Jana Nekvindova

List of Publications by Year in descending order

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414414 471509 1,327 34 17 32 citations h-index g-index papers 34 34 34 2770 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Role of miR-653 and miR-29c in downregulation of CYP1A2 expression in hepatocellular carcinoma. Pharmacological Reports, 2022, 74, 148-158.	3.3	O
2	Delimitation and phylogeny of <i> Dictyochaeta </i> , and introduction of <i> Achrochaeta </i> and <i> Tubulicolla </i> , genera nova. Mycologia, 2021, 113, 390-433.	1.9	13
3	Phylogeny, Global Biogeography and Pleomorphism of Zanclospora. Microorganisms, 2021, 9, 706.	3.6	10
4	Reflections on Menisporopsis, Multiguttulispora and Tainosphaeria Using Molecular and Morphological Data. Journal of Fungi (Basel, Switzerland), 2021, 7, 438.	3.5	7
5	Phylogeny and taxonomy of Catenularia and similar fungi with catenate conidia. MycoKeys, 2021, 81, 1-44.	1.9	11
6	Deregulation of signaling pathways controlling cell survival and proliferation in cancer cells alters induction of cytochrome P450 family 1 enzymes. Toxicology, 2021, 461, 152897.	4.2	5
7	Phylogenetic Reassessment, Taxonomy, and Biogeography of Codinaea and Similar Fungi. Journal of Fungi (Basel, Switzerland), 2021, 7, 1097.	3.5	11
8	Hepatocellular carcinoma: Gene expression profiling and regulation of xenobiotic-metabolizing cytochromes P450. Biochemical Pharmacology, 2020, 177, 113912.	4.4	24
9	New insights into the systematics of Bactrodesmium and its allies and introducing new genera, species and morphological patterns in the Pleurotheciales and Savoryellales (Sordariomycetes). Studies in Mycology, 2020, 95, 415-466.	7.2	25
10	Delimitation, new species and teleomorph-anamorph relationships in Codinaea, Dendrophoma, Paragaeumannomyces and Striatosphaeria (Chaetosphaeriaceae). MycoKeys, 2020, 74, 17-74.	1.9	14
11	Fungal diversity notes 1036–1150: taxonomic and phylogenetic contributions on genera and species of fungal taxa. Fungal Diversity, 2019, 96, 1-242.	12.3	148
12	Are haplotypes in a single methotrexate pathway more predictive for response in rheumatoid arthritis than in different pathways?. Pharmacogenomics, 2018, 19, 379-381.	1.3	0
13	Aberrant methylation of tumour suppressor genes <i>WT1</i> , <i>GATA5</i> and <i>PAX5</i> in hepatocellular carcinoma. Clinical Chemistry and Laboratory Medicine, 2016, 54, 1971-1980.	2.3	21
14	Methotrexate impact on radiographic progression in biologic-treated rheumatoid arthritis under clinical remission: A case report on monozygotic Caucasian twins. International Journal of Immunopathology and Pharmacology, 2016, 29, 790-795.	2.1	1
15	Circulating Serum MicroRNA-130a as a Novel Putative Marker of Extramedullary Myeloma. PLoS ONE, 2015, 10, e0137294.	2.5	16
16	The impact of C677T and A1298C MTHFR polymorphisms on methotrexate therapeutic response in East Bohemian region rheumatoid arthritis patients. Rheumatology International, 2015, 35, 1149-1161.	3.0	17
17	Combination of serum microRNAâ€320a and microRNAâ€320b as a marker for <scp>W</scp> aldenström macroglobulinemia. American Journal of Hematology, 2015, 90, E51-2.	4.1	8
18	Cervical Microbiota in Women with Preterm Prelabor Rupture of Membranes. PLoS ONE, 2015, 10, e0126884.	2.5	55

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19	Loss of PTEN Facilitates Rosiglitazone-Mediated Enhancement of Platinum(IV) Complex LA-12-Induced Apoptosis in Colon Cancer Cells. PLoS ONE, 2015, 10, e0141020.	2.5	6
20	Genetic polymorphisms in metabolic pathways of leflunomide in the treatment of rheumatoid arthritis. Clinical and Experimental Rheumatology, 2015, 33, 426-32.	0.8	5
21	Circulating serum microRNAs as novel diagnostic and prognostic biomarkers for multiple myeloma and monoclonal gammopathy of undetermined significance. Haematologica, 2014, 99, 511-518.	3.5	94
22	Acyclic nucleoside phosphonates: a study on cytochrome P450 gene expression. Xenobiotica, 2014, 44, 708-715.	1.1	1
23	Risk Score based on microRNA expression signature is independent prognostic classifier of glioblastoma patients. Carcinogenesis, 2014, 35, 2756-2762.	2.8	30
24	MiR-210 expression in tumor tissue and in vitro effects of its silencing in renal cell carcinoma. Tumor Biology, 2013, 34, 481-491.	1.8	70
25	Extension of microRNA expression pattern associated with high-risk neuroblastoma. Tumor Biology, 2013, 34, 2315-2319.	1.8	16
26	Circulating miR-378 and miR-451 in serum are potential biomarkers for renal cell carcinoma. Journal of Translational Medicine, 2012, 10, 55.	4.4	228
27	Identification and functional screening of micro <scp>RNA</scp> s highly deregulated in colorectal cancer. Journal of Cellular and Molecular Medicine, 2012, 16, 2655-2666.	3.6	127
28	MicroRNA expression profile associated with response to neoadjuvant chemoradiotherapy in locally advanced rectal cancer patients. Radiation Oncology, 2012, 7, 195.	2.7	111
29	Identification of MicroRNAs associated with early relapse after nephrectomy in renal cell carcinoma patients. Genes Chromosomes and Cancer, 2012, 51, 707-716.	2.8	97
30	The role of high-risk human papillomavirus infection in oral and oropharyngeal squamous cell carcinoma in non-smoking and non-drinking patients: a clinicopathological and molecular study of 46 cases. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2011, 458, 179-187.	2.8	49
31	IL-4-mediated transcriptional regulation of human CYP2E1 by two independent signaling pathways. Biochemical Pharmacology, 2010, 80, 1592-1600.	4.4	22
32	New Insights into the Regulation of CYP2C9 Gene Expression: The Role of the Transcription Factor GATA-4. Drug Metabolism and Disposition, 2010, 38, 415-421.	3.3	25
33	Colorectal Cancer-Specific Cytochrome P450 2W1: Intracellular Localization, Glycosylation, and Catalytic Activity. Molecular Pharmacology, 2010, 78, 1004-1011.	2.3	36
34	The transcription factor GATA-4 regulates cytochrome P4502C19 gene expression. Life Sciences, 2010, 86, 699-706.	4.3	24