

# Soo Ching Lee

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5007142/publications.pdf>

Version: 2024-02-01

20  
papers

1,179  
citations

687220

13  
h-index

794469

19  
g-index

23  
all docs

23  
docs citations

23  
times ranked

2148  
citing authors

#	ARTICLE	IF	CITATIONS
1	Helminth Colonization Is Associated with Increased Diversity of the Gut Microbiota. <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e2880.	1.3	353
2	Helminth infection promotes colonization resistance via type 2 immunity. <i>Science</i> , 2016, 352, 608-612.	6.0	347
3	Effect of ethnicity and socioeconomic variation to the gut microbiota composition among pre-adolescent in Malaysia. <i>Scientific Reports</i> , 2015, 5, 13338.	1.6	68
4	Reduced microbial diversity in adult survivors of childhood acute lymphoblastic leukemia and microbial associations with increased immune activation. <i>Microbiome</i> , 2017, 5, 35.	4.9	63
5	Enrichment of gut-derived <i>Fusobacterium</i> is associated with suboptimal immune recovery in HIV-infected individuals. <i>Scientific Reports</i> , 2018, 8, 14277.	1.6	57
6	Inferred metagenomic comparison of mucosal and fecal microbiota from individuals undergoing routine screening colonoscopy reveals similar differences observed during active inflammation. <i>Gut Microbes</i> , 2015, 6, 48-56.	4.3	55
7	Gastrointestinal parasites in rural dogs and cats in Selangor and Pahang states in Peninsular Malaysia. <i>Acta Parasitologica</i> , 2014, 59, 737-44.	0.4	33
8	Patterns and Risk Factors of Soil-Transmitted Helminthiasis Among Orang Asli Subgroups in Peninsular Malaysia. <i>American Journal of Tropical Medicine and Hygiene</i> , 2015, 93, 361-370.	0.6	33
9	Co-infection of <i>Haemonchus contortus</i> and <i>Trichostrongylus</i> spp. among livestock in Malaysia as revealed by amplification and sequencing of the internal transcribed spacer II DNA region. <i>BMC Veterinary Research</i> , 2014, 10, 38.	0.7	31
10	Prevalence and risk factors of intestinal parasitism among two indigenous sub-ethnic groups in Peninsular Malaysia. <i>Infectious Diseases of Poverty</i> , 2016, 5, 77.	1.5	28
11	Linking the effects of helminth infection, diet and the gut microbiota with human whole-blood signatures. <i>PLoS Pathogens</i> , 2019, 15, e1008066.	2.1	25
12	Neglected Tropical Diseases among Two Indigenous Subtribes in Peninsular Malaysia: Highlighting Differences and Co-Infection of Helminthiasis and Sarcocystosis. <i>PLoS ONE</i> , 2014, 9, e107980.	1.1	15
13	Effects of helminths on the human immune response and the microbiome. <i>Mucosal Immunology</i> , 2022, 15, 1224-1233.	2.7	15
14	Aquatic biomonitoring of <i>Giardia</i> cysts and <i>Cryptosporidium</i> oocysts in peninsular Malaysia. <i>Environmental Science and Pollution Research</i> , 2014, 21, 445-453.	2.7	14
15	Understanding <i>Giardia</i> infections among rural communities using the one health approach. <i>Acta Tropica</i> , 2017, 176, 349-354.	0.9	12
16	Loop-Mediated Isothermal Amplification Assay for Detection of Generic and Verocytotoxin-Producing <i>Escherichia coli</i> among Indigenous Individuals in Malaysia. <i>Scientific World Journal</i> , The, 2014, 2014, 1-6.	0.8	10
17	The study of seroprevalence of hepatitis E virus and an investigation into the lifestyle behaviours of the aborigines in Malaysia. <i>Zoonoses and Public Health</i> , 2020, 67, 263-270.	0.9	7
18	Seroprevalence of Nipah Virus Infection in Peninsular Malaysia. <i>Journal of Infectious Diseases</i> , 2020, 221, S370-S374.	1.9	6

#	ARTICLE	IF	CITATIONS
19	Tinea Imbricata among the Indigenous Communities: Current Global Epidemiology and Research Gaps Associated with Host Genetics and Skin Microbiota. <i>Journal of Fungi (Basel, Switzerland)</i> , 2022, 8, 202.	1.5	2
20	The Oral, Gut Microbiota and Cardiometabolic Health of Indigenous Orang Asli Communities. <i>Frontiers in Cellular and Infection Microbiology</i> , 2022, 12, 812345.	1.8	1