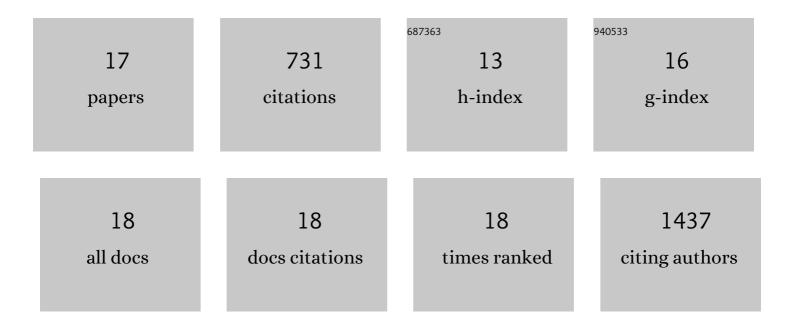
## Lesley A Ogilvie

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2295778/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	The human gut virome: a multifaceted majority. Frontiers in Microbiology, 2015, 6, 918.	3.5	171
2	Effects of selective digestive decontamination (SDD) on the gut resistome. Journal of Antimicrobial Chemotherapy, 2014, 69, 2215-2223.	3.0	90
3	Genome signature-based dissection of human gut metagenomes to extract subliminal viral sequences. Nature Communications, 2013, 4, 2420.	12.8	76
4	Bacterial Diversity of the Broadbalk â€~Classical' Winter Wheat Experiment in Relation to Long-Term Fertilizer Inputs. Microbial Ecology, 2008, 56, 525-537.	2.8	62
5	Comparative (Meta)genomic Analysis and Ecological Profiling of Human Gut-Specific Bacteriophage φB124-14. PLoS ONE, 2012, 7, e35053.	2.5	55
6	Dysbiosis modulates capacity for bile acid modification in the gut microbiomes of patients with inflammatory bowel disease: a mechanism and marker of disease?: Figure 1. Gut, 2012, 61, 1642-1643.	12.1	54
7	Identification of Aminoglycoside and β-Lactam Resistance Genes from within an Infant Gut Functional Metagenomic Library. PLoS ONE, 2014, 9, e108016.	2.5	48
8	High-throughput and single-cell imaging of NF-κB oscillations using monoclonal cell lines. BMC Cell Biology, 2010, 11, 21.	3.0	44
9	Evolutionary, ecological and biotechnological perspectives on plasmids resident in the human gut mobile metagenome. Bioengineered, 2012, 3, 13-31.	3.2	27
10	Activation of NF-κB by Neisseria gonorrhoeae is associated with microcolony formation and type Ⅳ pilus retraction. Cellular Microbiology, 2011, 13, 1168-1182.	2.1	25
11	Disruption of Escherichia coli Nissle 1917 K5 Capsule Biosynthesis, through Loss of Distinct kfi genes, Modulates Interaction with Intestinal Epithelial Cells and Impact on Cell Health. PLoS ONE, 2015, 10, e0120430.	2.5	22
12	Resolution of habitat-associated ecogenomic signatures in bacteriophage genomes and application to microbial source tracking. ISME Journal, 2018, 12, 942-958.	9.8	19
13	The potential of nanoflow liquid chromatography-nano electrospray ionisation-mass spectrometry for global profiling the faecal metabolome. Journal of Chromatography A, 2019, 1600, 127-136.	3.7	18
14	Genomic and Ecogenomic Characterization of Proteus mirabilis Bacteriophages. Frontiers in Microbiology, 2019, 10, 1783.	3.5	13
15	The human gut virome: form and function. Emerging Topics in Life Sciences, 2017, 1, 351-362.	2.6	5
16	Revisiting enterotypes: a viral perspective. Future Virology, 2017, 12, 391-394.	1.8	1
17	The Human Gut Mobile Metagenome: A Metazoan Perspective. , 2013, , 1-14.		1