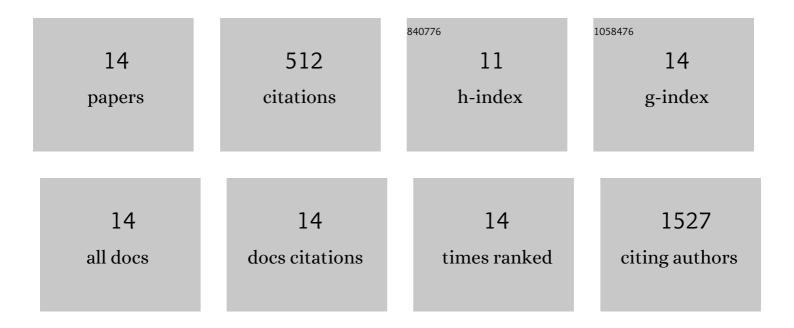
Laura Caleca

List of Publications by Year in descending order

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LALIDA CALECA

#	Article	IF	CITATIONS
1	<i>FANCM</i> c.5791C>T nonsense mutation (rs144567652) induces exon skipping, affects DNA repair activity and is a familial breast cancer risk factor. Human Molecular Genetics, 2015, 24, 5345-5355.	2.9	91
2	Mir-660 is downregulated in lung cancer patients and its replacement inhibits lung tumorigenesis by targeting MDM2-p53 interaction. Cell Death and Disease, 2014, 5, e1564-e1564.	6.3	73
3	Comparative In Vitro and In Silico Analyses of Variants in Splicing Regions of BRCA1 and BRCA2 Genes and Characterization of Novel Pathogenic Mutations. PLoS ONE, 2013, 8, e57173.	2.5	64
4	Circulating mirâ€320a promotes immunosuppressive macrophages M2 phenotype associated with lung cancer risk. International Journal of Cancer, 2019, 144, 2746-2761.	5.1	56
5	Unclassified variants in BRCA genes: guidelines for interpretation. Annals of Oncology, 2011, 22, i18-i23.	1.2	50
6	Characterization of the activation domain of the Rad53 checkpoint kinase. Cell Cycle, 2008, 7, 493-499.	2.6	38
7	c-Myc shuttled by tumour-derived extracellular vesicles promotes lung bronchial cell proliferation through miR-19b and miR-92a. Cell Death and Disease, 2019, 10, 759.	6.3	32
8	The FANCM:p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. Npj Breast Cancer, 2019, 5, 38.	5.2	28
9	Characterization of an Italian Founder Mutation in the RING-Finger Domain of BRCA1. PLoS ONE, 2014, 9, e86924.	2.5	24
10	The <i>BRCA2</i> c.68-7TÂ>ÂA variant is not pathogenic: A model for clinical calibration of spliceogenicity. Human Mutation, 2018, 39, 729-741.	2.5	19
11	The rs12975333 variant in the miR-125a and breast cancer risk in Germany, Italy, Australia and Spain. Journal of Medical Genetics, 2011, 48, 703-704.	3.2	13
12	Two Missense Variants Detected in Breast Cancer Probands Preventing BRCA2-PALB2 Protein Interaction. Frontiers in Oncology, 2018, 8, 480.	2.8	11
13	LKB1 Down-Modulation by miR-17 Identifies Patients With NSCLC Having Worse Prognosis Eligible for Energy-Stress–Based Treatments. Journal of Thoracic Oncology, 2021, 16, 1298-1311.	1.1	9
14	GFP-Fragment Reassembly Screens for the Functional Characterization of Variants of Uncertain Significance in Protein Interaction Domains of the BRCA1 and BRCA2 Genes. Cancers, 2019, 11, 151.	3.7	4