

# ValÃ©rie Borde

## List of Publications by Year in descending order

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42  
papers

2,943  
citations

257450

24  
h-index

265206

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53  
all docs

53  
docs citations

53  
times ranked

2069  
citing authors

#	ARTICLE	IF	CITATIONS
1	Histone H3 lysine 4 trimethylation marks meiotic recombination initiation sites. <i>EMBO Journal</i> , 2009, 28, 99-111.	7.8	329
2	Direct Coupling Between Meiotic DNA Replication and Recombination Initiation. <i>Science</i> , 2000, 290, 806-809.	12.6	231
3	Mapping Meiotic Single-Strand DNA Reveals a New Landscape of DNA Double-Strand Breaks in <i>Saccharomyces cerevisiae</i> . <i>PLoS Biology</i> , 2007, 5, e324.	5.6	202
4	Spp1, a Member of the Set1 Complex, Promotes Meiotic DSB Formation in Promoters by Tethering Histone H3K4 Methylation Sites to Chromosome Axes. <i>Molecular Cell</i> , 2013, 49, 43-54.	9.7	179
5	Association of Mre11p with Double-Strand Break Sites during Yeast Meiosis. <i>Molecular Cell</i> , 2004, 13, 389-401.	9.7	129
6	Genome-Wide Analysis of Heteroduplex DNA in Mismatch Repair-Deficient Yeast Cells Reveals Novel Properties of Meiotic Recombination Pathways. <i>PLoS Genetics</i> , 2011, 7, e1002305.	3.5	128
7	The multiple roles of the Mre11 complex for meiotic recombination. <i>Chromosome Research</i> , 2007, 15, 551-563.	2.2	118
8	Programmed induction of DNA double strand breaks during meiosis: setting up communication between DNA and the chromosome structure. <i>Current Opinion in Genetics and Development</i> , 2013, 23, 147-155.	3.3	116
9	Budding Yeast ATM/ATR Control Meiotic Double-Strand Break (DSB) Levels by Down-Regulating Rec114, an Essential Component of the DSB-machinery. <i>PLoS Genetics</i> , 2013, 9, e1003545.	3.5	115
10	Crossing and zipping: molecular duties of the ZMM proteins in meiosis. <i>Chromosoma</i> , 2019, 128, 181-198.	2.2	114
11	Use of a Recombination Reporter Insert To Define Meiotic Recombination Domains on Chromosome III of <i>Saccharomyces cerevisiae</i> . <i>Molecular and Cellular Biology</i> , 1999, 19, 4832-4842.	2.3	98
12	A meiotic XPF-ERCC1-like complex recognizes joint molecule recombination intermediates to promote crossover formation. <i>Genes and Development</i> , 2018, 32, 283-296.	5.9	98
13	The control of Spo11's interaction with meiotic recombination hotspots. <i>Genes and Development</i> , 2005, 19, 255-269.	5.9	97
14	Chromosome Synapsis Alleviates Mek1-Dependent Suppression of Meiotic DNA Repair. <i>PLoS Biology</i> , 2016, 14, e1002369.	5.6	95
15	Genome-Wide Redistribution of Meiotic Double-Strand Breaks in <i>Saccharomyces cerevisiae</i> . <i>Molecular and Cellular Biology</i> , 2007, 27, 1868-1880.	2.3	90
16	Differential Association of the Conserved SUMO Ligase Zip3 with Meiotic Double-Strand Break Sites Reveals Regional Variations in the Outcome of Meiotic Recombination. <i>PLoS Genetics</i> , 2013, 9, e1003416.	3.5	90
17	Regulation of the MLH1-MLH3 endonuclease in meiosis. <i>Nature</i> , 2020, 586, 618-622.	27.8	88
18	Concerted action of the MutL <sup>2</sup> heterodimer and Mer3 helicase regulates the global extent of meiotic gene conversion. <i>ELife</i> , 2017, 6, .	6.0	67

#	ARTICLE	IF	CITATIONS
19	The spatial regulation of meiotic recombination hotspots: Are all DSB hotspots crossover hotspots?. <i>Experimental Cell Research</i> , 2012, 318, 1347-1352.	2.6	65
20	Gel Electrophoresis Assays for Analyzing DNA Double-Strand Breaks in <i>Saccharomyces cerevisiae</i> at Various Spatial Resolutions. <i>Methods in Molecular Biology</i> , 2009, 557, 117-142.	0.9	49
21	Correlation between premeiotic DNA replication and chromatin transition at yeast recombination initiation sites. <i>Nucleic Acids Research</i> , 2003, 31, 4085-4090.	14.5	45
22	The PHD finger protein Spp1 has distinct functions in the Set1 and the meiotic DSB formation complexes. <i>PLoS Genetics</i> , 2018, 14, e1007223.	3.5	41
23	Interplay between modifications of chromatin and meiotic recombination hotspots. <i>Biology of the Cell</i> , 2012, 104, 51-69.	2.0	35
24	Crossover recombination and synapsis are linked by adjacent regions within the N terminus of the Zip1 synaptonemal complex protein. <i>PLoS Genetics</i> , 2019, 15, e1008201.	3.5	31
25	Genome-wide Expression Profiling, In Vivo DNA Binding Analysis, and Probabilistic Motif Prediction Reveal Novel Abf1 Target Genes during Fermentation, Respiration, and Sporulation in Yeast. <i>Molecular Biology of the Cell</i> , 2008, 19, 2193-2207.	2.1	29
26	Exo1 recruits Cdc5 polo kinase to MutL <sup>3</sup> to ensure efficient meiotic crossover formation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 30577-30588.	7.1	28
27	Excess Single-Stranded DNA Inhibits Meiotic Double-Strand Break Repair. <i>PLoS Genetics</i> , 2007, 3, e223.	3.5	25
28	Double functions for the Mre11 complex during DNA double-strand break repair and replication. <i>International Journal of Biochemistry and Cell Biology</i> , 2009, 41, 1249-1253.	2.8	23
29	The Zip4 protein directly couples meiotic crossover formation to synaptonemal complex assembly. <i>Genes and Development</i> , 2022, 36, 53-69.	5.9	22
30	Coupling DNA Damage and Repair: an Essential Safeguard during Programmed DNA Double-Strand Breaks?. <i>Trends in Cell Biology</i> , 2020, 30, 87-96.	7.9	20
31	A POLD3/BLM dependent pathway handles DSBs in transcribed chromatin upon excessive RNA:DNA hybrid accumulation. <i>Nature Communications</i> , 2022, 13, 2012.	12.8	20
32	Genetic evidence for the involvement of mismatch repair proteins, PMS2 and MLH3, in a late step of homologous recombination. <i>Journal of Biological Chemistry</i> , 2020, 295, 17460-17475.	3.4	18
33	Molecular basis of the dual role of the Mlh1-Mlh3 endonuclease in MMR and in meiotic crossover formation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	18
34	The Pif1 helicase is actively inhibited during meiotic recombination which restrains gene conversion tract length. <i>Nucleic Acids Research</i> , 2021, 49, 4522-4533.	14.5	16
35	The mapping of DNA topoisomerase sites in vivo: A tool to enlight the functions of topoisomerases. <i>Biochimie</i> , 1998, 80, 223-233.	2.6	14
36	The CAF-1 and Hir Histone Chaperones Associate with Sites of Meiotic Double-Strand Breaks in Budding Yeast. <i>PLoS ONE</i> , 2015, 10, e0125965.	2.5	13

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37	A Timeless but Timely Connection between Replication and Recombination. <i>Cell</i> , 2014, 158, 697-698.	28.9	9
38	Meiosis: Early DNA Double-Strand Breaks Pave the Way for Inter-Homolog Repair. <i>Developmental Cell</i> , 2015, 32, 663-664.	7.0	8
39	Methods to Map Meiotic Recombination Proteins in <i>Saccharomyces cerevisiae</i> . <i>Methods in Molecular Biology</i> , 2021, 2153, 295-306.	0.9	7
40	DNA topoisomerase II sites in the histone H4 gene during the highly synchronous cell cycle of <i>Physarum polycephalum</i> [published erratum appears in <i>Nucleic Acids Res</i> 1998 Oct 15;26(20):following 4789]. <i>Nucleic Acids Research</i> , 1998, 26, 2042-2049.	14.5	6
41	Special issue on "Recent advances in meiosis from DNA replication to chromosome segregation". <i>Chromosoma</i> , 2019, 128, 177-180.	2.2	0
42	Réplication et recombinaison vont de pair pendant la méiose.. <i>Medecine/Sciences</i> , 2001, 17, 482.	0.2	0