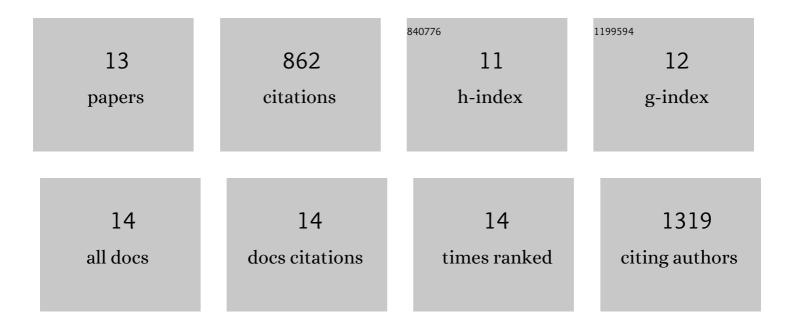
## Giuseppe Leuzzi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3454424/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Functional interrogation of DNA damage response variants with base editing screens. Cell, 2021, 184, 1081-1097.e19.	28.9	145
2	REV1-Polζ maintains the viability of homologous recombination-deficient cancer cells through mutagenic repair of PRIMPOL-dependent ssDNA gaps. Molecular Cell, 2021, 81, 4008-4025.e7.	9.7	78
3	HATtracting Nucleases to Stalled Forks. Molecular Cell, 2020, 80, 177-180.	9.7	1
4	Time for remodeling: SNF2-family DNA translocases in replication fork metabolism and human disease. DNA Repair, 2020, 95, 102943.	2.8	25
5	MCM8IP activates the MCM8-9 helicase to promote DNA synthesis and homologous recombination upon DNA damage. Nature Communications, 2020, 11, 2948.	12.8	28
6	Stimulation of CRISPR-mediated homology-directed repair by an engineered RAD18 variant. Nature Communications, 2019, 10, 3395.	12.8	85
7	The BRCT Domains of the BRCA1 and BARD1 Tumor Suppressors Differentially Regulate Homology-Directed Repair and Stalled Fork Protection. Molecular Cell, 2018, 72, 127-139.e8.	9.7	58
8	CSA and CSB play a role in the response to DNA breaks. Oncotarget, 2018, 9, 11581-11591.	1.8	23
9	Restoration of Replication Fork Stability in BRCA1- and BRCA2-Deficient Cells by Inactivation of SNF2-Family Fork Remodelers. Molecular Cell, 2017, 68, 414-430.e8.	9.7	295
10	Crosstalk between mismatch repair and base excision repair in human gastric cancer. Oncotarget, 2017, 8, 84827-84840.	1.8	13
11	WRNIP1: A new guardian of genome integrity at stalled replication forks. Molecular and Cellular Oncology, 2016, 3, e1215777.	0.7	3
12	<scp>WRNIP</scp> 1 protects stalled forks from degradation and promotes fork restart after replication stress. EMBO Journal, 2016, 35, 1437-1451.	7.8	78
13	Checkpoint-dependent and independent roles of the Werner syndrome protein in preserving genome integrity in response to mild replication stress. Nucleic Acids Research, 2014, 42, 12628-12639.	14.5	30