO Journal</i> , 2004, 23, 4527-4537.	7.8	273
6	Griscelli syndrome restricted to hypopigmentation results from a melanophilin defect (GS3) or a MYO5A F-exon deletion (GS1). <i>Journal of Clinical Investigation</i> , 2003, 112, 450-456.	8.2	251
7	Mechanism of Microtubule Stabilization by Doublecortin. <i>Molecular Cell</i> , 2004, 14, 833-839.	9.7	220
8	Structure of the regulatory domain of scallop myosin at 2 Å resolution: implications for regulation. <i>Structure</i> , 1996, 4, 21-32.	3.3	214
9	The structure of the myosin VI motor reveals the mechanism of directionality reversal. <i>Nature</i> , 2005, 435, 779-785.	27.8	206
10	Structures of four Ca ²⁺ -bound troponin C at 2.0 Å resolution: further insights into the Ca ²⁺ -switch in the calmodulin superfamily. <i>Structure</i> , 1997, 5, 1695-1711.	3.3	165
11	Mechanistic and structural basis for activation of cardiac myosin force production by omecamtiv mecarbil. <i>Nature Communications</i> , 2017, 8, 190.	12.8	153
12	Myosin motors: missing structures and hidden springs. <i>Current Opinion in Structural Biology</i> , 2001, 11, 182-194.	5.7	147
13	The structure of the rigor complex and its implications for the power stroke. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2004, 359, 1819-1828.	4.0	137
14	How Myosin Generates Force on Actin Filaments. <i>Trends in Biochemical Sciences</i> , 2016, 41, 989-997.	7.5	135
15	Oxidation of F-actin controls the terminal steps of cytokinesis. <i>Nature Communications</i> , 2017, 8, 14528.	12.8	130
16	Distinct roles of doublecortin modulating the microtubule cytoskeleton. <i>EMBO Journal</i> , 2006, 25, 4448-4457.	7.8	126
17	Crystal structure of apo-calmodulin bound to the first two IQ motifs of myosin V reveals essential recognition features. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 19326-19331.	7.1	125
18	Rab GTPases and their interacting protein partners: Structural insights into Rab functional diversity. <i>Small GTPases</i> , 2018, 9, 22-48.	1.6	122

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19	How Actin Initiates the Motor Activity of Myosin. <i>Developmental Cell</i> , 2015, 33, 401-412.	7.0	118
20	Template-free 13-protofilament microtubule-MAP assembly visualized at 8 Å... resolution. <i>Journal of Cell Biology</i> , 2010, 191, 463-470.	5.2	116
21	Molecular Basis for Specific Regulation of Neuronal Kinesin-3 Motors by Doublecortin Family Proteins. <i>Molecular Cell</i> , 2012, 47, 707-721.	9.7	116
22	Myosin VI Rewrites the Rules for Myosin Motors. <i>Cell</i> , 2010, 141, 573-582.	28.9	110
23	What can myosin VI do in cells?. <i>Current Opinion in Cell Biology</i> , 2007, 19, 57-66.	5.4	108
24	A model of Ca ²⁺ -free calmodulin binding to unconventional myosins reveals how calmodulin acts as a regulatory switch. <i>Structure</i> , 1996, 4, 1475-1490.	3.3	101
25	Conserved mechanisms of microtubule-stimulated ADP release, ATP binding, and force generation in transport kinesins. <i>ELife</i> , 2014, 3, e03680.	6.0	100
26	New insights into genotype-phenotype correlations for the doublecortin-related lissencephaly spectrum. <i>Brain</i> , 2013, 136, 223-244.	7.6	99
27	Rab35 GTPase couples cell division with initiation of epithelial apico-basal polarity and lumen opening. <i>Nature Communications</i> , 2016, 7, 11166.	12.8	97
28	Cadherin-23, myosin VIIa and harmonin, encoded by Usher syndrome type I genes, form a ternary complex and interact with membrane phospholipids. <i>Human Molecular Genetics</i> , 2010, 19, 3557-3565.	2.9	94
29	Hypertrophic cardiomyopathy disease results from disparate impairments of cardiac myosin function and auto-inhibition. <i>Nature Communications</i> , 2018, 9, 4019.	12.8	91
30	Force Generation by Myosin Motors: A Structural Perspective. <i>Chemical Reviews</i> , 2020, 120, 5-35.	47.7	91
31	Myosin VI Dimerization Triggers an Unfolding of a Three-Helix Bundle in Order to Extend Its Reach. <i>Molecular Cell</i> , 2009, 35, 305-315.	9.7	89
32	Biochemical and functional characterization of Rab27a mutations occurring in Griscelli syndrome patients. <i>Blood</i> , 2003, 101, 2736-2742.	1.4	87
33	Structural basis for ARF1-mediated recruitment of ARHGAP21 to Golgi membranes. <i>EMBO Journal</i> , 2007, 26, 1953-1962.	7.8	86
34	The unique insert at the end of the myosin VI motor is the sole determinant of directionality. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 778-783.	7.1	76
35	Force-producing ADP state of myosin bound to actin. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E1844-52.	7.1	76
36	The Structural Basis for the Large Powerstroke of Myosin VI. <i>Cell</i> , 2007, 131, 300-308.	28.9	75

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37	The myosin X motor is optimized for movement on actin bundles. <i>Nature Communications</i> , 2016, 7, 12456.	12.8	75
38	The unique insert in myosin VI is a structural calcium-calmodulin binding site. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 4787-4792.	7.1	73
39	Structural Basis for Recruitment of Rab6-Interacting Protein 1 to Golgi via a RUN Domain. <i>Structure</i> , 2009, 17, 21-30.	3.3	73
40	Magnesium Regulates ADP Dissociation from Myosin V. <i>Journal of Biological Chemistry</i> , 2005, 280, 6072-6079.	3.4	69
41	The structural basis of Arf effector specificity: the crystal structure of ARF6 in a complex with JIP4. <i>EMBO Journal</i> , 2009, 28, 2835-2845.	7.8	68
42	The motor mechanism of myosin V: insights for muscle contraction. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2004, 359, 1829-1842.	4.0	66
43	Emerging roles of MICAL family proteins “ from actin oxidation to membrane trafficking during cytokinesis. <i>Journal of Cell Science</i> , 2017, 130, 1509-1517.	2.0	63
44	Free Brick1 Is a Trimeric Precursor in the Assembly of a Functional Wave Complex. <i>PLoS ONE</i> , 2008, 3, e2462.	2.5	63
45	An Overview and Online Registry of Microvillus Inclusion Disease Patients and their MYO5B Mutations. <i>Human Mutation</i> , 2013, 34, 1597-1605.	2.5	62
46	Coupling fission and exit of RAB6 vesicles at Golgi hotspots through kinesin-myosin interactions. <i>Nature Communications</i> , 2017, 8, 1254.	12.8	55
47	Coordinated recruitment of Spir actin nucleators and myosin V motors to Rab11 vesicle membranes. <i>ELife</i> , 2016, 5, .	6.0	53
48	Myosin 7 and its adaptors link cadherins to actin. <i>Nature Communications</i> , 2017, 8, 15864.	12.8	49
49	Plasmodium myosin A drives parasite invasion by an atypical force generating mechanism. <i>Nature Communications</i> , 2019, 10, 3286.	12.8	49
50	MAPping out distribution routes for kinesin couriers. <i>Biology of the Cell</i> , 2013, 105, 465-487.	2.0	48
51	MYO5B, STX3, and STXBP2 mutations reveal a common disease mechanism that unifies a subset of congenital diarrheal disorders: A mutation update. <i>Human Mutation</i> , 2018, 39, 333-344.	2.5	48
52	Biochemical and Structural Characterization of the Gem GTPase. <i>Journal of Biological Chemistry</i> , 2007, 282, 1905-1915.	3.4	46
53	Myosin VI and branched actin filaments mediate membrane constriction and fission of melanosomal tubule carriers. <i>Journal of Cell Biology</i> , 2018, 217, 2709-2726.	5.2	46
54	Novel myosin mutations for hereditary hearing loss revealed by targeted genomic capture and massively parallel sequencing. <i>European Journal of Human Genetics</i> , 2014, 22, 768-775.	2.8	44

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55	The many roles of myosins in filopodia, microvilli and cilia. <i>Current Biology</i> , 2021, 31, R586-R602.	3.9	41
56	The divergent mitotic kinesin MKLP2 exhibits atypical structure and mechanochemistry. <i>ELife</i> , 2017, 6, .	6.0	39
57	Highly selective inhibition of myosin motors provides the basis of potential therapeutic application. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E7448-E7455.	7.1	34
58	The post-rigor structure of myosin VI and implications for the recovery stroke. <i>EMBO Journal</i> , 2008, 27, 244-252.	7.8	31
59	<i>Drosophila melanogaster</i> Myosin-18 Represents a Highly Divergent Motor with Actin Tethering Properties. <i>Journal of Biological Chemistry</i> , 2011, 286, 21755-21766.	3.4	28
60	Circularly Permuted Fluorogenic Proteins for the Design of Modular Biosensors. <i>ACS Chemical Biology</i> , 2018, 13, 2392-2397.	3.4	27
61	High-resolution structures of the actomyosin-V complex in three nucleotide states provide insights into the force generation mechanism. <i>ELife</i> , 2021, 10, .	6.0	27
62	Myosin VI Must Dimerize and Deploy Its Unusual Lever Arm in Order to Perform Its Cellular Roles. <i>Cell Reports</i> , 2014, 8, 1522-1532.	6.4	26
63	Eml1 loss impairs apical progenitor spindle length and soma shape in the developing cerebral cortex. <i>Scientific Reports</i> , 2017, 7, 17308.	3.3	26
64	Myosin Structures. <i>Advances in Experimental Medicine and Biology</i> , 2020, 1239, 7-19.	1.6	26
65	MyTH4-FERM myosins have an ancient and conserved role in filopod formation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E8059-E8068.	7.1	24
66	The actomyosin interface contains an evolutionary conserved core and an ancillary interface involved in specificity. <i>Nature Communications</i> , 2021, 12, 1892.	12.8	23
67	An intermediate along the recovery stroke of myosin VI revealed by X-ray crystallography and molecular dynamics. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 6213-6218.	7.1	22
68	Structural aspects of Rab6 effector complexes. <i>Biochemical Society Transactions</i> , 2009, 37, 1037-1041.	3.4	21
69	Single Residue Variation in Skeletal Muscle Myosin Enables Direct and Selective Drug Targeting for Spasticity and Muscle Stiffness. <i>Cell</i> , 2020, 183, 335-346.e13.	28.9	21
70	Structural Studies of the Doublecortin Family of MAPs. <i>Methods in Cell Biology</i> , 2013, 115, 27-48.	1.1	20
71	Myosin MyTH4-FERM structures highlight important principles of convergent evolution. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E2906-15.	7.1	20
72	Role of Insert-1 of Myosin VI in Modulating Nucleotide Affinity. <i>Journal of Biological Chemistry</i> , 2011, 286, 11716-11723.	3.4	19

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73	Metastasis-suppressor NME1 controls the invasive switch of breast cancer by regulating MT1-MMP surface clearance. <i>Oncogene</i> , 2021, 40, 4019-4032.	5.9	19
74	Full-length Plasmodium falciparum myosin A and essential light chain PfELC structures provide new anti-malarial targets. <i>ELife</i> , 2020, 9, .	6.0	19
75	Electrospray ionization mass spectrometry studies of noncovalent myosin VI complexes reveal a new specific calmodulin binding site. <i>Journal of the American Society for Mass Spectrometry</i> , 2005, 16, 1367-1376.	2.8	18
76	Optimized filopodia formation requires myosin tail domain cooperation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 22196-22204.	7.1	13
77	Dynein Swings into Action. <i>Cell</i> , 2009, 136, 395-396.	28.9	12
78	VASP-mediated actin dynamics activate and recruit a filopodia myosin. <i>ELife</i> , 2021, 10, .	6.0	11
79	Isolation and Characterization of an Aggresome Determinant in the NF2 Tumor Suppressor. <i>Journal of Biological Chemistry</i> , 2003, 278, 6235-6242.	3.4	10
80	Crystallization and Preliminary X-ray Diffraction Study of an Idiotope-Anti-Idiotope Fv-Fv Complex. <i>Journal of Molecular Biology</i> , 1994, 241, 739-743.	4.2	9
81	Kinesin-6 Klp9 orchestrates spindle elongation by regulating microtubule sliding and growth. <i>ELife</i> , 2021, 10, .	6.0	9
82	A double-take on MAPs. <i>Nature Structural and Molecular Biology</i> , 2003, 10, 314-316.	8.2	6
83	Improving Diffraction from 3 to 2 Å... for a Complex between a Small GTPase and Its Effector by Analysis of Crystal Contacts and Use of Reverse Screening. <i>Crystal Growth and Design</i> , 2007, 7, 2140-2146.	3.0	5
84	Griscelli syndrome restricted to hypopigmentation results from a melanophilin defect (GS3) or a MYO5A F-exon deletion (GS1). <i>Journal of Clinical Investigation</i> , 2005, 115, 1100-1100.	8.2	5
85	Cloning, bacterial expression and crystallization of Fv antibody fragments. <i>Journal of Crystal Growth</i> , 1992, 122, 337-343.	1.5	3
86	Biological nanomotors, driving forces of life. <i>Comptes Rendus - Biologies</i> , 2020, 343, 53-78.	0.2	2
87	Myosin VI Dimerizes And Walks Processively Along Actin. <i>Biophysical Journal</i> , 2009, 96, 142a.	0.5	0
88	Dimerization is Essential for the Large Step Size of Myosin VI. <i>Biophysical Journal</i> , 2010, 98, 724a.	0.5	0
89	Steered Molecular Dynamics Simulation of Unfolding of Myosin VI Proximal Tail Domain. <i>Biophysical Journal</i> , 2010, 98, 724a.	0.5	0
90	How Actin Initiates the Motor Activity of Myosin. <i>Biophysical Journal</i> , 2015, 108, 25a.	0.5	0

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91	The Force Producing ADP State of Myosin Bound to Actin. Biophysical Journal, 2016, 110, 13a.	0.5	0
92	A Structural Model of the Mitotic Kinesin-6 Mechanochemical Cycle. Biophysical Journal, 2016, 110, 192a-193a.	0.5	0