

# Stephan Schmitz-Esser

## List of Publications by Year in descending order

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124  
papers

4,591  
citations

87723

38  
h-index

114278

63  
g-index

131  
all docs

131  
docs citations

131  
times ranked

4905  
citing authors

#	ARTICLE	IF	CITATIONS
1	Illuminating the Evolutionary History of Chlamydiae. <i>Science</i> , 2004, 304, 728-730.	6.0	373
2	16S-23S rDNA intergenic spacer and 23S rDNA of anaerobic ammonium-oxidizing bacteria: implications for phylogeny and in situ detection. <i>Environmental Microbiology</i> , 2001, 3, 450-459.	1.8	227
3	Tn6188 - A Novel Transposon in <i>Listeria monocytogenes</i> Responsible for Tolerance to Benzalkonium Chloride. <i>PLoS ONE</i> , 2013, 8, e76835.	1.1	167
4	ATP/ADP Translocases: a Common Feature of Obligate Intracellular Amoebal Symbionts Related to Chlamydiae and Rickettsiae. <i>Journal of Bacteriology</i> , 2004, 186, 683-691.	1.0	162
5	The Genome of the Amoeba Symbiont <i>Candidatus</i> <i>Amoebophilus asiaticus</i> Reveals Common Mechanisms for Host Cell Interaction among Amoeba-Associated Bacteria. <i>Journal of Bacteriology</i> , 2010, 192, 1045-1057.	1.0	138
6	Comparative Genomics Suggests an Independent Origin of Cytoplasmic Incompatibility in <i>Cardinium hertigii</i> . <i>PLoS Genetics</i> , 2012, 8, e1003012.	1.5	135
7	Age-Related Differences in the Luminal and Mucosa-Associated Gut Microbiome of Broiler Chickens and Shifts Associated with <i>Campylobacter jejuni</i> Infection. <i>Frontiers in Cellular and Infection Microbiology</i> , 2016, 6, 154.	1.8	134
8	Grain-rich diets differently alter ruminal and colonic abundance of microbial populations and lipopolysaccharide in goats. <i>Anaerobe</i> , 2013, 20, 65-73.	1.0	121
9	Mucosa-Associated Bacterial Microbiome of the Gastrointestinal Tract of Weaned Pigs and Dynamics Linked to Dietary Calcium-Phosphorus. <i>PLoS ONE</i> , 2014, 9, e86950.	1.1	106
10	Environmental genomics reveals a functional chlorite dismutase in the nitrite-oxidizing bacterium <i>Candidatus</i> <i>Nitrospira defluvia</i> ™. <i>Environmental Microbiology</i> , 2008, 10, 3043-3056.	1.8	102
11	A candidate NAD <sup>+</sup> transporter in an intracellular bacterial symbiont related to Chlamydiae. <i>Nature</i> , 2004, 432, 622-625.	13.7	95
12	Diversity of Bacterial Endosymbionts of Environmental <i>Acanthamoeba</i> Isolates. <i>Applied and Environmental Microbiology</i> , 2008, 74, 5822-5831.	1.4	92
13	Age-associated microbiome shows the giant panda lives on hemicelluloses, not on cellulose. <i>ISME Journal</i> , 2018, 12, 1319-1328.	4.4	89
14	Genomes of sequence type 121 <i>Listeria monocytogenes</i> strains harbor highly conserved plasmids and prophages. <i>Frontiers in Microbiology</i> , 2015, 6, 380.	1.5	87
15	<i>Candidatus</i> <i>Thiobios zoothamnii</i> , an Ectosymbiotic Bacterium Covering the Giant Marine Ciliate <i>Zoothamnium niveum</i> . <i>Applied and Environmental Microbiology</i> , 2006, 72, 2014-2021.	1.4	84
16	The Plant Mitochondrial Carrier Family: Functional and Evolutionary Aspects. <i>Frontiers in Plant Science</i> , 2012, 3, 2.	1.7	83
17	Diatom plastids depend on nucleotide import from the cytosol. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 3621-3626.	3.3	80
18	Bacterial diversity of floor drain biofilms and drain waters in a <i>Listeria monocytogenes</i> contaminated food processing environment. <i>International Journal of Food Microbiology</i> , 2016, 223, 33-40.	2.1	79

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19	Pyrosequencing reveals diverse fecal microbiota in Simmental calves during early development. <i>Frontiers in Microbiology</i> , 2014, 5, 622.	1.5	78
20	The Plastidic Sugar Transporter pSuT Influences Flowering and Affects Cold Responses. <i>Plant Physiology</i> , 2019, 179, 569-587.	2.3	77
21	Transcriptome analysis of <i>Listeria monocytogenes</i> exposed to biocide stress reveals a multi-system response involving cell wall synthesis, sugar uptake, and motility. <i>Frontiers in Microbiology</i> , 2014, 5, 68.	1.5	75
22	Cultivation-independent analysis of microbial communities on Austrian raw milk hard cheese rinds. <i>International Journal of Food Microbiology</i> , 2014, 180, 88-97.	2.1	74
23	Recovery of an environmental chlamydia strain from activated sludge by co-cultivation with <i>Acanthamoeba</i> sp.. <i>Microbiology (United Kingdom)</i> , 2005, 151, 301-309.	0.7	73
24	Intestinal microbiota profiles associated with low and high residual feed intake in chickens across two geographical locations. <i>PLoS ONE</i> , 2017, 12, e0187766.	1.1	73
25	Tapping the nucleotide pool of the host: novel nucleotide carrier proteins of <i>Protochlamydia amoebophila</i> . <i>Molecular Microbiology</i> , 2006, 60, 1534-1545.	1.2	69
26	The <i>Listeria monocytogenes</i> transposon Tn6188 provides increased tolerance to various quaternary ammonium compounds and ethidium bromide. <i>FEMS Microbiology Letters</i> , 2014, 361, 166-173.	0.7	68
27	Changing Dietary Calcium-Phosphorus Level and Cereal Source Selectively Alters Abundance of Bacteria and Metabolites in the Upper Gastrointestinal Tracts of Weaned Pigs. <i>Applied and Environmental Microbiology</i> , 2013, 79, 7264-7272.	1.4	64
28	Fluoxetine-induced alteration of murine gut microbial community structure: evidence for a microbial endocrinology-based mechanism of action responsible for fluoxetine-induced side effects. <i>PeerJ</i> , 2019, 7, e6199.	0.9	62
29	Microbiota of newborn calves and their mothers reveals possible transfer routes for newborn calves' gastrointestinal microbiota. <i>PLoS ONE</i> , 2019, 14, e0220554.	1.1	61
30	Epimural bacterial community structure in the rumen of Holstein cows with different responses to a long-term subacute ruminal acidosis diet challenge. <i>Journal of Dairy Science</i> , 2017, 100, 1829-1844.	1.4	60
31	Adaptation of the Cecal Bacterial Microbiome of Growing Pigs in Response to Resistant Starch Type 4. <i>Applied and Environmental Microbiology</i> , 2015, 81, 8489-8499.	1.4	59
32	Genome Sequencing of <i>Listeria monocytogenes</i> "Quargel" Listeriosis Outbreak Strains Reveals Two Different Strains with Distinct In Vitro Virulence Potential. <i>PLoS ONE</i> , 2014, 9, e89964.	1.1	55
33	Metatranscriptome Sequencing Reveals Insights into the Gene Expression and Functional Potential of Rumen Wall Bacteria. <i>Frontiers in Microbiology</i> , 2018, 9, 43.	1.5	54
34	Plasmids contribute to food processing environment-associated stress survival in three <i>Listeria monocytogenes</i> ST121, ST8, and ST5 strains. <i>International Journal of Food Microbiology</i> , 2019, 299, 39-46.	2.1	52
35	Bacteriocyte-associated gammaproteobacterial symbionts of the <i>Adelges nordmannianae/piceae</i> complex (Hemiptera: Adelgidae). <i>ISME Journal</i> , 2012, 6, 384-396.	4.4	49
36	Pyrosequencing reveals shifts in the bacterial epimural community relative to dietary concentrate amount in goats. <i>Journal of Dairy Science</i> , 2015, 98, 5572-5587.	1.4	46

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37	New Aspects on <i>Listeria monocytogenes</i> ST5-ECVI Predominance in a Heavily Contaminated Cheese Processing Environment. <i>Frontiers in Microbiology</i> , 2018, 9, 64.	1.5	45
38	Comparative genomics of human and non-human <i>Listeria monocytogenes</i> sequence type 121 strains. <i>PLoS ONE</i> , 2017, 12, e0176857.	1.1	42
39	<i>Lawsonia intracellularis</i> Contains a Gene Encoding a Functional Rickettsia-Like ATP/ADP Translocase for Host Exploitation. <i>Journal of Bacteriology</i> , 2008, 190, 5746-5752.	1.0	37
40	High genetic similarity between two geographically distinct strains of the sulfur-oxidizing symbiont <i>Candidatus Thiobios zoothamnii</i> . <i>FEMS Microbiology Ecology</i> , 2009, 67, 229-241.	1.3	35
41	Transcriptome Sequencing Reveals Novel Candidate Genes for <i>Cardinium hertigii</i> -Caused Cytoplasmic Incompatibility and Host-Cell Interaction. <i>MSystems</i> , 2017, 2, .	1.7	35
42	<i>Brevibacterium</i> from Austrian hard cheese harbor a putative histamine catabolism pathway and a plasmid for adaptation to the cheese environment. <i>Scientific Reports</i> , 2019, 9, 6164.	1.6	35
43	Epimural Indicator Phylotypes of Transiently-Induced Subacute Ruminal Acidosis in Dairy Cattle. <i>Frontiers in Microbiology</i> , 2016, 7, 274.	1.5	34
44	Biphasic Metabolism and Host Interaction of a Chlamydial Symbiont. <i>MSystems</i> , 2017, 2, .	1.7	32
45	Microbial shifts in Minas artisanal cheeses from the Serra do Salitre region of Minas Gerais, Brazil throughout ripening time. <i>Food Microbiology</i> , 2019, 82, 349-362.	2.1	32
46	The genome of the amoeba symbiont " <i>Candidatus Amoebophilus asiaticus</i> " encodes an <i>afp</i> -like prophage possibly used for protein secretion. <i>Virulence</i> , 2010, 1, 541-545.	1.8	31
47	Autochthonous facility-specific microbiota dominates washed-rind Austrian hard cheese surfaces and its production environment. <i>International Journal of Food Microbiology</i> , 2018, 267, 54-61.	2.1	31
48	Transcriptome Sequencing of <i>Listeria monocytogenes</i> Reveals Major Gene Expression Changes in Response to Lactic Acid Stress Exposure but a Less Pronounced Response to Oxidative Stress. <i>Frontiers in Microbiology</i> , 2019, 10, 3110.	1.5	31
49	Exoproteome analysis reveals higher abundance of proteins linked to alkaline stress in persistent <i>Listeria monocytogenes</i> strains. <i>International Journal of Food Microbiology</i> , 2016, 218, 17-26.	2.1	30
50	Nucleotide Parasitism by <i>Simkania negevensis</i> ( <i>Chlamydiae</i> ). <i>Journal of Bacteriology</i> , 2011, 193, 225-235.	1.0	27
51	Microbiomes of Unreactive and Pathologically Altered Ileocecal Lymph Nodes of Slaughter Pigs. <i>Applied and Environmental Microbiology</i> , 2014, 80, 193-203.	1.4	27
52	Psychrophile spoilers dominate the bacterial microbiome in musculature samples of slaughter pigs. <i>Meat Science</i> , 2016, 117, 36-40.	2.7	27
53	Symbiont-driven sulfur crystal formation in a thiotrophic symbiosis from deep-sea hydrocarbon seeps. <i>Environmental Microbiology Reports</i> , 2014, 6, 364-372.	1.0	26
54	Microbial communities in dairy processing environment floor-drains are dominated by product-associated bacteria and yeasts. <i>Food Control</i> , 2016, 70, 210-215.	2.8	26

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55	Virulence characterization and comparative genomics of <i>Listeria monocytogenes</i> sequence type 155 strains. <i>BMC Genomics</i> , 2020, 21, 847.	1.2	26
56	The Metabolically Active Bacterial Microbiome of Tonsils and Mandibular Lymph Nodes of Slaughter Pigs. <i>Frontiers in Microbiology</i> , 2015, 6, 1362.	1.5	23
57	Rumen Epithelial Communities Share a Core Bacterial Microbiota: A Meta-Analysis of 16S rRNA Gene Illumina MiSeq Sequencing Datasets. <i>Frontiers in Microbiology</i> , 2021, 12, 625400.	1.5	23
58	A bacterial genome in transition - an exceptional enrichment of IS elements but lack of evidence for recent transposition in the symbiont <i>Amoebophilus asiaticus</i> . <i>BMC Evolutionary Biology</i> , 2011, 11, 270.	3.2	22
59	Comparison of intestinal permeability, morphology, and ileal microbial communities of commercial hens housed in conventional cages and cage-free housing systems. <i>Poultry Science</i> , 2021, 100, 1178-1191.	1.5	21
60	A Large-Scale Sequencing-Based Survey of Plasmids in <i>Listeria monocytogenes</i> Reveals Global Dissemination of Plasmids. <i>Frontiers in Microbiology</i> , 2021, 12, 653155.	1.5	21
61	High diversity of viable bacteria isolated from lymph nodes of slaughter pigs and its possible impacts for food safety. <i>Journal of Applied Microbiology</i> , 2015, 119, 1420-1432.	1.4	20
62	Shuttling of (deoxyâ€) purine nucleotides between compartments of the diatom <i>Phaeodactylum tricornutum</i>. <i>New Phytologist</i> , 2017, 213, 193-205.	3.5	20
63	Complete Genome Sequence of the Persistent <i>Listeria monocytogenes</i> Strain R479a. <i>Genome Announcements</i> , 2015, 3, .	0.8	18
64	Abundance and potential contribution of Gram-negative cheese rind bacteria from Austrian artisanal hard cheeses. <i>International Journal of Food Microbiology</i> , 2018, 266, 95-103.	2.1	17
65	The transcriptome of <i>Listeria monocytogenes</i> during co-cultivation with cheese rind bacteria suggests adaptation by induction of ethanolamine and 1,2-propanediol catabolism pathway genes. <i>PLoS ONE</i> , 2020, 15, e0233945.	1.1	17
66	Vaginal microbiota diverges in sows with low and high reproductive performance after porcine reproductive and respiratory syndrome vaccination. <i>Scientific Reports</i> , 2020, 10, 3046.	1.6	17
67	Xylanase modulates the microbiota of ileal mucosa and digesta of pigs fed corn-based arabinoxylans likely through both a stimbiotic and prebiotic mechanism. <i>PLoS ONE</i> , 2021, 16, e0246144.	1.1	17
68	Investigating the relationship between vaginal microbiota and host genetics and their impact on immune response and farrowing traits in commercial gilts. <i>Journal of Animal Breeding and Genetics</i> , 2020, 137, 84-102.	0.8	16
69	Beef cattle that respond differently to fescue toxicosis have distinct gastrointestinal tract microbiota. <i>PLoS ONE</i> , 2020, 15, e0229192.	1.1	16
70	Characterization of mucosa-associated bacterial communities in abomasal ulcers by pyrosequencing. <i>Veterinary Microbiology</i> , 2015, 177, 132-141.	0.8	15
71	Xylanase Supplementation Modulates the Microbiota of the Large Intestine of Pigs Fed Corn-Based Fiber by Means of a Stimbiotic Mechanism of Action. <i>Frontiers in Microbiology</i> , 2021, 12, 619970.	1.5	14
72	Dietary calcium concentration and cereals differentially affect mineral balance and tight junction proteins expression in jejunum of weaned pigs. <i>British Journal of Nutrition</i> , 2015, 113, 1019-1031.	1.2	13

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73	Austrian Raw-Milk Hard-Cheese Ripening Involves Successional Dynamics of Non-Inoculated Bacteria and Fungi. <i>Foods</i> , 2020, 9, 1851.	1.9	13
74	A clinical <i>Acanthamoeba</i> isolate harboring two distinct bacterial endosymbionts. <i>European Journal of Protistology</i> , 2016, 56, 21-25.	0.5	12
75	Development of a multi-locus sequence typing system helps reveal the evolution of <i>Cardinium hertigii</i> , a reproductive manipulator symbiont of insects. <i>BMC Microbiology</i> , 2019, 19, 266.	1.3	12
76	Ewe Vaginal Microbiota: Associations With Pregnancy Outcome and Changes During Gestation. <i>Frontiers in Microbiology</i> , 2021, 12, 745884.	1.5	12
77	Microbiota of the Gut-Lymph Node Axis: Depletion of Mucosa-Associated Segmented Filamentous Bacteria and Enrichment of <i>Methanobrevibacter</i> by Colistin Sulfate and Linco-Spectin in Pigs. <i>Frontiers in Microbiology</i> , 2019, 10, 599.	1.5	11
78	The Cobalamin-Dependent Gene Cluster of <i>Listeria monocytogenes</i> : Implications for Virulence, Stress Response, and Food Safety. <i>Frontiers in Microbiology</i> , 2020, 11, 601816.	1.5	11
79	The Endosymbiont <i>Amoebophilus asiaticus</i> Encodes an <i>S</i> -Adenosylmethionine Carrier That Compensates for Its Missing Methylation Cycle. <i>Journal of Bacteriology</i> , 2013, 195, 3183-3192.	1.0	9
80	High-Fat Diets Led to OTU-Level Shifts in Fecal Samples of Healthy Adult Dogs. <i>Frontiers in Microbiology</i> , 2020, 11, 564160.	1.5	9
81	<i>Cardinium</i> Localization During Its Parasitoid Wasp Host's Development Provides Insights Into Cytoplasmic Incompatibility. <i>Frontiers in Microbiology</i> , 2020, 11, 606399.	1.5	9
82	Vaginal microbiome and serum metabolite differences in late gestation commercial sows at risk for pelvic organ prolapse. <i>Scientific Reports</i> , 2021, 11, 6189.	1.6	9
83	Exogenous carbohydrases added to a starter diet reduced markers of systemic immune activation and decreased <i>Lactobacillus</i> in weaned pigs. <i>Journal of Animal Science</i> , 2019, 97, 1242-1253.	0.2	8
84	Beginning to offer drinking water at birth increases the species richness and the abundance of <i>Faecalibacterium</i> and <i>Bifidobacterium</i> in the gut of preweaned dairy calves. <i>Journal of Dairy Science</i> , 2020, 103, 4262-4274.	1.4	8
85	Vaginal microbiota differences associated with pelvic organ prolapse risk during late gestation in commercial sows. <i>Biology of Reproduction</i> , 2021, 105, 1545-1561.	1.2	6
86	Certain <i>Listeria monocytogenes</i> plasmids contribute to increased UVC ultraviolet light stress. <i>FEMS Microbiology Letters</i> , 2021, 368, .	0.7	5
87	Dietary Supplementation with Sugar Beet Fructooligosaccharides and Garlic Residues Promotes Growth of Beneficial Bacteria and Increases Weight Gain in Neonatal Lambs. <i>Biomolecules</i> , 2020, 10, 1179.	1.8	4
88	Quality over quantity: unraveling the contributions to cytoplasmic incompatibility caused by two coinfecting <i>Cardinium</i> symbionts. <i>Heredity</i> , 2022, , .	1.2	4
89	Metatranscriptomic Analyses Unravel Dynamic Changes in the Microbial and Metabolic Transcriptional Profiles in Artisanal Austrian Hard-Cheeses During Ripening. <i>Frontiers in Microbiology</i> , 2022, 13, 813480.	1.5	4
90	The validation of the existence of the entero-mammary pathway and the assessment of the differences of the pathway between first and third parity sows. <i>Translational Animal Science</i> , 2022, 6, .	0.4	4

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91	The Rumen Epithelial Microbiota: Possible Gatekeepers of the Rumen Epithelium and Its Potential Contributions to Epithelial Barrier Function and Animal Health and Performance. <i>Meat and Muscle Biology</i> , 2020, 4, .	0.7	3
92	Host immunity and the colon microbiota of mice infected with <i>Citrobacter rodentium</i> are beneficially modulated by lipid-soluble extract from late-cutting alfalfa in the early stages of infection. <i>PLoS ONE</i> , 2020, 15, e0236106.	1.1	2
93	Genome Sequencing of <i>Listeria monocytogenes</i> . <i>Methods in Molecular Biology</i> , 2014, 1157, 223-232.	0.4	2
94	Genome Sequences of Soft Rot-Causing <i>Pectobacterium</i> Isolates from Different Vegetables. <i>Microbiology Resource Announcements</i> , 2022, 11, e0106621.	0.3	2
95	PSII-16 Evidence for stratification of rumen wall microbial communities revealed by 16S rRNA based amplicon sequencing. <i>Journal of Animal Science</i> , 2019, 97, 226-227.	0.2	1
96	Virulence Pattern Analysis of Three <i>Listeria monocytogenes</i> Lineage I Epidemic Strains with Distinct Outbreak Histories. <i>Microorganisms</i> , 2021, 9, 1745.	1.6	1
97	PSVIII-3 Can the fermentation of insoluble corn fiber be improved in the pig? An investigation into the in vivo mode of action of xylanase. <i>Journal of Animal Science</i> , 2020, 98, 207-207.	0.2	1
98	The life cycle-dependent transcriptional profile of the obligate intracellular amoeba symbiont <i>Amoebophilus asiaticus</i> . <i>FEMS Microbiology Ecology</i> , 2022, 98, .	1.3	1
99	Cheese and Microbes. Edited by Catherine W. Donnelly. Washington (DC): ASM Press. \$150.00. xii + 333 p.; ill.; index. ISBN: 978-1-55581-586-8 (hc); 978-1-55581-859-3 (eb). 2014.. <i>Quarterly Review of Biology</i> , 2015, 90, 346-346.	0.0	0
100	385 Evidence for Stratification of Rumen Wall Microbial Communities Revealed By 16S rRNA Based Amplicon Sequencing.. <i>Journal of Animal Science</i> , 2018, 96, 207-207.	0.2	0
101	23 Effect of Genetic Response to Endophyte-Infected Fescue on Beef Cattle Gastrointestinal Tract Microbiota.. <i>Journal of Animal Science</i> , 2018, 96, 12-13.	0.2	0
102	300 Impact of Fiber-Degrading Enzymes on Microbial Composition and NSP Metabolites in Nursery Pigs Fed a Higher Fiber Diet.. <i>Journal of Animal Science</i> , 2018, 96, 161-162.	0.2	0
103	27 Composition and in Situ Gene Expression of Rumen Wall Microbial Communities.. <i>Journal of Animal Science</i> , 2018, 96, 14-15.	0.2	0
104	77 Responses to alfalfa supplementation in mice. <i>Journal of Animal Science</i> , 2019, 97, 45-46.	0.2	0
105	130 Young Scholar Presentation: Can exogenous carbohydrase supplementation to higher-fiber diets improve gut function, microbiota, and growth performance of weaned pigs?. <i>Journal of Animal Science</i> , 2019, 97, 76-76.	0.2	0
106	69 The ability of an artificial sweetener (Sucram®) to influence microbial community structure in the rumen papillae and content through the production of microbial-based neurochemicals. <i>Journal of Animal Science</i> , 2019, 97, 100-101.	0.2	0
107	219 Relationship between host-genetics and the vaginal microbiome in commercial gilts. <i>Journal of Animal Science</i> , 2019, 97, 43-44.	0.2	0
108	20 Vaginal microbiome composition is associated with sow longevity. <i>Journal of Animal Science</i> , 2020, 98, 23-24.	0.2	0

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109	399 Towards improving pig livability: Strategies to understand and mitigate sow mortality. Journal of Animal Science, 2020, 98, 183-184.	0.2	0
110	185 The Validation of the Existence of the Entero-Mammary Pathway and the Assessment of the Differences of the Pathway Between Gilts and Sows. Journal of Animal Science, 2022, 100, 88-88.	0.2	0
111	Title is missing!. , 2020, 15, e0229192.		0
112	Title is missing!. , 2020, 15, e0229192.		0
113	Title is missing!. , 2020, 15, e0229192.		0
114	Title is missing!. , 2020, 15, e0229192.		0
115	Title is missing!. , 2020, 15, e0229192.		0
116	Title is missing!. , 2020, 15, e0229192.		0
117	Title is missing!. , 2020, 15, e0236106.		0
118	Title is missing!. , 2020, 15, e0236106.		0
119	Title is missing!. , 2020, 15, e0236106.		0
120	Title is missing!. , 2020, 15, e0236106.		0
121	Title is missing!. , 2020, 15, e0233945.		0
122	Title is missing!. , 2020, 15, e0233945.		0
123	Title is missing!. , 2020, 15, e0233945.		0
124	Title is missing!. , 2020, 15, e0233945.		0