Jean-Mathieu Bart

List of Publications by Year in descending order

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47 1,778 papers citations

24 h-index 40 g-index

49 all docs 49 docs citations 49 times ranked 1759 citing authors

#	Article	IF	CITATIONS
1	Do Cryptic Reservoirs Threaten Gambiense-Sleeping Sickness Elimination?. Trends in Parasitology, 2018, 34, 197-207.	3.3	139
2	NUP-1 Is a Large Coiled-Coil Nucleoskeletal Protein in Trypanosomes with Lamin-Like Functions. PLoS Biology, 2012, 10, e1001287.	5.6	105
3	Genetic typing of Echinococcus granulosus in Romania. Parasitology Research, 2006, 98, 130-137.	1.6	86
4	Genotyping of human cystic echinococcosis in Xinjiang, PR China. Parasitology, 2006, 133, 571.	1.5	77
5	Genetic Diversity of the Cestode Echinococcus multilocularis in Red Foxes at a Continental Scale in Europe. PLoS Neglected Tropical Diseases, 2009, 3, e452.	3.0	74
6	Practical Approach for Typing Strains of Leishmania infantum by Microsatellite Analysis. Journal of Clinical Microbiology, 2002, 40, 3391-3397.	3.9	73
7	Cohesin regulates <i>VSG </i> monoallelic expression in trypanosomes. Journal of Cell Biology, 2009, 186, 243-254.	5.2	73
8	Third target of rapamycin complex negatively regulates development of quiescence in <i>Trypanosoma brucei</i> . Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 14399-14404.	7.1	70
9	Taxonomic position and geographical distribution of the common sheep G1 and camel G6 strains of Echinococcus granulosus in three African countries. Parasitology Research, 2006, 100, 495-503.	1.6	67
10	Cystic echinococcosis in Algeria: cattle act as reservoirs of a sheep strain and may contribute to human contamination. Veterinary Parasitology, 2003, 116, 35-44.	1.8	61
11	Assessment of Use of Microsatellite Polymorphism Analysis for Improving Spatial Distribution Tracking of <i>Echinococcus multilocularis</i> Journal of Clinical Microbiology, 2007, 45, 2943-2950.	3.9	60
12	EmsB, a tandem repeated multi-loci microsatellite, new tool to investigate the genetic diversity of Echinococcus multilocularis. Infection, Genetics and Evolution, 2006, 6, 390-400.	2.3	51
13	SUMOylation by the E3 Ligase TbSIZ1/PIAS1 Positively Regulates VSG Expression in Trypanosoma brucei. PLoS Pathogens, 2014, 10, e1004545.	4.7	48
14	Genetic diversity of Echinococcus multilocularis on a local scale. Infection, Genetics and Evolution, 2008, 8, 367-373.	2.3	46
15	Extravascular Dermal Trypanosomes in Suspected and Confirmed Cases of <i>gambiense</i> Human African Trypanosomiasis. Clinical Infectious Diseases, 2021, 73, 12-20.	5.8	46
16	Multi-locus microsatellite analysis supports the hypothesis of an autochthonous focus of Echinococcus multilocularis in northern Italy. International Journal for Parasitology, 2009, 39, 837-842.	3.1	44
17	The AMPKα1 Pathway Positively Regulates the Developmental Transition from Proliferation to Quiescence in Trypanosoma brucei. Cell Reports, 2016, 17, 660-670.	6.4	44
18	Using mitochondrial and nuclear markers to evaluate the degree of genetic cohesion among Echinococcus populations. Experimental Parasitology, 2008, 119, 453-459.	1.2	43

#	Article	IF	Citations
19	Echinococcus multilocularis in Svalbard, Norway: Microsatellite genotyping to investigate the origin of a highly focal contamination. Infection, Genetics and Evolution, 2012, 12, 1270-1274.	2.3	41
20	Echinococcus granulosus strain typing in North Africa: comparison of eight nuclear and mitochondrial DNA fragments. Parasitology, 2004, 128, 229-234.	1.5	39
21	Morphological and molecular characteristics of Echinococcus multilocularis and Echinococcus granulosus mixed infection in a dog from Xinjiang, China. Veterinary Parasitology, 2006, 139, 244-248.	1.8	31
22	Multiple-Locus Variable-Number Tandem-Repeat Analysis for Rapid Typing of Candida glabrata. Journal of Clinical Microbiology, 2007, 45, 3781-3784.	3.9	28
23	Assessment of Echinococcus granulosus polymorphism in Qinghai Province, People's Republic of China. Parasitology Research, 2008, 102, 1201-1206.	1.6	28
24	Trypa-NO! contributes to the elimination of gambiense human African trypanosomiasis by combining tsetse control with "screen, diagnose and treat―using innovative tools and strategies. PLoS Neglected Tropical Diseases, 2020, 14, e0008738.	3.0	28
25	Autophagic-related cell death of Trypanosoma brucei induced by bacteriocin AS-48. International Journal for Parasitology: Drugs and Drug Resistance, 2018, 8, 203-212.	3.4	27
26	Target of rapamycin (TOR) kinase in <i>Trypanosoma brucei</i> : an extended family. Biochemical Society Transactions, 2013, 41, 934-938.	3.4	26
27	Comparison of several commercial serologic kits and Em18 serology for detection of human alveolar echinococcosis. Diagnostic Microbiology and Infectious Disease, 2007, 59, 93-95.	1.8	25
28	The genomic Echinococcus microsatellite EmsB sequences: from a molecular marker to the epidemiological tool. Parasitology, 2010, 137, 439-449.	1.5	24
29	Defeating Leishmania resistance to Miltefosine (hexadecylphosphocholine) by peptide-mediated drug smuggling: A proof of mechanism for trypanosomatid chemotherapy. Journal of Controlled Release, 2012, 161, 835-842.	9.9	24
30	Co-dependence between trypanosome nuclear lamina components in nuclear stability and control of gene expression. Nucleic Acids Research, 2016, 44, 10554-10570.	14.5	23
31	A receptor for the complement regulator factor H increases transmission of trypanosomes to tsetse flies. Nature Communications, 2020, 11, 1326.	12.8	23
32	<i>Echinococcus vogeli</i> Infection in a Hunter, French Guiana. Emerging Infectious Diseases, 2009, 15, 2029-2031.	4.3	22
33	Detection of Echinococcus multilocularis in wild boars in France using PCR techniques against larval form. Veterinary Parasitology, 2005, 129, 259-266.	1.8	21
34	Molecular characterization of a novel gene encoding an 8-kDa-subunit of antigen B from Echinococcus granulosus genotypes 1 and 6. Parasitology International, 2007, 56, 313-316.	1.3	21
35	Increased uracil insertion in DNA is cytotoxic and increases the frequency of mutation, double strand break formation and VSG switching in Trypanosoma brucei. DNA Repair, 2012, 11, 986-995.	2.8	21
36	Molecular evidence of a Trypanosoma brucei gambiense sylvatic cycle in the human african trypanosomiasis foci of Equatorial Guinea. Frontiers in Microbiology, 2015, 6, 765.	3.5	20

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37	Genotyping Echinococcus multilocularis in Human Alveolar Echinococcosis Patients: An EmsB Microsatellite Analysis. Pathogens, 2020, 9, 282.	2.8	17
38	Localization of serum resistance-associated protein in <i>Trypanosoma brucei rhodesiense</i> and transgenic <i>Trypanosoma brucei brucei</i> . Cellular Microbiology, 2015, 17, 1523-1535.	2.1	13
39	Practices in research, surveillance and control of neglected tropical diseases by One Health approaches: A survey targeting scientists from French-speaking countries. PLoS Neglected Tropical Diseases, 2021, 15, e0009246.	3.0	13
40	Free-ranging pigs identified as a multi-reservoir of Trypanosoma brucei and Trypanosoma congolense in the Vavoua area, a historical sleeping sickness focus of Cà te d'Ivoire. PLoS Neglected Tropical Diseases, 2021, 15, e0010036.	3.0	11
41	Glossina palpalis palpalis populations from Equatorial Guinea belong to distinct allopatric clades. Parasites and Vectors, 2014, 7, 31.	2.5	10
42	Seroprevalence of brucellosis, Q fever and Rift Valley fever in domestic ruminants in Guinea in 2017–2019. BMC Veterinary Research, 2022, 18, 64.	1.9	9
43	Trypanosoma brucei gambiense Adaptation to Different Mammalian Sera Is Associated with VSG Expression Site Plasticity. PLoS ONE, 2013, 8, e85072.	2.5	8
44	Metabarcoding: A Powerful Yet Still Underestimated Approach for the Comprehensive Study of Vector-Borne Pathogen Transmission Cycles and Their Dynamics. , 0, , .		7
45	Involvement in surface antigen expression by a moonlighting FG-repeat nucleoporin in trypanosomes. Molecular Biology of the Cell, 2018, 29, 1100-1110.	2.1	5
46	SARS-CoV-2 Circulation, Guinea, March 2020–July 2021. Emerging Infectious Diseases, 2022, 28, 457-460.	4.3	4
47	Cohesin regulates <i>VSG </i> monoallelic expression in trypanosomes. Journal of Experimental Medicine, 2009, 206, i17-i17.	8.5	0