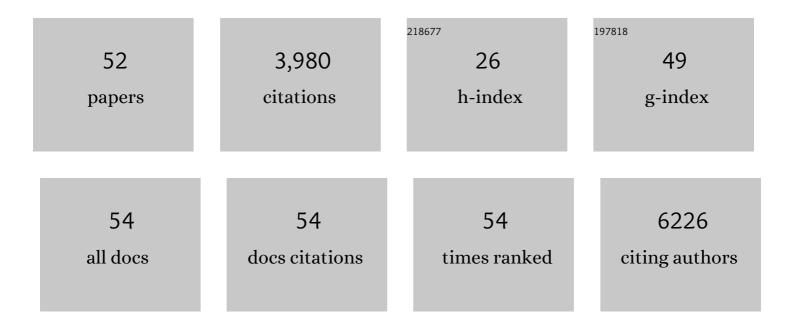
Isabelle Rouiller

List of Publications by Year in descending order

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ISARELLE ROULLER

#	Article	IF	CITATIONS
1	NMMD: Efficient Cryo-EM Flexible Fitting Based on Simultaneous Normal Mode and Molecular Dynamics atomic displacements. Journal of Molecular Biology, 2022, 434, 167483.	4.2	26
2	The AAA+ ATPase p97 as a novel parasite and tuberculosis drug target. Trends in Parasitology, 2022, 38, 572-590.	3.3	4
3	Broad and ultra-potent cross-clade neutralization of HIV-1 by a vaccine-induced CD4 binding site bovine antibody. Cell Reports Medicine, 2022, 3, 100635.	6.5	3
4	A "Drug Sweeping―State of the TriABC Triclosan Efflux Pump from Pseudomonas aeruginosa. Structure, 2021, 29, 261-274.e6.	3.3	11
5	Plant-derived virus-like particle vaccines drive cross-presentation of influenza A hemagglutinin peptides by human monocyte-derived macrophages. Npj Vaccines, 2019, 4, 17.	6.0	23
6	An Asymmetric Opening of HIV-1 Envelope Mediates Antibody-Dependent Cellular Cytotoxicity. Cell Host and Microbe, 2019, 25, 578-587.e5.	11.0	93
7	Morphological characterization of a plant-made virus-like particle vaccine bearing influenza virus hemagglutinins by electron microscopy. Vaccine, 2018, 36, 2147-2154.	3.8	37
8	Mfn2 ubiquitination by PINK1/parkin gates the p97-dependent release of ER from mitochondria to drive mitophagy. ELife, 2018, 7, .	6.0	261
9	The Peptidisc, a simple method for stabilizing membrane proteins in detergent-free solution. ELife, 2018, 7, .	6.0	119
10	Negative Stain Single-particle EM of the Maltose Transporter in Nanodiscs Reveals Asymmetric Closure of MalK2 and Catalytic Roles of ATP, MalE, and Maltose. Journal of Biological Chemistry, 2017, 292, 5457-5464.	3.4	14
11	Molecular mechanism of DRP1 assembly studied in vitro by cryo-electron microscopy. PLoS ONE, 2017, 12, e0179397.	2.5	44
12	Structure of anthrax lethal toxin prepore complex suggests a pathway for efficient cell entry. Journal of General Physiology, 2016, 148, 313-324.	1.9	16
13	Structure–Function Profile of MmpL3, the Essential Mycolic Acid Transporter from <i>Mycobacterium tuberculosis</i> . ACS Infectious Diseases, 2016, 2, 702-713.	3.8	86
14	Molecular assembly of lethal factor enzyme and pre-pore heptameric protective antigen in early stage of translocation. Journal of Molecular Modeling, 2016, 22, 7.	1.8	4
15	Cryo-EM of the pathogenic VCP variant R155P reveals long-range conformational changes in the D2 ATPase ring. Biochemical and Biophysical Research Communications, 2015, 468, 636-641.	2.1	7
16	Fibril formation pH controls intrafibrillar collagen biomineralization inÂvitro and inÂvivo. Biomaterials, 2015, 37, 252-259.	11.4	40
17	Quaternary structure of <scp>W</scp> zzB and <scp>W</scp> zzE polysaccharide copolymerases. Protein Science, 2015, 24, 58-69.	7.6	19
18	Regional differences in the expression of K+–Clâ^ 2 cotransporter in the developing rat cortex. Brain Structure and Function, 2014, 219, 527-538.	2.3	14

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19	Coordinated Rearrangements between Cytoplasmic and Periplasmic Domains of the Membrane Protein Complex ExbB-ExbD of Escherichia coli. Structure, 2014, 22, 791-797.	3.3	19
20	Study of Membrane Proteins by Single Particles Electron Microscopy Using Detergent, Liposomes and Nanodiscs. Microscopy and Microanalysis, 2014, 20, 1212-1213.	0.4	0
21	Family of phenylacetyl-CoA monooxygenases differs in subunit organization from other monooxygenases. Journal of Structural Biology, 2013, 184, 147-154.	2.8	5
22	The maltose ABC transporter: Action of membrane lipids on the transporter stability, coupling and ATPase activity. Biochimica Et Biophysica Acta - Biomembranes, 2013, 1828, 1723-1730.	2.6	26
23	Three-dimensional reconstructions of Arp2/3 complex with bound nucleation promoting factors. EMBO Journal, 2012, 31, 236-247.	7.8	67
24	A Stretch of Polybasic Residues Mediates Cdc42 GTPase-activating Protein (CdGAP) Binding to Phosphatidylinositol 3,4,5-Trisphosphate and Regulates Its GAP Activity. Journal of Biological Chemistry, 2012, 287, 19610-19621.	3.4	13
25	Protein-Protein Interactions in the β-Oxidation Part of the Phenylacetate Utilization Pathway. Journal of Biological Chemistry, 2012, 287, 37986-37996.	3.4	12
26	Full-length structural model of RET3 and SEC21 in COPI: identification of binding sites on the appendage for accessory protein recruitment motifs. Journal of Molecular Modeling, 2012, 18, 3199-3212.	1.8	5
27	Label-Free Visualization of Ultrastructural Features of Artificial Synapses via Cryo-EM. ACS Chemical Neuroscience, 2011, 2, 700-704.	3.5	5
28	Unravelling Electronic Energy Transfer in Single Conjugated Polyelectrolytes Encapsulated in Lipid Vesicles. Biophysical Journal, 2011, 100, 138a.	0.5	0
29	Growth-Arrest-Specific Protein 2 Inhibits Cell Division in Xenopus Embryos. PLoS ONE, 2011, 6, e24698.	2.5	17
30	Mutant huntingtin binds the mitochondrial fission GTPase dynamin-related protein-1 and increases its enzymatic activity. Nature Medicine, 2011, 17, 377-382.	30.7	467
31	Unraveling electronic energy transfer in single conjugated polyelectrolytes encapsulated in lipid vesicles. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 17480-17485.	7.1	38
32	<i>Desulfovibrio magneticus</i> RS-1 contains an iron- and phosphorus-rich organelle distinct from its bullet-shaped magnetosomes. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 12263-12268.	7.1	74
33	Lipid Bilayer Membrane-Triggered Presynaptic Vesicle Assembly. ACS Chemical Neuroscience, 2010, 1, 86-94.	3.5	17
34	Hereditary Inclusion Body Myopathy-Linked p97/VCP Mutations in the NH ₂ Domain and the D1 Ring Modulate p97/VCP ATPase Activity and D2 Ring Conformation. Molecular and Cellular Biology, 2009, 29, 4484-4494.	2.3	102
35	Role of capsid sequence and immature nucleocapsid proteins p9 and p15 in Human Immunodeficiency Virus type 1 genomic RNA dimerization. Virology, 2009, 385, 233-244.	2.4	16
36	Metal–nucleic acid cages. Nature Chemistry, 2009, 1, 390-396.	13.6	151

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37	"Breathing―Vesicles. Journal of the American Chemical Society, 2009, 131, 10557-10566.	13.7	217
38	Supported Bilayers Formed from Different Phospholipids on Spherical Silica Substrates. Langmuir, 2009, 25, 5455-5458.	3.5	55
39	Gigantism in unique biogenic magnetite at the Paleocene–Eocene Thermal Maximum. Proceedings of the United States of America, 2008, 105, 17648-17653.	7.1	69
40	The structural basis of actin filament branching by the Arp2/3 complex. Journal of Cell Biology, 2008, 180, 887-895.	5.2	270
41	Reconstituted NALP1 Inflammasome Reveals Two-Step Mechanism of Caspase-1 Activation. Molecular Cell, 2007, 25, 713-724.	9.7	610
42	Mechanistic Differences in Actin Bundling Activity of Two Mammalian Formins, FRL1 and mDia2. Journal of Biological Chemistry, 2006, 281, 14383-14392.	3.4	152
43	In Vitro Reconstitution of the NALP1 Inflammasome Reveals Requirements for Caspase Activation Blood, 2006, 108, 3658-3658.	1.4	0
44	Mechanism of Filament Nucleation and Branch Stability Revealed by the Structure of the Arp2/3 Complex at Actin Branch Junctions. PLoS Biology, 2005, 3, e383.	5.6	84
45	Effects of Arp2 and Arp3 nucleotide-binding pocket mutations on Arp2/3 complex function. Journal of Cell Biology, 2005, 168, 315-328.	5.2	73
46	Prepore to pore transition of a cholesterol-dependent cytolysin visualized by electron microscopy. Journal of Structural Biology, 2005, 150, 100-108.	2.8	72
47	The Subcellular Distribution of Multigene Family 110 Proteins of African Swine Fever Virus Is Determined by Differences in C-Terminal KDEL Endoplasmic Reticulum Retention Motifs. Journal of Virology, 2004, 78, 3710-3721.	3.4	41
48	Model-based particle picking for cryo-electron microscopy. Journal of Structural Biology, 2004, 145, 157-167.	2.8	19
49	Conformational changes of the multifunction p97 AAA ATPase during its ATPase cycle. Nature Structural Biology, 2002, 9, 950-957.	9.7	189
50	Automated Image Acquisition for Single-Particle Reconstruction Using p97 as the Biological Sample. Journal of Structural Biology, 2001, 133, 102-107.	2.8	12
51	A Major Conformational Change in p97 AAA ATPase upon ATP Binding. Molecular Cell, 2000, 6, 1485-1490.	9.7	146
52	African Swine Fever Virus Is Wrapped by the Endoplasmic Reticulum. Journal of Virology, 1998, 72, 2373-2387.	3.4	111