

Martin A M Reijns

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4812511/publications.pdf>

Version: 2024-02-01

38
papers

4,349
citations

201385

27
h-index

315357

38
g-index

49
all docs

49
docs citations

49
times ranked

6968
citing authors

#	ARTICLE	IF	CITATIONS
1	Signatures of TOP1 transcription-associated mutagenesis in cancer and germline. <i>Nature</i> , 2022, 602, 623-631.	13.7	38
2	DNA damage contributes to neurotoxic inflammation in Aicardi-Goutières syndrome astrocytes. <i>Journal of Experimental Medicine</i> , 2022, 219, .	4.2	35
3	User acceptability of saliva and gargle samples for identifying COVID-19 positive high-risk workers and household contacts. <i>Diagnostic Microbiology and Infectious Disease</i> , 2022, , 115732.	0.8	1
4	Novel <i>Escherichia coli</i> active site <i>dnaE</i> alleles with altered base and sugar selectivity. <i>Molecular Microbiology</i> , 2021, 116, 909-925.	1.2	3
5	cGAS-mediated induction of type I interferon due to inborn errors of histone pre-mRNA processing. <i>Nature Genetics</i> , 2020, 52, 1364-1372.	9.4	105
6	PRIM1 deficiency causes a distinctive primordial dwarfism syndrome. <i>Genes and Development</i> , 2020, 34, 1520-1533.	2.7	20
7	A sensitive and affordable multiplex RT-qPCR assay for SARS-CoV-2 detection. <i>PLoS Biology</i> , 2020, 18, e3001030.	2.6	32
8	Biallelic variants in <i>DNA2</i> cause microcephalic primordial dwarfism. <i>Human Mutation</i> , 2019, 40, 1063-1070.	1.1	16
9	Epithelial RNase H2 Maintains Genome Integrity and Prevents Intestinal Tumorigenesis in Mice. <i>Gastroenterology</i> , 2019, 156, 145-159.e19.	0.6	46
10	Gain-of-function DNMT3A mutations cause microcephalic dwarfism and hypermethylation of Polycomb-regulated regions. <i>Nature Genetics</i> , 2019, 51, 96-105.	9.4	110
11	DNA Polymerase Epsilon Deficiency Causes IMAGE Syndrome with Variable Immunodeficiency. <i>American Journal of Human Genetics</i> , 2018, 103, 1038-1044.	2.6	71
12	Ribonucleotide Excision Repair Is Essential to Prevent Squamous Cell Carcinoma of the Skin. <i>Cancer Research</i> , 2018, 78, 5917-5926.	0.4	40
13	CRISPR screens identify genomic ribonucleotides as a source of PARP-trapping lesions. <i>Nature</i> , 2018, 559, 285-289.	13.7	297
14	RNase H2, mutated in Aicardi-Goutières syndrome, promotes LINE1 retrotransposition. <i>EMBO Journal</i> , 2018, 37, .	3.5	67
15	Mutations in DONSON disrupt replication fork stability and cause microcephalic dwarfism. <i>Nature Genetics</i> , 2017, 49, 537-549.	9.4	81
16	Rare ADAR and RNASEH2B variants and a type I interferon signature in glioma and prostate carcinoma risk and tumorigenesis. <i>Acta Neuropathologica</i> , 2017, 134, 905-922.	3.9	12
17	cGAS surveillance of micronuclei links genome instability to innate immunity. <i>Nature</i> , 2017, 548, 461-465.	13.7	1,158
18	Aberrant ribonucleotide incorporation and multiple deletions in mitochondrial DNA of the murine MPV17 disease model. <i>Nucleic Acids Research</i> , 2017, 45, 12808-12815.	6.5	43

#	ARTICLE	IF	CITATIONS
19	Ribonuclease H2 mutations induce a cGAS/STING-dependent innate immune response. <i>EMBO Journal</i> , 2016, 35, 831-844.	3.5	200
20	TRAIIP promotes DNA damage response during genome replication and is mutated in primordial dwarfism. <i>Nature Genetics</i> , 2016, 48, 36-43.	9.4	74
21	Defective removal of ribonucleotides from DNA promotes systemic lupus erythematosus. <i>Pediatric Rheumatology</i> , 2015, 13, .	0.9	1
22	Defective removal of ribonucleotides from DNA promotes systemic autoimmunity. <i>Journal of Clinical Investigation</i> , 2015, 125, 413-424.	3.9	190
23	Lagging-strand replication shapes the mutational landscape of the genome. <i>Nature</i> , 2015, 518, 502-506.	13.7	213
24	Genome-wide mapping of embedded ribonucleotides and other noncanonical nucleotides using emRiboSeq and EndoSeq. <i>Nature Protocols</i> , 2015, 10, 1433-1444.	5.5	42
25	RNA:DNA hybrids are a novel molecular pattern sensed by TLR9. <i>EMBO Journal</i> , 2014, 33, 542-558.	3.5	133
26	Ribonuclease H2 in health and disease. <i>Biochemical Society Transactions</i> , 2014, 42, 717-725.	1.6	37
27	Ribonucleotides Misincorporated into DNA Act as Strand-Discrimination Signals in Eukaryotic Mismatch Repair. <i>Molecular Cell</i> , 2013, 50, 323-332.	4.5	139
28	Synonymous Mutations in <i>RNASEH2A</i> Create Cryptic Splice Sites Impairing RNase H2 Enzyme Function in Aicardi-Goutières Syndrome. <i>Human Mutation</i> , 2013, 34, 1066-1070.	1.1	16
29	Enzymatic Removal of Ribonucleotides from DNA Is Essential for Mammalian Genome Integrity and Development. <i>Cell</i> , 2012, 149, 1008-1022.	13.5	397
30	Human and Mouse Mutations in WDR35 Cause Short-Rib Polydactyly Syndromes Due to Abnormal Ciliogenesis. <i>American Journal of Human Genetics</i> , 2011, 88, 508-515.	2.6	122
31	PCNA directs type 2 RNase H activity on DNA replication and repair substrates. <i>Nucleic Acids Research</i> , 2011, 39, 3652-3666.	6.5	112
32	The Structure of the Human RNase H2 Complex Defines Key Interaction Interfaces Relevant to Enzyme Function and Human Disease. <i>Journal of Biological Chemistry</i> , 2011, 286, 10530-10539.	1.6	94
33	Analysis of Lsm1p and Lsm8p domains in the cellular localization of Lsm complexes in budding yeast. <i>FEBS Journal</i> , 2009, 276, 3602-3617.	2.2	10
34	A role for Q/N-rich aggregation-prone regions in P-body localization. <i>Journal of Cell Science</i> , 2008, 121, 2463-2472.	1.2	191
35	Requirements for nuclear localization of the Lsm2-8p complex and competition between nuclear and cytoplasmic Lsm complexes. <i>Journal of Cell Science</i> , 2007, 120, 4310-4320.	1.2	25
36	The Lsm2-8 complex determines nuclear localization of the spliceosomal U6 snRNA. <i>Nucleic Acids Research</i> , 2007, 35, 923-929.	6.5	30

#	ARTICLE	IF	CITATIONS
37	Mutagenesis of PepA suggests a new model for the Xer/ceI synaptic complex. <i>Molecular Microbiology</i> , 2005, 57, 927-941.	1.2	41
38	Identification of VP19 and VP15 of white spot syndrome virus (WSSV) and glycosylation status of the WSSV major structural proteins. <i>Journal of General Virology</i> , 2002, 83, 257-265.	1.3	105