PaweÅ, Åaniewski

List of Publications by Year
in descending order

20 papers	**966** citations	759233 **12** h-index	839539 **18** g-index
21 all docs	**21** docs citations	**21** times ranked	**850** citing authors

#	Article	IF	Citations
1	The microbiome and gynaecological cancer development, prevention and therapy. Nature Reviews Urology, 2020, 17, 232-250.	3.8	194
2	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
3	Deciphering the complex interplay between microbiota, HPV, inflammation and cancer through cervicovaginal metabolic profiling. EBioMedicine, 2019, 44, 675-690.	6.1	142
4	Host–vaginal microbiota interactions in the pathogenesis of bacterial vaginosis. Current Opinion in Infectious Diseases, 2020, 33, 59-65.	3.1	97
5	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
6	Features of the cervicovaginal microenvironment drive cancer biomarker signatures in patients across cervical carcinogenesis. Scientific Reports, 2019, 9, 7333.	3.3	70
7	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
8	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
9	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
10	Interleukin-36γ 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
11	Members of <i>Prevotella</i> 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	Multi-omics data integration reveals metabolome as the top predictor of the cervicovaginal microenvironment. PLoS Computational Biology, 2022, 18, e1009876.	3.2	21
13	Vaginal microbiota, genital inflammation, and neoplasia impact immune checkpoint protein profiles in the cervicovaginal microenvironment. Npj Precision Oncology, 2020, 4, 22.	5.4	18
14	Cervicovaginal DNA Virome Alterations Are Associated with Genital Inflammation and Microbiota Composition. MSystems, 2022, 7, e0006422.	3.8	14
15	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
16	Connecting microbiome and menopause for healthy ageing. Nature Microbiology, 2022, 7, 354-358.	13.3	11
17	Vagina. , 2018, , 353-359.		8
18	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

#	Article	IF	Citations
19	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
20	O05.6â€…Cervicovaginal metabolic profiling reveals the interplay between HPV, microbiota and inflammation in cervical carcinogenesis. , 2019, , .		0