

Xiaochun Yu

List of Publications by Year in descending order

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67
papers

5,315
citations

185998

28
h-index

110170

64
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69
all docs

69
docs citations

69
times ranked

6336
citing authors

#	ARTICLE	IF	CITATIONS
1	RNF8 Transduces the DNA-Damage Signal via Histone Ubiquitylation and Checkpoint Protein Assembly. <i>Cell</i> , 2007, 131, 901-914.	13.5	906
2	The BRCT Domain Is a Phospho-Protein Binding Domain. <i>Science</i> , 2003, 302, 639-642.	6.0	770
3	DNA Damage-Induced Cell Cycle Checkpoint Control Requires CtIP, a Phosphorylation-Dependent Binding Partner of BRCA1 C-Terminal Domains. <i>Molecular and Cellular Biology</i> , 2004, 24, 9478-9486.	1.1	355
4	Function of BRCA1 in the DNA Damage Response Is Mediated by ADP-Ribosylation. <i>Cancer Cell</i> , 2013, 23, 693-704.	7.7	261
5	BRCA1 ubiquitinates its phosphorylation-dependent binding partner CtIP. <i>Genes and Development</i> , 2006, 20, 1721-1726.	2.7	254
6	Functions of PARylation in DNA Damage Repair Pathways. <i>Genomics, Proteomics and Bioinformatics</i> , 2016, 14, 131-139.	3.0	215
7	Phosphopeptide Binding Specificities of BRCA1 COOH-terminal (BRCT) Domains. <i>Journal of Biological Chemistry</i> , 2003, 278, 52914-52918.	1.6	206
8	RNF8-Dependent Histone Modifications Regulate Nucleosome Removal during Spermatogenesis. <i>Developmental Cell</i> , 2010, 18, 371-384.	3.1	200
9	Chfr is required for tumor suppression and Aurora A regulation. <i>Nature Genetics</i> , 2005, 37, 401-406.	9.4	199
10	The role of poly ADP-ribosylation in the first wave of DNA damage response. <i>Nucleic Acids Research</i> , 2017, 45, 8129-8141.	6.5	157
11	ADP-ribosyltransferases, an update on function and nomenclature. <i>FEBS Journal</i> , 2022, 289, 7399-7410.	2.2	150
12	The FHA and BRCT domains recognize ADP-ribosylation during DNA damage response. <i>Genes and Development</i> , 2013, 27, 1752-1768.	2.7	132
13	PARP2 mediates branched poly ADP-ribosylation in response to DNA damage. <i>Nature Communications</i> , 2018, 9, 3233.	5.8	114
14	Super-resolution imaging identifies PARP1 and the Ku complex acting as DNA double-strand break sensors. <i>Nucleic Acids Research</i> , 2018, 46, 3446-3457.	6.5	88
15	Structural Basis of BACH1 Phosphopeptide Recognition by BRCA1 Tandem BRCT Domains. <i>Structure</i> , 2004, 12, 1137-1146.	1.6	87
16	53BP1 Cooperates with p53 and Functions as a Haploinsufficient Tumor Suppressor in Mice. <i>Molecular and Cellular Biology</i> , 2005, 25, 10079-10086.	1.1	80
17	ADP-Ribosyltransferases and Poly ADP-Ribosylation. <i>Current Protein and Peptide Science</i> , 2015, 16, 491-501.	0.7	76
18	SIRT1 Activation Disrupts Maintenance of Myelodysplastic Syndrome Stem and Progenitor Cells by Restoring TET2 Function. <i>Cell Stem Cell</i> , 2018, 23, 355-369.e9.	5.2	68

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19	Poly-ADP ribosylation in DNA damage response and cancer therapy. <i>Mutation Research - Reviews in Mutation Research</i> , 2019, 780, 82-91.	2.4	61
20	Targeting dePARylation selectively suppresses DNA repair in defective and PARP inhibitor-resistant malignancies. <i>Science Advances</i> , 2019, 5, eaav4340.	4.7	57
21	The oligonucleotide/oligosaccharide-binding fold motif is a poly(ADP-ribose)-binding domain that mediates DNA damage response. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 7278-7283.	3.3	55
22	The PIN domain of EXO1 recognizes poly(ADP-ribose) in DNA damage response. <i>Nucleic Acids Research</i> , 2015, 43, 10782-10794.	6.5	53
23	NAD ⁺ is an endogenous PARP inhibitor in DNA damage response and tumor suppression. <i>Nature Communications</i> , 2019, 10, 693.	5.8	45
24	Selective targeting of TET catalytic domain promotes somatic cell reprogramming. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 3621-3626.	3.3	44
25	Poly(ADP-Ribose) Mediates the BRCA2-Dependent Early DNA Damage Response. <i>Cell Reports</i> , 2015, 13, 678-689.	2.9	43
26	OGT restrains the expansion of DNA damage signaling. <i>Nucleic Acids Research</i> , 2016, 44, gkw663.	6.5	40
27	Chemopreventive Effects of ROS Targeting in a Murine Model of BRCA1-Deficient Breast Cancer. <i>Cancer Research</i> , 2017, 77, 448-458.	0.4	40
28	The zinc finger proteins ZNF644 and WIZ regulate the G9a/GLP complex for gene repression. <i>ELife</i> , 2015, 4, .	2.8	40
29	Double-strand break repair on sex chromosomes: challenges during male meiotic prophase. <i>Cell Cycle</i> , 2015, 14, 516-525.	1.3	37
30	ADP-ribosylation of histone variant H2AX promotes base excision repair. <i>EMBO Journal</i> , 2021, 40, e104542.	3.5	32
31	Poly(ADP-ribosyl)ation mediates early phase histone eviction at DNA lesions. <i>Nucleic Acids Research</i> , 2020, 48, 3001-3013.	6.5	29
32	Regulation of the DNA damage response on male meiotic sex chromosomes. <i>Nature Communications</i> , 2013, 4, 2105.	5.8	28
33	CTCF participates in DNA damage response via poly(ADP-ribosyl)ation. <i>Scientific Reports</i> , 2017, 7, 43530.	1.6	25
34	Human DNA ligase IV is able to use NAD ⁺ as an alternative adenylation donor for DNA ends ligation. <i>Nucleic Acids Research</i> , 2019, 47, 1321-1334.	6.5	22
35	LGR5 regulates gastric adenocarcinoma cell proliferation and invasion via activating Wnt signaling pathway. <i>Oncogenesis</i> , 2018, 7, 57.	2.1	20
36	Targeting dePARylation for cancer therapy. <i>Cell and Bioscience</i> , 2020, 10, 7.	2.1	20

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37	Truncated PARP1 mediates ADP-ribosylation of RNA polymerase III for apoptosis. <i>Cell Discovery</i> , 2022, 8, 3.	3.1	20
38	GALNT14 Involves the Regulation of Multidrug Resistance in Breast Cancer Cells. <i>Translational Oncology</i> , 2018, 11, 786-793.	1.7	19
39	Zinc Finger Protein 618 Regulates the Function of UHRF2 (Ubiquitin-like with PHD and Ring Finger) Tj ETQq1 1 0.784314 rgBT /Overlo 13679-13688.	1.6	17
40	Structureâ€“function analyses reveal the mechanism of the ARH3-dependent hydrolysis of ADP-ribosylation. <i>Journal of Biological Chemistry</i> , 2018, 293, 14470-14480.	1.6	17
41	Molecular basis for the inhibition of the methyl-lysine binding function of 53BP1 by TIRR. <i>Nature Communications</i> , 2018, 9, 2689.	5.8	17
42	The RNF20/40 complex regulates p53-dependent gene transcription and mRNA splicing. <i>Journal of Molecular Cell Biology</i> , 2020, 12, 113-124.	1.5	16
43	Topoisomerase II Regulates the Maintenance of DNA Methylation. <i>Journal of Biological Chemistry</i> , 2015, 290, 851-860.	1.6	15
44	EFEMP2 Mediates GALNT14-Dependent Breast Cancer Cell Invasion. <i>Translational Oncology</i> , 2018, 11, 346-352.	1.7	15
45	Molecular basis for the MacroD1-mediated hydrolysis of ADP-ribosylation. <i>DNA Repair</i> , 2020, 94, 102899.	1.3	15
46	Pre-ribosomal RNA reorganizes DNA damage repair factors in nucleus during meiotic prophase and DNA damage response. <i>Cell Research</i> , 2022, 32, 254-268.	5.7	15
47	UHRF2 regulates local 5-methylcytosine and suppresses spontaneous seizures. <i>Epigenetics</i> , 2017, 12, 551-560.	1.3	14
48	ATR/ATM-Mediated Phosphorylation of BRCA1 T1394 Promotes Homologous Recombinational Repair and G2â€“M Checkpoint Maintenance. <i>Cancer Research</i> , 2021, 81, 4676-4684.	0.4	14
49	8â€“chloroâ€“adenosine activity in FLT3â€“TD acute myeloid leukemia. <i>Journal of Cellular Physiology</i> , 2019, 234, 16295-16303.	2.0	12
50	Targeting reactive nitrogen species suppresses hereditary pancreatic cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 7106-7111.	3.3	11
51	Tissue-Specific Carcinogens as Soil to Seed BRCA1/2-Mutant Hereditary Cancers. <i>Trends in Cancer</i> , 2020, 6, 559-568.	3.8	11
52	OGA is associated with deglycosylation of NONO and the KU complex during DNA damage repair. <i>Cell Death and Disease</i> , 2021, 12, 622.	2.7	11
53	Poly(ADP-ribose) protects vascular smooth muscle cells from oxidative DNA damage. <i>BMB Reports</i> , 2015, 48, 354-359.	1.1	9
54	Functional deficiency of DNA repair gene EXO5 results in androgen-induced genomic instability and prostate tumorigenesis. <i>Oncogene</i> , 2020, 39, 1246-1259.	2.6	8

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55	AI26 inhibits the ADP-ribosylhydrolase ARH3 and suppresses DNA damage repair. <i>Journal of Biological Chemistry</i> , 2020, 295, 13838-13849.	1.6	8
56	The role of dePARylation in DNA damage repair and cancer suppression. <i>DNA Repair</i> , 2019, 76, 20-29.	1.3	7
57	Silencing of CHFR Sensitizes Gastric Carcinoma to PARP Inhibitor Treatment. <i>Translational Oncology</i> , 2020, 13, 113-121.	1.7	7
58	Cancer-associated 53BP1 mutations induce DNA damage repair defects. <i>Cancer Letters</i> , 2021, 501, 43-54.	3.2	7
59	Ribosomal RNA regulates chromosome clustering during mitosis. <i>Cell Discovery</i> , 2022, 8, .	3.1	7
60	CHFR is important for the survival of male premeiotic germ cells. <i>Cell Cycle</i> , 2015, 14, 3454-3460.	1.3	5
61	ATR prevents Ca ²⁺ overload-induced necrotic cell death through phosphorylation-mediated inactivation of PARP1 without DNA damage signaling. <i>FASEB Journal</i> , 2021, 35, e21373.	0.2	4
62	ADP-ribosylhydrolases: from DNA damage repair to COVID-19. <i>Journal of Zhejiang University: Science B</i> , 2021, 22, 21-30.	1.3	2
63	Functional defects of cancer-associated MDC1 mutations in DNA damage repair. <i>DNA Repair</i> , 2022, 114, 103330.	1.3	2
64	A special issue on the DNA damage response and genomic instability. <i>Acta Biochimica Et Biophysica Sinica</i> , 2016, 48, 593-593.	0.9	0
65	TET2 Activity Is Modulated By SIRT1-Mediated Protein Deacetylation: A Potential Therapeutic Target in Myelodysplastic Syndrome. <i>Blood</i> , 2016, 128, 1053-1053.	0.6	0
66	Antileukemic Activity of 8-Chloro-Adenosine (8-Cl-Ado) Is Mediated By Mir-155 Degradation and ErbB3 Binding Protein (Ebp1)-Dependent p53 Activation: A Novel Therapeutic Approach for FLT3-ITD Acute Myeloid Leukemia (AML). <i>Blood</i> , 2018, 132, 3938-3938.	0.6	0
67	The impact of TOPBP1 mutations in human cancers on the DNA damage response. <i>Genome Instability & Disease</i> , 0, , .	0.5	0