Raja Mokdad-Gargouri

List of Publications by Year in descending order

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394421 61 928 19 citations h-index papers

g-index 62 62 62 1568 docs citations times ranked citing authors all docs

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#	Article	IF	CITATIONS
1	Whole and Purified Aqueous Extracts of Nigella sativa L. Seeds Attenuate Apoptosis and the Overproduction of Reactive Oxygen Species Triggered by p53 Over-Expression in the Yeast Saccharomyces cerevisiae. Cells, 2022, 11, 869.	4.1	1
2	Novel and recurrent BRCA1/BRCA2 germline mutations in patients with breast/ovarian cancer: a series from the south of Tunisia. Journal of Translational Medicine, 2021, 19, 108.	4.4	14
3	Establishment of primary cell culture of Ruditapes decussatus haemocytes for metal toxicity assessment. In Vitro Cellular and Developmental Biology - Animal, 2021, 57, 477-484.	1.5	4
4	Lower p66Shc promoter methylation in subjects with chronic renal failure. PLoS ONE, 2021, 16, e0257176.	2.5	2
5	Identification of novel candidate genes by exome sequencing in Tunisian familial male breast cancer patients. Molecular Biology Reports, 2020, 47, 6507-6516.	2.3	4
6	FOXA1 Expression in Nasopharyngeal Carcinoma: Association with Clinicopathological Characteristics and EMT Markers. BioMed Research International, 2020, 2020, 1-9.	1.9	10
7	B1.12: a novel peptide interacting with the extracellular loop of the EBV oncoprotein LMP1. Scientific Reports, 2019, 9, 4389.	3.3	3
8	Association of FOXA1 and EMT markers (Twist1 and E-cadherin) in breast cancer. Molecular Biology Reports, 2019, 46, 3247-3255.	2.3	17
9	Serine protease inhibitors and human wellbeing interplay: new insights for old friends. PeerJ, 2019, 7, e7224.	2.0	20
10	Overexpression of the Oncogenic Variant (KLF6-SV1) in Young NPC Patients and Correlation with Lack of E-Cadherin. Analytical Cellular Pathology, 2018, 2018, 1-7.	1.4	3
11	Overexpression of miR-10b in colorectal cancer patients: Correlation with <i>TWIST-1</i> and E-cadherin expression. Tumor Biology, 2017, 39, 101042831769591.	1.8	32
12	CpG methylation of APC promoter 1A in sporadic and familial breast cancer patients. Cancer Biomarkers, 2017, 18, 133-141.	1.7	13
13	Clinical and prognosis value of the CIMP status combined with MLH1 or p16 INK4a methylation in colorectal cancer. Medical Oncology, 2017, 34, 147.	2.5	10
14	RIP140 and LCoR expression in gastrointestinal cancers. Oncotarget, 2017, 8, 111161-111175.	1.8	7
15	Phage-display screening identifies LMP1-binding peptides targeting the C-terminus region of the EBV oncoprotein. Peptides, 2016, 85, 73-79.	2.4	3
16	Clinical and prognosis relevance of COX-2 expression in Tunisian patients with primary gastric adenocarcinoma. Cancer Biomarkers, 2016, 17, 67-73.	1.7	4
17	Quantitative measurement of iNOS expression in melanoma, nasopharyngeal, colorectal, and breast tumors of Tunisian patients: comparative study and clinical significance. Tumor Biology, 2016, 37, 5153-5164.	1.8	3
18	CpG methylation of ubiquitin carboxyl-terminal hydrolase 1 (UCHL1) and P53 mutation pattern in sporadic colorectal cancer. Tumor Biology, 2016, 37, 1707-1714.	1.8	17

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19	Overexpression of yeast thioredoxin TRX2 reduces p53-mediated cell death in yeast. Applied Microbiology and Biotechnology, 2015, 99, 8619-8628.	3.6	5
20	Over-expression of miR-10b in NPC patients: correlation with LMP1 and Twist1. Tumor Biology, 2015, 36, 3807-3814.	1.8	20
21	Expression and Mutation Pattern of \hat{l}^2 -Catenin and Adenomatous Polyposis Coli in Colorectal Cancer Patients. Archives of Medical Research, 2015, 46, 54-62.	3.3	20
22	Characterization of C69R variant HBsAg: effect on binding to anti-HBs and the structure of virus-like particles. Archives of Virology, 2015, 160, 2427-2433.	2.1	8
23	Identification of a Novel Methylated Gene in Nasopharyngeal Carcinoma: TTC40. BioMed Research International, 2014, 2014, 1-7.	1.9	5
24	Negative control glucose dependent mediated by the PreS2 region on the translation efficiency of the reporter <i>Sh-bleomycin</i> gene in <i>Saccharomyces cerevisiae</i> . FEMS Yeast Research, 2014, 14, 357-363.	2.3	0
25	Expression of APC, \hat{l}^2 -catenin and E-cadherin in Tunisian patients with gastric adenocarcinoma: clinical significance. Tumor Biology, 2014, 35, 1775-1783.	1.8	14
26	Clinicopathological and prognostic significance of p53, Ki-67, and Bcl-2 expression in Tunisian gastric adenocarcinomas. Acta Histochemica, 2014, 116, 1244-1250.	1.8	13
27	Downregulation of WIF-1 and Wnt5a in patients with colorectal carcinoma: clinical significance. Tumor Biology, 2014, 35, 7975-7982.	1.8	14
28	Loss of WIF-1 and Wnt5a expression is related to aggressiveness of sporadic breast cancer in Tunisian patients. Tumor Biology, 2013, 34, 1625-1633.	1.8	23
29	Secreted recombinant P53 protein from Pichia pastoris is a useful antigen for detection of serum p53: autoantibody in patients with advanced colorectal adenocarcinoma. Molecular Biology Reports, 2013, 40, 3865-3872.	2.3	2
30	Quantitative expression analysis and prognostic significance of the BCL2-associated Xgene in nasopharyngeal carcinoma: a retrospective cohort study. BMC Cancer, 2013, 13, 293.	2.6	24
31	Extraction and purification of hepatitis B virus-like M particles from a recombinant Saccharomyces cerevisiae strain using alumina powder. Journal of Virological Methods, 2013, 187, 132-137.	2.1	18
32	Frequent CpG methylation of ubiquitin carboxyl-terminal hydrolase 1 (UCHL1) in sporadic and hereditary Tunisian breast cancer patients: clinical significance. Medical Oncology, 2013, 30, 418.	2.5	11
33	Yeasts as a Tool for Heterologous Gene Expression. Methods in Molecular Biology, 2012, 824, 359-370.	0.9	22
34	High prevalence of the c.1227_1228dup (p.Glu410GlyfsX43) mutation in Tunisian families affected with MUTYH-associated-polyposis. Familial Cancer, 2012, 11, 503-508.	1.9	11
35	Expression of COX-2 and E-cadherin in Tunisian patients with colorectal adenocarcinoma. Acta Histochemica, 2012, 114, 577-581.	1.8	22
36	Expression of the human tumor suppressor p53 induces cell death in Pichia pastoris. FEMS Yeast Research, 2012, 12, 2-8.	2.3	5

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37	Negative/low HER2 expression alone or combined with E-cadherin positivity is predictive of better prognosis in patients with breast carcinoma. Histology and Histopathology, 2012, 27, 377-85.	0.7	4
38	Aberrant methylation of hMLH1 and p16INK4a in Tunisian patients with sporadic colorectal adenocarcinoma. Bioscience Reports, 2011, 31, 257-264.	2.4	17
39	BCL2L12 is a Novel Biomarker for the Prediction of Short-Term Relapse in Nasopharyngeal Carcinoma. Molecular Medicine, 2011, 17, 163-171.	4.4	39
40	Expression of p16INK4a, Alone or Combined With p53, is Predictive of Better Prognosis in Colorectal Adenocarcinoma in Tunisian Patients. Applied Immunohistochemistry and Molecular Morphology, 2011, 19, 562-568.	1.2	9
41	Characteristics of epstein barr virus variants associated with gastric carcinoma in Southern Tunisia. Virology Journal, 2011, 8, 500.	3.4	18
42	A novel pathogenic germline mutation in the adenomatous polyposis coli gene in a Tunisian family with FAP. Familial Cancer, 2011, 10, 567-571.	1.9	5
43	Cellular localization of human p53 expressed in the yeast Saccharomyces cerevisiae: effect of NLSI deletion. Apoptosis: an International Journal on Programmed Cell Death, 2011, 16, 746-756.	4.9	9
44	Methylation status and overexpression of COX-2 in Tunisian patients with ductal invasive breast carcinoma. Tumor Biology, 2011, 32, 461-468.	1.8	8
45	Hypermethylation of tumorâ€related genes in tunisian patients with gastric carcinoma: Clinical and biological significance. Journal of Surgical Oncology, 2011, 103, 687-694.	1.7	32
46	Aberrant methylation of RASSF1A is associated with poor survival in Tunisian breast cancer patients. Journal of Cancer Research and Clinical Oncology, 2010, 136, 203-210.	2.5	36
47	Hypermethylation of $RAR\hat{I}^2$ 2 correlates with high COX-2 expression and poor prognosis in patients with colorectal carcinoma. Tumor Biology, 2010, 31, 503-511.	1.8	19
48	Epigenetic Alteration of the Wnt Inhibitory Factor-1 Promoter Is Common and Occurs in Advanced Stage of Tunisian Nasopharyngeal Carcinoma. Cancer Investigation, 2010, 28, 896-903.	1.3	30
49	Clinical Significance of Epigenetic Inactivation of hMLH1 and BRCA1 in Tunisian Patients with Invasive Breast Carcinoma. Journal of Biomedicine and Biotechnology, 2009, 2009, 1-7.	3.0	23
50	Inactivation of RASSF1A, RAR \hat{i}^2 2 and DAP-kinase by promoter methylation correlates with lymph node metastasis in nasopharyngeal carcinoma. Cancer Biology and Therapy, 2009, 8, 444-451.	3.4	48
51	Cloning and characterization of cDNA probes for the analysis of metallothionein gene expression in the Mediterranean bivalves: Ruditapes decussatus and Cerastoderma glaucum. Molecular Biology Reports, 2009, 36, 1007-1014.	2.3	14
52	Selection of cell deathâ€deficient <i>p53</i> mutants in <i>Saccharomyces cerevisiae</i> Yeast, 2009, 26, 441-450.	1.7	7
53	PIK3CA amplification is predictive of poor prognosis in Tunisian patients with nasopharyngeal carcinoma. Cancer Science, 2009, 100, 2034-2039.	3.9	37
54	Expression of HBsAg and preS2-S protein in different yeast based system: A comparative analysis. Protein Expression and Purification, 2009, 66, 131-137.	1.3	14

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55	Human p53 induces cell death and downregulates thioredoxin expression in (i>Saccharomyces cerevisiae . FEMS Yeast Research, 2008, 8, 1254-1262.	2.3	28
56	Overexpression of COX-2 and LMP1 are correlated with lymph node in Tunisian NPC patients. Oral Oncology, 2008, 44, 710-715.	1.5	13
57	High-level expression of human tumour suppressor P53 in the methylotrophic yeast: Pichia pastoris. Protein Expression and Purification, 2007, 54, 283-288.	1.3	12
58	Various 30 and 69bp deletion variants of the Epstein-Barr virus LMP1 may arise by homologous recombination in nasopharyngeal carcinoma of Tunisian patients. Virus Research, 2006, 115, 24-30.	2.2	32
59	Glucose dependant negative translational control of the heterologous expression of the preS2 HBV antigen in yeast. Gene, 2003, 311, 165-170.	2.2	8
60	Translational control of human p53 expression in yeast mediated by 5'-UTR-ORF structural interaction. Nucleic Acids Research, 2001, 29, 1222-1227.	14.5	23
61	Genotyping of Tunisian hepatitis B virus isolates based on the sequencing of preS2 and S regions. Microbes and Infection, 2000, 2, 607-612.	1.9	39