

Angelika Feldmann

List of Publications by Year in descending order

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7
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1478505

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1720034

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docs citations

9
times ranked

1737
citing authors

#	ARTICLE	IF	CITATIONS
1	Transcription Factor Occupancy Can Mediate Active Turnover of DNA Methylation at Regulatory Regions. PLoS Genetics, 2013, 9, e1003994.	3.5	194
2	PRC1 Catalytic Activity Is Central to Polycomb System Function. Molecular Cell, 2020, 77, 857-874.e9.	9.7	184
3	Cohesin Disrupts Polycomb-Dependent Chromosome Interactions in Embryonic Stem Cells. Cell Reports, 2020, 30, 820-835.e10.	6.4	129
4	A genome-scale map of DNA methylation turnover identifies site-specific dependencies of DNMT and TET activity. Nature Communications, 2020, 11, 2680.	12.8	97
5	The SET1 Complex Selects Actively Transcribed Target Genes via Multivalent Interaction with CpG Island Chromatin. Cell Reports, 2017, 20, 2313-2327.	6.4	86
6	FBXL19 recruits CDK-Mediator to CpG islands of developmental genes priming them for activation during lineage commitment. ELife, 2018, 7, .	6.0	22
7	CDK-Mediator and FBXL19 prime developmental genes for activation by promoting atypical regulatory interactions. Nucleic Acids Research, 2020, 48, 2942-2955.	14.5	9