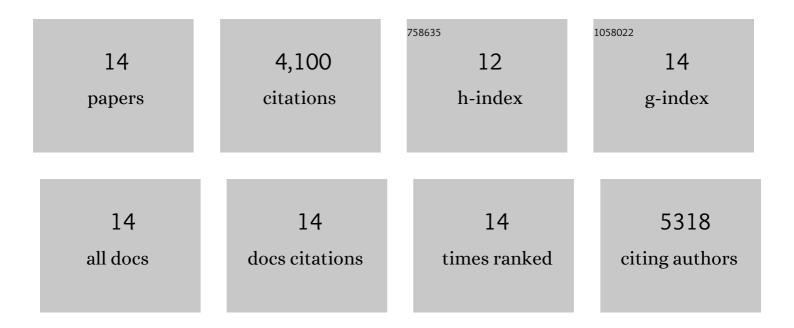
Sara Zaccara

List of Publications by Year in descending order

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SARA ZACCARA

#	Article	IF	CITATIONS
1	Reading, writing and erasing mRNA methylation. Nature Reviews Molecular Cell Biology, 2019, 20, 608-624.	16.1	1,403
2	The N6-methyladenosine (m6A)-forming enzyme METTL3 controls myeloid differentiation of normal hematopoietic and leukemia cells. Nature Medicine, 2017, 23, 1369-1376.	15.2	971
3	m6A enhances the phase separation potential of mRNA. Nature, 2019, 571, 424-428.	13.7	460
4	A Unified Model for the Function of YTHDF Proteins in Regulating m6A-Modified mRNA. Cell, 2020, 181, 1582-1595.e18.	13.5	448
5	The need for transparency and good practices in the qPCR literature. Nature Methods, 2013, 10, 1063-1067.	9.0	251
6	Identification of the m6Am Methyltransferase PCIF1 Reveals the Location and Functions of m6Am in the Transcriptome. Molecular Cell, 2019, 75, 631-643.e8.	4.5	183
7	Identification of a core TP53 transcriptional program with highly distributed tumor suppressive activity. Genome Research, 2017, 27, 1645-1657.	2.4	123
8	Circulating cell-free DNA in plasma of melanoma patients: Qualitative and quantitative considerations. Clinica Chimica Acta, 2011, 412, 2141-2145.	0.5	82
9	Live imaging of mRNA using RNA-stabilized fluorogenic proteins. Nature Methods, 2019, 16, 862-865.	9.0	71
10	Whole-genome cartography of p53 response elements ranked on transactivation potential. BMC Genomics, 2015, 16, 464.	1.2	58
11	Cooperative interactions between p53 and NFκB enhance cell plasticity. Oncotarget, 2014, 5, 12111-12125.	0.8	28
12	Nutlin-Induced Apoptosis Is Specified by a Translation Program Regulated by PCBP2 and DHX30. Cell Reports, 2020, 30, 4355-4369.e6.	2.9	18
13	Translation control can shape TP53-dependent cell fate. Molecular and Cellular Oncology, 2020, 7, 1767483.	0.3	2
14	TranSNPs: A class of functional SNPs affecting mRNA translation potential revealed by fraction-based allelic imbalance. IScience, 2021, 24, 103531.	1.9	2