Fabio Tosini

List of Publications by Year in descending order

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471509 454955 1,146 32 17 30 citations h-index g-index papers 33 33 33 1304 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Prevalence and genetic characterization of Dientamoeba fragilis in asymptomatic children attending daycare centers. Revista Do Instituto De Medicina Tropical De Sao Paulo, 2021, 63, e39.	1.1	4
2	Isolation and Characterization of Mouse Monoclonal Antibodies That Neutralize SARS-CoV-2 and Its Variants of Concern Alpha, Beta, Gamma and Delta by Binding Conformational Epitopes of Glycosylated RBD With High Potency. Frontiers in Immunology, 2021, 12, 750386.	4.8	6
3	Delivery of SA35 and SA40 peptides in mice enhances humoral and cellular immune responses and confers protection against Cryptosporidium parvum infection. Parasites and Vectors, 2019, 12, 233.	2.5	6
4	Multilocus sequence typing of Dientamoeba fragilis identified a major clone with widespread geographical distribution. International Journal for Parasitology, 2016, 46, 793-798.	3.1	16
5	Molecular characterization of intestinal protozoa in two poor communities in the State of São Paulo, Brazil. Parasites and Vectors, 2015, 8, 103.	2.5	77
6	A real-time assemblage-specific PCR assay for the detection of Giardia duodenalis assemblages A, B and E in fecal samples. Veterinary Parasitology, 2015, 211, 28-34.	1.8	17
7	Myiasis of the Tracheostomy Wound Caused by <i>Sarcophaga </i> (i>Liopygia) <i>argyrostoma </i> (Diptera: Sarcophagidae): Molecular Identification Based on the Mitochondrial Cytochrome c Oxidase I Gene: Fig. 1 Journal of Medical Entomology. 2015. 52. 1357-1360.	1.8	14
8	P.03.11 MOLECULAR DIAGNOSIS OF INTESTINAL PROTOZOA IN PATIENTS WITH IRRITABLE BOWEL SYNDROME (IBS). Digestive and Liver Disease, 2014, 46, S63-S64.	0.9	0
9	A rare Cryptosporidium parvum genotype associated with infection of lambs and zoonotic transmission in Italy. Veterinary Parasitology, 2013, 191, 128-131.	1.8	40
10	Detection of Giardia duodenalis Assemblages A and B in Human Feces by Simple, Assemblage-Specific PCR Assays. PLoS Neglected Tropical Diseases, 2012, 6, e1776.	3.0	38
11	Pigs as Natural Hosts of <i>Dientamoeba fragilis </i> Cenotypes Found in Humans. Emerging Infectious Diseases, 2012, 18, 838-41.	4.3	40
12	Expression of Cryptosporidium parvum Cpa135/CpCCP1 chimeras in Giardia duodenalis: Organization of the protein domains affects the protein secretion pathway. Experimental Parasitology, 2011, 127, 680-686.	1.2	1
13	The CpA135 gene as a marker to identify Cryptosporidium species infecting humans. Parasitology International, 2010, 59, 606-609.	1.3	13
14	Biochemical characterization of MLC1 protein in astrocytes and its association with the dystrophin–glycoprotein complex. Molecular and Cellular Neurosciences, 2008, 37, 480-493.	2.2	38
15	Proteomics Analysis and Protein Expression during Sporozoite Excystation of Cryptosporidium parvum (Coccidia, Apicomplexa). Molecular and Cellular Proteomics, 2007, 6, 346-355.	3.8	68
16	The immunological selection of recombinant peptides from Cryptosporidium parvum reveals 14 proteins expressed at the sporozoite stage, 7 of which are conserved in other apicomplexaâ ⁻ †. Molecular and Biochemical Parasitology, 2007, 152, 159-169.	1.1	14
17	Cryptosporidium parvum -Specific CD4 Th1 Cells from Sensitized Donors Responding to Both Fractionated and Recombinant Antigenic Proteins. Infection and Immunity, 2004, 72, 1306-1310.	2.2	30
18	Cryptosporidium parvum at Different Developmental Stages Modulates Host Cell Apoptosis In Vitro. Infection and Immunity, 2004, 72, 6061-6067.	2.2	88

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19	A new modular protein of Cryptosporidium parvum, with ricin B and LCCL domains, expressed in the sporozoite invasive stage. Molecular and Biochemical Parasitology, 2004, 134, 137-147.	1.1	51
20	Indinavir reduces Cryptosporidium parvum infection in both in vitro and in vivo models. International Journal for Parasitology, 2003, 33, 757-764.	3.1	53
21	Identification and Characterisation of the Antigenic Cpa135 protein., 2003,, 87-88.		0
22	Detection and counting of <i>Cryptosporidium parvum </i> ii HCT-8 cells by flowcytometry. Parasite, 2003, 10, 297-302.	2.0	7
23	Composite Integron Array Generated by Insertion of an ORF341-Type Integron Within a Tn21-like Element. Microbial Drug Resistance, 2002, 8, 1-8.	2.0	31
24	Characterization of Plasmids Carrying CMY-2 from Expanded-Spectrum Cephalosporin-Resistant Salmonella Strains Isolated in the United States between 1996 and 1998. Antimicrobial Agents and Chemotherapy, 2002, 46, 1269-1272.	3.2	139
25	Integration of Integrons in res Sites. Antimicrobial Agents and Chemotherapy, 2002, 46, 2058-2058.	3.2	1
26	Multiple-Antibiotic Resistance Mediated by Structurally Related IncL/M Plasmids Carrying an Extended-Spectrum β-Lactamase Gene and a Class 1 Integron. Antimicrobial Agents and Chemotherapy, 2000, 44, 2911-2914.	3.2	87
27	Antibiotic Resistance Conferred by a Conjugative Plasmid and a Class I Integron in <i>Vibrio cholerae</i> O1 El Tor Strains Isolated in Albania and Italy. Antimicrobial Agents and Chemotherapy, 1999, 43, 693-696.	3.2	96
28	identification and characterisation of three antigenic proteins from Cryptosporidium parvum sporozoites using a DNA library expressing poly-histidine tagged peptides1Note: Nucleotide sequences reported in this paper are available in the GenBankâ,,¢, EMBL and DDBJ databases under the accession numbers AJ006592 (sa20), AJ006593 (sa35), and AJ132769 (sa40).1. International Journal for Parasitology,	3.1	19
29	1999, 29, 1925-1933. The uvp1 gene on the R46 plasmid encodes a resolvase that catalyzes site-specific resolution involving the 5′-conserved segment of the adjacent integron In1. Molecular Genetics and Genomics, 1998, 258, 404-411.	2.4	13
30	Multidrug-ResistantSalmonella entericaSerotype Typhimurium Infections. New England Journal of Medicine, 1998, 339, 921-922.	27.0	7
31	Class 1 Integron-Borne Multiple-Antibiotic Resistance Carried by IncFI and IncL/M Plasmids in <1>Salmonella enterica 1 Serotype Typhimurium. Antimicrobial Agents and Chemotherapy, 1998, 42, 3053-3058.	3.2	129
32	The gene coding for proteins HC and HI-30 of inter-alpha-trypsin inhibitor maps to 9q22.3â†'q33. Cytogenetic and Genome Research, 1989, 50, 46-48.	1.1	3