Radoslav S Davidović

List of Publications by Year in descending order

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18 papers	978 citations	840776 11 h-index	17 g-index
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19 all docs	19 docs citations	19 times ranked	1496 citing authors

#	Article	IF	CITATIONS
1	The CAFA challenge reports improved protein function prediction and new functional annotations for hundreds of genes through experimental screens. Genome Biology, 2019, 20, 244.	8.8	261
2	DisProt 7.0: a major update of the database of disordered proteins. Nucleic Acids Research, 2017, 45, D219-D227.	14.5	242
3	DisProt: intrinsic protein disorder annotation in 2020. Nucleic Acids Research, 2020, 48, D269-D276.	14.5	141
4	DisProt in 2022: improved quality and accessibility of protein intrinsic disorder annotation. Nucleic Acids Research, 2022, 50, D480-D487.	14.5	117
5	MicroRNA in breast cancer: The association with BRCA1/2. Cancer Biomarkers, 2017, 19, 119-128.	1.7	47
6	The impact of PTEN tumor suppressor gene on acquiring resistance to tamoxifen treatment in breast cancer patients. Cancer Biology and Therapy, 2012, 13, 1165-1174.	3.4	36
7	Prognostic significance of epigenetic inactivation of p16, p15, MGMT and DAPK genes in follicular lymphoma. Medical Oncology, 2013, 30, 441.	2.5	18
8	Skewed X-Chromosome Inactivation in Women Affected by Alzheimer's Disease. Journal of Alzheimer's Disease, 2014, 43, 1251-1259.	2.6	17
9	Recurrence of giant cell tumour of bone: role of p53, cyclin D1, \hat{l}^2 -catenin and Ki67. International Orthopaedics, 2016, 40, 2393-2399.	1.9	13
10	Changes in miR-221/222 Levels in Invasive and In Situ Carcinomas of the Breast: Differences in Association with Estrogen Receptor and TIMP3 Expression Levels. Molecular Diagnosis and Therapy, 2016, 20, 603-615.	3.8	13
11	Mapping of Protein-Protein Interactions: Web-Based Resources for Revealing Interactomes. Current Medicinal Chemistry, 2019, 26, 3890-3910.	2.4	11
12	p14ARF methylation is a common event in the pathogenesis and progression of myxoid and pleomorphic liposarcoma. Medical Oncology, 2013, 30, 682.	2.5	10
13	Methylation-specific PCR: four steps in primer design. Open Life Sciences, 2014, 9, 1127-1139.	1.4	8
14	DiNGO: standalone application for Gene Ontology and Human Phenotype Ontology term enrichment analysis. Bioinformatics, 2020, 36, 1981-1982.	4.1	2
15	Immunohistochemical expression of cyclin-dependent kinase inhibitors p16 and p57 in rhabdomyosarcoma. Pathology Research and Practice, 2021, 225, 153558.	2.3	1
16	Methylation status of p16 and p14 genes in locally advanced rectal cancer: Potential clinical implication. Archives of Biological Sciences, 2018, 70, 681-690.	0.5	1
17	Significance of \hat{I}^2 -catenin expression for the incidence of pathological fractures in giant cell tumors of bone. Polish Journal of Pathology, 2016, 4, 345-350.	0.3	O
18	Alignment-free method for functional annotation of amino acid substitutions: Application on epigenetic factors involved in hematologic malignancies. PLoS ONE, 2021, 16, e0244948.	2.5	0