## Itay Onn

## List of Publications by Year in descending order

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567281 580821 1,212 25 26 15 citations h-index g-index papers 26 26 26 1372 citing authors docs citations times ranked all docs

#	Article	IF	CITATIONS
1	A Molecular Determinant for the Establishment of Sister Chromatid Cohesion. Science, 2008, 321, 566-569.	12.6	414
2	Sister Chromatid Cohesion: A Simple Concept with a Complex Reality. Annual Review of Cell and Developmental Biology, 2008, 24, 105-129.	9.4	295
3	Reconstitution and subunit geometry of human condensin complexes. EMBO Journal, 2007, 26, 1024-1034.	7.8	85
4	Redox Potential Regulates Binding of Universal Minicircle Sequence Binding Protein at the Kinetoplast DNA Replication Origin. Eukaryotic Cell, 2004, 3, 277-287.	3.4	57
5	A Conserved Domain in the Scc3 Subunit of Cohesin Mediates the Interaction with Both Mcd1 and the Cohesin Loader Complex. PLoS Genetics, $2015$ , $11$ , $e1005036$ .	3.5	49
6	Genetic Evidence that the Acetylation of the Smc3p Subunit of Cohesin Modulates Its ATP-Bound State to Promote Cohesion Establishment in <i>Saccharomyces cerevisiae</i> . Genetics, 2010, 185, 1249-1256.	2.9	32
7	Assigning functions to genes: identification of S-phase expressed genes in Leishmania major based on post-transcriptional control elements. Nucleic Acids Research, 2005, 33, 4235-4242.	14.5	29
8	Binding of the Universal Minicircle Sequence Binding Protein at the Kinetoplast DNA Replication Origin*. Journal of Biological Chemistry, 2006, 281, 37468-37476.	3.4	25
9	A new twist in the coil: functions of the coiled-coil domain of structural maintenance of chromosome (SMC) proteins. Current Genetics, 2018, 64, 109-116.	1.7	24
10	In vitro assembly of physiological cohesin/DNA complexes. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 12198-12205.	7.1	23
11	The emerging roles for the chromatin structure regulators CTCF and cohesin in neurodevelopment and behavior. Cellular and Molecular Life Sciences, 2018, 75, 1205-1214.	5.4	23
12	DNA motif associated with meiotic doubleâ€strand break regions in Saccharomyces cerevisiae. EMBO Reports, 2000, 1, 232-238.	4.5	21
13	The zinc finger of Eco1 enhances its acetyltransferase activity during sister chromatid cohesion. Nucleic Acids Research, 2009, 37, 6126-6134.	14.5	20
14	Monomeric cohesin state revealed by liveâ€cell singleâ€molecule spectroscopy. EMBO Reports, 2020, 21, e48211.	4.5	20
15	The chromatin remodeler Chd1 regulates cohesin in budding yeast and humans. Scientific Reports, 2019, 9, 8929.	3.3	18
16	Hit the brakes $\hat{a}\in$ a new perspective on the loop extrusion mechanism of cohesin and other SMC complexes. Journal of Cell Science, 2021, 134, .	2.0	17
17	Identification of a region in the coiled-coil domain of Smc3 that is essential for cohesin activity. Nucleic Acids Research, 2016, 44, 6309-6317.	14.5	15
18	Structural maintenance of chromosome complexes and bone development: the beginning of a wonderful relationship?. BoneKEy Reports, 2013, 2, 388.	2.7	9

#	Article	IF	CITATIONS
19	Chromosome loading of cohesin depends on conserved residues in Scc3. Current Genetics, 2021, 67, 447-459.	1.7	9
20	Identification of Functional Domains in the Cohesin Loader Subunit Scc4 by a Random Insertion/Dominant Negative Screen. G3: Genes, Genomes, Genetics, 2016, 6, 2655-2663.	1.8	8
21	Dysregulation of the cohesin subunit RAD21 by Hepatitis C virus mediates host–virus interactions. Nucleic Acids Research, 2019, 47, 2455-2471.	14.5	7
22	Analyzing chromosome condensation in yeast by second-harmonic generation microscopy. Current Genetics, 2020, 66, 437-443.	1.7	4
23	Disrupting the MAD2L2-Rev1 Complex Enhances Cell Death upon DNA Damage. Molecules, 2022, 27, 636.	3.8	4
24	Alternative Functional rad21 Paralogs in Fusarium oxysporum. Frontiers in Microbiology, 2019, 10, 1370.	3.5	3
25	Identifying Functional Domains in Subunits of Structural Maintenance of Chromosomes (SMC) Complexes by Transposon Mutagenesis Screen in Yeast. Methods in Molecular Biology, 2019, 2004, 63-78.	0.9	1
26	Redox Potential Regulates Binding of Universal Minicircle Sequence Binding Protein at the Kinetoplast DNA Replication Origin. Eukaryotic Cell, 2010, 9, 477-477.	3.4	0