Matthew J Neale

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Meiosis and beyond – understanding the mechanistic and evolutionary processes shaping the germline genome. Biological Reviews, 2021, 96, 822-841.	10.4	25
2	Concerted cutting by Spo11 illuminates meiotic DNA break mechanics. Nature, 2021, 594, 572-576.	27.8	34
3	Telomerase subunit Est2 marks internal sites that are prone to accumulate DNA damage. BMC Biology, 2021, 19, 247.	3.8	4
4	Convergent genes shape budding yeast pericentromeres. Nature, 2020, 582, 119-123.	27.8	50
5	Principles of meiotic chromosome assembly revealed in S. cerevisiae. Nature Communications, 2019, 10, 4795.	12.8	88
6	A nucleotide resolution map of Top2-linked DNA breaks in the yeast and human genome. Nature Communications, 2019, 10, 4846.	12.8	64
7	Regulatory control of DNA end resection by Sae2 phosphorylation. Nature Communications, 2018, 9, 4016.	12.8	64
8	Meiotic DSB patterning: A multifaceted process. Cell Cycle, 2016, 15, 13-21.	2.6	53
9	Tel1ATM-mediated interference suppresses clustered meiotic double-strand-break formation. Nature, 2015, 520, 114-118.	27.8	150
10	DNA Double-Strand Break Repair Pathway Choice Is Directed by Distinct MRE11 Nuclease Activities. Molecular Cell, 2014, 53, 7-18.	9.7	466
11	Homeostatic regulation of meiotic DSB formation by ATM/ATR. Experimental Cell Research, 2014, 329, 124-131.	2.6	64
12	Positive regulation of meiotic DNA double-strand break formation by activation of the DNA damage checkpoint kinase Mec1(ATR). Open Biology, 2013, 3, 130019.	3.6	65
13	A Hierarchical Combination of Factors Shapes the Genome-wide Topography of Yeast Meiotic Recombination Initiation. Cell, 2011, 144, 719-731.	28.9	520
14	Bidirectional resection of DNA double-strand breaks by Mre11 and Exo1. Nature, 2011, 479, 241-244.	27.8	373
15	Evidence that MEK1 positively promotes interhomologue double-strand break repair. Nucleic Acids Research, 2010, 38, 4349-4360.	14.5	19
16	PRDM9 points the zinc finger at meiotic recombination hotspots. Genome Biology, 2010, 11, 104.	9.6	19
17	Distinct Requirements for the Rad32Mre11 Nuclease and Ctp1CtlP in the Removal of Covalently Bound Topoisomerase I and II from DNA. Molecular Cell, 2009, 33, 117-123.	9.7	170
18	End-Labeling and Analysis of Spo11-Oligonucleotide Complexes in Saccharomyces cerevisiae. Methods in Molecular Biology, 2009, 557, 183-195.	0.9	29

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19	Excess Single-Stranded DNA Inhibits Meiotic Double-Strand Break Repair. PLoS Genetics, 2007, 3, e223.	3.5	25
20	Interactions between Mei4, Rec114, and other proteins required for meiotic DNA double-strand break formation in Saccharomyces cerevisiae. Chromosoma, 2007, 116, 471-486.	2.2	126
21	Clarifying the mechanics of DNA strand exchange in meiotic recombination. Nature, 2006, 442, 153-158.	27.8	383
22	Endonucleolytic processing of covalent protein-linked DNA double-strand breaks. Nature, 2005, 436, 1053-1057.	27.8	536
23	Wild-Type Levels of Spo11-Induced DSBs Are Required for Normal Single-Strand Resection during Meiosis. Molecular Cell, 2002, 9, 835-846.	9.7	58