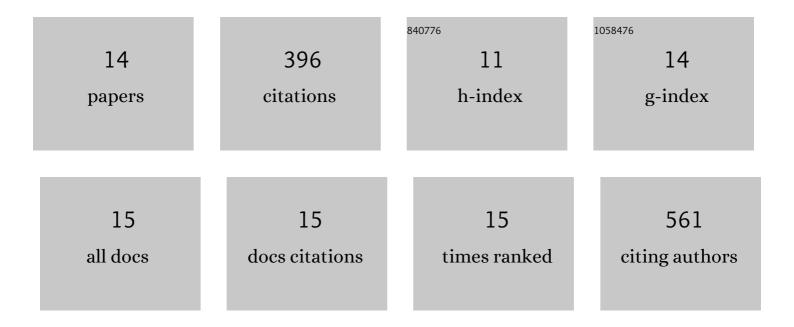
## Elena Jiménez-Ruiz

List of Publications by Year in descending order

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FLENA LIMÃONEZ-PULZ

#	Article	IF	CITATIONS
1	A splitCas9 phenotypic screen in Toxoplasma gondii identifies proteins involved in host cell egress and invasion. Nature Microbiology, 2022, 7, 882-895.	13.3	18
2	An endocytic-secretory cycle participates in Toxoplasma gondii in motility. PLoS Biology, 2019, 17, e3000060.	5.6	49
3	Characterization of the <i>Neospora caninum</i> NcROP40 and NcROP2Fam-1 rhoptry proteins during the tachyzoite lytic cycle. Parasitology, 2016, 143, 97-113.	1.5	12
4	UAP56 is a conserved crucial component of a divergent mRNA export pathway in <i>Toxoplasma gondii</i> . Molecular Microbiology, 2016, 102, 672-689.	2.5	22
5	Vacuolar protein sorting mechanisms in apicomplexan parasites. Molecular and Biochemical Parasitology, 2016, 209, 18-25.	1.1	33
6	Gliding Associated Proteins Play Essential Roles during the Formation of the Inner Membrane Complex of Toxoplasma gondii. PLoS Pathogens, 2016, 12, e1005403.	4.7	59
7	Conditional U1 Gene Silencing in Toxoplasma gondii. PLoS ONE, 2015, 10, e0130356.	2.5	44
8	Advantages and disadvantages of conditional systems for characterization of essential genes in <i>Toxoplasma gondii</i> . Parasitology, 2014, 141, 1390-1398.	1.5	20
9	Clinical outcome and vertical transmission variability among canine <i>Neospora caninum</i> isolates in a pregnant mouse model of infection. Parasitology, 2014, 141, 356-366.	1.5	22
10	Mice congenitally infected with low-to-moderate virulence Neospora caninum isolates exhibited clinical reactivation during the mating period without transmission to the next generation. Experimental Parasitology, 2013, 134, 244-248.	1.2	9
11	Specific antibody responses against Neospora caninum recombinant rNcGRA7, rNcSAG4, rNcBSR4 and rNcSRS9 proteins are correlated with virulence in mice. Parasitology, 2013, 140, 569-579.	1.5	10
12	Low efficacy of NcGRA7, NcSAG4, NcBSR4 and NcSRS9 formulated in poly-É›-caprolactone against Neospora caninum infection in mice. Vaccine, 2012, 30, 4983-4992.	3.8	22
13	Identification of a gene cluster for cell-surface genes of the SRS superfamily in <i>Neospora caninum</i> and characterization of the novel <i>SRS9</i> gene. Parasitology, 2011, 138, 1832-1842.	1.5	13
14	Influence of <i>Neospora caninum</i> intra-specific variability in the outcome of infection in a pregnant BALB/c mouse model. Veterinary Research, 2010, 41, 52.	3.0	63