Supplementary Figure S3. Replication origin activation profiles for all chromosomes (I – XVI) in wild-type, *mec1-1* and *rad53-1* cells. Chromosomes I-XVI are shown as cartoons at the top of each figure. The distribution of replication origins on the chromosome is indicated by solid black bars. The letter E indicates early-firing origins (Raghuraman *et al*, 2001). Each point on the graph represents the average of three independent experiments. Standard error is indicated by error bars. Significance levels 1 and 2 correspond to one standard deviation and one half of a standard deviation above background, respectively. Negative controls used for normalization are marked by a black square in wild-type profiles. CEOs are indicated by their ARS number in wild-type profiles and are circled in *mec1-1* and/or *rad53-1* profiles.

#### **Supplementary Figure S3**















