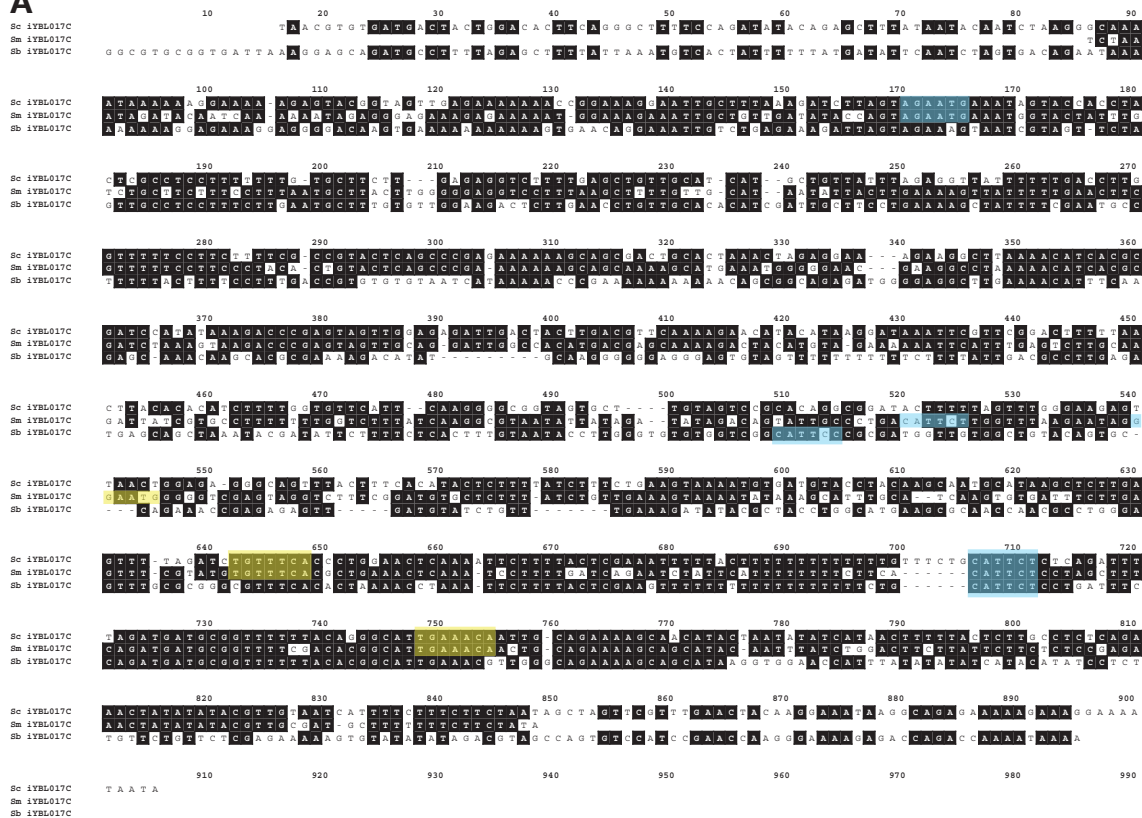
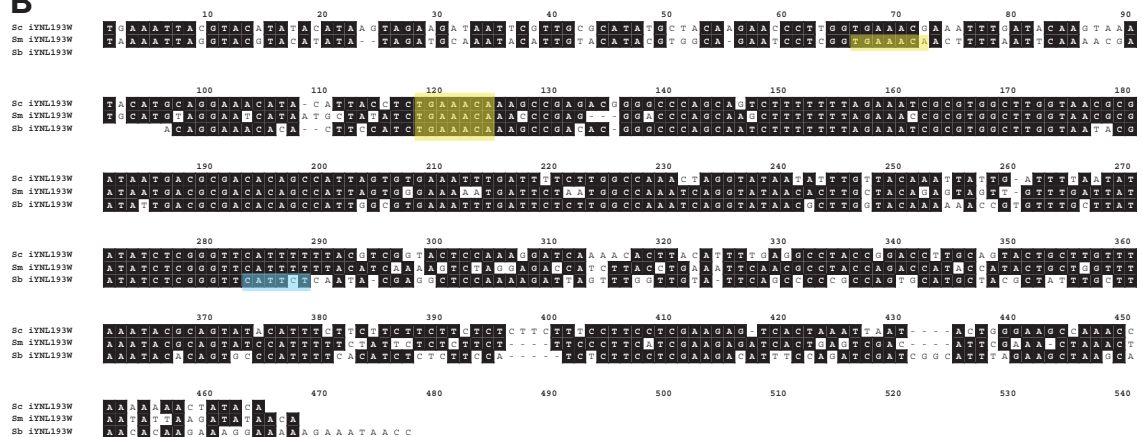


Supplemental Figure 3. Borneman et al.

A



B



Supplemental Figure 3. Sequence alignments of **(A)** the YBL017C/YBL016C and **(B)** the YNL193W/YNL192W intergenic region from *S. cerevisiae*, *S. mikatae* and *S. bayanus*. The position of Ste12 (yellow) and Tec1 (blue) consensus sequences are also highlighted.