Ute H Willhoeft

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8766735/publications.pdf

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21 papers 2,874 citations

567281 15 h-index 713466 21 g-index

21 all docs

21 docs citations

times ranked

21

4838 citing authors

#	Article	IF	CITATIONS
1	Fine-grained annotation and classification of de novo predicted LTR retrotransposons. Nucleic Acids Research, 2009, 37, 7002-7013.	14.5	276
2	LTRharvest, an efficient and flexible software for de novo detection of LTR retrotransposons. BMC Bioinformatics, 2008, 9, 18.	2.6	1,110
3	The genome of the protist parasite Entamoeba histolytica. Nature, 2005, 433, 865-868.	27.8	783
4	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
5	Characterization of Chitin Synthases from Entamoeba. Protist, 2004, 155, 323-330.	1.5	38
6	Pig complement regulator factor H: molecular cloning and functional characterization. Immunogenetics, 2003, 55, 462-471.	2.4	11
7	The Abundant Polyadenylated Transcript 2 DNA Sequence of the Pathogenic Protozoan Parasite Entamoeba histolytica Represents a Nonautonomous Non-Long-Terminal-Repeat Retrotransposon- Like Element Which Is Absent in the Closely Related Nonpathogenic Species Entamoeba dispar. Infection and Immunity, 2002, 70, 6798-6804.	2.2	40
8	Introns of Entamoeba histolytica and Entamoeba dispar. Protist, 2001, 152, 149-156.	1.5	40
9	Genetic Differences Between Entamoeba histolytica and Entamoeba dispar. Archives of Medical Research, 2000, 31, S254.	3.3	4
10	Fluorescence microscopy and fluorescence in situ hybridization of Entamoeba histolytica nuclei to analyse mitosis and the localization of repetitive DNA. Molecular and Biochemical Parasitology, 2000, 105, 291-296.	1.1	17
11	DNA sequences corresponding to the ariel gene family of Entamoeba histolytica are not present in E . dispar. Parasitology Research, 1999, 85, 787-789.	1.6	25
12	The electrophoretic karyotype of Entamoeba histolytica. Molecular and Biochemical Parasitology, 1999, 99, 41-53.	1.1	75
13	Analysis of cDNA Expressed Sequence Tags from Entamoeba histolytica: Identification of two Highly Abundant Polyadenylated Transcripts with no Overt Open Reading Frames. Protist, 1999, 150, 61-70.	1.5	23
14	A DNA Sequence Corresponding to the Gene Encoding Cysteine Proteinase 5 in Entamoeba histolytica Is Present and Positionally Conserved but Highly Degenerated in Entamoeba dispar. Infection and Immunity, 1999, 67, 5925-5929.	2.2	58
15	Analysis of the sex chromosomes of the Mediterranean fruit fly by microdissected DNA probes. Genome, 1998, 41, 74-78.	2.0	9
16	The Hermes element from Musca domestica can transpose in four families of cyclorrhaphan flies. Genetica, 1997, 99, 15-29.	1.1	14
17	Fluorescence in situ hybridization of ribosomal DNA to mitotic chromosomes of tsetse flies (Diptera:) Tj ETQq1	1 0. <u>78</u> 431	4 rgBT /Ovedo
18	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

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#	Article	IF	CITATIONS
19	Identification of the Sex-Determining Region of the <i>Ceratitis capitata Y</i> Chromosome by Deletion Mapping. Genetics, 1996, 144, 737-745.	2.9	106
20	A jumping sex determining factor in the flyMegaselia scalaris. Chromosoma, 1990, 99, 407-412.	2.2	74
21	Molecular differentiation of the homomorphic sex chromosomes inMegaselia scalaris (Diptera) detected by random DNA probes. Chromosoma, 1990, 99, 237-242.	2.2	31