

# Bo Zeng

## List of Publications by Year in descending order

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Version: 2024-02-01

43  
papers

1,251  
citations

471061

17  
h-index

395343

33  
g-index

44  
all docs

44  
docs citations

44  
times ranked

1669  
citing authors

#	ARTICLE	IF	CITATIONS
1	Dynamic transcriptome and chromatin architecture in granulosa cells during chicken folliculogenesis. <i>Nature Communications</i> , 2022, 13, 131.	5.8	24
2	Corrigendum to "Transcriptome Profiling across Five Tissues of Giant Panda". <i>BioMed Research International</i> , 2022, 2022, 1-1.	0.9	0
3	Effects of Dietary Alteration on the Gut Microbiome and Metabolome of the Rescued Bengal Slow Loris. <i>Frontiers in Microbiology</i> , 2021, 12, 650991.	1.5	6
4	Gut microbiota in healthy and unhealthy long-living people. <i>Gene</i> , 2021, 779, 145510.	1.0	17
5	A pig BodyMap transcriptome reveals diverse tissue physiologies and evolutionary dynamics of transcription. <i>Nature Communications</i> , 2021, 12, 3715.	5.8	60
6	Dihydromyricetin promotes longevity and activates the transcription factors FOXO and AOP in <i>Drosophila</i> . <i>Aging</i> , 2021, 13, 460-476.	1.4	15
7	Effects of anti-aging interventions on intestinal microbiota. <i>Gut Microbes</i> , 2021, 13, 1994835.	4.3	32
8	Inbreeding Alters the Gut Microbiota of the Banna Minipig. <i>Animals</i> , 2020, 10, 2125.	1.0	8
9	Transcriptome Profiling across Five Tissues of Giant Panda. <i>BioMed Research International</i> , 2020, 2020, 1-13.	0.9	8
10	The near complete mitochondrial genome of <i>Oreolalax schmidtii</i> (Anura: Megophryidae). <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 3536-3537.	0.2	1
11	AFB1 Induced Transcriptional Regulation Related to Apoptosis and Lipid Metabolism in Liver of Chicken. <i>Toxins</i> , 2020, 12, 290.	1.5	32
12	The Use of RNAi Technology to Interfere with Zfx Gene Increases the Male Rates of Red Deer ( <i>Cervus</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 059		1
13	The carnivorous digestive system and bamboo diet of giant pandas may shape their low gut bacterial diversity. , 2020, 8, coz104.		17
14	The complete mitogenome of the large toothed toad, <i>Oreolalax major</i> (Anura: Megophryidae) with phylogenetic analysis. <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 1117-1118.	0.2	3
15	Gut microbiota of Tibetans and Tibetan pigs varies between high and low altitude environments. <i>Microbiological Research</i> , 2020, 235, 126447.	2.5	39
16	Gut microbiota in reintroduction of giant panda. <i>Ecology and Evolution</i> , 2020, 10, 1012-1028.	0.8	18
17	The Regulatory Functions of Circular RNAs in Digestive System Cancers. <i>Cancers</i> , 2020, 12, 770.	1.7	18
18	Variation in Gut Microbiota of Captive Bengal Slow Lorises. <i>Current Microbiology</i> , 2020, 77, 2623-2632.	1.0	5

#	ARTICLE	IF	CITATIONS
19	Transplant of microbiota from long-living people to mice reduces aging-related indices and transfers beneficial bacteria. <i>Aging</i> , 2020, 12, 4778-4793.	1.4	38
20	Characterization of the Rumen Microbiota and Volatile Fatty Acid Profiles of Weaned Goat Kids under Shrub-Grassland Grazing and Indoor Feeding. <i>Animals</i> , 2020, 10, 176.	1.0	14
21	Deubiquitinase USP7 regulates aging through ubiquitination and autophagy. <i>Aging</i> , 2020, 12, 23082-23095.	1.4	1
22	The Complete Mitochondrial Genome of <i>Platysternon megacephalum peguense</i> and Molecular Phylogenetic Analysis. <i>Genes</i> , 2019, 10, 487.	1.0	11
23	The complete mitogenome of the splendid japalure <i>Japalura splendida</i> (Squamata, Agamidae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 2641-2642.	0.2	3
24	The complete mitogenome of the granular torrent frog, <i>Amolops granulosus</i> (Anura: Ranidae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 2643-2644.	0.2	1
25	Comparative Study of Gut Microbiota in Wild and Captive Giant Pandas ( <i>Ailuropoda melanoleuca</i> ). <i>Genes</i> , 2019, 10, 827.	1.0	69
26	Complete mitochondrial genome of the webbed-toed gecko <i>Gekko subpalmatus</i> (Squamata: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	0.2	0
27	The complete mitochondrial genome of the tree frog, <i>Polypedates braueri</i> (Anura,) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 5	0.2	4
28	Genetic diversity of chemokine XCL1 and its receptor XCR1 in murine rodents. <i>Developmental and Comparative Immunology</i> , 2019, 98, 80-88.	1.0	3
29	The revised complete mitogenome sequence of the tree frog <i>Polypedatesmegacephalus</i> (Anura,) Tj ETQq1 1 0.784314 rgBT /Over	0.9	5
30	Diarrhea-Associated Intestinal Microbiota in Captive Sichuan Golden Snub-Nosed Monkeys (&lt;i>Rhinopithecus roxellana&lt;/i>). <i>Microbes and Environments</i> , 2018, 33, 249-256.	0.7	14
31	Metagenomic Study Suggests That the Gut Microbiota of the Giant Panda ( <i>Ailuropoda melanoleuca</i> ) May Not Be Specialized for Fiber Fermentation. <i>Frontiers in Microbiology</i> , 2018, 9, 229.	1.5	70
32	Microbiome of Total Versus Live Bacteria in the Gut of Rex Rabbits. <i>Frontiers in Microbiology</i> , 2018, 9, 733.	1.5	30
33	Conservation implications of primate trade in China over 18 years based on web news reports of confiscations. <i>PeerJ</i> , 2018, 6, e6069.	0.9	18
34	Correlations between gut microbiota community structures of Tibetans and geography. <i>Scientific Reports</i> , 2017, 7, 16982.	1.6	65
35	High-Altitude Living Shapes the Skin Microbiome in Humans and Pigs. <i>Frontiers in Microbiology</i> , 2017, 8, 1929.	1.5	25
36	Gut microbiota signatures of longevity. <i>Current Biology</i> , 2016, 26, R832-R833.	1.8	265

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37	Molecular Evolution of PTEN Pseudogenes in Mammals. PLoS ONE, 2016, 11, e0167851.	1.1	23
38	The evolution of the gut microbiota in the giant and the red pandas. Scientific Reports, 2015, 5, 10185.	1.6	71
39	The primary structure of COMT gene is not involved in the diet shift of the giant or the red pandas. Gene, 2015, 562, 244-246.	1.0	19
40	The bacterial communities associated with fecal types and body weight of rex rabbits. Scientific Reports, 2015, 5, 9342.	1.6	115
41	Characterization of the Gut Microbiota in the Red Panda ( <i>Ailurus fulgens</i> ). PLoS ONE, 2014, 9, e87885.	1.1	70
42	Alterations in cecal microbiota of Jinhua piglets fostered by a Yorkshire sow. Science Bulletin, 2014, 59, 4304-4311.	1.7	9
43	Gut Microbiota Composition and Metabolic Potential of Long-Living People in China. Frontiers in Aging Neuroscience, 0, 14, .	1.7	2