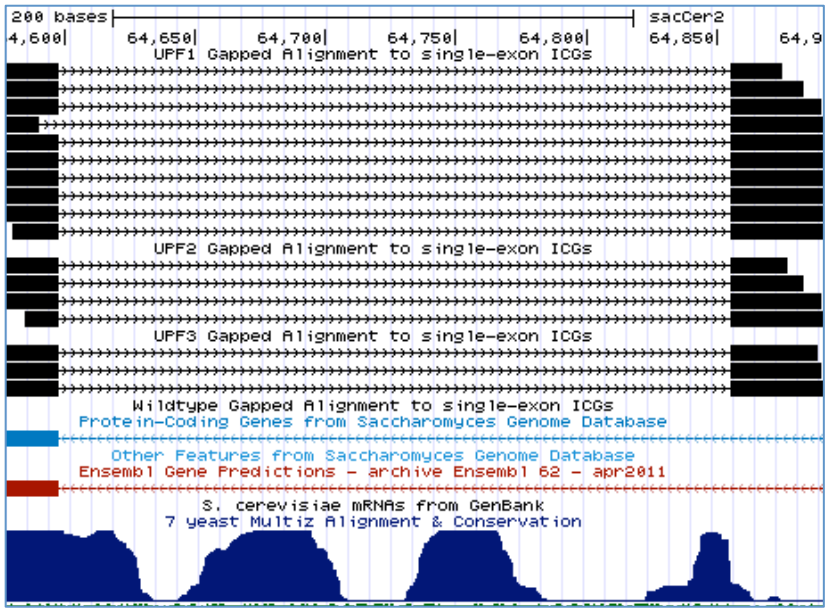
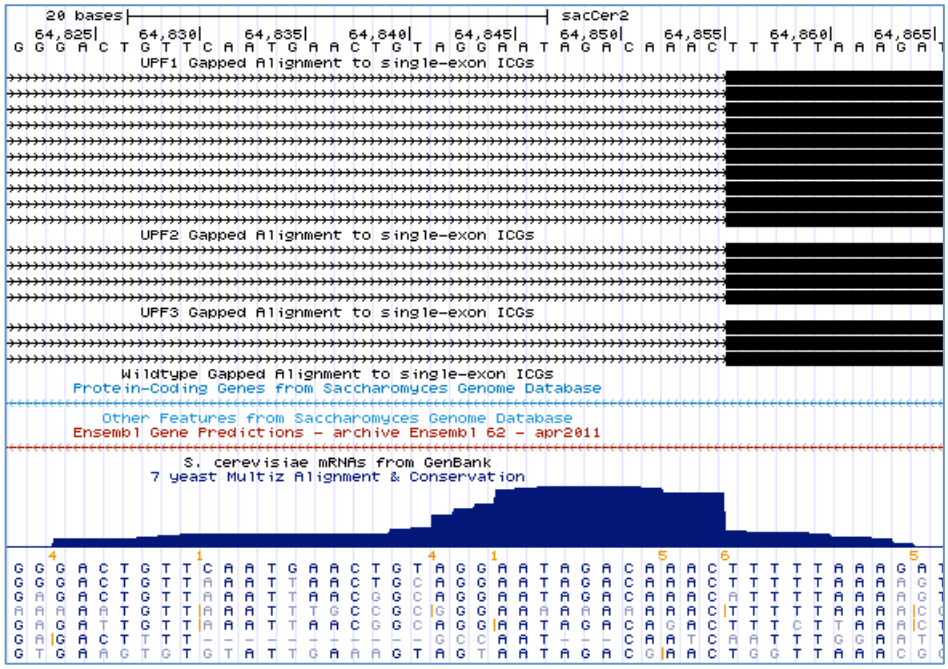
**A**


200 bases
sacCer2
4,600
64,650
64,700
64,750
64,800
64,850
64,9
UPF1 Gapped Alignment to single-exon ICGs
UPF2 Gapped Alignment to single-exon ICGs
UPF3 Gapped Alignment to single-exon ICGs
Wildtype Gapped Alignment to single-exon ICGs
Protein-Coding Genes from Saccharomyces Genome Database
Other Features from Saccharomyces Genome Database
Ensembl Gene Predictions - archive Ensembl 62 - apr2011
S. cerevisiae mRNAs from GenBank
7 yeast Multiz Alignment & Conservation

**B**


20 bases
sacCer2
64,825
64,830
64,835
64,840
64,845
64,850
64,855
64,860
64,865
UPF1 Gapped Alignment to single-exon ICGs
UPF2 Gapped Alignment to single-exon ICGs
UPF3 Gapped Alignment to single-exon ICGs
Wildtype Gapped Alignment to single-exon ICGs
Protein-Coding Genes from Saccharomyces Genome Database
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