## Manuel-Alfonso Patarroyo

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Cervical cancer screening programme attendance and compliance predictors regarding Colombia's Amazon region. PLoS ONE, 2022, 17, e0262069.	1.1	0
2	Comparing Class II MHC DRB3 Diversity in Colombian Simmental and Simbrah Cattle Across Worldwide Bovine Populations. Frontiers in Genetics, 2022, 13, 772885.	1.1	2
3	The prevalence of Chlamydia psittaci in confiscated Psittacidae in Colombia. Preventive Veterinary Medicine, 2022, 200, 105591.	0.7	5
4	The Cellular and Molecular Interaction Between Erythrocytes and Plasmodium falciparum Merozoites. Frontiers in Cellular and Infection Microbiology, 2022, 12, 816574.	1.8	6
5	Diagnostic Potential of the Serological Response to Synthetic Peptides from Mycobacterium tuberculosis Antigens for Discrimination Between Active and Latent Tuberculosis Infections. International Journal of Peptide Research and Therapeutics, 2022, 28, 98.	0.9	2
6	SM-COLSARSPROT: Highly Immunogenic Supramutational Synthetic Peptides Covering the World's Population. Frontiers in Immunology, 2022, 13, .	2.2	0
7	A novel platform for peptide-mediated affinity capture and LC-MS/MS identification of host receptors involved in Plasmodium invasion. Journal of Proteomics, 2021, 231, 104002.	1.2	7
8	The molecular basis for peptide-based antimalarial vaccine development targeting erythrocyte invasion by P.Âfalciparum. Biochemical and Biophysical Research Communications, 2021, 534, 86-93.	1.0	3
9	Babesia Bovis Ligand-Receptor Interaction: AMA-1 Contains Small Regions Governing Bovine Erythrocyte Binding. International Journal of Molecular Sciences, 2021, 22, 714.	1.8	4
10	T Cell Peptides Derived from Invasive Stages of Schistosoma mansoni as Potential Schistosomiasis Vaccine. Journal of Clinical Medicine, 2021, 10, 445.	1.0	1
11	Updating changes in human gut microbial communities associated with <i>Clostridioides difficile</i> infection. Gut Microbes, 2021, 13, 1966277.	4.3	5
12	Characterising four Sarconesiopsis magellanica (Diptera: Calliphoridae) larval fat body-derived antimicrobial peptides. Memorias Do Instituto Oswaldo Cruz, 2021, 116, e200587.	0.8	2
13	A comparative analysis of SLA-DRB1 genetic diversity in Colombian (creoles and commercial line) and worldwide swine populations. Scientific Reports, 2021, 11, 4340.	1.6	3
14	Two 20-Residue-Long Peptides Derived from Plasmodium vivax Merozoite Surface Protein 10 EGF-Like Domains Are Involved in Binding to Human Reticulocytes. International Journal of Molecular Sciences, 2021, 22, 1609.	1.8	2
15	Gut microbiota profiles in diarrheic patients with co-occurrence of Clostridioides difficile and Blastocystis. PLoS ONE, 2021, 16, e0248185.	1.1	19
16	MHCBI: a pipeline for calculating peptide-MHC binding energy using semi-empirical quantum mechanical methods with explicit/implicit solvent models. Briefings in Bioinformatics, 2021, 22, .	3.2	2
17	Gut microbiota composition in health-care facility-and community-onset diarrheic patients with Clostridioides difficile infection. Scientific Reports, 2021, 11, 10849.	1.6	8
18	Co-Circulation of Bovine Leukemia Virus Haplotypes among Humans, Animals, and Food Products: New Insights of Its Zoonotic Potential. International Journal of Environmental Research and Public Health, 2021, 18, 4883.	1.2	8

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19	Identifying HLA DRB1-DQB1 alleles associated with Chlamydia trachomatis infection and in silico prediction of potentially-related peptides. Scientific Reports, 2021, 11, 12837.	1.6	4
20	Mechanisms Associated with Trypanosoma cruzi Host Target Cell Adhesion, Recognition and Internalization. Life, 2021, 11, 534.	1.1	7
21	ESAT-6 and Ag85A Synthetic Peptides as Candidates for an Immunodiagnostic Test in Children with a Clinical Suspicion of Tuberculosis. Disease Markers, 2021, 2021, 1-9.	0.6	1
22	Genetic diversity and population structure of Rhipicephalus sanguineus sensu lato across different regions of Colombia. Parasites and Vectors, 2021, 14, 424.	1.0	13
23	Risk factor for breast cancer development under exposure to bovine leukemia virus in Colombian women: A case-control study. PLoS ONE, 2021, 16, e0257492.	1.1	5
24	The First Chemically-Synthesised, Highly Immunogenic Anti-SARS-CoV-2 Peptides in DNA Genotyped Aotus Monkeys for Human Use. Frontiers in Immunology, 2021, 12, 724060.	2.2	5
25	Nanovaccines against Animal Pathogens: The Latest Findings. Vaccines, 2021, 9, 988.	2.1	15
26	Molecular characterisation of parvorder Platyrrhini IgG sub-classes. Molecular Immunology, 2021, 139, 23-31.	1.0	2
27	Mycobacterium tuberculosis Rv0292 Protein Peptides Could be Included in a Synthetic Anti-tuberculosis Vaccine. International Journal of Peptide Research and Therapeutics, 2021, 27, 2823.	0.9	1
28	Trichomonas vaginalis follow-up and persistence in Colombian women. Scientific Reports, 2021, 11, 22597.	1.6	2
29	Structural Modelling of KCNQ1 and KCNH2 Double Mutant Proteins, Identified in Two Severe Long QT Syndrome Cases, Reveals New Insights into Cardiac Channelopathies. International Journal of Molecular Sciences, 2021, 22, 12861.	1.8	2
30	How to Combat Gram-Negative Bacteria Using Antimicrobial Peptides: A Challenge or an Unattainable Goal?. Antibiotics, 2021, 10, 1499.	1.5	19
31	NHP-immunome: A translational research-oriented database of non-human primate immune system proteins. Cellular Immunology, 2020, 347, 103999.	1.4	1
32	Robust, Comprehensive Molecular, and Phenotypical Characterisation of Atypical Candida albicans Clinical Isolates From BogotÃ <sub>i</sub> , Colombia. Frontiers in Cellular and Infection Microbiology, 2020, 10, 571147.	1.8	0
33	Molecular Markers for Detecting Schistosoma Species by Loop-Mediated Isothermal Amplification. Disease Markers, 2020, 2020, 1-11.	0.6	8
34	A Comprehensive Review of the Immunological Response against Foot-and-Mouth Disease Virus Infection and Its Evasion Mechanisms. Vaccines, 2020, 8, 764.	2.1	8
35	Loop-Mediated Isothermal Amplification as Point-of-Care Diagnosis for Neglected Parasitic Infections. International Journal of Molecular Sciences, 2020, 21, 7981.	1.8	29
36	Shorter Antibacterial Peptide Having High Selectivity for E. coli Membranes and Low Potential for Inducing Resistance. Microorganisms, 2020, 8, 867.	1.6	7

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37	<i>Strong</i> -LAMP Assay Based on a <i>Strongyloides</i> sppDerived Partial Sequence in the 18S rRNA as Potential Biomarker for Strongyloidiasis Diagnosis in Human Urine Samples. Disease Markers, 2020, 2020, 1-10.	0.6	13
38	Antibodies targeting Mycobacterium tuberculosis peptides inhibit mycobacterial entry to infection target cells. International Journal of Biological Macromolecules, 2020, 161, 712-720.	3.6	3
39	Identifying the HLA DRB1-DQB1 molecules and predicting epitopes associated with high-risk HPV infection clearance and redetection. Scientific Reports, 2020, 10, 7306.	1.6	9
40	Enhanced virulence of Beauveria bassiana against Diatraea saccharalis using a soluble recombinant enzyme with endo- and exochitinase activity. Biological Control, 2020, 144, 104211.	1.4	5
41	Plasmodium vivax Cell Traversal Protein for Ookinetes and Sporozoites (CelTOS) Functionally Restricted Regions Are Involved in Specific Host-Pathogen Interactions. Frontiers in Cellular and Infection Microbiology, 2020, 10, 119.	1.8	6
42	Hotspots in Plasmodium and RBC Receptor-Ligand Interactions: Key Pieces for Inhibiting Malarial Parasite Invasion. International Journal of Molecular Sciences, 2020, 21, 4729.	1.8	11
43	Sexual forms obtained in a continuous in vitro cultured Colombian strain of Plasmodium falciparum (FCB2). Malaria Journal, 2020, 19, 57.	0.8	2
44	Inferring Plasmodium vivax protein biology by using omics data. Journal of Proteomics, 2020, 218, 103719.	1.2	1
45	From a basic to a functional approach for developing a blood stage vaccine against Plasmodium vivax. Expert Review of Vaccines, 2020, 19, 195-207.	2.0	4
46	A <i>Trypanosoma cruzi</i> Genome Tandem Repetitive Satellite DNA Sequence as a Molecular Marker for a LAMP Assay for Diagnosing Chagas' Disease. Disease Markers, 2020, 2020, 1-8.	0.6	8
47	Major Histocompatibility Complex Class II (DRB3) Genetic Diversity in Spanish Morucha and Colombian Normande Cattle Compared to Taurine and Zebu Populations. Frontiers in Genetics, 2020, 10, 1293.	1.1	16
48	Plasmodium falciparum pre-erythrocytic stage vaccine development. Malaria Journal, 2020, 19, 56.	0.8	36
49	Occurrence of Blastocystis in Patients with Clostridioides difficile Infection. Pathogens, 2020, 9, 283.	1.2	13
50	Evaluating the immunogenicity of chemically-synthesised peptides derived from foot-and-mouth disease VP1, VP2 and VP3 proteins as vaccine candidates. Vaccine, 2020, 38, 3942-3951.	1.7	6
51	Malaria: Paving the way to developing peptide-based vaccines against invasion in infectious diseases. Biochemical and Biophysical Research Communications, 2020, 527, 1021-1026.	1.0	5
52	Una hoja de ruta para la Vacuna COVID 19 en Colombia, un reto posible. Infectio, 2020, 25, 7.	0.4	0
53	Integrated genomic epidemiology and phenotypic profiling of Clostridium difficile across intra-hospital and community populations in Colombia. Scientific Reports, 2019, 9, 11293.	1.6	12
54	Comparative genomics identifies potential virulence factors in <i>Clostridium tertium</i> and <i>C. paraputrificum</i> . Virulence, 2019, 10, 657-676.	1.8	13

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55	Using nextâ€generation sequencing for characterising HLA―DRB1 and DQB1 loci in a cohort of Colombian women. Hla, 2019, 94, 425-434.	0.4	1
56	Complement Receptor 1 availability on red blood cell surface modulates Plasmodium vivax invasion of human reticulocytes. Scientific Reports, 2019, 9, 8943.	1.6	14
57	High frequency of toxigenic Clostridium difficile and Clostridium perfringens coinfection among diarrheic patients at health care facility-onset (HCFO) and community-onset (CO) centers in BogotÃ;, Colombia. Gut Pathogens, 2019, 11, 27.	1.6	5
58	Sarconesin II, a New Antimicrobial Peptide Isolated from Sarconesiopsis magellanica Excretions and Secretions. Molecules, 2019, 24, 2077.	1.7	17
59	Specific Binding Peptides from Rv3632: A Strategy for BlockingMycobacterium tuberculosisEntry to Target Cells?. BioMed Research International, 2019, 2019, 1-13.	0.9	3
60	An epidemiological and molecular study regarding the spread of vancomycin-resistant Enterococcus faecium in a teaching hospital in BogotÃ;, Colombia 2016. BMC Infectious Diseases, 2019, 19, 258.	1.3	6
61	Behavior and abundance of Anopheles darlingi in communities living in the Colombian Amazon riverside. PLoS ONE, 2019, 14, e0213335.	1.1	9
62	Experimental models used in evaluating anti-tuberculosis vaccines: the latest advances in the field. Expert Review of Vaccines, 2019, 18, 365-377.	2.0	5
63	Peptides Derived of Kunitz-Type Serine Protease Inhibitor as Potential Vaccine Against Experimental Schistosomiasis. Frontiers in Immunology, 2019, 10, 2498.	2.2	21
64	Assessing Peptide Binding to MHC II: An Accurate Semiempirical Quantum Mechanics Based Proposal. Journal of Chemical Information and Modeling, 2019, 59, 5148-5160.	2.5	5
65	Immunocompetent patient with a brain abscess caused by Nocardia beijingensis in Latin America. Medicine (United States), 2019, 98, e14879.	0.4	16
66	Plasmodium falciparum Blood Stage Antimalarial Vaccines: An Analysis of Ongoing Clinical Trials and New Perspectives Related to Synthetic Vaccines. Frontiers in Microbiology, 2019, 10, 2712.	1.5	17
67	Parasite-Related Genetic and Epigenetic Aspects and Host Factors Influencing Plasmodium falciparum Invasion of Erythrocytes. Frontiers in Cellular and Infection Microbiology, 2019, 8, 454.	1.8	1
68	Abstract 4220: Evidence of bovine leukemia virus genes detected in Colombian women with and without breast cancer: A zoonotic infection. , 2019, , .		0
69	Preliminary Evaluation of the Safety and Immunogenicity of an Antimalarial Vaccine Candidate Modified Peptide (IMPIPS) Mixture in a Murine Model. Journal of Immunology Research, 2019, 2019, 1-12.	0.9	2
70	Abstract 4220: Evidence of bovine leukemia virus genes detected in Colombian women with and without breast cancer: A zoonotic infection. , 2019, , .		0
71	Quantifying intracellular <i>Mycobacterium tuberculosis</i> : An essential issue for in vitro assays. MicrobiologyOpen, 2018, 7, e00588.	1.2	15
72	Receptor-ligand and parasite protein-protein interactions in <i>Plasmodium vivax</i> : Analysing rhoptry neck proteins 2 and 4. Cellular Microbiology, 2018, 20, e12835.	1.1	15

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73	Comparison of healing of full-thickness skin wounds grafted with multidirectional or unidirectional autologous artificial dermis: differential delivery of healing biomarkers. Drug Delivery and Translational Research, 2018, 8, 1014-1024.	3.0	8
74	Micro-epidemiology of mixed-species malaria infections in a rural population living in the Colombian Amazon region. Scientific Reports, 2018, 8, 5543.	1.6	19
75	Towards designing a synthetic antituberculosis vaccine: The Rv3587c peptide inhibits mycobacterial entry to host cells. Bioorganic and Medicinal Chemistry, 2018, 26, 2401-2409.	1.4	13
76	Easy and fast method for expression and native extraction of Plasmodium vivax Duffy binding protein fragments. Malaria Journal, 2018, 17, 76.	0.8	1
77	Simultaneous detection of Plasmodium vivax dhfr, dhps, mdr1 and crt-o resistance-associated mutations in the Colombian Amazonian region. Malaria Journal, 2018, 17, 130.	0.8	12
78	Evaluating the anti-leishmania activity of Lucilia sericata and Sarconesiopsis magellanica blowfly larval excretions/secretions in an in vitro model. Acta Tropica, 2018, 177, 44-50.	0.9	19
79	Unveiling the Multilocus Sequence Typing (MLST) Schemes and Core Genome Phylogenies for Genotyping Chlamydia trachomatis. Frontiers in Microbiology, 2018, 9, 1854.	1.5	23
80	On the Evolution and Function of Plasmodium vivax Reticulocyte Binding Surface Antigen (pvrbsa). Frontiers in Genetics, 2018, 9, 372.	1.1	12
81	Cryptosporidium spp. CP15 and CSL protein-derived synthetic peptides' immunogenicity and in vitro seroneutralisation capability. Vaccine, 2018, 36, 6703-6710.	1.7	6
82	Sarconesin: Sarconesiopsis magellanica Blowfly Larval Excretions and Secretions With Antibacterial Properties. Frontiers in Microbiology, 2018, 9, 2249.	1.5	18
83	Plasmodium vivax Pv12 B-cell epitopes and HLA-DRβ1*-dependent T-cell epitopes in vitro antigenicity. PLoS ONE, 2018, 13, e0203715.	1.1	3
84	In silico and in vitro analysis of boAP3d1 protein interaction with bovine leukaemia virus gp51. PLoS ONE, 2018, 13, e0199397.	1.1	13
85	The in Vitro Antigenicity of Plasmodium vivax Rhoptry Neck Protein 2 (PvRON2) B- and T-Epitopes Selected by HLA-DRB1 Binding Profile. Frontiers in Cellular and Infection Microbiology, 2018, 8, 156.	1.8	9
86	Identifying Potential Plasmodium vivax Sporozoite Stage Vaccine Candidates: An Analysis of Genetic Diversity and Natural Selection. Frontiers in Genetics, 2018, 9, 10.	1.1	15
87	New Insights into Clostridium difficile (CD) Infection in Latin America: Novel Description of Toxigenic Profiles of Diarrhea-Associated to CD in BogotÃį, Colombia. Frontiers in Microbiology, 2018, 9, 74.	1.5	14
88	Mycobacterium tuberculosis H37Rv LpqG Protein Peptides Can Inhibit Mycobacterial Entry through Specific Interactions. Molecules, 2018, 23, 526.	1.7	5
89	Self-assembling functional programmable protein array for studying protein–protein interactions in malaria parasites. Malaria Journal, 2018, 17, 270.	0.8	10
90	Specific β-Turns Precede PPIIL Structures Binding to Allele-Specific HLA-DRβ1* PBRs in Fully-Protective Malaria Vaccine Components. Frontiers in Chemistry, 2018, 6, 106.	1.8	3

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91	Association of HIV status with infection by multiple HPV types. Tropical Medicine and International Health, 2018, 23, 1259-1268.	1.0	15
92	Plasmodium vivax in vitro continuous culture: the spoke in the wheel. Malaria Journal, 2018, 17, 301.	0.8	57
93	Diagnostic accuracy of combinations of serological biomarkers for identifying clinical tuberculosis. Journal of Infection in Developing Countries, 2018, 12, 429-441.	0.5	3
94	Transcriptome profiling of gene expression during immunisation trial against Fasciola hepatica: identification of genes and pathways involved in conferring immunoprotection in a murine model. BMC Infectious Diseases, 2017, 17, 94.	1.3	15
95	Identifying and characterising PPE7 (Rv0354c) high activity binding peptides and their role in inhibiting cell invasion. Molecular and Cellular Biochemistry, 2017, 430, 149-160.	1.4	6
96	pELMO, an optimised in-house cloning vector. AMB Express, 2017, 7, 26.	1.4	16
97	Multiple high-risk HPV genotypes are grouped by type and are associated with viral load and risk factors. Epidemiology and Infection, 2017, 145, 1479-1490.	1.0	11
98	New mutations in non-syndromic primary ovarian insufficiency patients identified via whole-exome sequencing. Human Reproduction, 2017, 32, 1512-1520.	0.4	65
99	PvGAMA reticulocyte binding activity: predicting conserved functional regions by natural selection analysis. Parasites and Vectors, 2017, 10, 251.	1.0	36
100	Critical role of HLA-DRβ* binding peptides' peripheral flanking residues in fully-protective malaria vaccine development. Biochemical and Biophysical Research Communications, 2017, 489, 339-345.	1.0	6
101	Semi-empirical quantum evaluation of peptide – MHC class II binding. Chemical Physics Letters, 2017, 668, 29-34.	1.2	12
102	Bovine leukaemia virus DNA in fresh milk and raw beef for human consumption. Epidemiology and Infection, 2017, 145, 3125-3130.	1.0	35
103	Purification of Trypanosoma cruzi metacyclic trypomastigotes by ion exchange chromatography in sepharose-DEAE, a novel methodology for host-pathogen interaction studies. Journal of Microbiological Methods, 2017, 142, 27-32.	0.7	15
104	Plasmodium vivax ligand-receptor interaction: PvAMA-1 domain I contains the minimal regions for specific interaction with CD71+ reticulocytes. Scientific Reports, 2017, 7, 9616.	1.6	29
105	Structural analysis of owl monkey MHC-DR shows that fully-protective malaria vaccine components can be readily used in humans. Biochemical and Biophysical Research Communications, 2017, 491, 1062-1069.	1.0	20
106	Determining Clostridium difficile intra-taxa diversity by mining multilocus sequence typing databases. BMC Microbiology, 2017, 17, 62.	1.3	44
107	Characterising PvRBSA: an exclusive protein from Plasmodium species infecting reticulocytes. Parasites and Vectors, 2017, 10, 243.	1.0	18
108	The Aotus nancymaae erythrocyte proteome and its importance for biomedical research. Journal of Proteomics, 2017, 152, 131-137.	1.2	6

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109	What Is Known about the Immune Response Induced by Plasmodium vivax Malaria Vaccine Candidates?. Frontiers in Immunology, 2017, 8, 126.	2.2	42
110	Novel genes and mutations in patients affected by recurrent pregnancy loss. PLoS ONE, 2017, 12, e0186149.	1.1	55
111	A novel loop-mediated isothermal amplification-based test for detecting Neospora caninum DNA. Parasites and Vectors, 2017, 10, 590.	1.0	5
112	Community-acquired infection with hypervirulent Clostridium difficile isolates that carry different toxin and antibiotic resistance loci: a case report. Gut Pathogens, 2017, 9, 63.	1.6	4
113	Far from the Madding Crowd: the Molecular Basis for Immunological Escape ofPlasmodium falciparum. Current Issues in Molecular Biology, 2017, 22, 65-78.	1.0	7
114	Size polymorphism and low sequence diversity in the locus encoding the Plasmodium vivaxÂrhoptry neck protein 4 (PvRON4) in Colombian isolates. Malaria Journal, 2016, 15, 501.	0.8	23
115	The Prevalence of High-Risk HPV Types and Factors Determining Infection in Female Colombian Adolescents. PLoS ONE, 2016, 11, e0166502.	1.1	14
116	The effect of Lucilia sericata - and Sarconesiopsis magellanica -derived larval therapy on Leishmania panamensis. Acta Tropica, 2016, 164, 280-289.	0.9	15
117	Evaluating the Biological Cycle and Reproductive and Population Parameters of <i>Calliphora vicina</i> (Diptera: Calliphoridae) Reared on Three Different Diets. Journal of Medical Entomology, 2016, 53, 1268-1275.	0.9	9
118	Plasmodium malariae in the Colombian Amazon region: you don't diagnose what you don't suspect. Malaria Journal, 2016, 15, 576.	0.8	20
119	Evidence of functional divergence in MSP7 paralogous proteins: a molecular-evolutionary and phylogenetic analysis. BMC Evolutionary Biology, 2016, 16, 256.	3.2	17
120	Cellâ€₽eptide Specific Interaction Can Inhibit <i>Mycobacterium tuberculosis H37Rv</i> Infection. Journal of Cellular Biochemistry, 2016, 117, 946-958.	1.2	6
121	TCR-contacting residues orientation and HLA-DRβ* binding preference determine long-lasting protective immunity against malaria. Biochemical and Biophysical Research Communications, 2016, 477, 654-660.	1.0	7
122	Mycobacterium tuberculosis PE9 protein has high activity binding peptides which inhibit target cell invasion. International Journal of Biological Macromolecules, 2016, 86, 646-655.	3.6	5
123	Evaluating Sarconesiopsis magellanica blowfly-derived larval therapy and comparing it to Lucilia sericata-derived therapy in an animal model. Acta Tropica, 2016, 154, 34-41.	0.9	12
124	Chlamydia trachomatis Frequency in a Cohort of HPV-Infected Colombian Women. PLoS ONE, 2016, 11, e0147504.	1.1	12
125	High Plasmodium malariae Prevalence in an Endemic Area of the Colombian Amazon Region. PLoS ONE, 2016, 11, e0159968.	1.1	31
126	The Malaria Parasite's Achilles' Heel: Functionally-relevant Invasion Structures. Current Issues in Molecular Biology, 2016, 18, 11-9.	1.0	5

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127	Malaria Parasite Survival Depends on Conserved Binding Peptides' Critical Biological Functions. Current Issues in Molecular Biology, 2016, 18, 57-78.	1.0	11
128	Immune protection-inducing protein structures (IMPIPS) against malaria: the weapons needed for beating Odysseus. Vaccine, 2015, 33, 7525-7537.	1.7	14
129	Evidence of recent interspecies horizontal gene transfer regarding nucleopolyhedrovirus infection of Spodoptera frugiperda. BMC Genomics, 2015, 16, 1008.	1.2	15
130	Characterising atypical Candida albicans clinical isolates from six third-level hospitals in BogotÃį, Colombia. BMC Microbiology, 2015, 15, 199.	1.3	17
131	IMPIPS: The Immune Protection-Inducing Protein Structure Concept in the Search for Steric-Electron and Topochemical Principles for Complete Fully-Protective Chemically Synthesised Vaccine Development. PLoS ONE, 2015, 10, e0123249.	1.1	25
132	Evaluating the Effect of <i>Sarconesiopsis magellanica</i> (Diptera: Calliphoridae) Larvae-Derived Haemolymph and Fat Body Extracts on Chronic Wounds in Diabetic Rabbits. Journal of Diabetes Research, 2015, 2015, 1-10.	1.0	7
133	Terapia larval en la curación de heridas. Infectio, 2015, 19, 1-2.	0.4	1
134	Determining the Plasmodium vivax VCG-1 strain blood stage proteome. Journal of Proteomics, 2015, 113, 268-280.	1.2	32
135	Recent advances in the development of a chemically synthesised anti-malarial vaccine. Expert Opinion on Biological Therapy, 2015, 15, 1567-1581.	1.4	13
136	New echinocandin susceptibility patterns for nosocomial Candida albicans in BogotÃ <sub>i</sub> , Colombia, in ten tertiary care centres: an observational study. BMC Infectious Diseases, 2015, 15, 108.	1.3	5
137	The DNA load of six high-risk human papillomavirus types and its association with cervical lesions. BMC Cancer, 2015, 15, 100.	1.1	36
138	The Plasmodium vivax rhoptry neck protein 5 is expressed in the apical pole of Plasmodium vivax VCG-1 strain schizonts and binds to human reticulocytes. Malaria Journal, 2015, 14, 106.	0.8	29
139	Inferring natural selection signals in Plasmodium vivax -encoded proteins having a potential role in merozoite invasion. Infection, Genetics and Evolution, 2015, 33, 182-188.	1.0	21
140	Mce4F Mycobacterium tuberculosis protein peptides can inhibit invasion of human cell lines. Pathogens and Disease, 2015, 73, .	0.8	17
141	The effects of Sarconesiopsis magellanica larvae (Diptera: Calliphoridae) excretions and secretions on fibroblasts. Acta Tropica, 2015, 142, 26-33.	0.9	7
142	Using the PfEMP1 Head Structure Binding Motif to Deal a Blow at Severe Malaria. PLoS ONE, 2014, 9, e88420.	1.1	8
143	Characterising a Microsatellite for DRB Typing in Aotus vociferans and Aotus nancymaae (Platyrrhini). PLoS ONE, 2014, 9, e96973.	1.1	15
144	In Vitro and In Vivo Studies for Assessing the Immune Response and Protection-Inducing Ability Conferred by Fasciola hepatica-Derived Synthetic Peptides Containing B- and T-Cell Epitopes. PLoS ONE, 2014. 9. e105323.	1.1	32

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145	Heterogeneous genetic diversity pattern in Plasmodium vivax genes encoding merozoite surface proteins (MSP) -7E, â^'7F and -7L. Malaria Journal, 2014, 13, 495.	0.8	24
146	Low genetic diversity in the locus encoding the Plasmodium vivax P41 protein in Colombia's parasite population. Malaria Journal, 2014, 13, 388.	0.8	22
147	Gauche+ side-chain orientation as a key factor in the search for an immunogenic peptide mixture leading to a complete fully protective vaccine. Vaccine, 2014, 32, 2117-2126.	1.7	20
148	Plasmodium falciparum rhoptry neck protein 5 peptides bind to human red blood cells and inhibit parasite invasion. Peptides, 2014, 53, 210-217.	1.2	9
149	Low genetic diversity and functional constraint in loci encoding Plasmodium vivax P12 and P38 proteins in the Colombian population. Malaria Journal, 2014, 13, 58.	0.8	28
150	Th1 and Th2 immune response to P30 and ROP18 peptides in human toxoplasmosis. Medical Microbiology and Immunology, 2014, 203, 315-322.	2.6	46
151	Specific Interaction between <i><scp>M</scp>ycobacterium tuberculosis</i> Lipoproteinâ€derived Peptides and Target Cells Inhibits Mycobacterial Entry <i>In Vitro</i> . Chemical Biology and Drug Design, 2014, 84, 626-641.	1.5	16
152	Persistence, clearance and reinfection regarding six high risk human papillomavirus types in Colombian women: a follow-up study. BMC Infectious Diseases, 2014, 14, 395.	1.3	18
153	Human papillomavirus detection in women with and without human immunodeficiency virus infection in Colombia. BMC Cancer, 2014, 14, 451.	1.1	14
154	Sarconesiopsis magellanica (Diptera: Calliphoridae) excretions and secretions have potent antibacterial activity. Acta Tropica, 2014, 136, 37-43.	0.9	21
155	Functional, biochemical and 3D studies of <i>Mycobacterium tuberculosis</i> protein peptides for an effective anti-tuberculosis vaccine. Critical Reviews in Microbiology, 2014, 40, 117-145.	2.7	14
156	Characterizing PvARP, a novel Plasmodium vivax antigen. Malaria Journal, 2013, 12, 165.	0.8	17
157	Rv1268c protein peptide inhibiting Mycobacterium tuberculosis H37Rv entry to target cells. Bioorganic and Medicinal Chemistry, 2013, 21, 6650-6656.	1.4	6
158	Sarconesiopsis magellanica (Diptera: Calliphoridae) life-cycle, reproductive and population parameters using different diets under laboratory conditions. Forensic Science International, 2013, 233, 380-386.	1.3	15
159	Classical Molecular Tests Using Urine Samples as a Potential Screening Tool for Human Papillomavirus Detection in Human Immunodeficiency Virus-Infected Women. Journal of Clinical Microbiology, 2013, 51, 3688-3693.	1.8	8
160	Annotation and characterization of the Plasmodium vivax rhoptry neck protein 4 (Pv RON4). Malaria Journal, 2013, 12, 356.	0.8	27
161	Proteolytic activity regarding Sarconesiopsis magellanica (Diptera: Calliphoridae) larval excretions and secretions. Acta Tropica, 2013, 128, 686-691.	0.9	12
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