























Keys:

-  = Represents the chromosome you are viewing
-  = Watson strand ORF
-  = Crick strand ORF
-  = tRNA_gene
-  = ncRNA_gene
-  = rRNA_gene
-  = snRNA_gene
-  = snoRNA gene
-  = LTR_retrotransposon
-  = ARS
-  = Dubious ORF
-  = gene_group
-  = long_terminal_repeat
-  = blocked_reading_frame
-  = origin_of_replication
-  = matrix_attachment_site
-  = mating_type_region
-  = pseudogene
-  = silent_mating_type_cassette_array
-  = telomere
-  = telomerase_RNA_gene
-  = transposable_element_gene

Notes:

1. Click on light blue bar to retrieve features left & right
2. Click on any gene or ORF name, any little colorful bar, or centromere to retrieve the corresponding LOCUS entry in SGD
3. A white box in the middle of a blue or red bar indicates that there is an intron in the ORF. The size of the white box represents the length of the intron.
4. For a definition of the features listed in the key please see the [glossary](#).