

Patrick Varga-Weisz

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10982073/publications.pdf>

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13
papers

1,638
citations

933447

10
h-index

1281871

11
g-index

13
all docs

13
docs citations

13
times ranked

2667
citing authors

#	ARTICLE	IF	CITATIONS
1	SATB1 targets chromatin remodelling to regulate genes over long distances. <i>Nature</i> , 2002, 419, 641-645.	27.8	456
2	Microbiota derived short chain fatty acids promote histone crotonylation in the colon through histone deacetylases. <i>Nature Communications</i> , 2018, 9, 105.	12.8	326
3	Butyrate Protects Mice from <i>Clostridium difficile</i> -Induced Colitis through an HIF-1-Dependent Mechanism. <i>Cell Reports</i> , 2019, 27, 750-761.e7.	6.4	212
4	WSTF-ISWI chromatin remodeling complex targets heterochromatic replication foci. <i>EMBO Journal</i> , 2002, 21, 2231-2241.	7.8	201
5	Maintenance of Silent Chromatin through Replication Requires SWI/SNF-like Chromatin Remodeler SMARCAD1. <i>Molecular Cell</i> , 2011, 42, 285-296.	9.7	156
6	ATP-dependent chromatin remodeling factors: Nucleosome shufflers with many missions. <i>Oncogene</i> , 2001, 20, 3076-3085.	5.9	109
7	The SNF2-Family Member Fun30 Promotes Gene Silencing in Heterochromatic Loci. <i>PLoS ONE</i> , 2009, 4, e8111.	2.5	59
8	SWI/SNF-Like Chromatin Remodeling Factor Fun30 Supports Point Centromere Function in <i>S. cerevisiae</i> . <i>PLoS Genetics</i> , 2012, 8, e1002974.	3.5	38
9	Chromatin dynamics and histone modifications in intestinal microbiota-host crosstalk. <i>Molecular Metabolism</i> , 2020, 38, 100925.	6.5	38
10	Chromatin Remodeling Factors and DNA Replication. , 2005, 38, 1-30.		22
11	Smardc1 mediates microbiota-induced inflammation in mouse and coordinates gene expression in the intestinal epithelium. <i>Genome Biology</i> , 2020, 21, 64.	8.8	13
12	The Roles of Chromatin Remodelling Factors in Replication. , 2006, 41, 91-107.		5
13	Functional Analysis of ISWI Complexes in Mammalian Cells. <i>Methods in Enzymology</i> , 2003, 377, 376-389.	1.0	3