## Frances V Fuller-Pace

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	DDX5 plays essential transcriptional and post-transcriptional roles in the maintenance and function of spermatogonia. Nature Communications, 2019, 10, 2278.	12.8	74
2	DDX17 nucleocytoplasmic shuttling promotes acquired gefitinib resistance in non-small cell lung cancer cells via activation of β-catenin. Cancer Letters, 2017, 400, 194-202.	7.2	57
3	p53 controls expression of the DNA deaminase APOBEC3B to limit its potential mutagenic activity in cancer cells. Nucleic Acids Research, 2017, 45, 11056-11069.	14.5	70
4	TP53 drives invasion through expression of its $\hat{I}$ "133p53 $\hat{I}$ 2 variant. ELife, 2016, 5, .	6.0	44
5	The loop structure and the RNA helicase p72/DDX17 influence the processing efficiency of the mice miR-132. Scientific Reports, 2016, 6, 22848.	3.3	15
6	LRH-1 drives colon cancer cell growth by repressing the expression of the <i>CDKN1A</i> gene in a p53-dependent manner. Nucleic Acids Research, 2016, 44, 582-594.	14.5	46
7	The miRNA biogenesis factors, p72/DDX17 and KHSRP regulate the protein level of Ago2 in human cells. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2016, 1859, 1299-1305.	1.9	16
8	Expression of CDK7, Cyclin H, and MAT1 Is Elevated in Breast Cancer and Is Prognostic in Estrogen Receptor–Positive Breast Cancer. Clinical Cancer Research, 2016, 22, 5929-5938.	7.0	66
9	APOBEC3B-Mediated Cytidine Deamination Is Required for Estrogen Receptor Action in Breast Cancer. Cell Reports, 2015, 13, 108-121.	6.4	105
10	Expression profiling of nuclear receptors in breast cancer identifies TLX as a mediator of growth and invasion in triple-negative breast cancer. Oncotarget, 2015, 6, 21685-21703.	1.8	24
11	The RNA helicase/transcriptional co-regulator, p68 (DDX5), stimulates expression of oncogenic protein kinase, Polo-like kinase-1 (PLK1), and is associated with elevated PLK1 levels in human breast cancers. Cell Cycle, 2014, 13, 1413-1423.	2.6	28
12	DEAD box RNA helicase functions in cancer. RNA Biology, 2013, 10, 121-132.	3.1	153
13	DEAD-Box RNA Helicases as Transcription Cofactors. Methods in Enzymology, 2012, 511, 347-367.	1.0	18
14	RNA helicases p68 and p72: multifunctional proteins with important implications for cancer development. Future Oncology, 2011, 7, 239-251.	2.4	101
15	The DEAD box RNA helicases p68 (Ddx5) and p72 (Ddx17): novel transcriptional co-regulators. Biochemical Society Transactions, 2008, 36, 609-612.	3.4	93
16	DExD/H box RNA helicases: multifunctional proteins with important roles in transcriptional regulation. Nucleic Acids Research, 2006, 34, 4206-4215.	14.5	360
17	Overexpression and poly-ubiquitylation of the DEAD-box RNA helicase p68 in colorectal tumours. Oncogene, 2001, 20, 7734-7743.	5.9	134
18	A chicken embryo protein related to the mammalian DEAD box protein p68 is tightly associated with the highly purified protein-RNA complex of 5-MeC-DNA glycosylase. Nucleic Acids Research, 1999, 27, 3245-3252.	14.5	58

#	Article	IF	CITATIONS
19	Expression of the â€~DEAD box' RNA helicase p68 is developmentally and growth regulated and correlates with organ differentiation/maturation in the fetus. , 1998, 184, 351-359.		99