

Marianna Borecká

List of Publications by Year in descending order

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Version: 2024-02-01

13
papers

221
citations

1163117

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1125743

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#	ARTICLE	IF	CITATIONS
1	Novel presentation of the c.1856A>>G (p.Asp619Gly) TSHR gene-activating variant: relapsing hyperthyroidism in three subsequent generations manifesting in early childhood and an in vitro functional study. <i>Hormones</i> , 2021, 20, 803-812.	1.9	1
2	Prevalence of Germline Pathogenic Variants in Cancer Predisposing Genes in Czech and Belgian Pancreatic Cancer Patients. <i>Cancers</i> , 2021, 13, 4430.	3.7	8
3	Multigene Panel Germline Testing of 1333 Czech Patients with Ovarian Cancer. <i>Cancers</i> , 2020, 12, 956.	3.7	19
4	Truncated PPM1D impairs stem cell response to genotoxic stress and promotes growth of APC-deficient tumors in the mouse colon. <i>Cell Death and Disease</i> , 2019, 10, 818.	6.3	12
5	Genetic analysis of subsequent second primary malignant neoplasms in long-term pancreatic cancer survivors suggests new potential hereditary genetic alterations. <i>Cancer Management and Research</i> , 2019, Volume 11, 599-609.	1.9	4
6	Estrogen Receptor Status Oppositely Modifies Breast Cancer Prognosis in BRCA1/BRCA2 Mutation Carriers Versus Non-Carriers. <i>Cancers</i> , 2019, 11, 738.	3.7	22
7	Identification of deleterious germline CHEK2 mutations and their association with breast and ovarian cancer. <i>International Journal of Cancer</i> , 2019, 145, 1782-1797.	5.1	62
8	Germline CHEK2 Gene Mutations in Hereditary Breast Cancer Predisposition – Mutation Types and their Biological and Clinical Relevance. <i>Klinicka Onkologie</i> , 2019, 32, 36-50.	0.3	2
9	Contribution of Massive Parallel Sequencing to Diagnosis of Hereditary Ovarian Cancer in the Czech Republic. <i>Klinicka Onkologie</i> , 2019, 32, 72-78.	0.3	1
10	BRCA1 and BRCA2 5' noncoding region variants identified in breast cancer patients alter promoter activity and protein binding. <i>Human Mutation</i> , 2018, 39, 2025-2039.	2.5	15
11	Validation of CZEKANCA (CZEch CAncer paNel for Clinical Application) for targeted NGS-based analysis of hereditary cancer syndromes. <i>PLoS ONE</i> , 2018, 13, e0195761.	2.5	31
12	The c.657del5 variant in the NBN gene predisposes to pancreatic cancer. <i>Gene</i> , 2016, 587, 169-172.	2.2	13
13	Mutation analysis of the PALB2 gene in unselected pancreatic cancer patients in the Czech Republic. <i>Cancer Genetics</i> , 2016, 209, 199-204.	0.4	27