Esther Orozco

List of Publications by Year in descending order

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218677 2,349 98 26 h-index citations papers

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#	Article	IF	CITATIONS
1	EhVps23, an ESCRT-I Member, Is a Key Factor in Secretion, Motility, Phagocytosis and Tissue Invasion by Entamoeba histolytica. Frontiers in Cellular and Infection Microbiology, 2022, 12, 835654.	3.9	4
2	The atypical protein arginine methyltrasferase of Entamoeba histolytica (EhPRMTA) is involved in cell proliferation, heat shock response and in vitro virulence. Experimental Parasitology, 2021, 222, 108077.	1.2	4
3	Protein Sumoylation Is Crucial for Phagocytosis in Entamoeba histolytica Trophozoites. International Journal of Molecular Sciences, 2021, 22, 5709.	4.1	3
4	EhVps23: A Component of ESCRT-I That Participates in Vesicular Trafficking and Phagocytosis of Entamoeba histolytica. Frontiers in Cellular and Infection Microbiology, 2021, 11, 770759.	3.9	4
5	A noncanonical GATA transcription factor of Entamoeba histolytica modulates genes involved in phagocytosis. Molecular Microbiology, 2020, 114, 1019-1037.	2.5	2
6	Lipids in Entamoeba histolytica: Host-Dependence and Virulence Factors. Frontiers in Cellular and Infection Microbiology, 2020, 10, 75.	3.9	18
7	Host Invasion by Pathogenic Amoebae: Epithelial Disruption by Parasite Proteins. Genes, 2019, 10, 618.	2.4	40
8	EhRabB mobilises the EhCPADH complex through the actin cytoskeleton during phagocytosis of <i>Entamoeba histolytica</i> . Cellular Microbiology, 2019, 21, e13071.	2.1	10
9	Identification of the gene encoding the TATA box-binding protein-associated factor 1 (TAF1) and its putative role in the heat shock response in the protozoan parasite Entamoeba histolytica. Parasitology Research, 2019, 118, 517-538.	1.6	6
10	Telomeric Repeat-Binding Factor Homologs in Entamoeba histolytica: New Clues for Telomeric Research. Frontiers in Cellular and Infection Microbiology, 2018, 8, 341.	3.9	4
11	Epithelial Cells Expressing EhADH, An Entamoeba histolytica Adhesin, Exhibit Increased Tight Junction Proteins. Frontiers in Cellular and Infection Microbiology, 2018, 8, 340.	3.9	9
12	A Calcium/Cation Exchanger Participates in the Programmed Cell Death and in vitro Virulence of Entamoeba histolytica. Frontiers in Cellular and Infection Microbiology, 2018, 8, 342.	3.9	7
13	The Sole DNA Ligase in Entamoeba histolytica Is a High-Fidelity DNA Ligase Involved in DNA Damage Repair. Frontiers in Cellular and Infection Microbiology, 2018, 8, 214.	3.9	8
14	The Conserved ESCRT-III Machinery Participates in the Phagocytosis of Entamoeba histolytica. Frontiers in Cellular and Infection Microbiology, 2018, 8, 53.	3.9	40
15	The Entamoeba histolytica TBP and TRF1 transcription factors are GAAC-box binding proteins, which display differential gene expression under different stress stimuli and during the interaction with mammalian cells. Parasites and Vectors, 2018, 11, 153.	2.5	8
16	Heterodimerization of the <i>Entamoeba histolytica</i> EhCPADH virulence complex through molecular dynamics and protein–protein docking. Journal of Biomolecular Structure and Dynamics, 2017, 35, 486-503.	3.5	14
17	Adherens junctions and desmosomes are damaged by <i>Entamoeba histolytica<i><i><i>><i i="">>Participation of EhCPADH complex and EhCP112 protease. Cellular Microbiology, 2017, 19, e12761.</i></i></i></i></i>	2.1	15
18	The Tudor Staphylococcal Nuclease Protein of Entamoeba histolytica Participates in Transcription Regulation and Stress Response. Frontiers in Cellular and Infection Microbiology, 2017, 7, 52.	3.9	7

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19	Entamoeba histolytica EhCP112 Dislocates and Degrades Claudin-1 and Claudin-2 at Tight Junctions of the Intestinal Epithelium. Frontiers in Cellular and Infection Microbiology, 2017, 7, 372.	3.9	30
20	A Novel Heat Shock Element (HSE) in Entamoeba histolytica that Regulates the Transcriptional Activation of the EhPgp5 Gene in the Presence of Emetine Drug. Frontiers in Cellular and Infection Microbiology, 2017, 7, 492.	3.9	5
21	Resveratrol Induces Apoptosis-Like Death and Prevents In Vitro and In Vivo Virulence of Entamoeba histolytica. PLoS ONE, 2016, 11, e0146287.	2.5	25
22	EhNPC1 and EhNPC2 Proteins Participate in Trafficking of Exogenous Cholesterol in Entamoeba histolytica Trophozoites: Relevance for Phagocytosis. PLoS Pathogens, 2016, 12, e1006089.	4.7	24
23	Identification and functional characterization of lysine methyltransferases of <i>Entamoeba histolytica</i> . Molecular Microbiology, 2016, 101, 351-365.	2.5	6
24	Data on docking and dynamics simulation of Entamoeba histolytica EhADH (an ALIX protein) and lysobisphosphatidic acid. Data in Brief, 2016, 7, 457-459.	1.0	6
25	The knockdown of each component of the cysteine proteinase-adhesin complex of (i>Entamoeba histolytica (i) (EhCPADH) affects the expression of the other complex element as well as the (i>in vitro (i>and (i>in vivo (i) virulence. Parasitology, 2016, 143, 50-59.	1.5	10
26	Identification of the phospholipid lysobisphosphatidic acid in the protozoan Entamoeba histolytica: An active molecule in endocytosis. Biochemistry and Biophysics Reports, 2016, 5, 224-236.	1.3	16
27	EhVps32 Is a Vacuole-Associated Protein Involved in Pinocytosis and Phagocytosis of Entamoeaba histolytica. PLoS Pathogens, 2015, 11, e1005079.	4.7	32
28	A recombinant influenza virus vaccine expressing the F protein of respiratory syncytial virus. Archives of Virology, 2014, 159, 1067-1077.	2.1	12
29	Effect of the silencing of the Ehcp112 gene on the in vitro virulence of Entamoeba histolytica. Parasites and Vectors, 2013, 6, 248.	2.5	16
30	The small GTPase EhRabB of Entamoeba histolytica is differentially expressed during phagocytosis. Parasitology Research, 2013, 112, 1631-1640.	1.6	11
31	The EhCPADH112 Complex of Entamoeba histolytica Interacts with Tight Junction Proteins Occludin and Claudin-1 to Produce Epithelial Damage. PLoS ONE, 2013, 8, e65100.	2.5	43
32	The 25 kDa Subunit of Cleavage Factor Im Is a RNA-Binding Protein That Interacts with the Poly(A) Polymerase in Entamoeba histolytica. PLoS ONE, 2013, 8, e67977.	2.5	15
33	EhADH112 is a Bro1 Domain-Containing Protein Involved in the <i>Entamoeba histolytica </i> Multivesicular Bodies Pathway. Journal of Biomedicine and Biotechnology, 2012, 2012, 1-15.	3.0	26
34	A high-throughput drug screen for Entamoeba histolytica identifies a new lead and target. Nature Medicine, 2012, 18, 956-960.	30.7	290
35	mRNA Decay Proteins Are Targeted to poly(A)+ RNA and dsRNA-Containing Cytoplasmic Foci That Resemble P-Bodies in Entamoeba histolytica. PLoS ONE, 2012, 7, e45966.	2.5	17
36	Identification of a polypeptide containing Tudor and staphyloccocal nuclease-like domains as the sequence-specific binding protein to the upstream regulatory element 1 of Entamoeba histolytica. International Journal for Parasitology, 2011, 41, 775-782.	3.1	9

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37	Detection of the Endosomal Sorting Complex Required for Transport inEntamoeba histolyticaand Characterization of the EhVps4 Protein. Journal of Biomedicine and Biotechnology, 2010, 2010, 1-15.	3.0	23
38	The R2R3 Myb protein family in Entamoeba histolytica. Gene, 2010, 455, 32-42.	2.2	38
39	Entamoeba histolytica: A unicellular organism containing two active genes encoding for members of the TBP family. Protein Expression and Purification, 2010, 70, 48-59.	1.3	12
40	DNA repair mechanisms in eukaryotes: Special focus in Entamoeba histolytica and related protozoan parasites. Infection, Genetics and Evolution, 2009, 9, 1051-1056.	2.3	28
41	A pcDNA-Ehcpadh vaccine against Entamoeba histolytica elicits a protective Th1-like response in hamster liver. Vaccine, 2009, 27, 4176-4186.	3.8	19
42	Purification, refolding and autoactivation of the recombinant cysteine proteinase EhCP112 from Entamoeba histolytica. Protein Expression and Purification, 2009, 63, 26-32.	1.3	24
43	Drug Resistance Mechanisms in Entamoeba histolytica, Giardia lamblia, Trichomonas vaginalis, and Opportunistic Anaerobic Protozoa. , 2009, , 549-559.		7
44	In silico analysis of EST and genomic sequences allowed the prediction of cis-regulatory elements for Entamoeba histolytica mRNA polyadenylation. Computational Biology and Chemistry, 2008, 32, 256-263.	2.3	24
45	Entamoeba histolytica EhDEAD1 is a conserved DEAD-box RNA helicase with ATPase and ATP-dependent RNA unwinding activities. Gene, 2008, 414, 19-31.	2.2	13
46	Putative DEAD and DExH-box RNA helicases families in Entamoeba histolytica. Gene, 2008, 424, 1-10.	2.2	16
47	Structural and functional analysis of the Entamoeba histolytica EhrabB gene promoter. BMC Molecular Biology, 2007, 8, 82.	3.0	21
48	A phagocytosis mutant of Entamoeba histolytica is less virulent due to deficient proteinase expression and release. Experimental Parasitology, 2007, 115, 192-199.	1.2	47
49	Molecular characterization of the Entamoeba histolytica enolase gene and modelling of the predicted protein. FEMS Microbiology Letters, 2006, 148, 123-129.	1.8	13
50	Identification and functional characterization of EhClC-A, an Entamoeba histolytica ClC chloride channel located at plasma membrane. Molecular Microbiology, 2006, 59, 1249-1261.	2.5	17
51	EhCP112 is an Entamoeba histolytica secreted cysteine protease that may be involved in the parasite-virulence. Cellular Microbiology, 2005, 7, 221-232.	2.1	53
52	Entamoeba histolytica TATA-box binding protein binds to different TATA variants in vitro. FEBS Journal, 2005, 272, 1354-1366.	4.7	18
53	Entamoeba histolytica: Identification of EhGPCR-1, a novel putative G protein-coupled receptor that binds to EhRabB. Experimental Parasitology, 2005, 110, 253-258.	1.2	17
54	Entamoeba histolytica: Cloning and expression of the poly(A) polymerase EhPAP. Experimental Parasitology, 2005, 110, 226-232.	1.2	12

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55	Entamoeba histolytica: Comparative genomics of the pre-mRNA $3\hat{a}\in^2$ end processing machinery. Experimental Parasitology, 2005, 110, 184-190.	1.2	27
56	Entamoeba histolytica: Alterations in EhRabB protein in a phagocytosis deficient mutant correlate with the Entamoeba dispar RabB sequence. Experimental Parasitology, 2005, 110, 259-264.	1.2	10
57	Entamoeba histolytica EhPgp5 transcriptional activation depends on putative emetine response elements. Experimental Parasitology, 2005, 110, 233-237.	1.2	8
58	Expression in fibroblasts and in live animals of Entamoeba histolytica polypeptides EhCP112 and EhADH112. Microbiology (United Kingdom), 2004, 150, 1251-1260.	1.8	23
59	The EhADH112 recombinant polypeptide inhibits cell destruction and liver abscess formation by Entamoeba histolytica trophozoites. Cellular Microbiology, 2004, 6, 367-376.	2.1	35
60	Entamoeba histolytica: expression and DNA binding of CCAAT/enhancer-binding proteins are regulated through the cell cycle. Experimental Parasitology, 2003, 103, 82-87.	1.2	8
61	EhPgp5 mRNA Stability Is a Regulatory Event in theEntamoeba histolytica Multidrug Resistance Phenotype. Journal of Biological Chemistry, 2003, 278, 11273-11280.	3.4	45
62	TheEntamoeba histolyticaEhPgp5 (MDR-like) Protein Induces Swelling of the Trophozoites and Alters Chloride-Dependent Currents inXenopus laevisOocytes. Microbial Drug Resistance, 2002, 8, 15-26.	2.0	13
63	Cellular Location and Function of the P-Glycoproteins (EhPgps) inEntamoeba histolyticaMultidrug-Resistant Trophozoites. Microbial Drug Resistance, 2002, 8, 291-300.	2.0	17
64	Two CCAAT/enhancer binding protein sites arecis-activator elements of theEntamoeba histolytica EhPgp1 (mdr-like) gene expression. Cellular Microbiology, 2002, 4, 725-737.	2.1	35
65	Specific detection of Entamoeba histolytica DNA by hemolysin gene targeted PCR. Acta Tropica, 2001, 78, 117-125.	2.0	20
66	Centromeric Structure Identification in Entamoeba histolytica by Anticentromeric/Kinetochore Antibodies Obtained from Patients with the CREST Syndrome. Archives of Medical Research, 2000, 31, S207-S209.	3.3	1
67	DNA Binding Activity and Predicted Tertiary Structure of the TATA Binding Protein of Entamoeba histolytica. Archives of Medical Research, 2000, 31, S299-S300.	3.3	1
68	Presence of Sequences Homologous to the Universal Minicircle Sequence (UMS) of Trypanosomatids in the DNA Circles of Entamoeba histolytica. Archives of Medical Research, 2000, 31, S296-S298.	3.3	0
69	An Initial Characterization of the 3′ Untranslated Region of the EhPgp5 mRNA in Entamoeba histolytica. Archives of Medical Research, 2000, 31, S282-S284.	3.3	3
70	Expression of Entamoeba histolytica EhPgp5 Gene in Xenopus laevis Oocytes. Archives of Medical Research, 2000, 31, S285-S287.	3.3	2
71	Algorithm to Predict MiniCHEF Electrophoresis Patterns of Entamoeba histolytica DNA. Archives of Medical Research, 2000, 31, S279-S281.	3.3	0
72	Expression and Immunodetection of a P-Glycoprotein in Emetine-Resistant Trophozoites of Entamoeba histolytica. Archives of Medical Research, 2000, 31, S288-S290.	3.3	1

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73	L6. Archives of Medical Research, 2000, 31, S237-S238.	3.3	6
74	An Entamoeba histolytica Rab-like encoding gene and protein: function and cellular location. Molecular and Biochemical Parasitology, 2000, 108, 199-206.	1.1	45
75	Entamoeba histolytica: Signaling through G Proteins. Experimental Parasitology, 1999, 91, 170-175.	1.2	12
76	Physiology and molecular genetics of multidrug resistance in Entamoeba histolytica. Drug Resistance Updates, 1999, 2, 188-197.	14.4	11
77	Chromatin Organization during the Nuclear Division Stages of LiveEntamoeba histolyticaTrophozoites. Experimental Parasitology, 1998, 89, 122-124.	1.2	10
78	The pyruvate:ferredoxin oxidoreductase enzyme is located in the plasma membrane and in a cytoplasmic structure inEntamoeba. Microbial Pathogenesis, 1998, 25, 1-10.	2.9	38
79	Transcriptional Analysis of the EhPgp1 Promoter of Entamoeba histolytica Multidrug-resistant Mutant. Journal of Biological Chemistry, 1998, 273, 7277-7284.	3.4	45
80	Transcriptional Analysis of the EhPgp5 Promoter of Entamoeba histolytica Multidrug-resistant Mutant. Journal of Biological Chemistry, 1998, 273, 7285-7292.	3.4	40
81	Cloning and characterization of the Entamoeba histolytica pyruvate: ferredoxin oxidoreductase gene. Molecular and Biochemical Parasitology, 1996, 78, 273-277.	1.1	24
82	Increase in mRNA of multiple Eh pgp genes encoding P-glycoprotein homologues in emetine-resistant Entamoeba histolytica parasites. Gene, 1995, 164, 179-184.	2.2	47
83	Invasion and metastasis mechanisms in Entamoeba histolytica and cancer cells. Some common cellular and molecular features. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 1994, 305, 229-239.	1.0	13
84	Cloning, genomic organization and transcription of the Entamoeba histolytica \hat{l} ±-tubulin-encodmg gene. Gene, 1994, 146, 239-244.	2.2	38
85	A variable DNA region of Entamoeba histolytica is expressed in several transcripts which differ in genetically related clones. Molecular Genetics and Genomics, 1993, 241-241, 271-279.	2.4	5
86	Molecular karyotype of related clones of Entamoeba histolytica. Molecular and Biochemical Parasitology, 1993, 59, 29-40.	1.1	19
87	Entamoeba histolytica: Generation and characterization of hybrid clones. Experimental Parasitology, 1991, 72, 236-242.	1.2	2
88	Non-pathogenic Entamoeba histolytica: functional and biochemical characterization of a monoxenic strain. Molecular and Biochemical Parasitology, 1990, 40, 193-201.	1.1	12
89	Entamoeba histolytica: Physiology of multidrug resistance. Experimental Parasitology, 1990, 71, 169-175.	1.2	38
90	Entamoeba histolytica: Correlation of the cytopathic effect of virulent trophozoites with secretion of a cysteine proteinase. Experimental Parasitology, 1990, 71, 199-206.	1.2	114

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91	Entamoeba histolytica: molecules involved in the target cell-parasite relationship. Molecular and Biochemical Parasitology, 1989, 37, 87-99.	1.1	37
92	Entamoeba histolytica: Cell cycle and nuclear division. Experimental Parasitology, 1988, 67, 85-95.	1.2	61
93	Entamoeba histolytica: Cytopathogenicity and lectin activity of avirulent mutants. Experimental Parasitology, 1987, 63, 157-165.	1.2	19
94	Localization and identification of an Entamoeba histolytica adhesin. Molecular and Biochemical Parasitology, 1987, 23, 151-158.	1.1	90
95	Structural Bases of the Cytolytic Mechanisms of Entamoeba histolytica1. Journal of Protozoology, 1985, 32, 166-175.	0.8	84
96	Isolation and characterization of Entamoeba histolytica mutants resistant to emetine. Molecular and Biochemical Parasitology, 1985, 15, 49-59.	1.1	55
97	Differences in adhesion, phagocytosis and virulence of clones from Entamoeba histolytica, strain HM1: IMSS. International Journal for Parasitology, 1985, 15, 655-660.	3.1	34
98	A Bioinformatical Approach to Study the Endosomal Sorting Complex Required for Transport (ESCRT) Machinery in Protozoan Parasites: The Entamoeba histolytica Case., 0,,.		2