

Reviewer Report

Title: Chromatin conformation capture (Hi-C) sequencing of patient-derived xenografts: analysis guidelines

Version: Revision 1 **Date: 1/31/2021**

Reviewer name: Joanna Achinger-Kawecka

Reviewer Comments to Author:

The authors answered all my questions to my complete satisfaction. Analysis of loops and TAD overlaps provide more in-depth and biologically relevant results that better support their conclusions.

Methods

Are the methods appropriate to the aims of the study, are they well described, and are necessary controls included? Choose an item.

Conclusions

Are the conclusions adequately supported by the data shown? Choose an item.

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