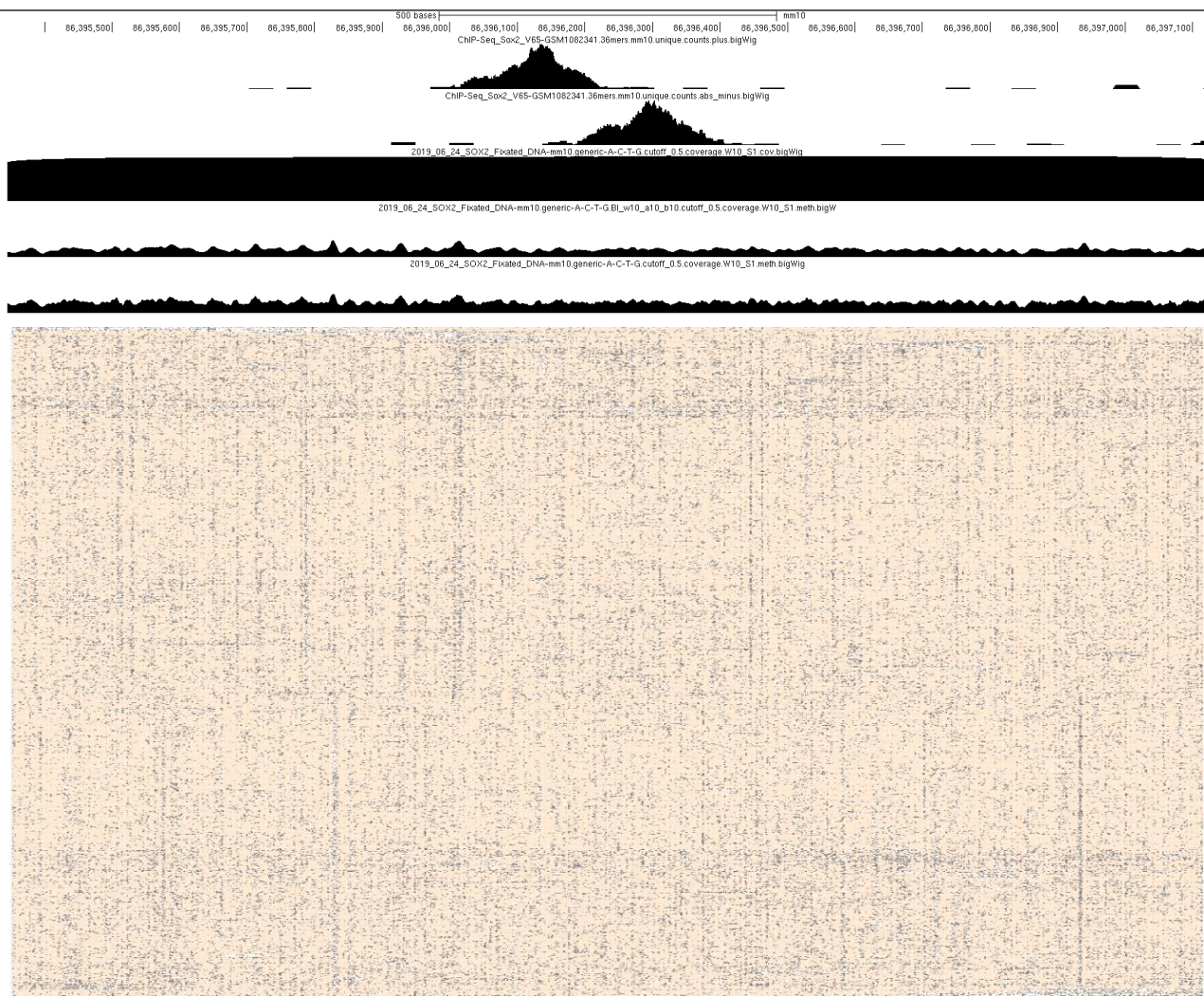


Mapping stats

Species	Assay	#	Mapping	Base Contexts	Reads	Total Bases	Mean Read Length	Median Read Length
mm10	CLAPP	2019_07_19_Mouse_Cerevisiae-Sox2_125nM-mm10	Tombo 1.3 denovo	generic-A-C-T-G	18,038	14,674,546	814	583

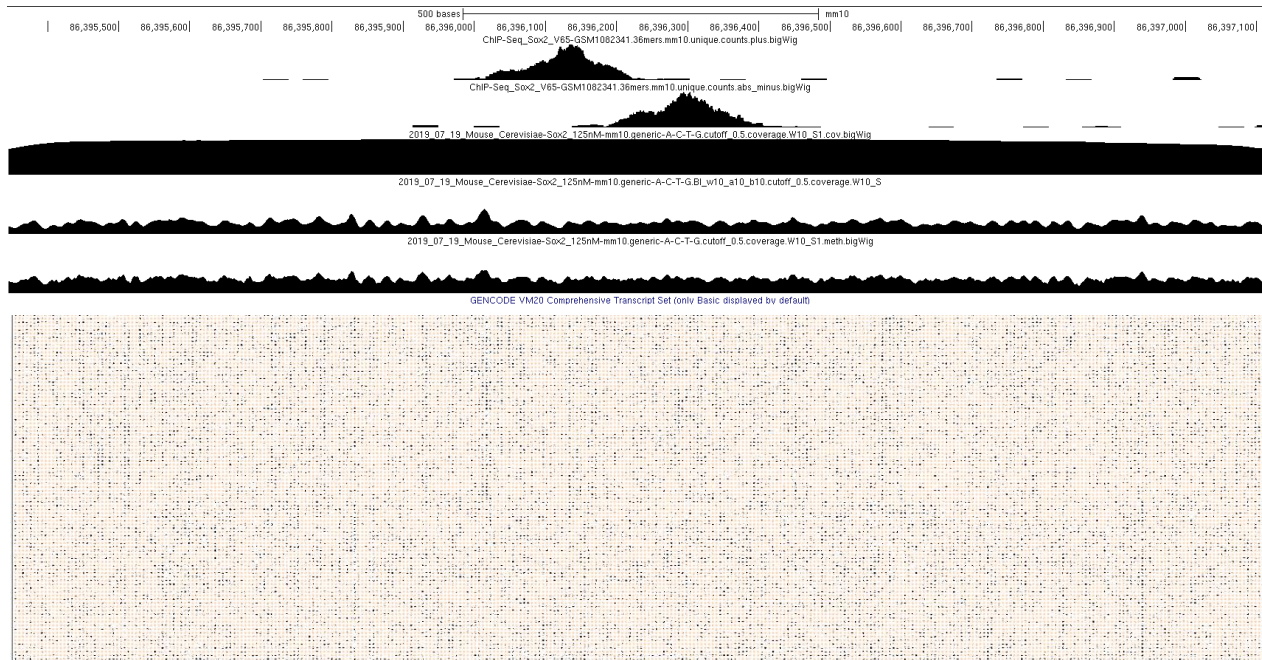
Single-molecule (1000
reads subsample)
 $p=0.3$ binarization,

500nM sample



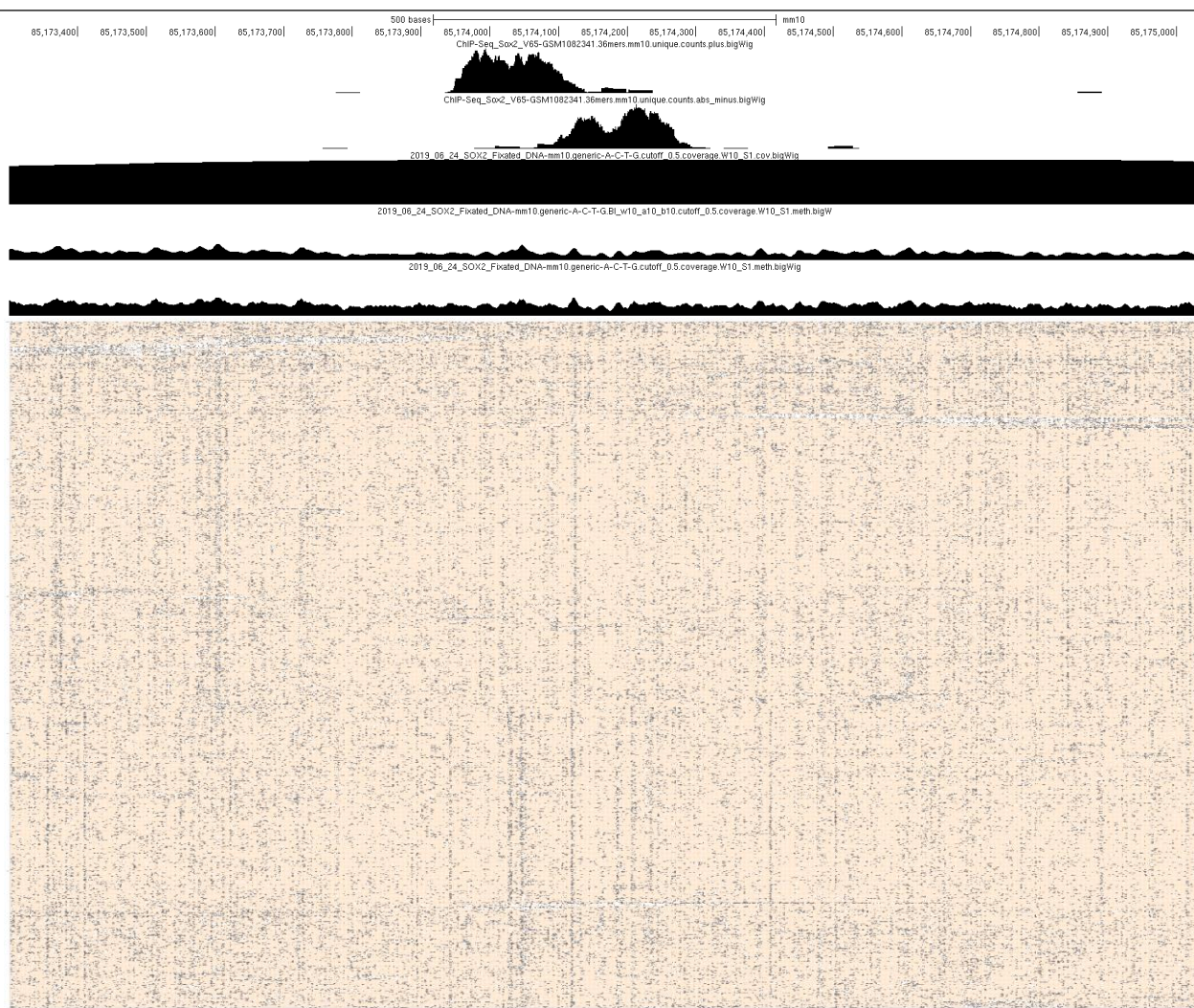
Single-molecule (1000
reads subsample)
 $p=0.3$ binarization,

125nM sample



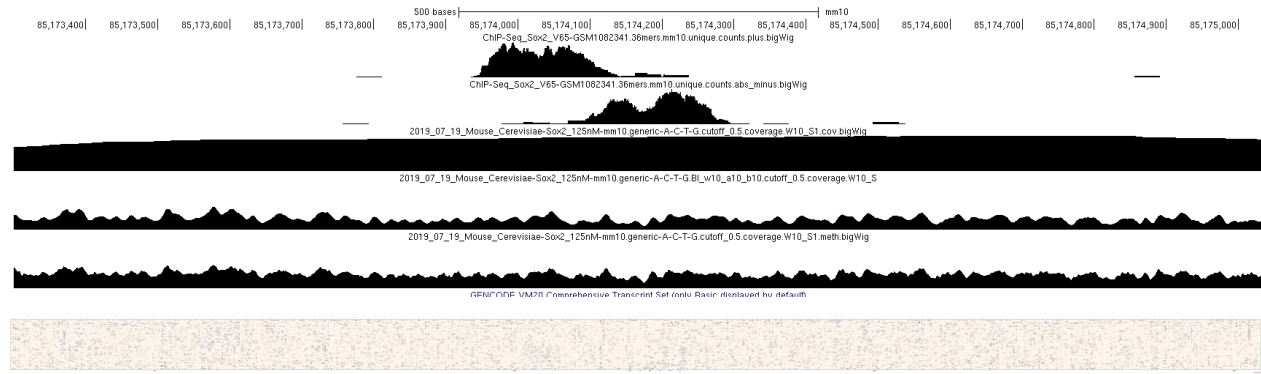
Single-molecule (1000
reads subsample)
 $p=0.3$ binarization

500nM sample



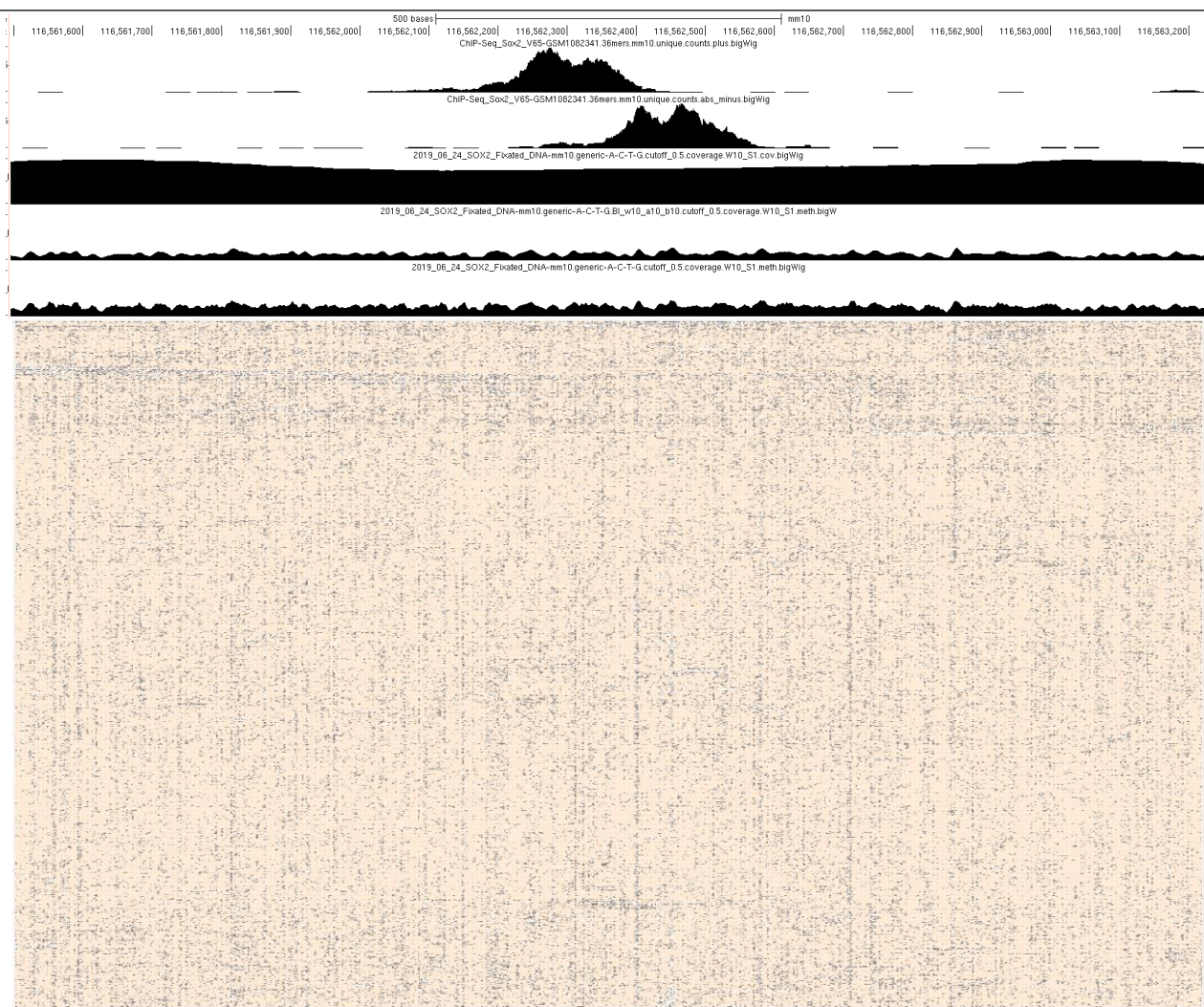
Single-molecule (1000
reads subsample)
 $p=0.3$ binarization

125nM sample



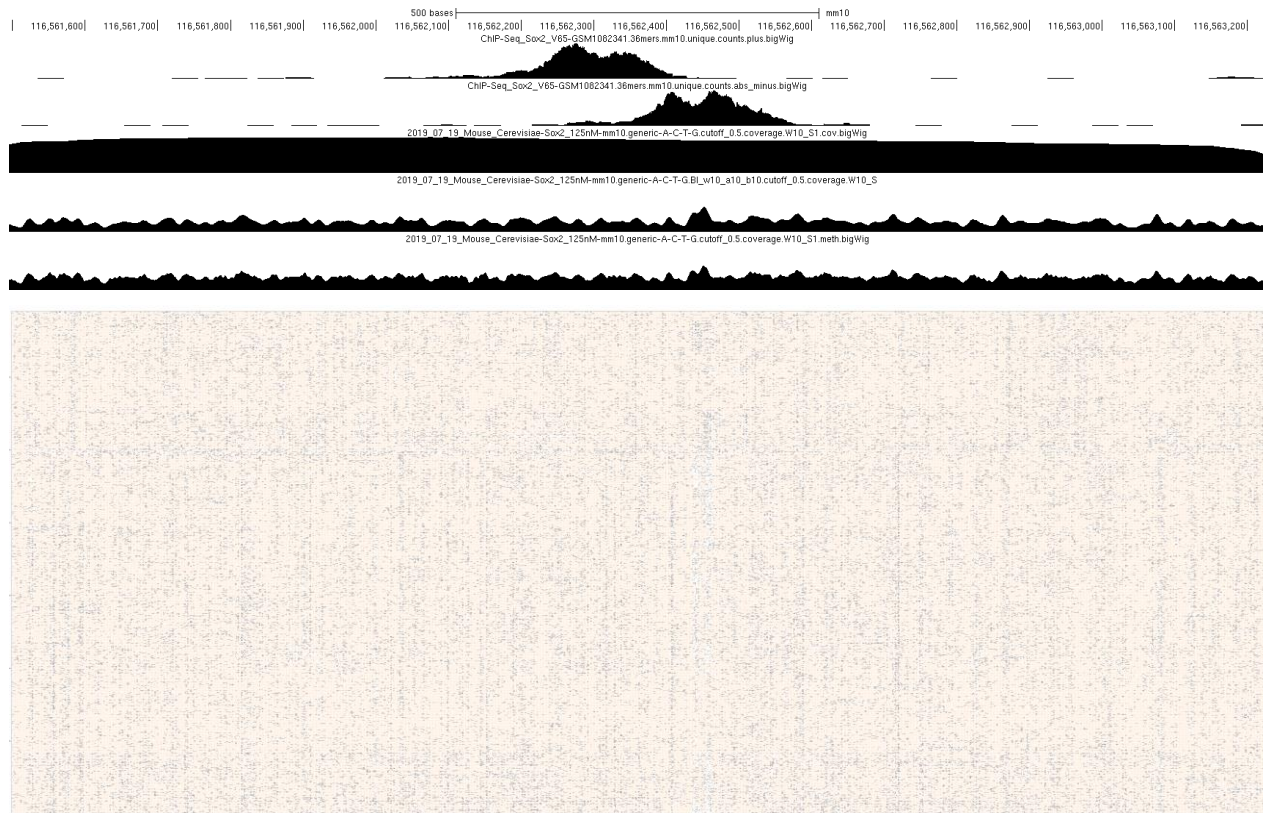
Single-molecule (1000
reads subsample)
 $p=0.3$ binarization

500nM sample



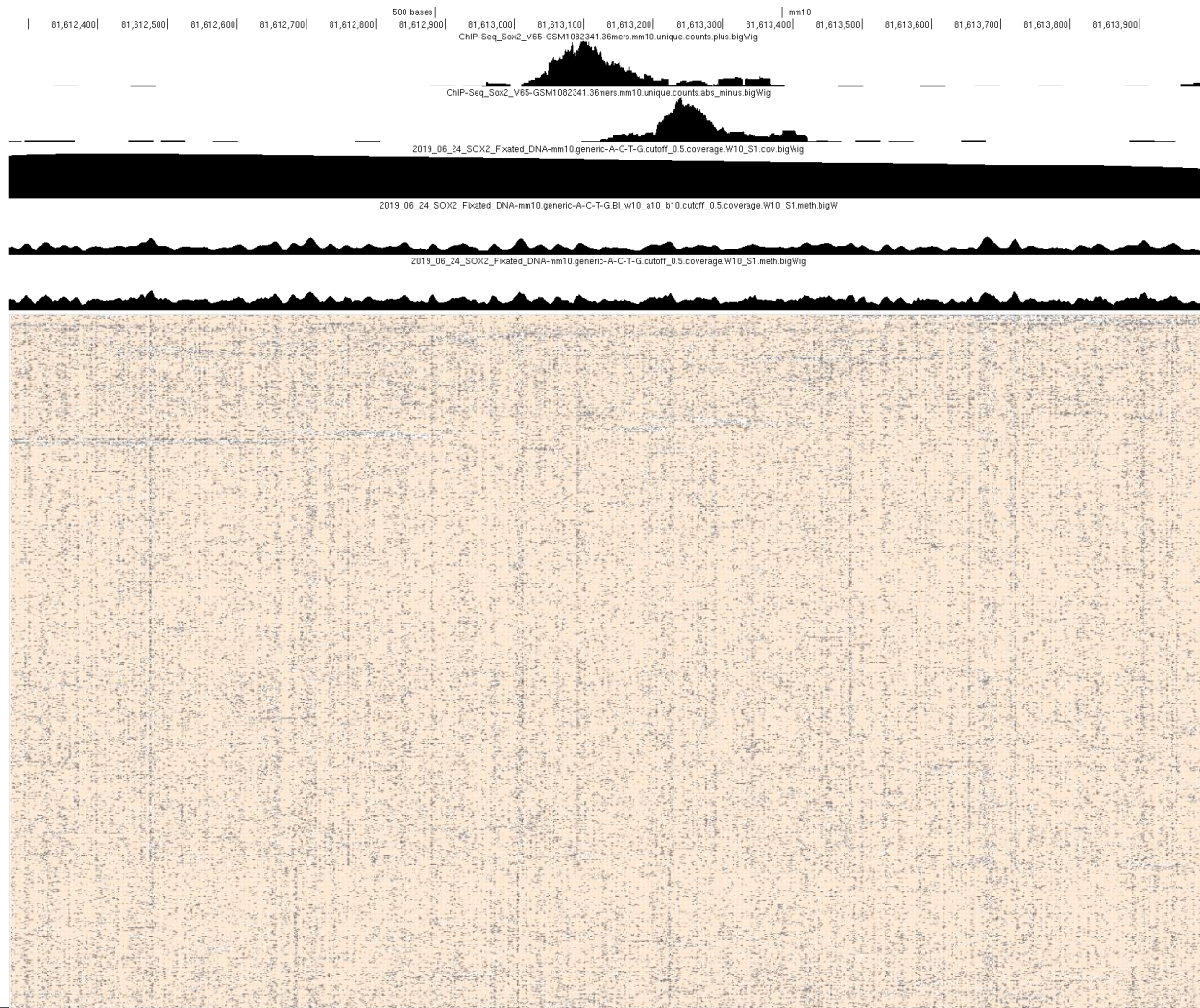
Single-molecule (1000
reads subsample)
 $p=0.3$ binarization

125nM sample



Single-molecule (1000
reads subsample)
 $p=0.3$ binarization

500nM sample



Single-molecule (1000
reads subsample)
 $p=0.3$ binarization

125nM sample

