

Agilent Webinar Series

Ask the Experts

Webinar 4:
Improving resolution through Capture HiC



Register now for the free Agilent Webinar Series

Want to enhance your skills in the Genomics world?

Have an hour to spare?

Join us on Tuesdays for a series of science topics presented by Agilent experts.

Presented by our field scientists, you'll learn tips and tricks to be more successful in your work.



Webinar 4: Improving resolution through Capture HiC

Date: January 11, 2022

Time: 11:00 CET

Speaker: Justin Lenhart, Agilent Technologies

High throughput chromatin conformation capture (HiC) provides a means to generate genome-wide chromatin architecture maps.

Despite this utility, HiC is a genome-wide sequencing assay that can be limited due to relatively high costs while simultaneously failing to attain high-enough resolution to identify individual chromatin loops. To address these problems, Agilent has partnered with Arima Genomics to generate a streamlined target enrichment workflow for HiC (capture HiC).

During this webinar, this streamlined capture HiC workflow will be presented, highlighting how it allows researchers to focus HiC reads on targeted areas of the genome, reducing sequencing read requirements (and, hence, cost) for high-resolution HiC workflows.

Please, register to the event [here](#)

This information is subject to change without notice
PR7000-8286

© Agilent Technologies, Inc 2021
Published in the Germany, October, 2021
DGG_FY22_SCITUES_EM-04