Figure S1 Spatial accuracy evaluation using all the binding events

A) Fraction of predicted GABP binding events with a motif within the given distance with event discovery by GEM, GPS, SISSRs, MACS, cisGenome, QuEST and PeakRanger. Events shown were all the events predicted by each of the seven methods and had a GABP motif within 100bp. **B**) Fraction of predicted CTCF binding events with a motif within the given distance with event discovery by GEM, GPS, SISSRs, MACS, cisGenome, QuEST, FindPeaks, spp-wtd and spp-mtc. Events shown were all the events predicted by each of the nine methods and had a CTCF motif within 100bp.



