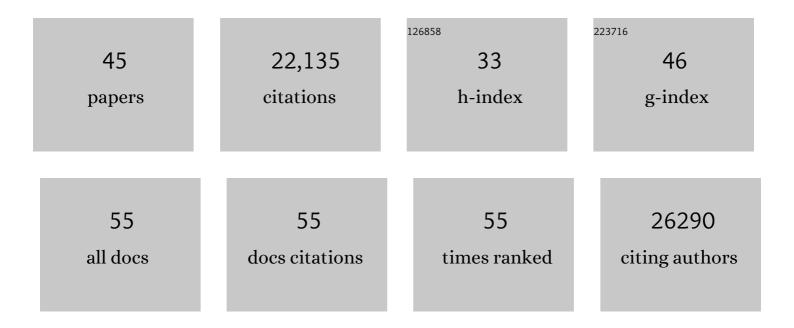
Se Jin Song

List of Publications by Year in descending order

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SELIN SONC

| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Reproducible, interactive, scalable and extensible microbiome data science using QIIME 2. Nature Biotechnology, 2019, 37, 852-857. | 9.4 | 11,167 |
| 2 | A communal catalogue reveals Earth's multiscale microbial diversity. Nature, 2017, 551, 457-463. | 13.7 | 1,942 |
| 3 | Cohabiting family members share microbiota with one another and with their dogs. ELife, 2013, 2, e00458. | 2.8 | 801 |
| 4 | Partial restoration of the microbiota of cesarean-born infants via vaginal microbial transfer. Nature Medicine, 2016, 22, 250-253. | 15.2 | 736 |
| 5 | The microbiome of uncontacted Amerindians. Science Advances, 2015, 1, . | 4.7 | 721 |
| 6 | Microbiome analyses of blood and tissues suggest cancer diagnostic approach. Nature, 2020, 579, 567-574. | 13.7 | 691 |
| 7 | American Gut: an Open Platform for Citizen Science Microbiome Research. MSystems, 2018, 3, . | 1.7 | 604 |
| 8 | Advancing Our Understanding of the Human Microbiome Using QIIME. Methods in Enzymology, 2013, 531, 371-444. | 0.4 | 553 |
| 9 | Microbial community assembly and metabolic function during mammalian corpse decomposition. Science, 2016, 351, 158-162. | 6.0 | 381 |
| 10 | Preservation Methods Differ in Fecal Microbiome Stability, Affecting Suitability for Field Studies. MSystems, 2016, 1, . | 1.7 | 367 |
| 11 | Convergence of gut microbiomes in myrmecophagous mammals. Molecular Ecology, 2014, 23, 1301-1317. | 2.0 | 311 |
| 12 | The Effects of Captivity on the Mammalian Gut Microbiome. Integrative and Comparative Biology, 2017, 57, 690-704. | 0.9 | 301 |
| 13 | Balance Trees Reveal Microbial Niche Differentiation. MSystems, 2017, 2, . | 1.7 | 284 |
| 14 | The human microbiome in evolution. BMC Biology, 2017, 15, 127. | 1.7 | 243 |
| 15 | Evolutionary trends in host physiology outweigh dietary niche in structuring primate gut microbiomes. ISME Journal, 2019, 13, 576-587. | 4.4 | 236 |
| 16 | The Bee Microbiome: Impact on Bee Health and Model for Evolution and Ecology of Host-Microbe Interactions. MBio, 2016, 7, e02164-15. | 1.8 | 215 |
| 17 | Comparative Analyses of Vertebrate Gut Microbiomes Reveal Convergence between Birds and Bats. MBio, 2020, 11, . | 1.8 | 204 |
| 18 | Learning representations of microbe–metabolite interactions. Nature Methods, 2019, 16, 1306-1314. | 9.0 | 184 |

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|----|---|-----|-----------|
| 19 | Tiny microbes, enormous impacts: what matters in gut microbiome studies?. Genome Biology, 2016, 17, 217. | 3.8 | 128 |
| 20 | Evaluating the impact of domestication and captivity on the horse gut microbiome. Scientific Reports, 2017, 7, 15497. | 1.6 | 112 |
| 21 | Home chemical and microbial transitions across urbanization. Nature Microbiology, 2020, 5, 108-115. | 5.9 | 83 |
| 22 | Consumption of Fermented Foods Is Associated with Systematic Differences in the Gut Microbiome and Metabolome. MSystems, 2020, 5, . | 1.7 | 81 |
| 23 | Using the gut microbiota as a novel tool for examining colobine primate GI health. Global Ecology and Conservation, 2016, 7, 225-237. | 1.0 | 76 |
| 24 | Walls talk: Microbial biogeography of homes spanning urbanization. Science Advances, 2016, 2, e1501061. | 4.7 | 72 |
| 25 | Engineering the microbiome for animal health and conservation. Experimental Biology and Medicine, 2019, 244, 494-504. | 1.1 | 65 |
| 26 | The Oral and Skin Microbiomes of Captive Komodo Dragons Are Significantly Shared with Their Habitat. MSystems, 2016, 1, . | 1.7 | 61 |
| 27 | How delivery mode and feeding can shape the bacterial community in the infant gut. Cmaj, 2013, 185, 373-374. | 0.9 | 54 |
| 28 | From Sample to Multi-Omics Conclusions in under 48 Hours. MSystems, 2016, 1, . | 1.7 | 53 |
| 29 | Major shifts in gut microbiota during development and its relationship to growth in ostriches. Molecular Ecology, 2019, 28, 2653-2667. | 2.0 | 53 |
| 30 | Stress response, gut microbial diversity and sexual signals correlate with social interactions. Biology Letters, 2016, 12, 20160352. | 1.0 | 47 |
| 31 | Prevalence and genetic diversity of Blastocystis in family units living in the United States. Infection, Genetics and Evolution, 2016, 45, 95-97. | 1.0 | 40 |
| 32 | Trace Evidence Potential in Postmortem Skin Microbiomes: From Death Scene to Morgue. Journal of Forensic Sciences, 2019, 64, 791-798. | 0.9 | 40 |
| 33 | Naturalization of the microbiota developmental trajectory of Cesarean-born neonates after vaginal seeding. Med, 2021, 2, 951-964.e5. | 2.2 | 37 |
| 34 | EMPress Enables Tree-Guided, Interactive, and Exploratory Analyses of Multi-omic Data Sets. MSystems, 2021, 6, . | 1.7 | 36 |
| 35 | Early-life gut dysbiosis linked to juvenile mortality in ostriches. Microbiome, 2020, 8, 147. | 4.9 | 30 |
| 36 | A posteriori dietary patterns better explain variations of the gut microbiome than individual markers in the American Gut Project. American Journal of Clinical Nutrition, 2022, 115, 432-443. | 2.2 | 28 |

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|----|---|-----|-----------|
| 37 | Are microbiome studies ready for hypothesis-driven research?. Current Opinion in Microbiology, 2018, 44, 61-69. | 2.3 | 27 |
| 38 | Evaluation of the Effect of Storage Methods on Fecal, Saliva, and Skin Microbiome Composition. MSystems, 2021, 6, . | 1.7 | 22 |
| 39 | Is there convergence of gut microbes in blood-feeding vertebrates?. Philosophical Transactions of the Royal Society B: Biological Sciences, 2019, 374, 20180249. | 1.8 | 21 |
| 40 | Reproducibility, stability, and accuracy of microbial profiles by fecal sample collection method in three distinct populations. PLoS ONE, 2019, 14, e0224757. | 1.1 | 19 |
| 41 | Coinfection and infection duration shape how pathogens affect the African buffalo gut microbiota. ISME Journal, 2021, 15, 1359-1371. | 4.4 | 17 |
| 42 | Reply to: Examining microbe–metabolite correlations by linear methods. Nature Methods, 2021, 18, 40-41. | 9.0 | 6 |
| 43 | Compositionally Aware Phylogenetic Beta-Diversity Measures Better Resolve Microbiomes Associated with Phenotype. MSystems, 2022, 7, e0005022. | 1.7 | 4 |
| 44 | Multiomic Analyses of Nascent Preterm Infant Microbiomes Differentiation Suggest Opportunities for Targeted Intervention. Advanced Biology, 2022, 6, . | 1.4 | 4 |
| 45 | The impact of maternal asthma on the preterm infants' gut metabolome and microbiome (MAP study). Scientific Reports, 2022, 12, 6437. | 1.6 | 3 |