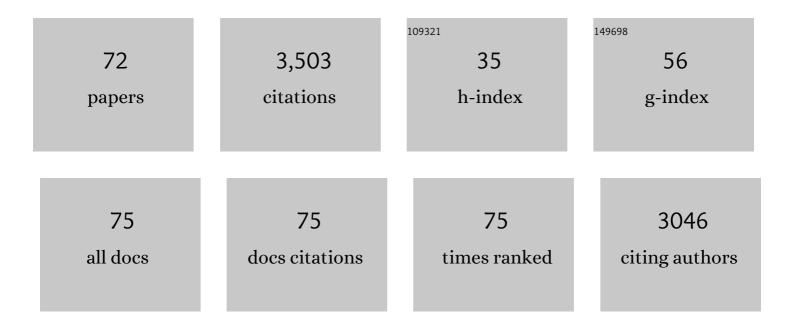
List of Publications by Year in descending order

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ΙΙΛΝ SΗΛ

#	Article	IF	CITATIONS
1	A new generation needle- and adjuvant-free trivalent plague vaccine utilizing adenovirus-5 nanoparticle platform. Npj Vaccines, 2021, 6, 21.	6.0	14
2	A universal bacteriophage T4 nanoparticle platform to design multiplex SARS-CoV-2 vaccine candidates by CRISPR engineering. Science Advances, 2021, 7, eabh1547.	10.3	44
3	Combinatorial Viral Vector-Based and Live Attenuated Vaccines without an Adjuvant to Generate Broader Immune Responses to Effectively Combat Pneumonic Plague. MBio, 2021, 12, e0322321.	4.1	6
4	New Host-Directed Therapeutics for the Treatment of Clostridioides difficile Infection. MBio, 2020, 11,	4.1	8
5	A prokaryotic-eukaryotic hybrid viral vector for delivery of large cargos of genes and proteins into human cells. Science Advances, 2019, 5, eaax0064.	10.3	28
6	T6SS and ExoA of flesh-eating <i>Aeromonas hydrophila</i> in peritonitis and necrotizing fasciitis during mono- and polymicrobial infections. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 24084-24092.	7.1	19
7	Combating Multidrug-Resistant Pathogens with Host-Directed Nonantibiotic Therapeutics. Antimicrobial Agents and Chemotherapy, 2018, 62, .	3.2	22
8	A Bacteriophage T4 Nanoparticle-Based Dual Vaccine against Anthrax and Plague. MBio, 2018, 9, .	4.1	62
9	Identification of New Virulence Factors and Vaccine Candidates for Yersinia pestis. Frontiers in Cellular and Infection Microbiology, 2017, 7, 448.	3.9	23
10	Protective Immunity Elicited by Oral Immunization of Mice with Salmonella enterica Serovar Typhimurium Braun Lipoprotein (Lpp) and Acetyltransferase (MsbB) Mutants. Frontiers in Cellular and Infection Microbiology, 2016, 6, 148.	3.9	13
11	Immunisation of two rodent species with new live-attenuated mutants of Yersinia pestis CO92 induces protective long-term humoral- and cell-mediated immunity against pneumonic plague. Npj Vaccines, 2016, 1, 16020.	6.0	17
12	New Insights into Autoinducer-2 Signaling as a Virulence Regulator in a Mouse Model of Pneumonic Plague. MSphere, 2016, 1, .	2.9	15
13	New Role for FDA-Approved Drugs in Combating Antibiotic-Resistant Bacteria. Antimicrobial Agents and Chemotherapy, 2016, 60, 3717-3729.	3.2	38
14	A Replication-Defective Human Type 5 Adenovirus-Based Trivalent Vaccine Confers Complete Protection against Plague in Mice and Nonhuman Primates. Vaccine Journal, 2016, 23, 586-600.	3.1	21
15	Cross-talk among flesh-eating <i>Aeromonas hydrophila</i> strains in mixed infection leading to necrotizing fasciitis. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 722-727.	7.1	113
16	Combinational Deletion of Three Membrane Protein-Encoding Genes Highly Attenuates Yersinia pestis while Retaining Immunogenicity in a Mouse Model of Pneumonic Plague. Infection and Immunity, 2015, 83, 1318-1338.	2.2	20
17	Further characterization of a highly attenuated Yersinia pestis CO92 mutant deleted for the genes encoding Braun lipoprotein and plasminogen activator protease in murine alveolar and primary human macrophages. Microbial Pathogenesis, 2015, 80, 27-38.	2.9	9
18	Intramuscular Immunization of Mice with a Live-Attenuated Triple Mutant of Yersinia pestis CO92 Induces Robust Humoral and Cell-Mediated Immunity To Completely Protect Animals against Pneumonic Plague. Vaccine Journal, 2015, 22, 1255-1268.	3.1	15

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19	High-Throughput, Signature-Tagged Mutagenic Approach To Identify Novel Virulence Factors of Yersinia pestis CO92 in a Mouse Model of Infection. Infection and Immunity, 2015, 83, 2065-2081.	2.2	19
20	Deletion of Braun Lipoprotein and Plasminogen-Activating Protease-Encoding Genes Attenuates Yersinia pestis in Mouse Models of Bubonic and Pneumonic Plague. Infection and Immunity, 2014, 82, 2485-2503.	2.2	22
21	Functional Genomic Characterization of Virulence Factors from Necrotizing Fasciitis-Causing Strains of Aeromonas hydrophila. Applied and Environmental Microbiology, 2014, 80, 4162-4183.	3.1	54
22	The Effects of Modeled Microgravity on Growth Kinetics, Antibiotic Susceptibility, Cold Growth, and the Virulence Potential of a <i>Yersinia pestis ymoA</i> -Deficient Mutant and Its Isogenic Parental Strain. Astrobiology, 2013, 13, 821-832.	3.0	24
23	A non-invasive inÂvivo imaging system to study dissemination of bioluminescent Yersinia pestis CO92 in a mouse model of pneumonic plague. Microbial Pathogenesis, 2013, 55, 39-50.	2.9	25
24	Evaluation of the roles played by Hcp and VgrG type 6 secretion system effectors in Aeromonas hydrophila SSU pathogenesis. Microbiology (United Kingdom), 2013, 159, 1120-1135.	1.8	55
25	Characterization of Aeromonas hydrophilaÂWound Pathotypes by Comparative Genomic and Functional Analyses of Virulence Genes. MBio, 2013, 4, e00064-13.	4.1	71
26	Mutated and Bacteriophage T4 Nanoparticle Arrayed F1-V Immunogens from Yersinia pestis as Next Generation Plague Vaccines. PLoS Pathogens, 2013, 9, e1003495.	4.7	56
27	Deletion of the Braun Lipoprotein-Encoding Gene and Altering the Function of Lipopolysaccharide Attenuate the Plague Bacterium. Infection and Immunity, 2013, 81, 815-828.	2.2	27
28	Evaluation of Protective Potential of Yersinia pestis Outer Membrane Protein Antigens as Possible Candidates for a New-Generation Recombinant Plague Vaccine. Vaccine Journal, 2013, 20, 227-238.	3.1	32
29	The two-component QseBC signalling system regulates in vitro and in vivo virulence of Aeromonas hydrophila. Microbiology (United Kingdom), 2012, 158, 259-271.	1.8	60
30	DNA adenine methyltransferase (Dam) controls the expression of the cytotoxic enterotoxin (act) gene of Aeromonas hydrophila via tRNA modifying enzyme-glucose-inhibited division protein (GidA). Gene, 2012, 498, 280-287.	2.2	15
31	Actin cross-linking domain of Aeromonas hydrophila repeat in toxin A (RtxA) induces host cell rounding and apoptosis. Gene, 2012, 506, 369-376.	2.2	39
32	Quorum sensing and c-di-GMP-dependent alterations in gene transcripts and virulence-associated phenotypes in a clinical isolate of Aeromonas hydrophila. Microbial Pathogenesis, 2011, 50, 213-223.	2.9	42
33	Progress on plague vaccine development. Applied Microbiology and Biotechnology, 2011, 91, 265-286.	3.6	40
34	Cethromycin-Mediated Protection against the Plague Pathogen Yersinia pestis in a Rat Model of Infection and Comparison with Levofloxacin. Antimicrobial Agents and Chemotherapy, 2011, 55, 5034-5042.	3.2	17
35	Characterization of an F1 Deletion Mutant of Yersinia pestis CO92, Pathogenic Role of F1 Antigen in Bubonic and Pneumonic Plague, and Evaluation of Sensitivity and Specificity of F1 Antigen Capture-Based Dipsticks. Journal of Clinical Microbiology, 2011, 49, 1708-1715.	3.9	42
36	Distribution of Virulence Factors and Molecular Fingerprinting of <i>Aeromonas</i> Species Isolates from Water and Clinical Samples: Suggestive Evidence of Water-to-Human Transmission. Applied and Environmental Microbiology, 2010, 76, 2313-2325.	3.1	124

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37	Comparative Global Gene Expression Profiles of Wild-Type <i>Yersinia pestis</i> CO92 and Its Braun Lipoprotein Mutant at Flea and Human Body Temperatures. Comparative and Functional Genomics, 2010, 2010, 1-11.	2.0	13
38	Deletion of Braun lipoprotein gene (lpp) attenuates Yersinia pestis KIM/D27 strain: Role of Lpp in modulating host immune response, NF-IºB activation and cell death. Microbial Pathogenesis, 2010, 48, 42-52.	2.9	14
39	Unraveling the mechanism of action of a new type III secretion system effector AexU from Aeromonas hydrophila. Microbial Pathogenesis, 2010, 49, 122-134.	2.9	34
40	Protection Afforded by Fluoroquinolones in Animal Models of Respiratory Infections with Bacillus anthracis, Yersinia pestis, and Francisella tularensis. Open Microbiology Journal, 2010, 4, 34-46.	0.7	35
41	Comparative Analyses of Transcriptional Profiles in Mouse Organs Using a Pneumonic Plague Model after Infection with Wild-Type <i>Yersinia pestis</i> CO92 and Its Braun Lipoprotein Mutant. Comparative and Functional Genomics, 2009, 2009, 1-16.	2.0	14
42	Phospholipase A2-activating protein (PLAA) enhances cisplatin-induced apoptosis in HeLa cells. Cellular Signalling, 2009, 21, 1085-1099.	3.6	29
43	Characterization of the rat pneumonic plague model: infection kinetics following aerosolization of Yersinia pestis CO92. Microbes and Infection, 2009, 11, 205-214.	1.9	50
44	N-Acylhomoserine lactones involved in quorum sensing control the type VI secretion system, biofilm formation, protease production, and in vivo virulence in a clinical isolate of Aeromonas hydrophila. Microbiology (United Kingdom), 2009, 155, 3518-3531.	1.8	124
45	Surface-Expressed Enolase Contributes to the Pathogenesis of Clinical Isolate SSU of <i>Aeromonas hydrophila</i> . Journal of Bacteriology, 2009, 191, 3095-3107.	2.2	93
46	Deletion of Braun lipoprotein gene (lpp) and curing of plasmid pPCP1 dramatically alter the virulence of Yersinia pestis CO92 in a mouse model of pneumonic plague. Microbiology (United Kingdom), 2009, 155, 3247-3259.	1.8	27
47	Alteration in the activation state of new inflammation-associated targets by phospholipase A2-activating protein (PLAA). Cellular Signalling, 2008, 20, 844-861.	3.6	18
48	Immunological responses against Salmonella enterica serovar Typhimurium Braun lipoprotein and lipid A mutant strains in Swiss-Webster mice: Potential use as live-attenuated vaccines. Microbial Pathogenesis, 2008, 44, 224-237.	2.9	26
49	Molecular characterization of a functional type VI secretion system from a clinical isolate of Aeromonas hydrophila. Microbial Pathogenesis, 2008, 44, 344-361.	2.9	193
50	Mutation in the S-ribosylhomocysteinase (luxS) gene involved in quorum sensing affects biofilm formation and virulence in a clinical isolate of Aeromonas hydrophila. Microbial Pathogenesis, 2008, 45, 343-354.	2.9	52
51	Braun Lipoprotein (Lpp) Contributes to Virulence of Yersiniae: Potential Role of Lpp in Inducing Bubonic and Pneumonic Plague. Infection and Immunity, 2008, 76, 1390-1409.	2.2	75
52	Characterization of a mouse model of plague after aerosolization of Yersinia pestis CO92. Microbiology (United Kingdom), 2008, 154, 1939-1948.	1.8	72
53	Cold Shock Exoribonuclease R (VacB) Is Involved in <i>Aeromonas hydrophila</i> Pathogenesis. Journal of Bacteriology, 2008, 190, 3467-3474.	2.2	54
54	Interaction between innate immune cells and a bacterial type III secretion system in mutualistic and pathogenic associations. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 9481-9486.	7.1	79

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55	Biological characterization of a new type III secretion system effector from a clinical isolate of Aeromonas hydrophila—Part II. Microbial Pathogenesis, 2007, 43, 147-160.	2.9	37
56	Global transcriptional responses of wild-type Aeromonas hydrophila and its virulence-deficient mutant in a murine model of infection. Microbial Pathogenesis, 2007, 42, 193-203.	2.9	15
57	Further characterization of a type III secretion system (T3SS) and of a new effector protein from a clinical isolate of Aeromonas hydrophila—Part I. Microbial Pathogenesis, 2007, 43, 127-146.	2.9	65
58	Identification of a new hemolysin from diarrheal isolate SSU of <i>Aeromonas hydrophila</i> . FEMS Microbiology Letters, 2007, 275, 301-311.	1.8	30
59	Genome Sequence of Aeromonas hydrophila ATCC 7966 T : Jack of All Trades. Journal of Bacteriology, 2006, 188, 8272-8282.	2.2	317
60	Deletion of the genes encoding the type III secretion system and cytotoxic enterotoxin alters host responses to Aeromonas hydrophila infection. Microbial Pathogenesis, 2006, 40, 198-210.	2.9	19
61	Host Immune Responses to Aeromonas Virulence Factors. Current Immunology Reviews, 2006, 2, 13-26.	1.2	63
62	Molecular and Functional Characterization of a ToxR-Regulated Lipoprotein from a Clinical Isolate of Aeromonas hydrophila. Infection and Immunity, 2006, 74, 3742-3755.	2.2	25
63	DNA Adenine Methyltransferase Influences the Virulence of Aeromonas hydrophila. Infection and Immunity, 2006, 74, 410-424.	2.2	61
64	Mutations within the Catalytic Motif of DNA Adenine Methyltransferase (Dam) of Aeromonas hydrophila Cause the Virulence of the Dam-Overproducing Strain To Revert to That of the Wild-Type Phenotype. Infection and Immunity, 2006, 74, 5763-5772.	2.2	19
65	Microarray and Proteomics Analyses of Human Intestinal Epithelial Cells Treated with the Aeromonas hydrophila Cytotoxic Enterotoxin. Infection and Immunity, 2005, 73, 2628-2643.	2.2	20
66	The Type III Secretion System and Cytotoxic Enterotoxin Alter the Virulence of Aeromonas hydrophila. Infection and Immunity, 2005, 73, 6446-6457.	2.2	84
67	Molecular Characterization of a Glucose-Inhibited Division Gene, gidA , That Regulates Cytotoxic Enterotoxin of Aeromonas hydrophila. Infection and Immunity, 2004, 72, 1084-1095.	2.2	71
68	Aeromonas hydrophila Cytotoxic Enterotoxin Activates Mitogen-activated Protein Kinases and Induces Apoptosis in Murine Macrophages and Human Intestinal Epithelial Cells. Journal of Biological Chemistry, 2004, 279, 37597-37612.	3.4	94
69	Differential expression of the enolase gene under in vivo versus in vitro growth conditions of Aeromonas hydrophila. Microbial Pathogenesis, 2003, 34, 195-204.	2.9	39
70	Identification of Aeromonas hydrophila Cytotoxic Enterotoxin-induced Genes in Macrophages Using Microarrays. Journal of Biological Chemistry, 2003, 278, 40198-40212.	3.4	47
71	Role of Various Enterotoxins in Aeromonas hydrophila-Induced Gastroenteritis: Generation of Enterotoxin Gene-Deficient Mutants and Evaluation of Their Enterotoxic Activity. Infection and Immunity, 2002, 70, 1924-1935.	2.2	203
72	Regulation of the Cytotoxic Enterotoxin Gene in Aeromonas hydrophila : Characterization of an Iron Uptake Regulator. Infection and Immunity, 2001, 69, 6370-6381.	2.2	37