

Dace Pjanova

List of Publications by Year in descending order

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36
papers

799
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567281

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501196

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all docs

39
docs citations

39
times ranked

1449
citing authors

#	ARTICLE	IF	CITATIONS
1	Role of the Circadian Clock "Death-Loop" in the DNA Damage Response Underpinning Cancer Treatment Resistance. <i>Cells</i> , 2022, 11, 880.	4.1	8
2	Effect of Bacteriophage-Derived Double Stranded RNA on Rat Peritoneal Macrophages and Microglia in Normoxia and Hypoxia. <i>Proceedings of the Latvian Academy of Sciences</i> , 2021, 75, 343-349.	0.1	2
3	Association of HERV-K and LINE-1 hypomethylation with reduced disease-free survival in melanoma patients. <i>Epigenomics</i> , 2020, 12, 1689-1706.	2.1	11
4	"Mitotic Slippage" and Extranuclear DNA in Cancer Chemoresistance: A Focus on Telomeres. <i>International Journal of Molecular Sciences</i> , 2020, 21, 2779.	4.1	36
5	Features associated with melanoma metastasis in Latvia. <i>Oncology Letters</i> , 2020, 20, 1-1.	1.8	0
6	Estimating CDKN2A mutation carrier probability among global familial melanoma cases using GenoMELPREDICT. <i>Journal of the American Academy of Dermatology</i> , 2019, 81, 386-394.	1.2	17
7	Comparison of the effects of bacteriophage-derived dsRNA and poly(I:C) on ex vivo cultivated peripheral blood mononuclear cells. <i>Immunology Letters</i> , 2019, 212, 114-119.	2.5	8
8	The Cancer Aneuploidy Paradox: In the Light of Evolution. <i>Genes</i> , 2019, 10, 83.	2.4	41
9	The complementary effect of rs1042522 in TP53 and rs1805007 in MC1R is associated with an elevated risk of cutaneous melanoma in Latvian population. <i>Oncology Letters</i> , 2019, 18, 5225-5234.	1.8	2
10	Oncolytic viruses sensitize human tumor cells for NY-ESO-1 tumor antigen recognition by CD4+ effector T cells. <i>Onc Immunology</i> , 2018, 7, e1407897.	4.6	22
11	Association of the 16q24.3 region gene variants rs1805007 and rs4785763 with heightened risk of melanoma in Latvian population. <i>Meta Gene</i> , 2018, 18, 87-92.	0.6	1
12	The effect of intranasally administered TLR3 agonist larifan on metabolic profile of microglial cells in rat with C6 glioma. <i>Ukrainian Biochemical Journal</i> , 2018, 90, 110-119.	0.5	2
13	Germline Variation at CDKN2A and Associations with Nevus Phenotypes among Members of Melanoma Families. <i>Journal of Investigative Dermatology</i> , 2017, 137, 2606-2612.	0.7	18
14	Phenotypic and Histopathological Tumor Characteristics According to CDKN2A Mutation Status among Affected Members of Melanoma Families. <i>Journal of Investigative Dermatology</i> , 2016, 136, 1066-1069.	0.7	13
15	The lack of E318K MITF germline mutation in Latvian melanoma patients. <i>Cancer Genetics</i> , 2015, 208, 355-356.	0.4	1
16	Replacement of short segments within transmembrane domains of MC2R disrupts retention signal. <i>Journal of Molecular Endocrinology</i> , 2014, 53, 201-215.	2.5	10
17	Somatic BRAF and NRAS Mutations in Familial Melanomas with Known Germline CDKN2A Status: A GenoMEL Study. <i>Journal of Investigative Dermatology</i> , 2014, 134, 287-290.	0.7	18
18	Inherited variation in the PARP1 gene and survival from melanoma. <i>International Journal of Cancer</i> , 2014, 135, 1625-1633.	5.1	24

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19	<i>Ex vivo</i> cytokine production in peripheral blood mononuclear cells after their stimulation with dsRNA of natural origin. <i>Biotechnology and Applied Biochemistry</i> , 2014, 61, 65-73.	3.1	9
20	An inherited variant in the gene coding for vitamin D-binding protein and survival from cutaneous melanoma: a BioGenoMEL study. <i>Pigment Cell and Melanoma Research</i> , 2014, 27, 234-243.	3.3	25
21	Melanoma risk associated with MC1R gene variants in Latvia and the functional analysis of rare variants. <i>Cancer Genetics</i> , 2013, 206, 81-91.	0.4	7
22	Melanoma prone families with CDK4 germline mutation: phenotypic profile and associations with MC1R variants. <i>Journal of Medical Genetics</i> , 2013, 50, 264-270.	3.2	112
23	Analysis of Latvian familial melanoma patients shows novel variants in the noncoding regions of CDKN2A and that the CDK4 mutation R24H is a founder mutation. <i>Melanoma Research</i> , 2013, 23, 221-226.	1.2	13
24	Melanoma epidemiology, prognosis and trends in Latvia. <i>Journal of the European Academy of Dermatology and Venereology</i> , 2013, 27, 1352-1359.	2.4	14
25	Skin Examination Behavior. <i>Archives of Dermatology</i> , 2012, 148, 1142.	1.4	36
26	Inherited variants in the MC1R gene and survival from cutaneous melanoma: a BioGenoMEL study. <i>Pigment Cell and Melanoma Research</i> , 2012, 25, 384-394.	3.3	61
27	Abstract 867: Somatic BRAF and NRAS mutations in familial melanoma: a GenoMEL study. , 2011, , .		0
28	Abstract 869: Characterization of malignant melanoma families with CDK4 germ-line mutation. , 2011, , .		0
29	Prognostic factors and epidemiological characteristics of cutaneous and mucosal head and neck melanoma. <i>Stomatologija</i> , 2011, 13, 49-54.	0.3	4
30	Melanoma risk factors, perceived threat and intentional tanning: an international online survey. <i>European Journal of Cancer Prevention</i> , 2010, 19, 216-226.	1.3	47
31	Predictors of Sun Protection Behaviors and Severe Sunburn in an International Online Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 2199-2210.	2.5	106
32	Identification of a CDK4 R24H mutation-positive melanoma family by analysis of early-onset melanoma patients in Latvia. <i>Melanoma Research</i> , 2009, 19, 119-122.	1.2	16
33	Genetic alteration in melanoma development. <i>Proceedings of the Latvian Academy of Sciences</i> , 2009, 63, 73-80.	0.1	0
34	CDKN2A and CDK4 variants in Latvian melanoma patients: analysis of a clinic-based population. <i>Melanoma Research</i> , 2007, 17, 185-191.	1.2	34
35	Toluidine blue test for sperm DNA integrity and elaboration of image cytometry algorithm. <i>Cytometry</i> , 2003, 52A, 19-27.	1.8	74
36	The Role of the Meiotic Component in Reproduction of B-RAF-Mutated Melanoma: A Review and Brainstorming Session. , 0, , .		2