

# Dimitris Tsakogiannis

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/878408/publications.pdf>

Version: 2024-02-01

28  
papers

312  
citations

933447

10  
h-index

888059

17  
g-index

29  
all docs

29  
docs citations

29  
times ranked

416  
citing authors

#	ARTICLE	IF	CITATIONS
1	Serum Concentration of Selected Angiogenesis-Related Molecules Differs among Molecular Subtypes, Body Mass Index and Menopausal Status in Breast Cancer Patients. <i>Journal of Clinical Medicine</i> , 2022, 11, 4079.	2.4	1
2	WarmStart colorimetric RT-LAMP for the rapid, sensitive and specific detection of Enteroviruses targeting the 5'UTR region. <i>Journal of Applied Microbiology</i> , 2021, 130, 292-301.	3.1	8
3	A colorimetric IsoPCR for the rapid and sensitive visual detection of high-risk HPV16 in clinical samples with hydroxynaphthol blue. <i>Journal of Virological Methods</i> , 2021, 290, 114072.	2.1	3
4	HPV16-Genotyper: A Computational Tool for Risk-Assessment, Lineage Genotyping and Recombination Detection in HPV16 Sequences, Based on a Large-Scale Evolutionary Analysis. <i>Diversity</i> , 2021, 13, 497.	1.7	3
5	Genetic variability of the HPV16 early genes and LCR. Present and future perspectives. <i>Expert Reviews in Molecular Medicine</i> , 2021, 23, e19.	3.9	10
6	WarmStart colorimetric LAMP for the specific and rapid detection of HPV16 and HPV18 DNA. <i>Journal of Virological Methods</i> , 2019, 270, 87-94.	2.1	20
7	2-stage, nested-like nucleic acid amplification method (IsoPCR) for the highly sensitive detection of HPV16 and HPV18 DNA. <i>Molecular and Cellular Probes</i> , 2019, 45, 1-7.	2.1	3
8	Large-scale genomic analysis reveals recurrent patterns of intertypic recombination in human enteroviruses. <i>Virology</i> , 2019, 526, 72-80.	2.4	36
9	Intra- and inter-serotypic recombinations in the 5'UTR-VP4 region of Echovirus 30 strains. <i>Archives of Virology</i> , 2018, 163, 365-375.	2.1	4
10	Association of p16 (CDKN2A) polymorphisms with the development of HPV16-related precancerous lesions and cervical cancer in the Greek population. <i>Journal of Medical Virology</i> , 2018, 90, 965-971.	5.0	10
11	Polymorphic variability in the exon 19 of the RB1 gene and its flanking intronic sequences in HPV16-associated precancerous lesions in the Greek population. <i>Journal of Medical Microbiology</i> , 2018, 67, 1638-1644.	1.8	4
12	Molecular approaches for HPV genotyping and HPV-DNA physical status. <i>Expert Reviews in Molecular Medicine</i> , 2017, 19, e1.	3.9	45
13	Association of codon 72 polymorphism of p53 with the severity of cervical dysplasia, E6-T350G and HPV16 variant lineages in HPV16-infected women. <i>Journal of Medical Microbiology</i> , 2017, 66, 1358-1365.	1.8	8
14	Development of a multiplex RT-PCR assay for the identification of recombination types at different genomic regions of vaccine-derived polioviruses. <i>Virus Genes</i> , 2016, 52, 453-462.	1.6	0
15	Identification of rearranged sequences of HPV16 DNA in precancerous and cervical cancer cases. <i>Molecular and Cellular Probes</i> , 2016, 30, 6-12.	2.1	7
16	Sites of disruption within E1 and E2 genes of HPV16 and association with cervical dysplasia. <i>Journal of Medical Virology</i> , 2015, 87, 1973-1980.	5.0	29
17	Multiplex PCR assay for the rapid identification of human papillomavirus genotypes 16, 18, 45, 35, 66, 33, 51, 58, and 31 in clinical samples. <i>Archives of Virology</i> , 2015, 160, 207-214.	2.1	16
18	Molecular epidemiology and evolutionary dynamics of Echovirus 3 serotype. <i>Infection, Genetics and Evolution</i> , 2015, 32, 305-312.	2.3	10

#	ARTICLE	IF	CITATIONS
19	Duplex Real-time PCR assay and SYBR green I melting curve analysis for molecular identification of HPV genotypes 16, 18, 31, 35, 51 and 66. <i>Molecular and Cellular Probes</i> , 2015, 29, 13-18.	2.1	6
20	Prevalence of HPV16 E1â€¦1374^63nt variants in Greek women. <i>Journal of Medical Virology</i> , 2014, 86, 778-784.	5.0	2
21	Determination of human papillomavirus 16 physical status through E1/E6 and E2/E6 ratio analysis. <i>Journal of Medical Microbiology</i> , 2014, 63, 1716-1723.	1.8	16
22	Nucleotide polymorphisms of the human papillomavirus 16 E1 gene. <i>Archives of Virology</i> , 2014, 159, 51-63.	2.1	10
23	PCR assays for the identification of rare recombination types from VP1 to 3D genomic region of vaccine derived poliovirus strains. <i>Molecular and Cellular Probes</i> , 2014, 28, 107-112.	2.1	3
24	Characterization of novel intergenogroup and intergenotype recombinant noroviruses from central Greece. <i>Molecular and Cellular Probes</i> , 2014, 28, 204-210.	2.1	7
25	Molecular and evolutionary analysis of HPV16 E6 and E7 genes in Greek women. <i>Journal of Medical Microbiology</i> , 2013, 62, 1688-1696.	1.8	13
26	Identification of novel E6-E7 sequence variants of human papillomavirus 16. <i>Archives of Virology</i> , 2013, 158, 821-828.	2.1	7
27	Molecular and phylogenetic analysis of the HPV 16 E4 gene in cervical lesions from women in Greece. <i>Archives of Virology</i> , 2012, 157, 1729-1739.	2.1	12
28	Sequence variation analysis of the E2 gene of human papilloma virus type 16 in cervical lesions from women in Greece. <i>Archives of Virology</i> , 2012, 157, 825-832.	2.1	15