preciseTAD: A machine learning framework for precise 3D domain boundary prediction at base-level resolution

Additional File Legends

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# Additional File 1

An example script for applying Arrowhead to in situ Hi-C data (HIC001-HIC018) to obtain chromosome-specific TAD boundaries on the GM12878 cell line at 5kb, 10kb, 25kb, 50kb, 100kb resolutions. An example of deepTools commands to generate plotProfiles and plotHeatmaps is also provided.