Jan Hojný

List of Publications by Year in descending order

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933447 839539 19 350 10 18 citations h-index g-index papers 19 19 19 627 citing authors docs citations times ranked all docs

#	Article	IF	Citations
1	Identification of deleterious germline <i>CHEK2</i> mutations and their association with breast and ovarian cancer. International Journal of Cancer, 2019, 145, 1782-1797.	5.1	62
2	Multiplex PCR and NGS-based identification of mRNA splicing variants: Analysis of BRCA1 splicing pattern as a model. Gene, 2017, 637, 41-49.	2.2	43
3	A comprehensive evaluation of pathogenic mutations in primary cutaneous melanomas, including the identification of novel loss-of-function variants. Scientific Reports, 2019, 9, 17050.	3.3	42
4	Expression of human BRCA1î"17–19 alternative splicing variant with a truncated BRCT domain in MCF-7 cells results in impaired assembly of DNA repair complexes and aberrant DNA damage response. Cellular Signalling, 2013, 25, 1186-1193.	3.6	35
5	Ovarian mesonephric-like adenocarcinoma arising in serous borderline tumor: a case report with complex morphological and molecular analysis. Diagnostic Pathology, 2020, 15, 91.	2.0	35
6	Association of Germline CHEK2 Gene Variants with Risk and Prognosis of Non-Hodgkin Lymphoma. PLoS ONE, 2015, 10, e0140819.	2.5	31
7	Leiomyoma with Bizarre Nuclei: a Study of 108 Cases Focusing on Clinicopathological Features, Morphology, and Fumarate Hydratase Alterations. Pathology and Oncology Research, 2020, 26, 1527-1537.	1.9	22
8	Germline mutation in the TP53 gene in uveal melanoma. Scientific Reports, 2018, 8, 7618.	3.3	13
9	Synchronous endometrioid endometrial and ovarian carcinomas are biologically related: A clinico‑pathological and molecular (next generation sequencing) study of 22�cases. Oncology Letters, 2019, 17, 2207-2214.	1.8	12
10	Results of targeted next-generation sequencing in children with cystic kidney diseases often change the clinical diagnosis. PLoS ONE, 2020, 15, e0235071.	2.5	12
11	Uterine cellular leiomyomas are characterized by common HMGA2 aberrations, followed by chromosome 1p deletion and MED12 mutation: morphological, molecular, and immunohistochemical study of 52 cases. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2022, 480, 281-291.	2.8	10
12	Expression, Epigenetic, and Genetic Changes of HNF1B in Colorectal Lesions: an Analysis of 145 Cases. Pathology and Oncology Research, 2020, 26, 2337-2350.	1.9	7
13	HNF1B, EZH2 and ECI2 in prostate carcinoma. Molecular, immunohistochemical and clinico-pathological study. Scientific Reports, 2020, 10, 14365.	3.3	6
14	Analysis of expression, epigenetic, and genetic changes of HNF1B in 130 kidney tumours. Scientific Reports, 2020, 10, 17151.	3.3	5
15	Identification of novel HNF1B mRNA splicing variants and their qualitative and semi-quantitative profile in selected healthy and tumour tissues. Scientific Reports, 2020, 10, 6958.	3.3	5
16	A comprehensive analysis of the expression, epigenetic and genetic changes of HNF1B and ECI2 in 122Âcases of high†grade serous ovarian carcinoma. Oncology Letters, 2021, 21, 185.	1.8	4
17	Microscopic extraovarian sex cord proliferation: report of a case with bilateral Fallopian tube involvement and a comprehensive molecular analysis. Polish Journal of Pathology, 2020, 71, 175-180.	0.3	2
18	Comprehensive quantitative analysis of alternative splicing variants reveals the HNF1B mRNA splicing pattern in various tumour and non-tumour tissues. Scientific Reports, 2022, 12, 199.	3.3	2

#	Article	IF	CITATIONS
19	Desmoplastic Small Round Cell Tumor of the Uterus: A Report of Molecularly Confirmed Case with EWSR1-WT1 Fusion. Diagnostics, 2022, 12, 1184.	2.6	2