Elizabeth K Mallott

List of Publications by Year
in descending order

28 papers	589 citations	687363 13 h-index	677142 22 g-index
30 all docs	30 docs citations	30 times ranked	862 citing authors

#	Article	IF	Citations
1	Butyrate Production Pathway Abundances Are Similar in Human and Nonhuman Primate Gut Microbiomes. Molecular Biology and Evolution, 2022, 39, .	8.9	13
2	Ancient and modern genomics of the Ohlone Indigenous population of California. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e2111533119.	7.1	10
3	The faecal metabolome of black howler monkeys (Alouatta pigra) varies in response to seasonal dietary changes. Molecular Ecology, 2022, 31, 4146-4161.	3.9	4
4	Effects of anthropogenic habitat disturbance and Giardia duodenalis infection on a sentinel species' gut bacteria. Ecology and Evolution, 2021, 11, 45-57.	1.9	3
5	Fermented food consumption in wild nonhuman primates and its ecological drivers. American Journal of Physical Anthropology, 2021, 175, 513-530.	2.1	16
6	Host specificity of the gut microbiome. Nature Reviews Microbiology, 2021, 19, 639-653.	28.6	77
7	Predigestion as an Evolutionary Impetus for Human Use of Fermented Food. Current Anthropology, 2021, 62, S207-S219.	1.6	22
8	The relationship between pinworm (Trypanoxyuris) infection and gut bacteria in wild black howler monkeys (Alouatta pigra). American Journal of Primatology, 2021, 83, e23330.	1.7	7
9	Phylosymbiosis, diet and gut microbiome-associated metabolic disease. Evolution, Medicine and Public Health, 2020, 2020, 100-101.	2.5	1
10	Reproductive hormones mediate changes in the gut microbiome during pregnancy and lactation in Phayreâ€™s leaf monkeys. Scientific Reports, 2020, 10, 9961.	3.3	44
11	Convergence of human and Old World monkey gut microbiomes demonstrates the importance of human ecology over phylogeny. Genome Biology, 2019, 20, 201.	8.8	57
12	The effect of captivity on the primate gut microbiome varies with host dietary niche. American Journal of Primatology, 2019, 81, e23061.	1.7	56
13	Plasticity in the Human Gut Microbiome Defies Evolutionary Constraints. MSphere, 2019, 4, .	2.9	40
14	The gut microbiome and metabolome of saddleback tamarins (Leontocebus weddelli): Insights into the foraging ecology of a smallâ€bodied primate. American Journal of Primatology, 2019, 81, e23003.	1.7	10
15	Assessing the comparability of different DNA extraction and amplification methods in gut microbial community profiling. Access Microbiology, 2019, 1, e000060.	0.5	10
16	Species identification and mitochondrial genomes of ancient fish bones from the Riverine Kachemak tradition of the Kenai Peninsula, Alaska. Mitochondrial DNA Part B: Resources, 2018, 3, 409-411.	0.4	9
17	Influence of fruit and invertebrate consumption on the gut microbiota of wild whiteâ€faced capuchins (Cebus capucinus). American Journal of Physical Anthropology, 2018, 165, 576-588.	2.1	36
18	Patterns of Genetic Coding Variation in a Native American Population before and after European Contact. American Journal of Human Genetics, 2018, 102, 806-815.	6.2	33

#	Article	IF	Citations
19	trnL outperforms rbcL as a DNA metabarcoding marker when compared with the observed plant component of the diet of wild white-faced capuchins (Cebus capucinus, Primates). PLoS ONE, 2018, 13, e0199556.	2.5	32
20	The microbial reproductive ecology of whiteâ€faced capuchins (Cebus capucinus). American Journal of Primatology, 2018, 80, e22896.	1.7	36
21	Locus. , 2018, , 1-2.		0
22	Somatic Cells. , 2018, , 1-2.		0
23	Integrating feeding behavior, ecological data, and DNA barcoding to identify developmental differences in invertebrate foraging strategies in wild whiteâ€faced capuchins (Cebus capucinus). American Journal of Physical Anthropology, 2017, 162, 241-254.	2.1	25
24	Complete Mitochondrial Genome Sequencing of a Burial from a Romanoâ€“Christian Cemetery in the Dakhleh Oasis, Egypt: Preliminary Indications. Genes, 2017, 8, 262.	2.4	14
25	ANÃLISIS DEL GENOMA MITOCONDRIAL DE DOS INDIVIDUOS INHUMADOS EN EL SITIO ARQUEOLÃ“GICO CG14E01 â€œISLA LARGAâ€•(ROCHA, URUGUAY). Revista Argentina De Antropologia Biologica, 2017, 19, 17.	0.4	4
26	Highâ€throughput sequencing of fecal <scp>DNA</scp> to identify insects consumed by wild <scp>W</scp>eddell's saddleback tamarins (<scp>S</scp>aguinus weddelli,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 462 T Anthropology, 2015, 156, 474-481.	2.1	27
27	The ecology of trunk-to-trunk leaping in <i>Saguinus fuscicollis</i>: implications for understanding locomotor diversity in Callitrichines. Neotropical Primates, 2012, 19, 1-7.	0.1	2
28	Environmental Stress and the Primate Microbiome: Glucocorticoids Contribute to Structure Gut Bacterial Communities of Black Howler Monkeys in Anthropogenically Disturbed Forest Fragments. Frontiers in Ecology and Evolution, 0, 10, .	2.2	1