Miluse Hradilova

List of Publications by Year in descending order

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35 papers 1,326 citations

430874 18 h-index 35 g-index

37 all docs

37 docs citations

37 times ranked

2148 citing authors

#	Article	IF	CITATIONS
1	Exosomes produced by melanoma cells significantly influence the biological properties of normal and cancer-associated fibroblasts. Histochemistry and Cell Biology, 2022, 157, 153-172.	1.7	17
2	Desmoplastic Crosstalk in Pancreatic Ductal Adenocarcinoma Is Reflected by Different Responses of Panc-1, MIAPaCa-2, PaTu-8902, and CAPAN-2 Cell Lines to Cancer-associated/Normal Fibroblasts. Cancer Genomics and Proteomics, 2021, 18, 221-243.	2.0	8
3	The <i>Mastigamoeba balamuthi</i> Genome and the Nature of the Free-Living Ancestor of <i>Entamoeba</i> . Molecular Biology and Evolution, 2021, 38, 2240-2259.	8.9	14
4	Heterotrophic euglenid Rhabdomonas costata resembles its phototrophic relatives in many aspects of molecular and cell biology. Scientific Reports, $2021, 11, 13070$.	3.3	5
5	Comparison of Silks from Pseudoips prasinana and Bombyx mori Shows Molecular Convergence in Fibroin Heavy Chains but Large Differences in Other Silk Components. International Journal of Molecular Sciences, 2021, 22, 8246.	4.1	7
6	Predominant Biphenyl Dioxygenase From Legacy Polychlorinated Biphenyl (PCB)-Contaminated Soil Is a Part of Unusual Gene Cluster and Transforms Flavone and Flavanone. Frontiers in Microbiology, 2021, 12, 644708.	3.5	4
7	Anaeramoebae are a divergent lineage of eukaryotes that shed light on the transition from anaerobic mitochondria to hydrogenosomes. Current Biology, 2021, 31, 5605-5612.e5.	3.9	29
8	A rapid approach for in locus overexpression of Trypanosoma brucei proteins. Molecular and Biochemical Parasitology, 2020, 239, 111300.	1.1	3
9	Antiâ€angiogenic effects of the blueâ€green alga Arthrospira platensis on pancreatic cancer. Journal of Cellular and Molecular Medicine, 2020, 24, 2402-2415.	3.6	10
10	Analysis of HPV-Positive and HPV-Negative Head and Neck Squamous Cell Carcinomas and Paired Normal Mucosae Reveals Cyclin D1 Deregulation and Compensatory Effect of Cyclin D2. Cancers, 2020, 12, 792.	3.7	9
11	Ms1 RNA increases the amount of RNA polymerase in <i>Mycobacterium smegmatis</i> Microbiology, 2019, 111, 354-372.	2.5	26
12	Modular structure, sequence diversification and appropriate nomenclature of seroins produced in the silk glands of Lepidoptera. Scientific Reports, 2019, 9, 3797.	3.3	8
13	The Head and Neck Squamous Cell Carcinoma Microenvironment as a Potential Target for Cancer Therapy. Cancers, 2019, 11, 440.	3.7	43
14	The expansion of genes encoding soluble silk components in the greater wax moth, Galleria mellonella. Insect Biochemistry and Molecular Biology, 2019, 106, 28-38.	2.7	17
15	Diversity of root-associated microbial populations of Tamarix parviflora cultivated under various conditions. Applied Soil Ecology, 2018, 125, 264-272.	4.3	16
16	Complete genome sequence of Pseudomonas alcaliphila JAB1 (=DSM 26533), a versatile degrader of organic pollutants. Standards in Genomic Sciences, 2018, 13, 3.	1.5	36
17	Hybrid asexuality as a primary postzygotic barrier between nascent species: On the interconnection between asexuality, hybridization and speciation. Molecular Ecology, 2018, 27, 248-263.	3.9	64
18	Detection of Distinct Changes in Gene-expression Profiles in Specimens of Tumors and Transition Zones of Tenascinpositive/- negative Head and Neck Squamous Cell Carcinoma. Anticancer Research, 2018, 38, 1279-1290.	1.1	8

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19	Analysis of dermal fibroblasts isolated from neonatal and child cleft lip and adult skin: Developmental implications on reconstructive surgery. International Journal of Molecular Medicine, 2017, 40, 1323-1334.	4.0	17
20	Nuclear genetic codes with a different meaning of the UAG and the UAA codon. BMC Biology, 2017, 15, 8.	3.8	25
21	Mitochondrial Genome of Prasinophyte Alga <i>Pyramimonas parkeae</i> . Journal of Eukaryotic Microbiology, 2017, 64, 360-369.	1.7	3
22	Arginine deiminase pathway enzymes: evolutionary history in metamonads and other eukaryotes. BMC Evolutionary Biology, 2016, 16 , 197 .	3.2	40
23	A Eukaryote without a Mitochondrial Organelle. Current Biology, 2016, 26, 1274-1284.	3.9	302
24	A Comparative Analysis of Mitochondrial Genomes in Eustigmatophyte Algae. Genome Biology and Evolution, 2016, 8, 705-722.	2.5	33
25	First multigene analysis of Archamoebae (Amoebozoa: Conosa) robustly reveals its phylogeny and shows that Entamoebidae represents a deep lineage of the group. Molecular Phylogenetics and Evolution, 2016, 98, 41-51.	2.7	23
26	Characterization of three distinct metallothionein genes of the Ag-hyperaccumulating ectomycorrhizal fungus Amanita strobiliformis. Fungal Biology, 2016, 120, 358-369.	2.5	32
27	Pseudomonads Rule Degradation of Polyaromatic Hydrocarbons in Aerated Sediment. Frontiers in Microbiology, 2015, 6, 1268.	3.5	54
28	Combined Culture-Based and Culture-Independent Approaches Provide Insights into Diversity of Jakobids, an Extremely Plesiomorphic Eukaryotic Lineage. Frontiers in Microbiology, 2015, 6, 1288.	3.5	20
29	Microarray Analysis of Serum mRNA in Patients with Head and Neck Squamous Cell Carcinoma at Whole-Genome Scale. BioMed Research International, 2014, 2014, 1-10.	1.9	6
30	Plant secondary metabolite-induced shifts in bacterial community structure and degradative ability in contaminated soil. Applied Microbiology and Biotechnology, 2013, 97, 9245-9256.	3.6	56
31	Diversity, Phylogeny and Expression Patterns of Pou and Six Homeodomain Transcription Factors in Hydrozoan Jellyfish Craspedacusta sowerbyi. PLoS ONE, 2012, 7, e36420.	2.5	20
32	Smooth muscle actinâ \in expressing stromal fibroblasts in head and neck squamous cell carcinoma: Increased expression of galectinâ \in 1 and induction of poor prognosis factors. International Journal of Cancer, 2012, 131, 2499-2508.	5.1	67
33	Identification of Bacteria Utilizing Biphenyl, Benzoate, and Naphthalene in Long-Term Contaminated Soil. PLoS ONE, 2012, 7, e40653.	2.5	124
34	Matrix-Assisted Laser Desorption Ionization (MALDI)-Time of Flight Mass Spectrometry- and MALDI Biotyper-Based Identification of Cultured Biphenyl-Metabolizing Bacteria from Contaminated Horseradish Rhizosphere Soil. Applied and Environmental Microbiology, 2011, 77, 6858-6866.	3.1	77
35	Biphenyl-Metabolizing Bacteria in the Rhizosphere of Horseradish and Bulk Soil Contaminated by Polychlorinated Biphenyls as Revealed by Stable Isotope Probing. Applied and Environmental Microbiology, 2009, 75, 6471-6477.	3.1	102

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