

# Joanna R Morris

## List of Publications by Year in descending order

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36  
papers

2,652  
citations

257429

24  
h-index

361001

35  
g-index

47  
all docs

47  
docs citations

47  
times ranked

4168  
citing authors

#	ARTICLE	IF	CITATIONS
1	The SUMO modification pathway is involved in the BRCA1 response to genotoxic stress. <i>Nature</i> , 2009, 462, 886-890.	27.8	377
2	Critical research gaps and translational priorities for the successful prevention and treatment of breast cancer. <i>Breast Cancer Research</i> , 2013, 15, R92.	5.0	320
3	BRCA1 and BARD1 induces the formation of conjugated ubiquitin structures, dependent on K6 of ubiquitin, in cells during DNA replication and repair. <i>Human Molecular Genetics</i> , 2004, 13, 807-817.	2.9	230
4	BRCA1 RING Function Is Essential for Tumor Suppression but Dispensable for Therapy Resistance. <i>Cancer Cell</i> , 2011, 20, 797-809.	16.8	228
5	Human BRCA1 and BARD1 ubiquitin ligase activity counteracts chromatin barriers to DNA resection. <i>Nature Structural and Molecular Biology</i> , 2016, 23, 647-655.	8.2	222
6	The proteasomal de-ubiquitinating enzyme POH1 promotes the double-strand DNA break response. <i>EMBO Journal</i> , 2012, 31, 3918-3934.	7.8	127
7	BRCA1 methylation: a significant role in tumour development?. <i>Seminars in Cancer Biology</i> , 2002, 12, 359-371.	9.6	104
8	Genetic analysis of BRCA1 ubiquitin ligase activity and its relationship to breast cancer susceptibility. <i>Human Molecular Genetics</i> , 2006, 15, 599-606.	2.9	96
9	Isomerization of BRCA1 and BARD1 promotes replication fork protection. <i>Nature</i> , 2019, 571, 521-527.	27.8	88
10	The deSUMOylase SENP7 promotes chromatin relaxation for homologous recombination DNA repair. <i>EMBO Reports</i> , 2013, 14, 975-983.	4.5	82
11	USP48 restrains resection by site-specific cleavage of the BRCA1 ubiquitin mark from H2A. <i>Nature Communications</i> , 2018, 9, 229.	12.8	76
12	PrimPol-dependent single-stranded gap formation mediates homologous recombination at bulky DNA adducts. <i>Nature Communications</i> , 2020, 11, 5863.	12.8	69
13	SUMO, a small, but powerful, regulator of double-strand break repair. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2017, 372, 20160281.	4.0	66
14	Two GC Boxes (Sp1 Sites) Are Involved in Regulation of the Activity of the Epithelium-specific MUC1 Promoter. <i>Journal of Biological Chemistry</i> , 1996, 271, 18140-18147.	3.4	54
15	The BRCA1 Ubiquitin ligase function sets a new trend for remodelling in DNA repair. <i>Nucleus</i> , 2017, 8, 116-125.	2.2	46
16	A fork in the road: Where homologous recombination and stalled replication fork protection part ways. <i>Seminars in Cell and Developmental Biology</i> , 2021, 113, 14-26.	5.0	44
17	Identification of Residues Required for the Interaction of BARD1 with BRCA1. <i>Journal of Biological Chemistry</i> , 2002, 277, 9382-9386.	3.4	38
18	Moving Mountains – The BRCA1 Promotion of DNA Resection. <i>Frontiers in Molecular Biosciences</i> , 2019, 6, 79.	3.5	37

#	ARTICLE	IF	CITATIONS
19	SUMO in the mammalian response to DNA damage. <i>Biochemical Society Transactions</i> , 2010, 38, 92-97.	3.4	32
20	The deSUMOylase SENP2 coordinates homologous recombination and nonhomologous end joining by independent mechanisms. <i>Genes and Development</i> , 2019, 33, 333-347.	5.9	32
21	Mouse ST6Gal sialyltransferase gene expression during mammary gland lactation. <i>Glycobiology</i> , 2001, 11, 407-412.	2.5	31
22	Lymphocyte radiosensitivity in BRCA1 and BRCA2 mutation carriers and implications for breast cancer susceptibility. <i>International Journal of Cancer</i> , 2007, 121, 1631-1636.	5.1	30
23	More Modifiers Move on DNA Damage. <i>Cancer Research</i> , 2010, 70, 3861-3863.	0.9	30
24	Discovery of peptide ligands targeting a specific ubiquitin-like domain binding site in the deubiquitinase USP11. <i>Journal of Biological Chemistry</i> , 2019, 294, 424-436.	3.4	28
25	SUMO in the DNA Double-Stranded Break Response: Similarities, Differences, and Cooperation with Ubiquitin. <i>Journal of Molecular Biology</i> , 2017, 429, 3376-3387.	4.2	27
26	GSK3 $\beta$ -SCFFBXW7 $\mu$ mediated phosphorylation and ubiquitination of IRF1 are required for its transcription-dependent turnover. <i>Nucleic Acids Research</i> , 2019, 47, 4476-4494.	14.5	21
27	The Sp1 Transcription Factor Regulates Cell Type-Specific Transcription of MUC1. <i>DNA and Cell Biology</i> , 2001, 20, 133-139.	1.9	20
28	Structural evolution of the BRCA1 genomic region in primates. <i>Genomics</i> , 2004, 84, 1071-1082.	2.9	20
29	Regulation of MUC1 Expression in Human Mammary Cell Lines by the c-ErbB2 and Ras Signaling Pathways. <i>DNA and Cell Biology</i> , 2001, 20, 265-274.	1.9	19
30	DNA methylation profiling to assess pathogenicity of BRCA1 unclassified variants in breast cancer. <i>Epigenetics</i> , 2015, 10, 1121-1132.	2.7	12
31	Up-regulation of MUC1 in mammary tumors generated in a double-transgenic mouse expressing human MUC1 cDNA, under the control of 1.4-kb 5' MUC1 promoter sequence and the middle T oncogene, expressed from the MMTV promoter. <i>International Journal of Cancer</i> , 2001, 92, 382-387.	5.1	11
32	Attenuation of the ubiquitin conjugate DNA damage signal by the proteasomal DUB POH1. <i>Cell Cycle</i> , 2012, 11, 4103-4104.	2.6	11
33	Imaging nanoscale nuclear structures with expansion microscopy. <i>Journal of Cell Science</i> , 2022, 135, .	2.0	6
34	Is it a wrap? Nucleosome interactions of the BRCA1-binding partner, BARD1, steal the scene. <i>Nature Structural and Molecular Biology</i> , 2021, 28, 708-710.	8.2	5
35	BRCA1-BARD1: the importance of being in shape. <i>Molecular and Cellular Oncology</i> , 2019, 6, e1656500.	0.7	4
36	Recent Advances in Understanding the Cellular Functions of BRCA1. , 2009, , 75-92.		1