Journal Club Abstract Research Center for Mathematics on Chromatin Live Dynamics (RcMcD)

A simple biophysical model emulates budding yeast chromosome condensation Soya Shinkai

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Abstract: The authors used a coarse-grained Brownian simulation of a budding yeast chromosome to explore chromatin behavior during chromosome condensation. Two-type intrachromosomal interactions by condensin were modeled. They compared the predictions from these simulations with 4C data on budding yeast chromosome 5. This study shows that stochastic pairwise interactions of a chromatin chain, mediated by condensin, provide a close fit to observed behavior in budding yeast.

References:

1. Tammy MK Cheng et al., <u>eLife (2015) 4 e05565</u>.