

Marcin M Gorski

List of Publications by Year in descending order

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Version: 2024-02-01

14
papers

1,197
citations

933447

10
h-index

1058476

14
g-index

14
all docs

14
docs citations

14
times ranked

2681
citing authors

#	ARTICLE	IF	CITATIONS
1	Selective transcriptional regulation by Myc in cellular growth control and lymphomagenesis. <i>Nature</i> , 2014, 511, 488-492.	27.8	411
2	SWI/SNF Mediates Polycomb Eviction and Epigenetic Reprogramming of the <i>INK4b-ARF-INK4a</i> Locus. <i>Molecular and Cellular Biology</i> , 2008, 28, 3457-3464.	2.3	251
3	Degradation dynamics of microRNAs revealed by a novel pulse-chase approach. <i>Genome Research</i> , 2016, 26, 554-565.	5.5	155
4	The <i>Drosophila</i> Mre11/Rad50 Complex Is Required to Prevent Both Telomeric Fusion and Chromosome Breakage. <i>Current Biology</i> , 2004, 14, 1360-1366.	3.9	108
5	Genome-wide mapping of Myc binding and gene regulation in serum-stimulated fibroblasts. <i>Oncogene</i> , 2012, 31, 1695-1709.	5.9	90
6	Disruption of <i>Drosophila</i> Rad50 causes pupal lethality, the accumulation of DNA double-strand breaks and the induction of apoptosis in third instar larvae. <i>DNA Repair</i> , 2004, 3, 603-615.	2.8	35
7	The <i>Drosophila melanogaster</i> DNA Ligase IV Gene Plays a Crucial Role in the Repair of Radiation-Induced DNA Double-Strand Breaks and Acts Synergistically With <i>Rad54</i> . <i>Genetics</i> , 2003, 165, 1929-1941.	2.9	34
8	The ISTH Bleeding Assessment Tool and the risk of future bleeding. <i>Journal of Thrombosis and Haemostasis</i> , 2018, 16, 125-130.	3.8	32
9	Whole-exome sequencing to identify genetic risk variants underlying inhibitor development in severe hemophilia A patients. <i>Blood</i> , 2016, 127, 2924-2933.	1.4	29
10	Lig4 and Rad54 Are Required for Repair of DNA Double-Strand Breaks Induced by P-Element Excision in <i>Drosophila</i> . This article is dedicated to the memory of our colleague and friend Dr. Jan C. J. Eeken, who died unexpectedly on May 24, 2002.. <i>Genetics</i> , 2005, 169, 795-806.	2.9	28
11	ImmunoChip analysis identifies novel susceptibility loci in the human leukocyte antigen region for acquired thrombotic thrombocytopenic purpura. <i>Journal of Thrombosis and Haemostasis</i> , 2016, 14, 2356-2367.	3.8	10
12	Single Nucleotide Variant rs2232710 in the Protein Z-Dependent Protease Inhibitor (ZPI, SERPINA10) Gene Is Not Associated with Deep Vein Thrombosis. <i>PLoS ONE</i> , 2016, 11, e0151347.	2.5	9
13	Next-generation DNA sequencing to identify novel genetic risk factors for cerebral vein thrombosis. <i>Thrombosis Research</i> , 2018, 169, 76-81.	1.7	4
14	Genetic variants at the chromosomal region 2q21.3 underlying inhibitor development in patients with severe haemophilia A. <i>Haemophilia</i> , 2022, 28, 270-277.	2.1	1