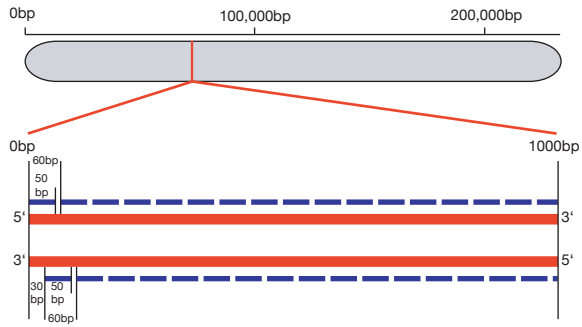


Supplemental Figure 1. Borneman et al.,



Supplemental Figure 1. Tiling array designs used in *S. mikatae*, *S. bayanus* and *C. albicans*. For each species, oligos were designed to tile the genome sequence contigs with a 50bp oligonucleotide every 60 bp on the Watson strand. For *S. mikatae* and *S. bayanus*, this same spacing was used on the Crick strand offset 30 bp compared to those oligonucleotide on the Watson strand. For *C. albicans*, limitations on the number a features allowed on the arrays resulted in a 50 bp oligonucleotide being spaced every 120 bp, offset 30 bp.