Relative copy number by sort-seq Mean BrdU content chrl 1.6 1.2 2.0 chrll 1.6 8.0 2.0 chrIII 1.6 0.8 2.0 chrlV 1.6 1.2 8.0 2.0 chrV 1.6 1.2 8.0 2.0 chrVI 1.6 1.2 0.8 2.0 chrVII 1.6 1.2 0.8 2.0 chrVIII 1.6 1.2 ₩ 0.8 c chrlX 1.6 1.2 8.0 2.0 chrX 1.6 1.2 8.0 2.0 chrXI 1.6 1.2 8.0 2.0 chrXII 1.6 0.8 2.0 chrXIII 1.6 8.0 2.0 chrXIV 1.6 0.8 2.0 chrXV 1.6 1.2 0.8 2.0 1.6 0.8 200 300 800 1000 1100 1200 1300 1400 1500 Genomic position (kb)