

# Jennifer M Fettweis

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/222855/publications.pdf>

Version: 2024-02-01

23  
papers

2,683  
citations

516710

16  
h-index

642732

23  
g-index

25  
all docs

25  
docs citations

25  
times ranked

3785  
citing authors

#	ARTICLE	IF	CITATIONS
1	The vaginal microbiome and preterm birth. <i>Nature Medicine</i> , 2019, 25, 1012-1021.	30.7	600
2	Differences in vaginal microbiome in African American women versus women of European ancestry. <i>Microbiology (United Kingdom)</i> , 2014, 160, 2272-2282.	1.8	390
3	The truth about metagenomics: quantifying and counteracting bias in 16S rRNA studies. <i>BMC Microbiology</i> , 2015, 15, 66.	3.3	388
4	Racioethnic diversity in the dynamics of the vaginal microbiome during pregnancy. <i>Nature Medicine</i> , 2019, 25, 1001-1011.	30.7	204
5	Does the human placenta delivered at term have a microbiota? Results of cultivation, quantitative real-time PCR, 16S rRNA gene sequencing, and metagenomics. <i>American Journal of Obstetrics and Gynecology</i> , 2019, 220, 267.e1-267.e39.	1.3	196
6	Reporting guidelines for human microbiome research: the STORMS checklist. <i>Nature Medicine</i> , 2021, 27, 1885-1892.	30.7	170
7	Species-level classification of the vaginal microbiome. <i>BMC Genomics</i> , 2012, 13, S17.	2.8	145
8	Genomic sequence analysis and characterization of <i>Sneathia amnii</i> sp. nov. <i>BMC Genomics</i> , 2012, 13, S4.	2.8	108
9	Effects of combined oral contraceptives, depot medroxyprogesterone acetate and the levonorgestrel-releasing intrauterine system on the vaginal microbiome. <i>Contraception</i> , 2017, 95, 405-413.	1.5	95
10	Changes in vaginal community state types reflect major shifts in the microbiome. <i>Microbial Ecology in Health and Disease</i> , 2017, 28, 1303265.	3.5	66
11	An Emerging <i>Mycoplasma</i> Associated with Trichomoniasis, Vaginal Infection and Disease. <i>PLoS ONE</i> , 2014, 9, e110943.	2.5	64
12	Identification of a gene in <i>Mycoplasma hominis</i> associated with preterm birth and microbial burden in intraamniotic infection. <i>American Journal of Obstetrics and Gynecology</i> , 2015, 212, 779.e1-779.e13.	1.3	64
13	Skin-to-Skin Care and the Development of the Preterm Infant Oral Microbiome. <i>American Journal of Perinatology</i> , 2015, 32, 1205-1216.	1.4	50
14	Relationship between vitamin D status and the vaginal microbiome during pregnancy. <i>Journal of Perinatology</i> , 2019, 39, 824-836.	2.0	40
15	Race, the Vaginal Microbiome, and Spontaneous Preterm Birth. <i>MSystems</i> , 2022, 7, e0001722.	3.8	24
16	Association between statin use, the vaginal microbiome, and <i>Gardnerella vaginalis</i> vaginolysin-mediated cytotoxicity. <i>PLoS ONE</i> , 2017, 12, e0183765.	2.5	21
17	The vaginal microbiome in women of reproductive age with healthy weight versus overweight/obesity. <i>Obesity</i> , 2022, 30, 142-152.	3.0	12
18	Is prenatal diet associated with the composition of the vaginal microbiome?. <i>Paediatric and Perinatal Epidemiology</i> , 2022, 36, 243-253.	1.7	11

#	ARTICLE	IF	CITATIONS
19	Two Different Species of <i>Mycoplasma</i> Endosymbionts Can Influence <i>Trichomonas vaginalis</i> Pathophysiology. <i>MBio</i> , 2022, 13, .	4.1	11
20	Multi-omic Microbiome Profiles in the Female Reproductive Tract in Early Pregnancy. <i>Infectious Microbes &amp; Diseases</i> , 2019, 1, 49-60.	1.3	9
21	Vaginal microbiome <i>Lactobacillus crispatus</i> is heritable among European American women. <i>Communications Biology</i> , 2021, 4, 872.	4.4	7
22	Unique roles of vaginal <i>Megasphaera</i> phylotypes in reproductive health. <i>Microbial Genomics</i> , 2021, 7, .	2.0	6
23	BOTUX: Bayesian-like operational taxonomic unit examiner. <i>International Journal of Computational Biology and Drug Design</i> , 2014, 7, 130.	0.3	1