## Devin A King

## List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Recognition of Histone Crotonylation by Taf14 Links Metabolic State to Gene Expression. Molecular Cell, 2019, 76, 909-921.e3.	9.7	83
2	Somatic mutation distributions in cancer genomes vary with three-dimensional chromatin structure. Nature Genetics, 2020, 52, 1178-1188.	21.4	79
3	Carcinogen susceptibility is regulated by genome architecture and predicts cancer mutagenesis. EMBO Journal, 2017, 36, 2829-2843.	7.8	71
4	Endothelial deletion of Ino80 disrupts coronary angiogenesis and causes congenital heart disease. Nature Communications, 2018, 9, 368.	12.8	71
5	The INO80 Complex Requires the Arp5-les6 Subcomplex for Chromatin Remodeling and Metabolic Regulation. Molecular and Cellular Biology, 2016, 36, 979-991.	2.3	48
6	Endocardial/endothelial angiocrines regulate cardiomyocyte development and maturation and induce features of ventricular non-compaction. European Heart Journal, 2021, 42, 4264-4276.	2.2	41
7	Set5 and Set1 cooperate to repress gene expression at telomeres and retrotransposons. Epigenetics, 2014, 9, 513-522.	2.7	28
8	INO80 Chromatin Remodeling Coordinates Metabolic Homeostasis with Cell Division. Cell Reports, 2018, 22, 611-623.	6.4	28
9	The INO80 chromatin remodeler sustains metabolic stability by promoting TOR signaling and regulating histone acetylation. PLoS Genetics, 2018, 14, e1007216.	3.5	23
10	Genome-wide profiles of UV lesion susceptibility, repair, and mutagenic potential in melanoma. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2021, 823, 111758.	1.0	4
11	Retinoblastoma protein regulates carcinogen susceptibility at heterochromatic cancer driver loci. Life Science Alliance, 2022, 5, e202101134.	2.8	4
12	Transcriptome profiling of Set5 and Set1 methyltransferases: Tools for visualization of gene expression. Genomics Data, 2014, 2, 216-218.	1.3	3