

Mireia Lopez-Siles

List of Publications by Year in descending order

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Version: 2024-02-01

25
papers

1,908
citations

687363

13
h-index

677142

22
g-index

25
all docs

25
docs citations

25
times ranked

3242
citing authors

#	ARTICLE	IF	CITATIONS
1	Cross-Recognition of SARS-CoV-2 B-Cell Epitopes with Other Betacoronavirus Nucleoproteins. <i>International Journal of Molecular Sciences</i> , 2022, 23, 2977.	4.1	4
2	Prevalence, Abundance, and Virulence of Adherent-Invasive <i>Escherichia coli</i> in Ulcerative Colitis, Colorectal Cancer, and Coeliac Disease. <i>Frontiers in Immunology</i> , 2022, 13, 748839.	4.8	12
3	Brief Research Report: Virus-Specific Humoral Immunity at Admission Predicts the Development of Respiratory Failure in Unvaccinated SARS-CoV-2 Patients. <i>Frontiers in Immunology</i> , 2022, 13, 878812.	4.8	3
4	Identification of Promoter Region Markers Associated With Altered Expression of Resistance-Nodulation-Division Antibiotic Efflux Pumps in <i>Acinetobacter baumannii</i> . <i>Frontiers in Microbiology</i> , 2022, 13, .	3.5	1
5	Vaccines for multidrug resistant Gram negative bacteria: lessons from the past for guiding future success. <i>FEMS Microbiology Reviews</i> , 2021, 45, .	8.6	18
6	Optimization of a Lambda-RED Recombination Method for Rapid Gene Deletion in Human Cytomegalovirus. <i>International Journal of Molecular Sciences</i> , 2021, 22, 10558.	4.1	1
7	Subinhibitory Concentrations of Clinically-Relevant Antimicrobials Affect Resistance-Nodulation-Division Family Promoter Activity in <i>Acinetobacter baumannii</i> . <i>Frontiers in Microbiology</i> , 2021, 12, 780201.	3.5	4
8	Healthy dietary pattern and their corresponding gut microbiota profile are linked to a lower risk of type 2 diabetes, independent of the presence of obesity. <i>Clinical Nutrition</i> , 2020, 39, 524-532.	5.0	25
9	Identification and Analysis of Unstructured, Linear B-Cell Epitopes in SARS-CoV-2 Virion Proteins for Vaccine Development. <i>Vaccines</i> , 2020, 8, 397.	4.4	17
10	Evaluation of bacterial biomarkers to aid in challenging inflammatory bowel diseases diagnostics and subtype classification. <i>World Journal of Gastrointestinal Pathophysiology</i> , 2020, 11, 64-77.	1.0	8
11	Gut microbiota imbalances in Tunisian participants with type 1 and type 2 diabetes mellitus. <i>Bioscience Reports</i> , 2019, 39, .	2.4	38
12	Genetic and Phenotypic Features to Screen for Putative Adherent-Invasive <i>Escherichia coli</i> . <i>Frontiers in Microbiology</i> , 2019, 10, 108.	3.5	23
13	Comparative genomics reveals new single-nucleotide polymorphisms that can assist in identification of adherent-invasive <i>Escherichia coli</i> . <i>Scientific Reports</i> , 2018, 8, 2695.	3.3	46
14	Alterations in the Abundance and Co-occurrence of <i>Akkermansia muciniphila</i> and <i>Faecalibacterium prausnitzii</i> in the Colonic Mucosa of Inflammatory Bowel Disease Subjects. <i>Frontiers in Cellular and Infection Microbiology</i> , 2018, 8, 281.	3.9	135
15	<i>Faecalibacterium prausnitzii</i> : from microbiology to diagnostics and prognostics. <i>ISME Journal</i> , 2017, 11, 841-852.	9.8	510
16	Changes in the Abundance of <i>Faecalibacterium prausnitzii</i> Phylogroups I and II in the Intestinal Mucosa of Inflammatory Bowel Disease and Patients with Colorectal Cancer. <i>Inflammatory Bowel Diseases</i> , 2016, 22, 28-41.	1.9	108
17	Anti-tumour Necrosis Factor Treatment with Adalimumab Induces Changes in the Microbiota of Crohn's Disease. <i>Journal of Crohn's and Colitis</i> , 2015, 9, 899-906.	1.3	59
18	Mucosa-Associated <i>Faecalibacterium prausnitzii</i> Phylotype Richness Is Reduced in Patients with Inflammatory Bowel Disease. <i>Applied and Environmental Microbiology</i> , 2015, 81, 7582-7592.	3.1	89

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19	Mucosa-associated <i>Faecalibacterium prausnitzii</i> and <i>Escherichia coli</i> co-abundance can distinguish Irritable Bowel Syndrome and Inflammatory Bowel Disease phenotypes. <i>International Journal of Medical Microbiology</i> , 2014, 304, 464-475.	3.6	114
20	Cultured Representatives of Two Major Phylogroups of Human Colonic <i>Faecalibacterium prausnitzii</i> Can Utilize Pectin, Uronic Acids, and Host-Derived Substrates for Growth. <i>Applied and Environmental Microbiology</i> , 2012, 78, 420-428.	3.1	341
21	A New Validated Real-Time PCR-Based Method for the Specific and Fast Detection of <i>Cronobacter</i> spp. in Infant Formula. <i>Food Analytical Methods</i> , 2012, 5, 179-187.	2.6	10
22	A validated simple and rapid method for the simultaneous detection of both <i>Cronobacter</i> spp. and <i>Salmonella</i> spp. for infant formula quality control. <i>Dairy Science and Technology</i> , 2012, 92, 151-166.	2.2	2
23	Gut Environmental Factors May Shape the Persistence of <i>Faecalibacterium Prausnitzii</i> in the Healthy and Diseased Large Intestine. <i>Gastroenterology</i> , 2011, 140, S-665.	1.3	1
24	W1261 Effect of Adalimumab Treatment on the Microbiota Recovery in the Intestinal Mucosa of Crohn's Disease Patients. <i>Gastroenterology</i> , 2010, 138, S-685-S-686.	1.3	0
25	Molecular diversity of <i>Escherichia coli</i> in the human gut: New ecological evidence supporting the role of adherent-invasive <i>E. coli</i> (AIEC) in Crohn's disease. <i>Inflammatory Bowel Diseases</i> , 2009, 15, 872-882.	1.9	339