Figure S1 Relative copy number by sort-seq 

Mean BrdU content chrl 1.6 1.2 8.0 2.0 chrll 1.6 1.2 0.8 2.0 chrIII 1.6 1.2 0.8 2.0 chrlV 1.6 1.2 8.0 2.0 chr∨ 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 1.2 0.8 2.0 chrXIII 1.6 1.2 0.8 2.0 chrXIV 1.6 1.2 0.8 2.0 chrXV 1.6 1.2 0.8 2.0 chrXVI 1.6 1.2 0.8

700

800

Genomic position (kb)

900

1000

1100

1200

1300

1400

1500

100

200

300