

Jennifer M Fettweis

List of Publications by Year in descending order

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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	Is prenatal diet associated with the composition of the vaginal microbiome?. Paediatric and Perinatal Epidemiology, 2022, 36, 243-253.	1.7	11
2	The vaginal microbiome in women of reproductive age with healthy weight versus overweight/obesity. Obesity, 2022, 30, 142-152.	3.0	12
3	Race, the Vaginal Microbiome, and Spontaneous Preterm Birth. MSystems, 2022, 7, e0001722.	3.8	24
4	Two Different Species of <i>Mycoplasma</i> Endosymbionts Can Influence <i>Trichomonas vaginalis</i> Pathophysiology. MBio, 2022, 13, .	4.1	11
5	Vaginal microbiome <i>Lactobacillus crispatus</i> is heritable among European American women. Communications Biology, 2021, 4, 872.	4.4	7
6	Reporting guidelines for human microbiome research: the STORMS checklist. Nature Medicine, 2021, 27, 1885-1892.	30.7	170
7	Unique roles of vaginal <i>Megasphaera</i> phylotypes in reproductive health. Microbial Genomics, 2021, 7, .	2.0	6
8	The vaginal microbiome and preterm birth. Nature Medicine, 2019, 25, 1012-1021.	30.7	600
9	Racioethnic diversity in the dynamics of the vaginal microbiome during pregnancy. Nature Medicine, 2019, 25, 1001-1011.	30.7	204
10	Does the human placenta delivered at term have a microbiota? Results of cultivation, quantitative real-time PCR, 16S rRNA gene sequencing, and metagenomics. American Journal of Obstetrics and Gynecology, 2019, 220, 267.e1-267.e39.	1.3	196
11	Relationship between vitamin D status and the vaginal microbiome during pregnancy. Journal of Perinatology, 2019, 39, 824-836.	2.0	40
12	Multi-omic Microbiome Profiles in the Female Reproductive Tract in Early Pregnancy. Infectious Microbes & Diseases, 2019, 1, 49-60.	1.3	9
13	Effects of combined oral contraceptives, depot medroxyprogesterone acetate and the levonorgestrel-releasing intrauterine system on the vaginal microbiome. Contraception, 2017, 95, 405-413.	1.5	95
14	Changes in vaginal community state types reflect major shifts in the microbiome. Microbial Ecology in Health and Disease, 2017, 28, 1303265.	3.5	66
15	Association between statin use, the vaginal microbiome, and <i>Gardnerella vaginalis</i> vaginolysin-mediated cytotoxicity. PLoS ONE, 2017, 12, e0183765.	2.5	21
16	The truth about metagenomics: quantifying and counteracting bias in 16S rRNA studies. BMC Microbiology, 2015, 15, 66.	3.3	388
17	Identification of a gene in <i>Mycoplasma hominis</i> associated with preterm birth and microbial burden in intraamniotic infection. American Journal of Obstetrics and Gynecology, 2015, 212, 779.e1-779.e13.	1.3	64
18	Skin-to-Skin Care and the Development of the Preterm Infant Oral Microbiome. American Journal of Perinatology, 2015, 32, 1205-1216.	1.4	50

#	ARTICLE	IF	CITATIONS
19	An Emerging Mycoplasma Associated with Trichomoniasis, Vaginal Infection and Disease. PLoS ONE, 2014, 9, e110943.	2.5	64
20	Differences in vaginal microbiome in African American women versus women of European ancestry. Microbiology (United Kingdom), 2014, 160, 2272-2282.	1.8	390
21	BOTUX: Bayesian-like operational taxonomic unit examiner. International Journal of Computational Biology and Drug Design, 2014, 7, 130.	0.3	1
22	Genomic sequence analysis and characterization of <i>Sneathia amnii</i> sp. nov. BMC Genomics, 2012, 13, S4.	2.8	108
23	Species-level classification of the vaginal microbiome. BMC Genomics, 2012, 13, S17.	2.8	145