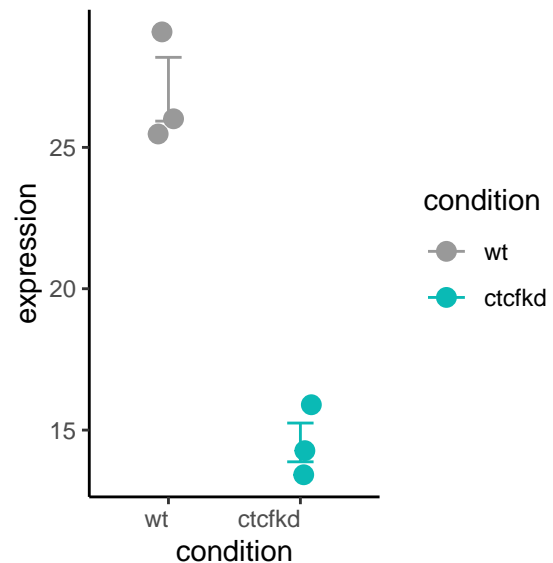
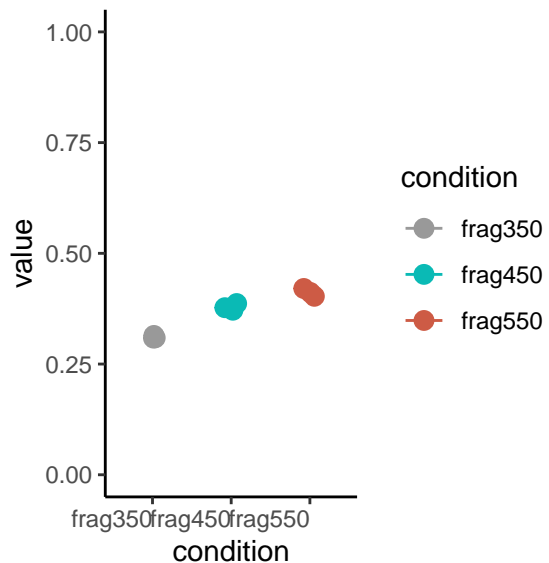


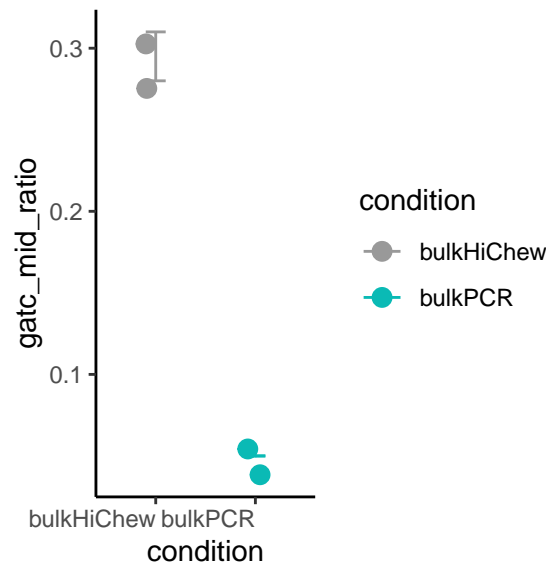
RNAseq CTCF expression
p-value: 0.00167



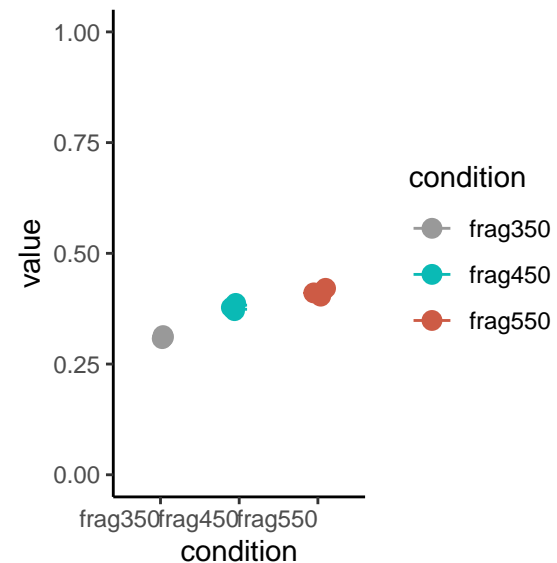
valid pair ratio
p-value: 0.0389



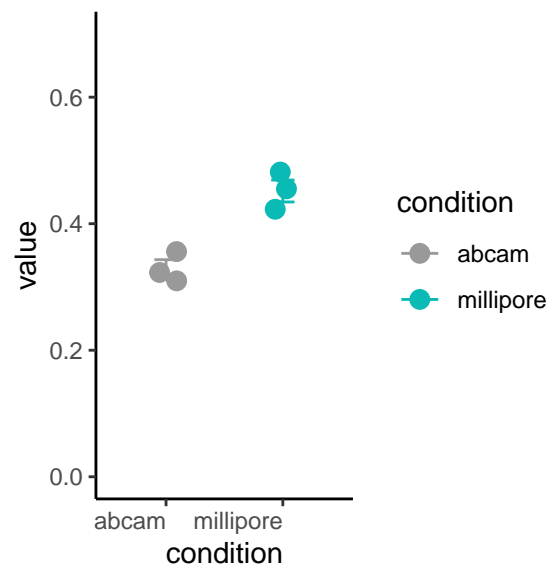
sfig 5e ab efficiency
p-value: 0.0389



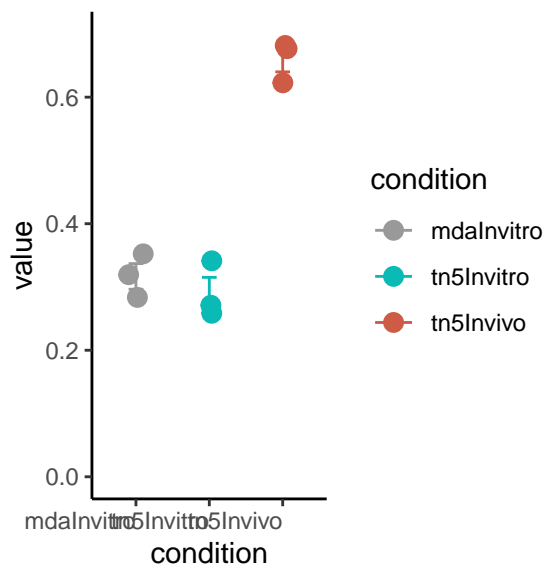
frag size valid pair ratio



sfig 5g frag size valid pair ratio
p-value: 0.00609



sfig 5j lib compare valid pair ratio
p-value: 0.00609



sfig 5d ab pulldown

