

Jens Mayer

List of Publications by Year in descending order

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56
papers

2,937
citations

147801

31
h-index

175258

52
g-index

60
all docs

60
docs citations

60
times ranked

3637
citing authors

#	ARTICLE	IF	CITATIONS
1	New insights into the Tyrolean Iceman's origin and phenotype as inferred by whole-genome sequencing. <i>Nature Communications</i> , 2012, 3, 698.	12.8	382
2	CpG Methylation Directly Regulates Transcriptional Activity of the Human Endogenous Retrovirus Family HERV-K(HML-2). <i>Journal of Virology</i> , 2005, 79, 876-883.	3.4	188
3	An almost-intact human endogenous retrovirus K on human chromosome 7. <i>Nature Genetics</i> , 1999, 21, 257-258.	21.4	139
4	Nomenclature for endogenous retrovirus (ERV) loci. <i>Retrovirology</i> , 2018, 15, 59.	2.0	103
5	Classification and nomenclature of endogenous retroviral sequences (ERVs). <i>Gene</i> , 2009, 448, 115-123.	2.2	101
6	Expression patterns of transcribed human endogenous retrovirus HERV-K(HML-2) loci in human tissues and the need for a HERV Transcriptome Project. <i>BMC Genomics</i> , 2008, 9, 354.	2.8	95
7	A revised nomenclature for transcribed human endogenous retroviral loci. <i>Mobile DNA</i> , 2011, 2, 7.	3.6	94
8	Transcriptional Profiling of Human Endogenous Retrovirus Group HERV-K(HML-2) Loci in Melanoma. <i>Genome Biology and Evolution</i> , 2013, 5, 307-328.	2.5	94
9	The human L1 promoter: Variable transcription initiation sites and a major impact of upstream flanking sequence on promoter activity. <i>Genome Research</i> , 2004, 14, 2253-2260.	5.5	92
10	Ortervirales: New Virus Order Unifying Five Families of Reverse-Transcribing Viruses. <i>Journal of Virology</i> , 2018, 92, .	3.4	79
11	Variable Transcriptional Activity of Endogenous Retroviruses in Human Breast Cancer. <i>Journal of Virology</i> , 2008, 82, 1808-1818.	3.4	78
12	Complex humoral immune response against a benign tumor: Frequent antibody response against specific antigens as diagnostic targets. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 9601-9606.	7.1	73
13	Human Endogenous Retrovirus Family HERV-K(HML-2) RNA Transcripts Are Selectively Packaged into Retroviral Particles Produced by the Human Germ Cell Tumor Line Tera-1 and Originate Mainly from a Provirus on Chromosome 22q11.21. <i>Journal of Virology</i> , 2008, 82, 10008-10016.	3.4	73
14	Human endogenous retrovirus HERV-K(HML-2) RNA causes neurodegeneration through Toll-like receptors. <i>JCI Insight</i> , 2020, 5, .	5.0	68
15	Analysis of transcribed human endogenous retrovirus W env loci clarifies the origin of multiple sclerosis-associated retrovirus env sequences. <i>Retrovirology</i> , 2009, 6, 37.	2.0	65
16	Human endogenous retroviruses in the primate lineage and their influence on host genomes. <i>Cytogenetic and Genome Research</i> , 2005, 110, 448-456.	1.1	62
17	HERV-K(HML-2) rec and np9 transcripts not restricted to disease but present in many normal human tissues. <i>Mobile DNA</i> , 2015, 6, 4.	3.6	62
18	Comprehensive Analysis of Human Endogenous Retrovirus Group HERV-W Locus Transcription in Multiple Sclerosis Brain Lesions by High-Throughput Amplicon Sequencing. <i>Journal of Virology</i> , 2013, 87, 13837-13852.	3.4	59

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19	HERV-K(OLD): Ancestor Sequences of the Human Endogenous Retrovirus Family HERV-K(HML-2). <i>Journal of Virology</i> , 2001, 75, 8917-8926.	3.4	51
20	Human Endogenous Retrovirus Family HERV-K(HML-5): Status, Evolution, and Reconstruction of an Ancient Betaretrovirus in the Human Genome. <i>Journal of Virology</i> , 2004, 78, 8788-8798.	3.4	51
21	Modulation of human endogenous retrovirus (HERV) transcription during persistent and de novo HIV-1 infection. <i>Retrovirology</i> , 2015, 12, 27.	2.0	48
22	Expression of Human Endogenous Retrovirus-W Including Syncytin-1 in Cutaneous T-Cell Lymphoma. <i>PLoS ONE</i> , 2013, 8, e76281.	2.5	47
23	Transcriptional profiling of HERV-K(HML-2) in amyotrophic lateral sclerosis and potential implications for expression of HML-2 proteins. <i>Molecular Neurodegeneration</i> , 2018, 13, 39.	10.8	47
24	Genomic Organization of the Human Endogenous Retrovirus HERV-K(HML-2.HOM) (ERV6) on Chromosome 7. <i>Genomics</i> , 2001, 72, 314-320.	2.9	46
25	Human endogenous retrovirus HERV-K(HML-2) activity in prostate cancer is dominated by a few loci. <i>Prostate</i> , 2015, 75, 1958-1971.	2.3	43
26	Identification of Protease Specificity by Combining Proteome-Derived Peptide Libraries and Quantitative Proteomics. <i>Molecular and Cellular Proteomics</i> , 2016, 15, 2515-2524.	3.8	43
27	Degradation and remobilization of endogenous retroviruses by recombination during the earliest stages of a germ-line invasion. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 8609-8614.	7.1	40
28	The Human Endogenous Retrovirus Family HERV-K(HML-3). <i>Genomics</i> , 2002, 80, 331-343.	2.9	39
29	The Hsp70 chaperones of the <i>Trityps</i> are characterized by unusual features and novel members. <i>Parasitology International</i> , 2010, 59, 497-505.	1.3	39
30	Properties of LINE-1 proteins and repeat element expression in the context of amyotrophic lateral sclerosis. <i>Mobile DNA</i> , 2018, 9, 35.	3.6	37
31	Human Endogenous Retrovirus K Homologous Sequences and Their Coding Capacity in Old World Primates. <i>Journal of Virology</i> , 1998, 72, 1870-1875.	3.4	37
32	Multiple human endogenous retrovirus (HERV-K) loci with <i>gag</i> open reading frames in the human genome. <i>Cytogenetic and Genome Research</i> , 1997, 78, 1-5.	1.1	35
33	Human endogenous retrovirus HERV-K(HML-2) proviruses with <i>Rec</i> protein coding capacity and transcriptional activity. <i>Virology</i> , 2004, 322, 190-198.	2.4	34
34	Chromosomal assignment of human endogenous retrovirus K (HERV-K) <i>env</i> open reading frames. <i>Cytogenetic and Genome Research</i> , 1997, 79, 157-161.	1.1	34
35	Human Endogenous Retrovirus HERV-K14 Families: Status, Variants, Evolution, and Mobilization of Other Cellular Sequences. <i>Journal of Virology</i> , 2005, 79, 2941-2949.	3.4	33
36	An N-terminally truncated envelope protein encoded by a human endogenous retrovirus W locus on chromosome Xq22.3. <i>Retrovirology</i> , 2010, 7, 69.	2.0	30

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37	HERV-E-Mediated Modulation of PLA2G4A Transcription in Urothelial Carcinoma. <i>PLoS ONE</i> , 2012, 7, e49341.	2.5	29
38	Human endogenous retrovirus HERV-K(HML-2) encodes a stable signal peptide with biological properties distinct from Rec. <i>Retrovirology</i> , 2009, 6, 17.	2.0	27
39	Activation of HERV-K(HML-2) disrupts cortical patterning and neuronal differentiation by increasing NTRK3. <i>Cell Stem Cell</i> , 2021, 28, 1566-1581.e8.	11.1	27
40	HERV-W group evolutionary history in non-human primates: characterization of ERV-W orthologs in Catarrhini and related ERV groups in Platyrrhini. <i>BMC Evolutionary Biology</i> , 2018, 18, 6.	3.2	26
41	ICTV Virus Taxonomy Profile: Retroviridae 2021. <i>Journal of General Virology</i> , 2021, 102, .	2.9	24
42	Haplotype Analysis of the Human Endogenous Retrovirus Locus HERV-K(HML-2.HOM) and Its Evolutionary Implications. <i>Journal of Molecular Evolution</i> , 2005, 61, 706-715.	1.8	23
43	Presence of dUTPase in the Various Human Endogenous Retrovirus K (HERV-K) Families. <i>Journal of Molecular Evolution</i> , 2003, 57, 642-649.	1.8	19
44	Expression pattern analysis of transcribed HERV sequences is complicated by ex vivo recombination. <i>Retrovirology</i> , 2007, 4, 39.	2.0	19
45	The mouse KrÄppel-like Factor 4 (Klf4) gene: Four functional polyadenylation sites which are used in a cell-specific manner as revealed by testicular transcript analysis and multiple processed pseudogenes. <i>Gene</i> , 2005, 361, 149-156.	2.2	18
46	Np9, a cellular protein of retroviral ancestry restricted to human, chimpanzee and gorilla, binds and regulates ubiquitin ligase MDM2. <i>Cell Cycle</i> , 2015, 14, 2619-2633.	2.6	16
47	On the origin of a pathogenic HERV-W envelope protein present in multiple sclerosis lesions. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 19791-19792.	7.1	13
48	Identification and characterization of ERV-W-like sequences in Platyrrhini species provides new insights into the evolutionary history of ERV-W in primates. <i>Mobile DNA</i> , 2020, 11, 6.	3.6	12
49	A novel endogenous betaretrovirus group characterized from polar bears (<i>Ursus maritimus</i>) and giant pandas (<i>Ailuropoda melanoleuca</i>). <i>Virology</i> , 2013, 443, 1-10.	2.4	11
50	A human endogenous retrovirus encoded protease potentially cleaves numerous cellular proteins. <i>Mobile DNA</i> , 2019, 10, 36.	3.6	9
51	PTC124 for cystic fibrosis. <i>Lancet, The</i> , 2009, 373, 1426.	13.7	8
52	An Evolutionarily Young Polar Bear (<i>Ursus maritimus</i>) Endogenous Retrovirus Identified from Next Generation Sequence Data. <i>Viruses</i> , 2015, 7, 6089-6107.	3.3	7
53	Endogenous murine leukemia retroviral variation across wild European and inbred strains of house mouse. <i>BMC Genomics</i> , 2015, 16, 613.	2.8	4
54	A protocol for CRISPR-mediated activation and repression of human endogenous retroviruses in human pluripotent stem cells. <i>STAR Protocols</i> , 2022, 3, 101281.	1.2	2

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55	Comparing PreXMRV-2 gag sequence diversity in laboratory and wild mice using deep sequencing. <i>Virus Research</i> , 2012, 169, 30-37.	2.2	1
56	Endogenous retroviruses. , 2005, , .		0