Chengping Lu

List of Publications by Year
in descending order

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130 papers	3,373 citations	126907 33 h-index	206112 48 g-index
135 all docs	135 docs citations	135 times ranked	3217 citing authors

#	Article	IF	Citations
1	The Hcp proteins fused with diverse extended-toxin domains represent a novel pattern of antibacterial effectors in type VI secretion systems. Virulence, 2017, 8, 1189-1202.	4.4	120
2	Novel insights into the pathogenicity of epidemic Aeromonas hydrophila ST251 clones from comparative genomics. Scientific Reports, 2015, 5, 9833.	3.3	110
3	PAARâ€Rhs proteins harbor various Câ€terminal toxins to diversify the antibacterial pathways of type VI secretion systems. Environmental Microbiology, 2017, 19, 345-360.	3.8	105
4	Functional analysis of luxS in Streptococcus suis reveals a key role in biofilm formation and virulence. Veterinary Microbiology, 2011, 152, 151-160.	1.9	97
5	Genomic and Epidemiological Characteristics Provide New Insights into the Phylogeographical and Spatiotemporal Spread of Porcine Epidemic Diarrhea Virus in Asia. Journal of Clinical Microbiology, 2015, 53, 1484-1492.	3.9	86
6	Effects of ibeA Deletion on Virulence and Biofilm Formation of Avian Pathogenic Escherichia coli. Infection and Immunity, 2011, 79, 279-287.	2.2	75
7	Reduced virulence is an important characteristic of biofilm infection of Streptococcus suis. FEMS Microbiology Letters, 2011, 316, 36-43.	1.8	74
8	Comparative genomics analysis of Streptococcus agalactiae reveals that isolates from cultured tilapia in China are closely related to the human strain A909. BMC Genomics, 2013, 14, 775.	2.8	73
9	Hepatoprotective and antioxidant effects of Glycyrrhiza glabra extract against carbon tetrachloride (CCl4)-induced hepatocyte damage in common carp (Cyprinus carpio). Fish Physiology and Biochemistry, 2011, 37, 209-216.	2.3	72
10	Immunoproteomic assay of surface proteins ofStreptococcus suisserotype 9. FEMS Immunology and Medical Microbiology, 2008, 53, 52-59.	2.7	66
11	Comparative proteome analysis of secreted proteins of Streptococcus suis serotype 9 isolates from diseased and healthy pigs. Microbial Pathogenesis, 2008, 45, 159-166.	2.9	66
12	Identification of Novel Laminin- and Fibronectin-binding Proteins by Far-Western Blot: Capturing the Adhesins of Streptococcus suis Type 2. Frontiers in Cellular and Infection Microbiology, 2015, 5, 82.	3.9	64
13	Transcriptome profiling of zebrafish infected with Streptococcus suis. Microbial Pathogenesis, 2010, 48, 178-187.	2.9	63
14	Two Functional Type VI Secretion Systems in Avian Pathogenic Escherichia coli Are Involved in Different Pathogenic Pathways. Infection and Immunity, 2014, 82, 3867-3879.	2.2	63
15	The Streptococcus suis transcriptional landscape reveals adaptation mechanisms in pig blood and cerebrospinal fluid. Rna, 2014, 20, 882-898.	3.5	59
16	Novel Variant Serotype of Streptococcus suis Isolated from Piglets with Meningitis. Applied and Environmental Microbiology, 2015, 81, 976-985.	3.1	57
17	Inhibition of Aeromonas hydrophila-induced intestinal inflammation and mucosal barrier function damage in crucian carp by oral administration of Lactococcus lactis. Fish and Shellfish Immunology, 2018, 83, 359-367.	3.6	51
18	Comparative Proteomic Analysis of Streptococcus suis Biofilms and Planktonic Cells That Identified Biofilm Infection-Related Immunogenic Proteins. PLoS ONE, 2012, 7, e33371.	2.5	50

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19	Biofilm Formation, Host-Cell Adherence, and Virulence Genes Regulation of Streptococcus suis in Response to Autoinducer-2 Signaling. Current Microbiology, 2014, 68, 575-580.	2.2	48
20	cas9 Enhances Bacterial Virulence by Repressing the regR Transcriptional Regulator in Streptococcus agalactiae. Infection and Immunity, 2018, 86, .	2.2	48
21	Comparative genomic analysis shows that Streptococcus suis meningitis isolate SC070731 contains a unique 105K genomic island. Gene, 2014, 535, 156-164.	2.2	45
22	Isolation and characterization of bacteriophages against virulent Aeromonas hydrophila. BMC Microbiology, 2020, 20, 141.	3.3	43
23	Streptococcus suis serotype 9 strain GZ0565 contains a type VII secretion system putative substrate EsxA that contributes to bacterial virulence and a vanZ- like gene that confers resistance to teicoplanin and dalbavancin in Streptococcus agalactiae. Veterinary Microbiology, 2017, 205, 26-33.	1.9	42
24	Diverse toxic effectors are harbored by vgrG islands for interbacterial antagonism in type VI secretion system. Biochimica Et Biophysica Acta - General Subjects, 2018, 1862, 1635-1643.	2.4	42
25	Analysis of synonymous codon usage patterns in torque teno sus virus 1 (TTSuV1). Archives of Virology, 2013, 158, 145-154.	2.1	41
26	ArcA Controls Metabolism, Chemotaxis, and Motility Contributing to the Pathogenicity of Avian Pathogenic Escherichia coli. Infection and Immunity, 2015, 83, 3545-3554.	2.2	41
27	Alterations in gp37 Expand the Host Range of a T4-Like Phage. Applied and Environmental Microbiology, 2017, 83, .	3.1	41
28	Contribution of Eukaryotic-Type Serine/Threonine Kinase to Stress Response and Virulence of Streptococcus suis. PLoS ONE, 2014, 9, e91971.	2.5	40
29	Complete Genome Sequence of Streptococcus agalactiae GD201008-001, Isolated in China from Tilapia with Meningoencephalitis. Journal of Bacteriology, 2012, 194, 6653-6653.	2.2	38
30	Genetic and pathobiologic characterization of H3N2 canine influenza viruses isolated in the Jiangsu Province of China in 2009â€“2010. Veterinary Microbiology, 2012, 158, 247-258.	1.9	38
31	Canine Distemper Virus Causes Apoptosis of Vero Cells. Zoonoses and Public Health, 2000, 47, 183-190.	1.4	36
32	Biological activity and identification of a peptide inhibitor of LuxS fromStreptococcus suisserotype 2. FEMS Microbiology Letters, 2009, 294, 16-23.	1.8	35
33	Characterization and genome sequencing of a novel bacteriophage infecting Streptococcus agalactiae with high similarity to a phage from Streptococcus pyogenes. Archives of Virology, 2013, 158, 1733-1741.	2.1	35
34	The novel virulence-related gene stp of Streptococcus suis serotype 9 strain contributes to a significant reduction in mouse mortality. Microbial Pathogenesis, 2011, 51, 442-453.	2.9	33
35	Use of in vivo-induced antigen technology (IVIAT) for the identification of Streptococcus suis serotype 2 in vivo-induced bacterial protein antigens. BMC Microbiology, 2009, 9, 201.	3.3	30
36	Isolation, genome sequencing and functional analysis of two T7-like coliphages of avian pathogenic Escherichia coli. Gene, 2016, 582, 47-58.	2.2	29

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37	Streptococcus suis small RNA rss04 contributes to the induction of meningitis by regulating capsule synthesis and by inducing biofilm formation in a mouse infection model. Veterinary Microbiology, 2017, 199, 111-119.	1.9	29
38	Immunoproteomics selection of cross-protective vaccine candidates from Riemerella anatipestifer serotypes 1 and 2. Veterinary Microbiology, 2013, 162, 850-857.	1.9	27
39	Crystal Structure and Identification of Two Key Amino Acids Involved in AI-2 Production and Biofilm Formation in Streptococcus suis LuxS. PLoS ONE, 2015, 10, e0138826.	2.5	27
40	Latest developments on Streptococcus suis: an emerging zoonotic pathogen: part 2. Future Microbiology, 2014, 9, 587-591.	2.0	26
41	Comparative genome analysis provides deep insights into Aeromonas hydrophila taxonomy and virulence-related factors. BMC Genomics, 2018, 19, 712.	2.8	26
42	Overexpression ofluxSCannot Increase Autoinducer-2 Production, Only Affect the Growth and Biofilm Formation inStreptococcus suis. Scientific World Journal, The, 2013, 2013, 1-6.	2.1	25
43	The non-conserved region of MRP is involved in the virulence ofStreptococcus suisserotype 2. Virulence, 2017, 8, 1274-1289.	4.4	25
44	Streptococcus suis synthesizes deoxyadenosine and adenosine by 5â€™-nucleotidase to dampen host immune responses. Virulence, 2018, 9, 1509-1520.	4.4	24
45	Adhesion activity of glyceraldehyde-3-phosphate dehydrogenase in a Chinese Streptococcus suis type 2 strain. Berliner Und Munchener Tierarztliche Wochenschrift, 2007, 120, 207-9.	0.7	24
46	Virulence genotyping and population analysis of Streptococcus suis serotype 2 isolates from China. Infection, Genetics and Evolution, 2015, 36, 483-489.	2.3	23
47	Lethal infection by a novel reassortant H5N1 avian influenza A virus in a zoo-housed tiger. Microbes and Infection, 2015, 17, 54-61.	1.9	23
48	Identification and Characterization of an Aeromonas hydrophila Oligopeptidase Gene pepF Negatively Related to Biofilm Formation. Frontiers in Microbiology, 2016, 7, 1497.	3.5	23
49	Diverse roles of Hcp family proteins in the environmental fitness and pathogenicity of Aeromonas hydrophila Chinese epidemic strain NJ-35. Applied Microbiology and Biotechnology, 2018, 102, 7083-7095.	3.6	23
50	Roles of three TonB systems in the iron utilization and virulence of the Aeromonas hydrophila Chinese epidemic strain NJ-35. Applied Microbiology and Biotechnology, 2019, 103, 4203-4215.	3.6	23
51	Pre-Absorbed Immunoproteomics: A Novel Method for the Detection of Streptococcus suis Surface Proteins. PLoS ONE, 2011, 6, e21234.	2.5	23
52	Development of Rapid Serotype-Specific PCR Assays for Eight Serotypes of Streptococcus suis. Journal of Clinical Microbiology, 2012, 50, 3329-3334.	3.9	22
53	Isolation and characterization of a T4â€like phage with a relatively wide host range within Escherichia coli. Journal of Basic Microbiology, 2016, 56, 405-421.	3.3	22
54	Identification of two mutation sites in spike and envelope proteins mediating optimal cellular infection of porcine epidemic diarrhea virus from different pathways. Veterinary Research, 2017, 48, 44.	3.0	22

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55	Characterization of Streptococcus suis Isolates from Slaughter Swine. Current Microbiology, 2013, 66, 344-349.	2.2	21
56	Identification of six novel capsular polysaccharide loci (<scp>NCL</scp>) from StreptococcusÂsuis multidrug resistant nonâ€typeable strains and the pathogenic characteristic of strains carrying new <scp>NCL</scp> s. Transboundary and Emerging Diseases, 2019, 66, 995-1003.	3.0	21
57	Genetic analysis of the capsular polysaccharide synthesis locus in 15 Streptococcus suis serotypes. FEMS Microbiology Letters, 2011, 324, 117-124.	1.8	20
58	Natural infection with torque teno sus virus 1 (TTSuV1) suppresses the immune response to porcine reproductive and respiratory syndrome virus (PRRSV) vaccination. Archives of Virology, 2012, 157, 927-933.	2.1	20
59	Fibronectin-/fibrinogen-binding protein (FBPS) is not a critical virulence factor for the Streptococcus suis serotype 2 strain ZY05719. Veterinary Microbiology, 2017, 208, 38-46.	1.9	20
60	Three Hcp homologs with divergent extended loop regions exhibit different functions in avian pathogenic Escherichia coli. Emerging Microbes and Infections, 2018, 7, 1-13.	6.5	20
61	Evidence of circulation of an epidemic strain of Pasteurella multocida in Jiangsu, China by multi-locus sequence typing (MLST). Infection, Genetics and Evolution, 2013, 20, 34-38.	2.3	19
62	Prophage Lysin Ply30 Protects Mice from Streptococcus suis and Streptococcus equi subsp. zooepidemicus Infections. Applied and Environmental Microbiology, 2015, 81, 7377-7384.	3.1	19
63	A Streptococcus suis LysM domain surface protein contributes to bacterial virulence. Veterinary Microbiology, 2016, 187, 64-69.	1.9	19
64	Protective efficacy of recombinant hemolysin co-regulated protein (Hcp) of Aeromonas hydrophila in common carp (Cyprinus carpio). Fish and Shellfish Immunology, 2015, 46, 297-304.	3.6	18
65	Identification of novel virulence-related genes in Aeromonas hydrophila by screening transposon mutants in a Tetrahymena infection model. Veterinary Microbiology, 2017, 199, 36-46.	1.9	18
66	Inducible Prophage Mutant of Escherichia coli Can Lyse New Host and the Key Sites of Receptor Recognition Identification. Frontiers in Microbiology, 2017, 8, 147.	3.5	18
67	IolR, a negative regulator of the myo-inositol metabolic pathway, inhibits cell autoaggregation and biofilm formation by downregulating RpmA in Aeromonas hydrophila. Npj Biofilms and Microbiomes, 2020, 6, 22.	6.4	18
68	Catecholamine-Stimulated Growth of Aeromonas hydrophila Requires the TonB2 Energy Transduction System but Is Independent of the Amonabactin Siderophore. Frontiers in Cellular and Infection Microbiology, 2016, 6, 183.	3.9	17
69	Down-regulating heat shock protein 27 is involved in porcine epidemic diarrhea virus escaping from host antiviral mechanism. Veterinary Microbiology, 2017, 205, 6-13.	1.9	17
70	The Two-Component Signaling System VraSR _{ss} Is Critical for Multidrug Resistance and Full Virulence in Streptococcus suis Serotype 2. Infection and Immunity, 2018, 86, .	2.2	17
71	Chaperonin GroEL: A novel phylogenetically conserved protein with strong immunoreactivity of Avian Pathogenic Escherichia coli isolates from duck identified by immunoproteomics. Vaccine, 2013, 31, 2947-2953.	3.8	16
72	SssP1, a Streptococcus suis Fimbria-Like Protein Transported by the SecY2/A2 System, Contributes to Bacterial Virulence. Applied and Environmental Microbiology, 2018, 84, .	3.1	16

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73	The identification of six novel proteins with fibronectin or collagen type I binding activity from Streptococcus suis serotype 2. Journal of Microbiology, 2014, 52, 963-969.	2.8	15
74	AutA and AutR, Two Novel Global Transcriptional Regulators, Facilitate Avian Pathogenic Escherichia coli Infection. Scientific Reports, 2016, 6, 25085.	3.3	15
75	Lysogenic Streptococcus suis Isolate SS2-4 Containing Prophage SMP Showed Increased Mortality in Zebra Fish Compared to the Wild-Type Isolate. PLoS ONE, 2013, 8, e54227.	2.5	15
76	Detection of canine coronaviruses genotype I and II in raised Canidae animals in China. Berliner Und Munchener Tierarztliche Wochenschrift, 2006, 119, 35-9.	0.7	15
77	The cps locus of Streptococcus suis serotype 16: Development of a serotype-specific PCR assay. Veterinary Microbiology, 2011, 153, 403-406.	1.9	14
78	Immunoproteomic analysis of bacterial proteins of Actinobacillus pleuropneumoniae serotype 1. Proteome Science, 2011, 9, 32.	1.7	14
79	Immunoproteomic identification of 11 novel immunoreactive proteins ofRiemerella anatipestiferserotype 2. FEMS Immunology and Medical Microbiology, 2012, 65, 84-95.	2.7	14
80	A streptococcal Fic domain-containing protein disrupts blood-brain barrier integrity by activating moesin in endothelial cells. PLoS Pathogens, 2019, 15, e1007737.	4.7	14
81	Identification of a new effector-immunity pair of Aeromonas hydrophila type VI secretion system. Veterinary Research, 2020, 51, 71.	3.0	14
82	Identification of immunoreactive proteins ofStreptococcus agalactiaeisolated from cultured tilapia in China. Pathogens and Disease, 2013, 69, 223-231.	2.0	13
83	Identification of Aeromonas hydrophila Genes Preferentially Expressed after Phagocytosis by Tetrahymena and Involvement of Methionine Sulfoxide Reductases. Frontiers in Cellular and Infection Microbiology, 2016, 6, 199.	3.9	13
84	Tetrahymena thermophila Predation Enhances Environmental Adaptation of the Carp Pathogenic Strain Aeromonas hydrophila NJ-35. Frontiers in Cellular and Infection Microbiology, 2018, 8, 76.	3.9	13
85	The Novel Streptococcal Transcriptional Regulator XtgS Negatively Regulates Bacterial Virulence and Directly Represses PseP Transcription. Infection and Immunity, 2020, 88, .	2.2	13
86	Molecular characterization of the 9.36Âkb C-terminal region of canine coronavirus 1-71 strain. Virus Genes, 2008, 36, 491-497.	1.6	12
87	A Novel Dual Vector Coexpressing PhiX174 Lysis E Gene and Staphylococcal Nuclease A Gene on the Basis of Lambda Promoter pR and pL, Respectively. Molecular Biotechnology, 2013, 54, 436-444.	2.4	12
88	Antibacterial effect of porcine PTX3 against Streptococcus suis type 2 infection. Microbial Pathogenesis, 2015, 89, 128-139.	2.9	12
89	Factor H specifically capture novel Factor H-binding proteins of Streptococcus suis and contribute to the virulence of the bacteria. Microbiological Research, 2017, 196, 17-25.	5.3	12
90	Infection and adaption-based proteomic changes of Streptococcus suis serotype 2 in a pig model. Journal of Proteomics, 2018, 180, 41-52.	2.4	12

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91	Immune responses and protection efficacy of a recombinant swinepox virus expressing HA1 against swine H3N2 influenza virus in mice and pigs. Virus Research, 2012, 167, 188-195.	2.2	11
92	Identification of a novel collagen type Ð†-binding protein from Streptococcus suis serotype 2. Veterinary Journal, 2013, 197, 406-414.	1.7	11
93	The Truncated Major Pilin Subunit Sbp2 of the srtBCD Pilus Cluster Still Contributes to Streptococcus suis Pathogenesis in the Absence of Pilus Shaft. Current Microbiology, 2014, 69, 703-707.	2.2	11
94	The effects of H3N2 swine influenza virus infection on TLRs and RLRs signaling pathways in porcine alveolar macrophages. Virology Journal, 2015, 12, 61.	3.4	11
95	IbeR Facilitates Stress-Resistance, Invasion and Pathogenicity of Avian Pathogenic Escherichia coli. PLoS ONE, 2015, 10, e0119698.	2.5	10
96	The role of regulator Eha in Edwardsiella tarda pathogenesis and virulence gene transcription. Microbial Pathogenesis, 2016, 95, 216-223.	2.9	10
97	Evaluation of the differences between biofilm and planktonic Brucella abortus via metabolomics and proteomics. Functional and Integrative Genomics, 2021, 21, 421-433.	3.5	10
98	Enhanced replication of avian-origin H3N2 canine influenza virus in eggs, cell cultures and mice by a two-amino acid insertion in neuraminidase stalk. Veterinary Research, 2016, 47, 53.	3.0	9
99	Quantitative assessment of the blood-brain barrier opening caused by Streptococcus agalactiae hyaluronidase in a BALB/c mouse model. Scientific Reports, 2017, 7, 13529.	3.3	9
100	EsR240, a non-coding sRNA, is required for the resistance of Edwardsiella tarda to stresses in macrophages and for virulence. Veterinary Microbiology, 2019, 231, 254-263.	1.9	9
101	Immunoproteomic assay of secreted proteins of Streptococcus suis serotype 9 with convalescent sera from pigs. Folia Microbiologica, 2011, 56, 423-430.	2.3	8
102	Immune responses and protective efficacy of a recombinant swinepox virus co-expressing HA1 genes of H3N2 and H1N1 swine influenza virus in mice and pigs. Veterinary Microbiology, 2013, 162, 259-264.	1.9	8
103	Fifteen novel immunoreactive proteins of Chinese virulent Haemophilus parasuis serotype 5 verified by an immunoproteomic assay. Folia Microbiologica, 2015, 60, 81-87.	2.3	8
104	Monoclonal antibody specific to HA2 glycopeptide protects mice from H3N2 influenza virus infection. Veterinary Research, 2015, 46, 33.	3.0	8
105	Intracranial Subarachnoidal Route of Infection for Investigating Roles of Streptococcus suis Biofilms in Meningitis in a Mouse Infection Model. Journal of Visualized Experiments, 2018, , .	0.3	8
106	Diverse effects of nitric oxide reductase NorV on Aeromonas hydrophila virulence-associated traits under aerobic and anaerobic conditions. Veterinary Research, 2019, 50, 67.	3.0	8
107	Immune responses and protective efficacy of a recombinant swinepox virus expressing HA1 against swine H1N1 influenza virus in mice and pigs. Vaccine, 2012, 30, 3119-3125.	3.8	7
108	Establishment and characterization of a telomerase-immortalized canine bronchiolar epithelial cell line. Applied Microbiology and Biotechnology, 2015, 99, 9135-9146.	3.6	7

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109	Discovery of lahS as a Global Regulator of Environmental Adaptation and Virulence in Aeromonas hydrophila. International Journal of Molecular Sciences, 2018, 19, 2709.	4.1	7
110	Role of luxS in immune evasion and pathogenicity of piscine Streptococcus agalactiae is not dependent on autoinducer-2. Fish and Shellfish Immunology, 2020, 99, 274-283.	3.6	7
111	CRISPR-dependent endogenous gene regulation is required for virulence in piscine Streptococcus agalactiae. Emerging Microbes and Infections, 2021, 10, 1-53.	6.5	7
112	Protection of guinea pigs by vaccination with a recombinant swinepox virus co-expressing HA1 genes of swine H1N1 and H3N2 influenza viruses. Archives of Virology, 2013, 158, 629-637.	2.1	6
113	Whole-Genome Sequence of Streptococcus suis Serotype 4 Reference Strain 6407. Genome Announcements, 2014, 2, .	0.8	6
114	Identification of a virulence-related surface protein XF in piscine Streptococcus agalactiaeby pre-absorbed immunoproteomics. BMC Veterinary Research, 2014, 10, 259.	1.9	6
115	Eha, a regulator ofEdwardsiella tarda, required for resistance to oxidative stress in macrophages. FEMS Microbiology Letters, 2016, 363, fnw192.	1.8	6
116	Mac Protein is not an Essential Virulence Factor for the Virulent Reference Strain Streptococcus suis P1/7. Current Microbiology, 2017, 74, 90-96.	2.2	6
117	SBP1 is an adhesion-associated factor without the involvement of virulence in Streptococcus suis serotype 2. Microbial Pathogenesis, 2018, 122, 90-97.	2.9	6
118	Comparative proteomic and genomic analyses of Brucella abortus biofilm and planktonic cells. Molecular Medicine Reports, 2020, 21, 731-743.	2.4	6
119	Characterization and complete genome sequence analysis of Staphylococcus aureus bacteriophage JS01. Virus Genes, 2015, 50, 345-348.	1.6	5
120	AroC, a chorismate synthase, is required for the formation of Edwardsiella tarda biofilms. Microbes and Infection, 2022, 24, 104955.	1.9	5
121	Coronavirus as an Agent of Neonatal Calf Diarrhea in a Chinese Dairy Cattle Farm. Zoonoses and Public Health, 1991, 38, 473-476.	1.4	4
122	Development and evaluation of a dot blot assay for rapid determination of invasion-associated gene ibeA directly in fresh bacteria cultures of E. coli. Folia Microbiologica, 2012, 57, 557-561.	2.3	4
123	Mutations in the C-terminal tail of NS1 protein facilitate the replication of classical swine H1N1 influenza A virus in mice. Folia Microbiologica, 2012, 57, 169-175.	2.3	4
124	Mitochondrial antiviral signaling adaptor mediated apoptosis in H3N2 swine influenza virus infection is inhibited by viral protein NS1 in vitro. Veterinary Immunology and Immunopathology, 2015, 165, 34-44.	1.2	3
125	Target genes directly regulated by Eha are required for Edwardsiella tarda survival within macrophages. Veterinary Microbiology, 2020, 247, 108739.	1.9	2
126	The TonB system in Aeromonas hydrophila NJ-35 is essential for MacA2B2 efflux pump-mediated macrolide resistance. Veterinary Research, 2021, 52, 63.	3.0	1

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127	Pre-absorbed Immunoproteomics: A Novel Method for the Detection of Bacterial Surface Proteins. Methods in Molecular Biology, 2013, 1061, 113-121.	0.9	1
128	Nonstructural proteins of Torque teno sus virus 2 from O2AUG: Prediction to experimental validation. Virus Research, 2013, 178, 272-280.	2.2	0
129	Identification and Detection of Serotype-Specific Genes: Effective Serotyping of Streptococcus suis. Current Clinical Microbiology Reports, 2017, 4, 29-35.	3.4	0
130	Transcriptional regulator XtgS is involved in iron transition and attenuates the virulence of Streptococcus agalactiae. Research in Veterinary Science, 2021, 138, 109-115.	1.9	0