Carlos F SuÃ;rez

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	SM-COLSARSPROT: Highly Immunogenic Supramutational Synthetic Peptides Covering the World's Population. Frontiers in Immunology, 2022, 13, .	4.8	0
2	The molecular basis for peptide-based antimalarial vaccine development targeting erythrocyte invasion by P.Âfalciparum. Biochemical and Biophysical Research Communications, 2021, 534, 86-93.	2.1	3
3	Babesia Bovis Ligand-Receptor Interaction: AMA-1 Contains Small Regions Governing Bovine Erythrocyte Binding. International Journal of Molecular Sciences, 2021, 22, 714.	4.1	4
4	A comparative analysis of SLA-DRB1 genetic diversity in Colombian (creoles and commercial line) and worldwide swine populations. Scientific Reports, 2021, 11, 4340.	3.3	3
5	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.	4.1	2
6	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, .	6.5	2
7	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.	4.8	5
8	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.	4.1	2
9	Robust, Comprehensive Molecular, and Phenotypical Characterisation of Atypical Candida albicans Clinical Isolates From Bogotá, Colombia. Frontiers in Cellular and Infection Microbiology, 2020, 10, 571147.	3.9	0
10	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.	3.9	6
11	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.	2.3	16
12	Malaria: Paving the way to developing peptide-based vaccines against invasion in infectious diseases. Biochemical and Biophysical Research Communications, 2020, 527, 1021-1026.	2.1	5
13	Fundamentos y aplicaciones biomédicas de las principales tecnologÃas de secuenciación: una revisión de literatura. Revista Investigación En Salud Universidad De Boyacá, 2020, 7, .	0.1	Ο
14	Assessing Peptide Binding to MHC II: An Accurate Semiempirical Quantum Mechanics Based Proposal. Journal of Chemical Information and Modeling, 2019, 59, 5148-5160.	5.4	5
15	Plasmodium vivax Pv12 B-cell epitopes and HLA-DRβ1*-dependent T-cell epitopes in vitro antigenicity. PLoS ONE, 2018, 13, e0203715.	2.5	3
16	Self-assembling functional programmable protein array for studying protein–protein interactions in malaria parasites. Malaria Journal, 2018, 17, 270.	2.3	10
17	New mutations in non-syndromic primary ovarian insufficiency patients identified via whole-exome sequencing. Human Reproduction, 2017, 32, 1512-1520.	0.9	65
18	Semi-empirical quantum evaluation of peptide – MHC class II binding. Chemical Physics Letters, 2017, 668, 29-34.	2.6	12

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19	Mass & secondary structure propensity of amino acids explain their mutability and evolutionary replacements. Scientific Reports, 2017, 7, 7717.	3.3	7
20	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.	2.1	20
21	Novel genes and mutations in patients affected by recurrent pregnancy loss. PLoS ONE, 2017, 12, e0186149.	2.5	55
22	How frequently do clusters occur in hierarchical clustering analysis? A graph theoretical approach to studying ties in proximity. Journal of Cheminformatics, 2016, 8, 4.	6.1	11
23	TCR-contacting residues orientation and HLA-DRÎ ² * binding preference determine long-lasting protective immunity against malaria. Biochemical and Biophysical Research Communications, 2016, 477, 654-660.	2.1	7
24	Characterising atypical Candida albicans clinical isolates from six third-level hospitals in BogotÃ _i , Colombia. BMC Microbiology, 2015, 15, 199.	3.3	17
25	Characterising a Microsatellite for DRB Typing in Aotus vociferans and Aotus nancymaae (Platyrrhini). PLoS ONE, 2014, 9, e96973.	2.5	15
26	Characterisation and comparative analysis of MHC-DPA1 exon 2 in the owl monkey (Aotus nancymaae). Gene, 2011, 470, 37-45.	2.2	9
27	High polymorphism in Plasmodium vivax merozoite surface protein-5 (MSP5). Parasitology, 2006, 133, 661.	1.5	31
28	Reference strand conformational analysis (RSCA) is a valuable tool in identifying MHC-DRB sequences in three species of Aotus monkeys. Immunogenetics, 2006, 58, 590-597.	2.4	15
29	Owl monkey MHC-DRB exon 2 reveals high similarity with several HLA-DRB lineages. Immunogenetics, 2006, 58, 542-558.	2.4	68
30	High level of conservation in Plasmodium vivax merozoite surface protein 4 (PvMSP4)â~†. Infection, Genetics and Evolution, 2005, 5, 354-361.	2.3	16
31	A comparative study of MHC Class-II HLA-DRβ1*0401-Col II and HLA-DRβ1*0101-HA complexes: a theoretical point of view. Journal of Structural Biology, 2005, 149, 38-52.	2.8	13
32	Identification of five different IGHV gene families in owl monkeys (Aotus nancymaae). Tissue Antigens, 2005, 66, 640-649.	1.0	7
33	MHC class I genes in the owl monkey: mosaic organisation, convergence and loci diversity. Immunogenetics, 2005, 56, 818-832.	2.4	18
34	The T-cell receptor in primates: identifying and sequencing new owl monkey TRBV gene sub-groups. Immunogenetics, 2005, 57, 42-52.	2.4	19
35	Quantum chemical analysis explains hemagglutinin peptide–MHC Class II molecule HLA-DRβ1*0101 interactions. Biochemical and Biophysical Research Communications, 2004, 323, 1265-1277	2.1	17
36	Plasmodium vivax Duffy binding protein: a modular evolutionary proposal. Parasitology, 2004, 128, 353-366.	1.5	24

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37	α1 and α2 domains of Aotus MHC Class I and Catarrhini MHC Class Ia share similar characteristics. Tissue Antigens, 2003, 61, 362-373.	1.0	13
38	Characterizing T-cell receptor gamma-variable gene in Aotus nancymaae owl monkey peripheral blood. Tissue Antigens, 2003, 62, 472-482.	1.0	19
39	Electronic Energy and Multipolar Moments Characterize Amino Acid Side Chains into Chemically Related Groups. Journal of Physical Chemistry A, 2003, 107, 10090-10097.	2.5	22
40	Identification, cloning, and sequencing of different cytokine genes in four species of owl monkey. Immunogenetics, 2002, 54, 645-653.	2.4	20