



Figure S6. Whole genome-resequencing of *TAO3* allele replacement strain (YAD331, (Deutschbauer and Davis 2005) in comparison to S288c reference strain. Segmentation plots where each chromosome is divided into bins (shown as blue dots) depicted on the x-axis. Copy number of each bin relative to S288c reference strain is shown on the y-axis. Red dotted lines show the position where a chromosomal region (bin) would appear if there is a duplication (above 1.5) or deletion (below 0.5).