

Ute H Willhoeft

List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

2,874
citations

567281

15
h-index

713466

21
g-index

21
all docs

21
docs citations

21
times ranked

4838
citing authors

#	ARTICLE	IF	CITATIONS
1	LTRharvest, an efficient and flexible software for de novo detection of LTR retrotransposons. BMC Bioinformatics, 2008, 9, 18.	2.6	1,110
2	The genome of the protist parasite <i>Entamoeba histolytica</i> . Nature, 2005, 433, 865-868.	27.8	783
3	Fine-grained annotation and classification of de novo predicted LTR retrotransposons. Nucleic Acids Research, 2009, 37, 7002-7013.	14.5	276
4	Identification of the Sex-Determining Region of the <i>Ceratitis capitata</i> Y Chromosome by Deletion Mapping. Genetics, 1996, 144, 737-745.	2.9	106
5	Analysis of 5' junctions of human LINE-1 and Alu retrotransposons suggests an alternative model for 5'-end attachment requiring microhomology-mediated end-joining. Genome Research, 2005, 15, 780-789.	5.5	101
6	The electrophoretic karyotype of <i>Entamoeba histolytica</i> . Molecular and Biochemical Parasitology, 1999, 99, 41-53.	1.1	75
7	A jumping sex determining factor in the fly <i>Megaselia scalaris</i> . Chromosoma, 1990, 99, 407-412.	2.2	74
8	A DNA Sequence Corresponding to the Gene Encoding Cysteine Proteinase 5 in <i>Entamoeba histolytica</i> Is Present and Positionally Conserved but Highly Degenerated in <i>Entamoeba dispar</i> . Infection and Immunity, 1999, 67, 5925-5929.	2.2	58
9	Introns of <i>Entamoeba histolytica</i> and <i>Entamoeba dispar</i> . Protist, 2001, 152, 149-156.	1.5	40
10	The Abundant Polyadenylated Transcript 2 DNA Sequence of the Pathogenic Protozoan Parasite <i>Entamoeba histolytica</i> Represents a Nonautonomous Non-Long-Terminal-Repeat Retrotransposon-Like Element Which Is Absent in the Closely Related Nonpathogenic Species <i>Entamoeba dispar</i> . Infection and Immunity, 2002, 70, 6798-6804.	2.2	40
11	Characterization of Chitin Synthases from <i>Entamoeba</i> . Protist, 2004, 155, 323-330.	1.5	38
12	Molecular differentiation of the homomorphic sex chromosomes in <i>Megaselia scalaris</i> (Diptera) detected by random DNA probes. Chromosoma, 1990, 99, 237-242.	2.2	31
13	DNA sequences corresponding to the ariel gene family of <i>Entamoeba histolytica</i> are not present in <i>E. dispar</i> . Parasitology Research, 1999, 85, 787-789.	1.6	25
14	Fluorescence in situ hybridization of ribosomal DNA to mitotic chromosomes of tsetse flies (Diptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf	2.2	23
15	Analysis of cDNA Expressed Sequence Tags from <i>Entamoeba histolytica</i> : Identification of two Highly Abundant Polyadenylated Transcripts with no Overt Open Reading Frames. Protist, 1999, 150, 61-70.	1.5	23
16	Fluorescence microscopy and fluorescence in situ hybridization of <i>Entamoeba histolytica</i> nuclei to analyse mitosis and the localization of repetitive DNA. Molecular and Biochemical Parasitology, 2000, 105, 291-296.	1.1	17
17	Comparison of the mitotic karyotypes of <i>Ceratitis capitata</i> , <i>Ceratitis rosa</i> , and <i>Trirhithrum coffeae</i> (Diptera: Tephritidae) by C-banding and FISH. Genome, 1996, 39, 884-889.	2.0	16
18	The Hermes element from <i>Musca domestica</i> can transpose in four families of cyclorrhaphan flies. Genetica, 1997, 99, 15-29.	1.1	14

#	ARTICLE	IF	CITATIONS
19	Pig complement regulator factor H: molecular cloning and functional characterization. Immunogenetics, 2003, 55, 462-471.	2.4	11
20	Analysis of the sex chromosomes of the Mediterranean fruit fly by microdissected DNA probes. Genome, 1998, 41, 74-78.	2.0	9
21	Genetic Differences Between Entamoeba histolytica and Entamoeba dispar. Archives of Medical Research, 2000, 31, S254.	3.3	4