PaweÅ, Åaniewski

List of Publications by Year
in descending order

20	966	759233 12	839539 18
papers	citations	h-index	g-index

21	21	21	850
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Multi-omics data integration reveals metabolome as the top predictor of the cervicovaginal microenvironment. PLoS Computational Biology, 2022, 18, e1009876.	3.2	21
2	Cervicovaginal DNA Virome Alterations Are Associated with Genital Inflammation and Microbiota Composition. MSystems, 2022, 7, e0006422.	3.8	14
3	Connecting microbiome and menopause for healthy ageing. Nature Microbiology, 2022, 7, 354-358.	13.3	11
4	Clinical and Personal Lubricants Impact the Growth of Vaginal Lactobacillus Species and Colonization of Vaginal Epithelial Cells: An in Vitro Study. Sexually Transmitted Diseases, 2021, 48, 63-70.	1.7	11
5	Veillonellaceae family members uniquely alter the cervical metabolic microenvironment in a human three-dimensional epithelial model. Npj Biofilms and Microbiomes, 2021, 7, 57.	6.4	25
6	Bacterial vaginosis and health-associated bacteria modulate the immunometabolic landscape in 3D model of human cervix. Npj Biofilms and Microbiomes, 2021, 7, 88.	6.4	42
7	Immunometabolic Analysis of Mobiluncus mulieris and Eggerthella sp. Reveals Novel Insights Into Their Pathogenic Contributions to the Hallmarks of Bacterial Vaginosis. Frontiers in Cellular and Infection Microbiology, 2021, 11, 759697.	3.9	6
8	Interleukin-36Î³ Is Elevated in Cervicovaginal Epithelial Cells in Women With Bacterial Vaginosis and In Vitro After Infection With Microbes Associated With Bacterial Vaginosis. Journal of Infectious Diseases, 2020, 221, 983-988.	4.0	24
9	Hostâ€“vaginal microbiota interactions in the pathogenesis of bacterial vaginosis. Current Opinion in Infectious Diseases, 2020, 33, 59-65.	3.1	97
10	Vaginal microbiota, genital inflammation, and neoplasia impact immune checkpoint protein profiles in the cervicovaginal microenvironment. Npj Precision Oncology, 2020, 4, 22.	5.4	18
11	Members of Prevotella Genus Distinctively Modulate Innate Immune and Barrier Functions in a Human Three-Dimensional Endometrial Epithelial Cell Model. Journal of Infectious Diseases, 2020, 222, 2082-2092.	4.0	21
12	The microbiome and gynaecological cancer development, prevention and therapy. Nature Reviews Urology, 2020, 17, 232-250.	3.8	194
13	Personal and Clinical Vaginal Lubricants: Impact on Local Vaginal Microenvironment and Implications for Epithelial Cell Host Response and Barrier Function. Journal of Infectious Diseases, 2019, 220, 2009-2018.	4.0	29
14	Analysis of Host Responses to Neisseria gonorrhoeae Using a Human Three-Dimensional Endometrial Epithelial Cell Model. Methods in Molecular Biology, 2019, 1997, 347-361.	0.9	5
15	Features of the cervicovaginal microenvironment drive cancer biomarker signatures in patients across cervical carcinogenesis. Scientific Reports, 2019, 9, 7333.	3.3	70
16	Deciphering the complex interplay between microbiota, HPV, inflammation and cancer through cervicovaginal metabolic profiling. EBioMedicine, 2019, 44, 675-690.	6.1	142
17	O05.6â€…Cervicovaginal metabolic profiling reveals the interplay between HPV, microbiota and inflammation in cervical carcinogenesis. , 2019, , .		0
18	Linking cervicovaginal immune signatures, HPV and microbiota composition in cervical carcinogenesis in non-Hispanic and Hispanic women. Scientific Reports, 2018, 8, 7593.	3.3	155

#	Article	IF	Citations
19	Vagina. , 2018, , 353-359.		8
20	Human Three-Dimensional Endometrial Epithelial Cell Model To Study Host Interactions with Vaginal Bacteria and Neisseria gonorrhoeae. Infection and Immunity, 2017, 85, .	2.2	72