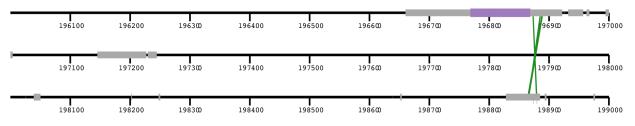
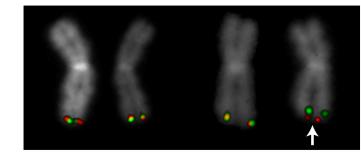
Figure S6. Inversion Identification and Validation. A schematic of clones used to identify 5 large inversions is shown (coloring as in Figure S3). Each of the depicted clones has an "inverted" orientation. Despite the presence of duplicated sequences (grey boxes), the clones have a single, best placement. For each locus, FISH validation is shown with the inverted allele indicated by the arrow. The 8q24 inversion was only observed in 3/20 nuclei and may be mosaic.

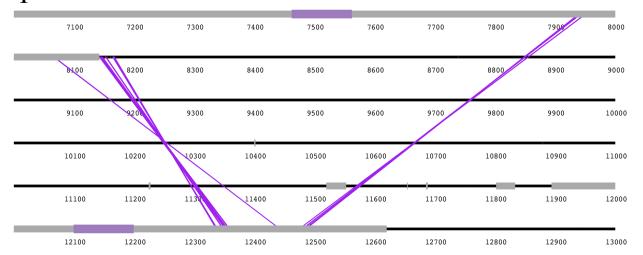
Figure S6

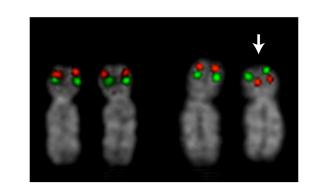
3q29



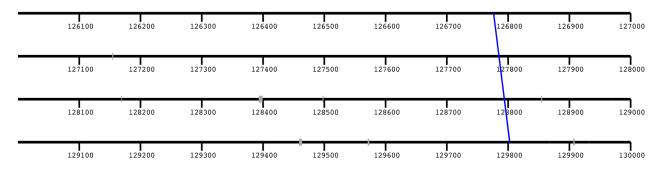


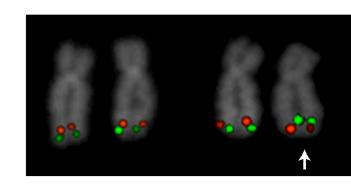
8p23



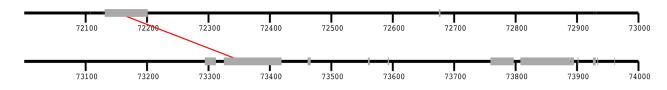


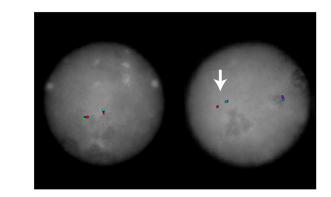
8q24





15q24





17q12

-	•	++							_	
	31100	31200	31300	31400	31500	31600	31700	31800	31900	32000
-										
	32100	32200	32300	32400	32500	32600	32700	32800	32900	33000
•										
	33100	33200	33300	33400	33500	33600	33700	33800	33900	34000

