

Supplemental Figure S4. Densities of top four enriched motifs at the end of the chromosomal *q* arm of the HG005 dataset (Chinese trio, son). Only the arms covered by at least 25 reads are displayed. Note that none of the chromosomal *p* arms pass this threshold in this dataset. Genomic coordinates are given in Kbp. Vertical red dashed lines denote the position of the boundary of the annotated telomeric tract.

