

# A (3D-Nuclear) Space Odyssey: Making Sense of Hi-C Maps

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Comparative Genomics and Transcriptomics To Analyze Fruiting Body Development in Filamentous Ascomycetes. <i>Genetics</i> , 2019, 213, 1545-1563.	2.9	14
2	Modern epigenetics methods in biological research. <i>Methods</i> , 2021, 187, 104-113.	3.8	97
3	Approaches for studying epigenetic aspects of the human genome. , 2021, , 155-209.		0
4	Identification of Enhancers and Promoters in the Genome by Multidimensional Scaling. <i>Genes</i> , 2021, 12, 1671.	2.4	1
5	OUP accepted manuscript. <i>Briefings in Bioinformatics</i> , 2021, , .	6.5	8
6	FISH Going Meso-Scale: A Microscopic Search for Chromatin Domains. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 753097.	3.7	11
8	Breaking boundaries: Pan BETi disrupt 3D chromatin structure, BD2-selective BETi are strictly epigenetic transcriptional regulators. <i>Biomedicine and Pharmacotherapy</i> , 2022, 152, 113230.	5.6	8
9	RNA polymerase II pausing regulates chromatin organization in erythrocytes. <i>Nature Structural and Molecular Biology</i> , 2023, 30, 1092-1104.	8.2	1