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	AroC, a chorismate synthase, is required for the formation of Edwardsiella tarda biofilms. Microbes and Infection, 2022, 24, 104955.	1.9	5
2	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
3	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
4	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
5	CRISPR-dependent endogenous gene regulation is required for virulence in piscine Streptococcus agalactiae. Emerging Microbes and Infections, 2021, 10, 1-53.	6.5	7
6	The Novel Streptococcal Transcriptional Regulator XtgS Negatively Regulates Bacterial Virulence and Directly Represses PsePTranscription. Infection and Immunity, 2020, 88, .	2.2	13
7	Target genes directly regulated by Eha are required for Edwardsiella tarda survival within macrophages. Veterinary Microbiology, 2020, 247, 108739.	1.9	2
8	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
9	Isolation and characterization of bacteriophages against virulent Aeromonas hydrophila. BMC Microbiology, 2020, 20, 141.	3.3	43
10	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
11	Identification of a new effector-immunity pair of Aeromonas hydrophila type VI secretion system. Veterinary Research, 2020, 51, 71.	3.0	14
12	Comparative proteomic and genomic analyses of Brucella abortus biofilm and planktonic cells. Molecular Medicine Reports, 2020, 21, 731-743.	2.4	6
13	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
14	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
15	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
16	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
17	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
18	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

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19	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
20	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
21	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
22	cas9 Enhances Bacterial Virulence by Repressing the regR Transcriptional Regulator in Streptococcus agalactiae. Infection and Immunity, 2018, 86, .	2.2	48
23	Comparative genome analysis provides deep insights into Aeromonas hydrophila taxonomy and virulence-related factors. BMC Genomics, 2018, 19, 712.	2.8	26
24	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
25	Streptococcus suis synthesizes deoxyadenosine and adenosine by 5â€™-nucleotidase to dampen host immune responses. Virulence, 2018, 9, 1509-1520.	4.4	24
26	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
27	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
28	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
29	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
30	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
31	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
32	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
33	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
34	Identification and Detection of Serotype-Specific Genes: Effective Serotyping of Streptococcus suis. Current Clinical Microbiology Reports, 2017, 4, 29-35.	3.4	0
35	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
36	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

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37	The non-conserved region of MRP is involved in the virulence ofStreptococcus suisserotype 2. Virulence, 2017, 8, 1274-1289.	4.4	25
38	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
39	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
40	Alterations in gp37 Expand the Host Range of a T4-Like Phage. Applied and Environmental Microbiology, 2017, 83, .	3.1	41
41	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
42	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
43	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
44	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
45	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
46	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
47	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
48	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
49	Identification and Characterization of an Aeromonas hydrophila Oligopeptidase Gene pepF Negatively Related to Biofilm Formation. Frontiers in Microbiology, 2016, 7, 1497.	3.5	23
50	A Streptococcus suis LysM domain surface protein contributes to bacterial virulence. Veterinary Microbiology, 2016, 187, 64-69.	1.9	19
51	Eha, a regulator ofEdwardsiella tarda, required for resistance to oxidative stress in macrophages. FEMS Microbiology Letters, 2016, 363, fnw192.	1.8	6
52	AutA and AutR, Two Novel Global Transcriptional Regulators, Facilitate Avian Pathogenic Escherichia coli Infection. Scientific Reports, 2016, 6, 25085.	3.3	15
53	The role of regulator Eha in Edwardsiella tarda pathogenesis and virulence gene transcription. Microbial Pathogenesis, 2016, 95, 216-223.	2.9	10
54	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

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55	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
56	Isolation, genome sequencing and functional analysis of two T7-like coliphages of avian pathogenic Escherichia coli. Gene, 2016, 582, 47-58.	2.2	29
57	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
58	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
59	IbeR Facilitates Stress-Resistance, Invasion and Pathogenicity of Avian Pathogenic Escherichia coli. PLoS ONE, 2015, 10, e0119698.	2.5	10
60	Novel insights into the pathogenicity of epidemic Aeromonas hydrophila ST251 clones from comparative genomics. Scientific Reports, 2015, 5, 9833.	3.3	110
61	Novel Variant Serotype of Streptococcus suis Isolated from Piglets with Meningitis. Applied and Environmental Microbiology, 2015, 81, 976-985.	3.1	57
62	Antibacterial effect of porcine PTX3 against Streptococcus suis type 2 infection. Microbial Pathogenesis, 2015, 89, 128-139.	2.9	12
63	Establishment and characterization of a telomerase-immortalized canine bronchiolar epithelial cell line. Applied Microbiology and Biotechnology, 2015, 99, 9135-9146.	3.6	7
64	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
65	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
66	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
67	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
68	Monoclonal antibody specific to HA2 glycopeptide protects mice from H3N2 influenza virus infection. Veterinary Research, 2015, 46, 33.	3.0	8
69	Characterization and complete genome sequence analysis of Staphylococcus aureus bacteriophage JS01. Virus Genes, 2015, 50, 345-348.	1.6	5
70	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
71	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
72	Virulence genotyping and population analysis of Streptococcus suis serotype 2 isolates from China. Infection, Genetics and Evolution, 2015, 36, 483-489.	2.3	23

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73	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
74	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
75	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
76	Latest developments on Streptococcus suis: an emerging zoonotic pathogen: part 2. Future Microbiology, 2014, 9, 587-591.	2.0	26
77	Whole-Genome Sequence of Streptococcus suis Serotype 4 Reference Strain 6407. Genome Announcements, 2014, 2, .	0.8	6
78	The Streptococcus suis transcriptional landscape reveals adaptation mechanisms in pig blood and cerebrospinal fluid. Rna, 2014, 20, 882-898.	3.5	59
79	Identification of a virulence-related surface protein XF in piscine Streptococcus agalactiaeby pre-absorbed immunoproteomics. BMC Veterinary Research, 2014, 10, 259.	1.9	6
80	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
81	Comparative genomic analysis shows that Streptococcus suis meningitis isolate SC070731 contains a unique 105K genomic island. Gene, 2014, 535, 156-164.	2.2	45
82	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
83	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
84	Contribution of Eukaryotic-Type Serine/Threonine Kinase to Stress Response and Virulence of Streptococcus suis. PLoS ONE, 2014, 9, e91971.	2.5	40
85	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
86	Characterization of Streptococcus suis Isolates from Slaughter Swine. Current Microbiology, 2013, 66, 344-349.	2.2	21
87	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
88	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
89	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
90	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

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91	Analysis of synonymous codon usage patterns in torque teno sus virus 1 (TTSuV1). Archives of Virology, 2013, 158, 145-154.	2.1	41
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	Nonstructural proteins of Torque teno sus virus 2 from O2AUG: Prediction to experimental validation. Virus Research, 2013, 178, 272-280.	2.2	0
94	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
95	Immunoproteomics selection of cross-protective vaccine candidates from Riemerella anatipestifer serotypes 1 and 2. Veterinary Microbiology, 2013, 162, 850-857.	1.9	27
96	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
97	Identification of immunoreactive proteins ofStreptococcus agalactiaeisolated from cultured tilapia in China. Pathogens and Disease, 2013, 69, 223-231.	2.0	13
98	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
99	Pre-absorbed Immunoproteomics: A Novel Method for the Detection of Bacterial Surface Proteins. Methods in Molecular Biology, 2013, 1061, 113-121.	0.9	1
100	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
101	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
102	Development of Rapid Serotype-Specific PCR Assays for Eight Serotypes of Streptococcus suis. Journal of Clinical Microbiology, 2012, 50, 3329-3334.	3.9	22
103	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
104	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
105	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
106	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
107	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
108	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

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109	Immunoproteomic identification of 11 novel immunoreactive proteins ofRiemerella anatipestiferserotype 2. FEMS Immunology and Medical Microbiology, 2012, 65, 84-95.	2.7	14
110	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
111	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
112	Reduced virulence is an important characteristic of biofilm infection of Streptococcus suis. FEMS Microbiology Letters, 2011, 316, 36-43.	1.8	74
113	Genetic analysis of the capsular polysaccharide synthesis locus in 15 Streptococcus suis serotypes. FEMS Microbiology Letters, 2011, 324, 117-124.	1.8	20
114	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
115	The cps locus of Streptococcus suis serotype 16: Development of a serotype-specific PCR assay. Veterinary Microbiology, 2011, 153, 403-406.	1.9	14
116	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
117	Immunoproteomic assay of secreted proteins of Streptococcus suis serotype 9 with convalescent sera from pigs. Folia Microbiologica, 2011, 56, 423-430.	2.3	8
118	Immunoproteomic analysis of bacterial proteins of Actinobacillus pleuropneumoniae serotype 1. Proteome Science, 2011, 9, 32.	1.7	14
119	Effects of ibeA Deletion on Virulence and Biofilm Formation of Avian Pathogenic Escherichia coli. Infection and Immunity, 2011, 79, 279-287.	2.2	75
120	Pre-Absorbed Immunoproteomics: A Novel Method for the Detection of Streptococcus suis Surface Proteins. PLoS ONE, 2011, 6, e21234.	2.5	23
121	Transcriptome profiling of zebrafish infected with Streptococcus suis. Microbial Pathogenesis, 2010, 48, 178-187.	2.9	63
122	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
123	Biological activity and identification of a peptide inhibitor of LuxS fromStreptococcus suisserotype 2. FEMS Microbiology Letters, 2009, 294, 16-23.	1.8	35
124	Molecular characterization of the 9.36Âkb C-terminal region of canine coronavirus 1-71 strain. Virus Genes, 2008, 36, 491-497.	1.6	12
125	Immunoproteomic assay of surface proteins ofStreptococcus suisserotype 9. FEMS Immunology and Medical Microbiology, 2008, 53, 52-59.	2.7	66
126	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

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127	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
128	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
129	Canine Distemper Virus Causes Apoptosis of Vero Cells. Zoonoses and Public Health, 2000, 47, 183-190.	1.4	36
130	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