Souhaila Al-Khodor

List of Publications by Year in descending order

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60 papers

2,337 citations

270111 25 h-index 263392 45 g-index

62 all docs

62 docs citations

62 times ranked 2969 citing authors

#	Article	IF	Citations
1	Microbial Dysbiosis Tunes the Immune Response Towards Allergic Disease Outcomes. Clinical Reviews in Allergy and Immunology, 2023, 65, 43-71.	2.9	14
2	Corneal confocal microscopy identifies a reduction in corneal keratocyte density and sub-basal nerves in children with type 1 diabetes mellitus. British Journal of Ophthalmology, 2022, 106, 1368-1372.	2.1	6
3	Tipping the Balance: Vitamin D Inadequacy in Children Impacts the Major Gut Bacterial Phyla. Biomedicines, 2022, 10, 278.	1.4	7
4	Microbiota medicine: towards clinical revolution. Journal of Translational Medicine, 2022, 20, 111.	1.8	87
5	The Impact of Nutritional Supplementation During Pregnancy on the Incidence of Gestational Diabetes and Glycaemia Control. Frontiers in Nutrition, 2022, 9, 867099.	1.6	3
6	Gut microbial influences on the adaptive immune system and the development of cow milk allergy. Qatar Medical Journal, 2022, 2022, .	0.2	0
7	Inflammatory Bowel Disease Treatments and Predictive Biomarkers of Therapeutic Response. International Journal of Molecular Sciences, 2022, 23, 6966.	1.8	32
8	Distinctive Microbial Signatures and Gut-Brain Crosstalk in Pediatric Patients with Coeliac Disease and Type 1 Diabetes Mellitus. International Journal of Molecular Sciences, 2021, 22, 1511.	1.8	10
9	Vaginal Microbiota and Cytokine Levels Predict Preterm Delivery in Asian Women. Frontiers in Cellular and Infection Microbiology, 2021, 11, 639665.	1.8	34
10	The Salivary miRNome: A Promising Biomarker of Disease. MicroRNA (Shariqah, United Arab Emirates), 2021, 10, 29-38.	0.6	4
11	Azithromycin Exhibits Activity Against Pseudomonas aeruginosa in Chronic Rat Lung Infection Model. Frontiers in Microbiology, 2021, 12, 603151.	1.5	11
12	Translational metagenomics. Journal of Translational Medicine, 2021, 19, 158.	1.8	0
13	COVID-19 Infection during Pregnancy: Risk of Vertical Transmission, Fetal, and Neonatal Outcomes. Journal of Personalized Medicine, 2021, 11, 483.	1.1	24
14	Immunomodulatory Effects of Vitamin D Supplementation in a Deficient Population. International Journal of Molecular Sciences, 2021, 22, 5041.	1.8	5
15	Oral microbiome and pregnancy: A bidirectional relationship. Journal of Reproductive Immunology, 2021, 145, 103293.	0.8	40
16	Omouma: a prospective mother and child cohort aiming to identify early biomarkers of pregnancy complications in women living in Qatar. BMC Pregnancy and Childbirth, 2021, 21, 570.	0.9	3
17	"Armed for the future Coronavirus pandemic― a promising use of the multimeric SARS-CoV-2 receptor binding domain nanoparticle as a new Pan-Coronavirus vaccine. Signal Transduction and Targeted Therapy, 2021, 6, 305.	7.1	1
18	Actionable genomic variants in 6045 participants from the Qatar Genome Program. Human Mutation, 2021, 42, 1584-1601.	1.1	13

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19	Exploring the Triple Interaction between the Host Genome, the Epigenome, and the Gut Microbiome in Type 1 Diabetes. International Journal of Molecular Sciences, 2021, 22, 125.	1.8	11
20	Thousands of Qatari genomes inform human migration history and improve imputation of Arab haplotypes. Nature Communications, 2021, 12, 5929.	5.8	18
21	Breast Milk: A Meal Worth Having. Frontiers in Nutrition, 2021, 8, 800927.	1.6	22
22	The Human Microbiome in Chronic Kidney Disease: A Double-Edged Sword. Frontiers in Medicine, 2021, 8, 790783.	1.2	31
23	Can the Salivary Microbiome Predict Cardiovascular Diseases? Lessons Learned From the Qatari Population. Frontiers in Microbiology, 2021, 12, 772736.	1.5	5
24	Corneal confocal microscopy demonstrates minimal evidence of distal neuropathy in children with celiac disease. PLoS ONE, 2020, 15, e0238859.	1.1	4
25	A literature-based approach for curating gene signatures in multifaceted diseases. Journal of Translational Medicine, 2020, 18, 279.	1.8	6
26	Annexin A3 in sepsis: novel perspectives from an exploration of public transcriptome data. Immunology, 2020, 161, 291-302.	2.0	32
27	Role of the gut microbiota in the pathogenesis of coeliac disease and potential therapeutic implications. European Journal of Nutrition, 2020, 59, 3369-3390.	1.8	42
28	Cohort profile: molecular signature in pregnancy (MSP): longitudinal high-frequency sampling to characterise cross-omic trajectories in pregnancy in a resource-constrained setting. BMJ Open, 2020, 10, e041631.	0.8	6
29	The Role of Polymorphisms in Vitamin D-Related Genes in Response to Vitamin D Supplementation. Nutrients, 2020, 12, 2608.	1.7	13
30	Pathophysiology and treatment strategies for COVID-19. Journal of Translational Medicine, 2020, 18, 353.	1.8	71
31	The potential role of vitamin D supplementation as a gut microbiota modifier in healthy individuals. Scientific Reports, 2020, 10, 21641.	1.6	100
32	Corneal nerve loss in children with typeÂ1 diabetes mellitus without retinopathy or microalbuminuria. Journal of Diabetes Investigation, 2020, 11, 1594-1601.	1.1	13
33	Profiling the Salivary microbiome of the Qatari population. Journal of Translational Medicine, 2020, 18, 127.	1.8	33
34	Microbiome as an Immunological Modifier. Methods in Molecular Biology, 2020, 2055, 595-638.	0.4	23
35	DS86760016, a Leucyl-tRNA Synthetase Inhibitor with Activity against Pseudomonas aeruginosa. Antimicrobial Agents and Chemotherapy, 2019, 63, .	1.4	9
36	Evaluation of Methods for the Extraction of Microbial DNA From Vaginal Swabs Used for Microbiome Studies. Frontiers in Cellular and Infection Microbiology, 2019, 9, 197.	1.8	27

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37	Vitamin D Deficiency in the Gulf Cooperation Council: Exploring the Triad of Genetic Predisposition, the Gut Microbiome and the Immune System. Frontiers in Immunology, 2019, 10, 1042.	2.2	31
38	A prospective cohort for the investigation of alteration in temporal transcriptional and microbiome trajectories preceding preterm birth: a study protocol. BMJ Open, 2019, 9, e023417.	0.8	15
39	Integrating omics for a better understanding of Inflammatory Bowel Disease: a step towards personalized medicine. Journal of Translational Medicine, 2019, 17, 419.	1.8	52
40	Gut microbiome and kidney disease: a bidirectional relationship. Pediatric Nephrology, 2017, 32, 921-931.	0.9	122
41	The Microbiome and Blood Pressure: Can Microbes Regulate Our Blood Pressure?. Frontiers in Pediatrics, 2017, 5, 138.	0.9	102
42	Human Microbiome and its Association With Health and Diseases. Journal of Cellular Physiology, 2016, 231, 1688-1694.	2.0	98
43	Activator of G-Protein Signaling 3–Induced Lysosomal Biogenesis Limits Macrophage Intracellular Bacterial Infection. Journal of Immunology, 2016, 196, 846-856.	0.4	31
44	Assay Development for Image-Based Quantification of Intracellular Bacterial Replication and Analysis of the Innate Immune Response to Infection. Assay and Drug Development Technologies, 2015, 13, 515-528.	0.6	5
45	Development of a cell system for siRNA screening of pathogen responses in human and mouse macrophages. Scientific Reports, 2015, 5, 9559.	1.6	21
46	<i>Burkholderia cenocepacia</i> â€J2315 escapes to the cytosol and actively subverts autophagy in human macrophages. Cellular Microbiology, 2014, 16, 378-395.	1.1	35
47	Indispensable Role for the Eukaryotic-Like Ankyrin Domains of the Ankyrin B Effector of Legionella pneumophila within Macrophages and Amoebae. Infection and Immunity, 2013, 81, 2660-2660.	1.0	0
48	Cell biology and molecular ecology of Francisella tularensis. Cellular Microbiology, 2010, 12, 129-139.	1.1	68
49	Triggering Ras signalling by intracellular Francisella tularensis through recruitment of PKC $\hat{l}\pm$ and \hat{l}^2 I to the SOS2/GrB2 complex is essential for bacterial proliferation in the cytosol. Cellular Microbiology, 2010, 12, 1604-1621.	1.1	18
50	Molecular Characterization of the Dot/Icm-Translocated AnkH and AnkJ Eukaryotic-Like Effectors of < i>Legionella pneumophila < / i>. Infection and Immunity, 2010, 78, 1123-1134.	1.0	36
51	Indispensable Role for the Eukaryotic-Like Ankyrin Domains of the Ankyrin B Effector of <i>Legionella pneumophila </i> within Macrophages and Amoebae. Infection and Immunity, 2010, 78, 2079-2088.	1.0	58
52	Functional diversity of ankyrin repeats in microbial proteins. Trends in Microbiology, 2010, 18, 132-139.	3.5	178
53	Temporal and differential regulation of expression of the eukaryoticâ€ike ankyrin effectors of <i>Legionella pneumophila</i> . Environmental Microbiology Reports, 2010, 2, 677-684.	1.0	12
54	Host Factors Required for Modulation of Phagosome Biogenesis and Proliferation of Francisella tularensis within the Cytosol. PLoS ONE, 2010, 5, e11025.	1.1	57

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55	Molecular Mimicry by an F-Box Effector of Legionella pneumophila Hijacks a Conserved Polyubiquitination Machinery within Macrophages and Protozoa. PLoS Pathogens, 2009, 5, e1000704.	2.1	156
56	The PmrA/PmrB Two-Component System of <i>Legionella pneumophila </i> Is a Global Regulator Required for Intracellular Replication within Macrophages and Protozoa. Infection and Immunity, 2009, 77, 374-386.	1.0	70
57	Role for the Ankyrin eukaryoticâ€like genes of <i>Legionella pneumophila</i> in parasitism of protozoan hosts and human macrophages. Environmental Microbiology, 2008, 10, 1460-1474.	1.8	108
58	A Dot/Icmâ€translocated ankyrin protein of <i>Legionella pneumophila</i> is required for intracellular proliferation within human macrophages and protozoa. Molecular Microbiology, 2008, 70, 908-923.	1.2	150
59	Chlamydial Entry Involves TARP Binding of Guanine Nucleotide Exchange Factors. PLoS Pathogens, 2008, 4, e1000014.	2.1	132
60	Infections and Pregnancy: Effects on Maternal and Child Health. Frontiers in Cellular and Infection Microbiology, 0, 12, .	1.8	22