



Figure 4 – figure supplement 1. Checkpoint restoration in *exo1*Δ *sgs1*Δ interchromosomal assay strains. **(A)** Survival frequency in the plating assay for the interchromosomal strains with the indicated genotypes. Bars represent mean values from at least 2 plating assays per strain. Error bars represent standard deviation. * $p < .05$, ** $p < .01$, *** $p < .001$, **** $p < .0001$ **(B)** HO cutting efficiency as measured by qPCR with primers flanking the HO cut site (top) and measurement of ssDNA at 640 bp from the HO DSB using the qPCR-based resection assay (bottom) as in Figure 3 – figure supplement 1. Each point represents the mean of 4 biological replicates and error bars represent standard deviation.