Ewa A Grzybowska

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Overall survival analysis of > 65-year-old patients with breast cancer based on their molecular, clinicopathological and laboratory factors. Archives of Medical Science, 2022, 18, 800-804.	0.9	0
2	Cell Death by Entosis: Triggers, Molecular Mechanisms and Clinical Significance. International Journal of Molecular Sciences, 2022, 23, 4985.	4.1	14
3	Clusters, Assemblies and Aggregates of Tumor Cells in the Blood of Breast Cancer Patients; Composition, Mode of Action, Detection and Impact on Metastasis and Survival. International Journal of Translational Medicine, 2021, 1, 55-68.	0.4	3
4	The type <scp>III</scp> secretion system effector <scp>EspO</scp> of enterohaemorrhagic <i>Escherichia coli</i> inhibits apoptosis through an interaction with <scp>HAX</scp> â€1. Cellular Microbiology, 2021, 23, e13366.	2.1	3
5	Protein Binding to Cis-Motifs in mRNAs Coding Sequence Is Common and Regulates Transcript Stability and the Rate of Translation. Cells, 2021, 10, 2910.	4.1	8
6	Editorial: The Role of Protein Post-Translational Modifications in Protein-RNA Interactions and RNP Assemblies. Frontiers in Molecular Biosciences, 2021, 8, 831810.	3.5	2
7	How to Predict Metastasis in Luminal Breast Cancer? Current Solutions and Future Prospects. International Journal of Molecular Sciences, 2020, 21, 8415.	4.1	16
8	Circulating Tumor Cells in Early and Advanced Breast Cancer; Biology and Prognostic Value. International Journal of Molecular Sciences, 2020, 21, 1671.	4.1	34
9	Genetic 3′UTR variations and clinical factors significantly contribute to survival prediction and clinical response in breast cancer patients. Scientific Reports, 2020, 10, 5736.	3.3	14
10	Impact of Medium pH on DOX Toxicity toward HeLa and A498 Cell Lines. ACS Omega, 2020, 5, 7979-7986.	3.5	16
11	Quantification of Cell-Substrate Adhesion Area and Cell Shape Distributions in MCF7 Cell Monolayers. Journal of Visualized Experiments, 2020, , .	0.3	2
12	The interactome of multifunctional HAX1 protein suggests its role in the regulation of energy metabolism, de-aggregation, cytoskeleton organization and RNA-processing. Bioscience Reports, 2020, 40, .	2.4	10
13	RNA–protein interactions: disorder, moonlighting and junk contribute to eukaryotic complexity. Open Biology, 2019, 9, 190096.	3.6	113
14	Cytoplasmic HAX1 Is an Independent Risk Factor for Breast Cancer Metastasis. Journal of Oncology, 2019, 1-13.	1.3	8
15	Calcium-Binding Proteins with Disordered Structure and Their Role in Secretion, Storage, and Cellular Signaling. Biomolecules, 2018, 8, 42.	4.0	27
16	CTC clusters in cancer progression and metastasis. Medical Oncology, 2017, 34, 12.	2.5	134
17	The calcium binding properties and structure prediction of the Hax-1 protein. Acta Biochimica Polonica, 2017, 64, 537-542.	0.5	7
18	The putative oncogene, <i>CRNDE,</i> is a negative prognostic factor in ovarian cancer patients. Oncotarget, 2015, 6, 43897-43910.	1.8	51

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19	HAX-1: A Novel P-Body Protein. DNA and Cell Biology, 2015, 34, 43-54.	1.9	9
20	Exploring the antiâ€apoptotic role of HAXâ€1 versus BCLâ€X _L in cytokineâ€dependent bone marrowâ€derived cells from mice. FEBS Letters, 2014, 588, 2921-2927.	2.8	16
21	<scp>HAX</scp> â€1 is a nucleocytoplasmic shuttling protein with a possible role in m <scp>RNA</scp> processing. FEBS Journal, 2013, 280, 256-272.	4.7	29
22	Comment on "HAX1 Augments Cell Proliferation, Migration, Adhesion, and Invasion Induced by Urokinase-Type Plasminogen Activator Receptor― Journal of Oncology, 2013, 2013, 1-1.	1.3	1
23	Human intronless genes: Functional groups, associated diseases, evolution, and mRNA processing in absence of splicing. Biochemical and Biophysical Research Communications, 2012, 424, 1-6.	2.1	85
24	HAX-1 overexpression, splicing and cellular localization in tumors. BMC Cancer, 2010, 10, 76.	2.6	63
25	Nα-Tosyl-l-phenylalanine Chloromethyl Ketone Induces Caspase-dependent Apoptosis in Transformed Human B Cell Lines with Transcriptional Down-regulation of Anti-apoptotic HS1-associated Protein X-1. Journal of Biological Chemistry, 2009, 284, 27827-27837.	3.4	26
26	Compound heterozygous <i>HAX1</i> mutations in a Swedish patient with severe congenital neutropenia and no neurodevelopmental abnormalities. Pediatric Blood and Cancer, 2009, 53, 1143-1146.	1.5	14
27	HAX-1: A multifunctional protein with emerging roles in human disease. Biochimica Et Biophysica Acta - General Subjects, 2009, 1790, 1139-1148.	2.4	97
28	Hairpin structure within the 3'UTR of DNA polymerase mRNA acts as a post-transcriptional regulatory element and interacts with Hax-1. Nucleic Acids Research, 2007, 35, 5499-5510.	14.5	43
29	Identification and expression analysis of alternative splice variants of the rat Hax-1 gene. Gene, 2006, 371, 84-92.	2.2	25
30	Low Expression of DNA Polymerase β in Human Testicular Germ Cell Tumours: Impact on Foetal Gonocytic Origin Theory. Acta Oncológica, 2002, 41, 188-191.	1.8	5
31	Saccharomyces cerevisiae Ferrochelatase Forms a Homodimer. Archives of Biochemistry and Biophysics, 2002, 398, 170-178.	3.0	16
32	Regulatory Functions of 3′UTRs. Biochemical and Biophysical Research Communications, 2001, 288, 291-295.	2.1	162
33	Probing the Active-site Residues in Saccharomyces cerevisiae Ferrochelatase by Directed Mutagenesis. Journal of Biological Chemistry, 1996, 271, 11810-11816.	3.4	51