Ville Nikolai Pimenoff

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Evidence of Still-Ongoing Convergence Evolution of the Lactase Persistence T-13910 Alleles in Humans. American Journal of Human Genetics, 2007, 81, 615-625.	2.6	135
2	Transmission between Archaic and Modern Human Ancestors during the Evolution of the Oncogenic Human Papillomavirus 16. Molecular Biology and Evolution, 2017, 34, 4-19.	3.5	103
3	Northwest Siberian Khanty and Mansi in the junction of West and East Eurasian gene pools as revealed by uniparental markers. European Journal of Human Genetics, 2008, 16, 1254-1264.	1.4	53
4	Whole-genome sequence analysis of a Pan African set of samples reveals archaic gene flow from an extinct basal population of modern humans into sub-Saharan populations. Genome Biology, 2019, 20, 77.	3.8	50
5	Gut microbiome diversity detected by high-coverage 16S and shotgun sequencing of paired stool and colon sample. Scientific Data, 2020, 7, 92.	2.4	37
6	Analysis of 16 Y STR loci in the Finnish population reveals a local reduction in the diversity of male lineages. Forensic Science International, 2004, 142, 37-43.	1.3	36
7	Finnish mitochondrial DNA HVS-I and HVS-II population data. Forensic Science International, 2007, 172, 171-178.	1.3	34
8	HPV16 variants distribution in invasive cancers of the cervix, vulva, vagina, penis, and anus. Cancer Medicine, 2016, 5, 2909-2919.	1.3	29
9	Vaccination With Moderate Coverage Eradicates Oncogenic Human Papillomaviruses If a Gender-Neutral Strategy Is Applied. Journal of Infectious Diseases, 2020, 222, 948-956.	1.9	29
10	Evaluation of HPV typeâ€replacement in unvaccinated and vaccinated adolescent females— <i>Postâ€hoc</i> analysis of a communityâ€randomized clinical trial (II). International Journal of Cancer, 2018, 142, 2491-2500.	2.3	28
11	Occurrence of human papillomavirus (HPV) type replacement by sexual riskâ€ŧaking behaviour group: Postâ€hoc analysis of a community randomized clinical trial up to nine years after vaccination (IV). International Journal of Cancer, 2019, 145, 785-796.	2.3	20
12	Disagreement in high-grade/low-grade intraepithelial neoplasia and high-risk/low-risk HPV infection: clinical implications for anal cancer precursor lesions in HIV-positive and HIV-negative MSM. Clinical Microbiology and Infection, 2015, 21, 605.e11-605.e19.	2.8	18
13	The Role of aDNA in Understanding the Coevolutionary Patterns of Human Sexually Transmitted Infections. Genes, 2018, 9, 317.	1.0	17
14	Differences in risk for SARS-CoV-2 infection among healthcare workers. Preventive Medicine Reports, 2021, 24, 101518.	0.8	17
15	Distinct geographic clustering of oncogenic human papillomaviruses multiple infections in cervical cancers: Results from a worldwide crossâ€sectional study. International Journal of Cancer, 2019, 144, 2478-2488.	2.3	14
16	Assessment of HV1 and HV2 mtDNA Variation for Forensic Purposes in an Uruguayan Population Sample. Journal of Forensic Sciences, 2005, 50, 1-4.	0.9	14
17	Longâ€ŧerm followâ€up of human papillomavirus type replacement among young pregnant Finnish females before and after a communityâ€randomised <scp>HPV</scp> vaccination trial with moderate coverage. International Journal of Cancer, 2020, 147, 3511-3522.	2.3	13
18	Similarity in recombination rate and linkage disequilibrium at CYP2C and CYP2D cytochrome P450 gene regions among Europeans indicates signs of selection and no advantage of using tagSNPs in population isolates. Pharmacogenetics and Genomics, 2012, 22, 846-857.	0.7	12

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19	Squamous intraepithelial lesions of the anal squamocolumnar junction: Histopathological classification and HPV genotyping. Papillomavirus Research (Amsterdam, Netherlands), 2017, 3, 11-17.	4.5	12
20	Phylogenetically related, clinically different: human papillomaviruses 6 and 11 variants distribution in genital warts and in laryngeal papillomatosis. Clinical Microbiology and Infection, 2014, 20, O406-O413.	2.8	9
21	High Amounts of SARS-CoV-2 Precede Sickness Among Asymptomatic Health Care Workers. Journal of Infectious Diseases, 2021, 224, 14-20.	1.9	8
22	Human papillomavirus seroprevalence in pregnant women following gender-neutral and girls-only vaccination programs in Finland: A cross-sectional cohort analysis following a cluster randomized trial. PLoS Medicine, 2021, 18, e1003588.	3.9	8
23	Human exposome assessment platform. Environmental Epidemiology, 2021, 5, e182.	1.4	7
24	Estimating Total Excess Mortality During a Coronavirus Disease 2019 Outbreak in Stockholm, Sweden. Clinical Infectious Diseases, 2021, 72, e890-e892.	2.9	5
25	Risk for SARS-CoV-2 infection in healthcare workers outside hospitals: A real-life immuno-virological study during the first wave of the COVID-19 epidemic. PLoS ONE, 2021, 16, e0257854.	1.1	5
26	Moral dilemma(s) in human papillomavirus vaccination – revisiting the role of the herd effect. Eurosurveillance, 2021, 26, .	3.9	3
27	Potential SARS-CoV-2 infectiousness among asymptomatic healthcare workers. PLoS ONE, 2021, 16, e0260453.	1.1	3
28	Head-to-Head Comparison of Bi- and Nonavalent Human Papillomavirus Vaccine-Induced Antibody Responses. Journal of Infectious Diseases, 2022, 226, 1195-1199.	1.9	3
29	Severe features during outbreak but low mortality observed immediately before and after a March–May 2020 COVID-19 outbreak in Stockholm, Sweden. International Journal of Infectious Diseases, 2021, 110, 433-435.	1.5	2
30	How infectious diseases arrived in the colonial Americas. ELife, 2021, 10, .	2.8	1
31	Population-Based Human Papillomavirus Serosurvey Reveals HPV16/18 Herd Effect But No Clear Type-Replacement in Unvaccinated Females 6 Years Post Gender-Neutral Vaccination in a Cluster Randomised Trial. SSRN Electronic Journal, 0, , .	0.4	0
32	Moderate Coverage Vaccination Eradicates Oncogenic Human Papillomaviruses if a Gender-Neutral Strategy is Applied. SSRN Electronic Journal, 0, , .	0.4	0