Enhancer-promoter communication in the 3D genome: is close enough, enough?

The large regulatory landscapes of developmental genes in mammals often correspond to topologically associating domains (TADs) that are formed through the process of cohesin-driven loop extrusion, and whose extent is constrained by orientation-dependent CTCF binding. I will discuss the potential role of this level of 3D genome organisation on the function of long-range enhancers.

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**Tuesday, March 22, 2022**  
11:30am - 12:30pm  

[Zoom Link](https://medicine.yale.edu/genetics/events/seminars.aspx)  
pw: 473124