



Figure S1 Quantitative analysis of the relative amounts of intact chromosomes in the homozygous S288C diploid strain analyzed by PFGE after the *cdc14-1* block-and-release. (A) The intensities of each ethidium-bromide-stained band in Fig. 2D and two more independent experiments were normalized to the sum of the band signals corresponding to chromosomes I, VI, and III, and then to the intensities at time 0 (time of the *cdc14* block). Error bars represent s.e.m (N=3). (B) In-gel and in-well signals of unsaturated exposures of the Southern blots shown in Fig. 2D were quantified and normalized to that of time 0' (*cdc14*-block). Overall is the sum of both in-gel and in-well signals.