



Figure 5 - figure supplement 2 | Retention of intergenic genomic segments in accessible- and inaccessible- chromatin is robust to changes in sampled region size and sequence identity threshold.

a, b, Repeating the procedure used in **Fig. 5c** and **Fig. 5d** sampling segments of different length (150 bp) and **c,d** increasing the LiftOver minMatch (0.5).