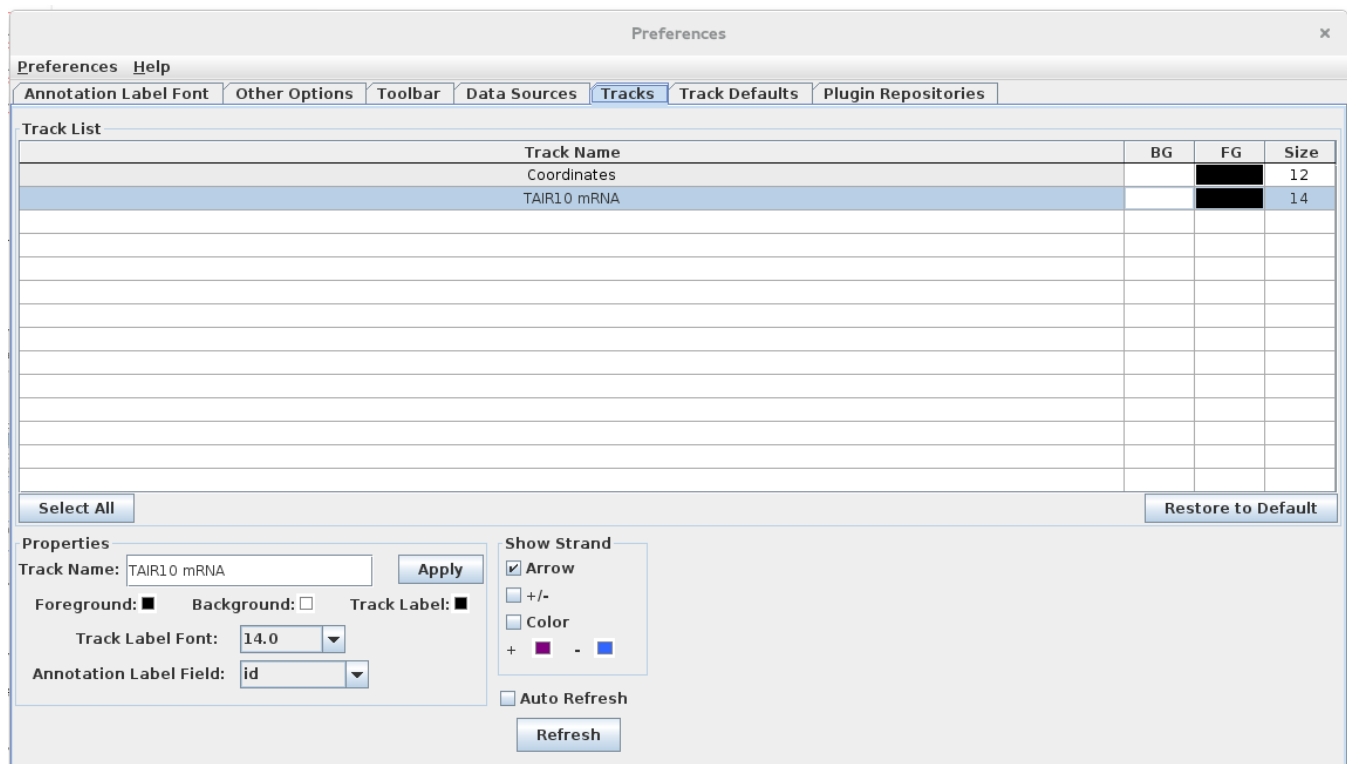


## General Function Checklist

1. Open the **A\_thaliana\_Jun\_2009** genome.
2. Navigate to the following coordinates: **Chr1:635,037-637,360**
3. Right click on the track label and select **Optimize Stack Height**.
4. Right click on the white space within a track.
5. Select **Customize...**



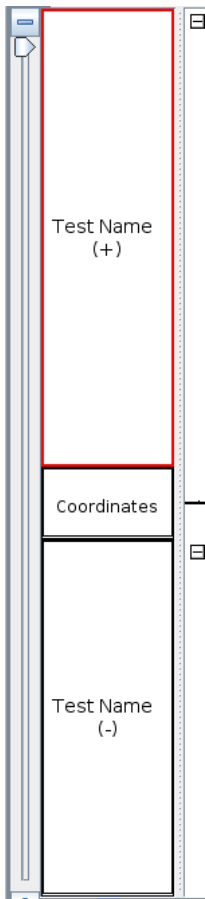
- This action opened the **Track Preferences** tab with the selected track highlighted.

- ☐ mac
- ☐ linux
- ☐ windows

Change the **Track Name** to **Test Name**, then select **Apply**.

**Properties**

Track Name:

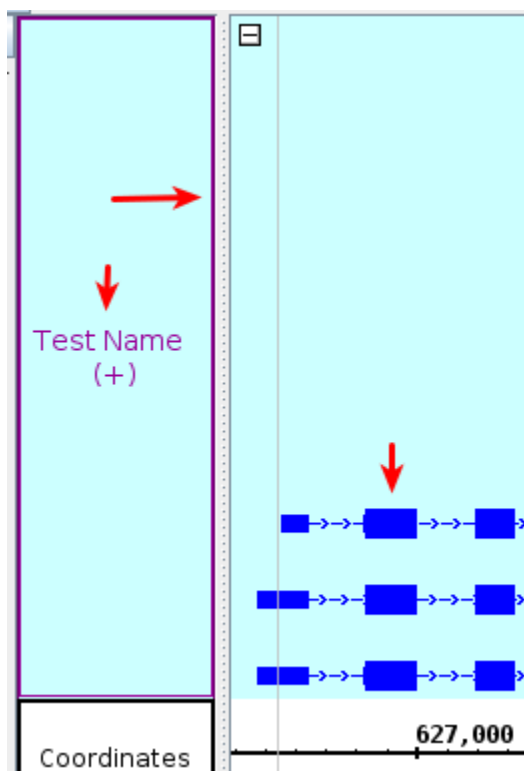


- The track names were updated.

- ☐ mac
- ☐ linux
- ☐ windows

Change the foreground, background, and track label colors to match the following image:

Foreground:  Background:  Track Label:



- The main view of IGB is now colored correctly with the three colors you chose.

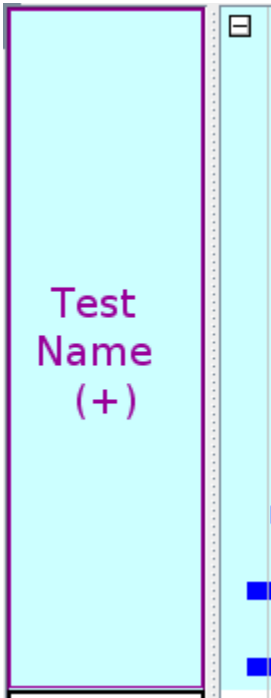
- ☐ mac
- ☐ linux
- ☐ windows

Change the track label font size to **20.0**.

Track Label Font:

20.0

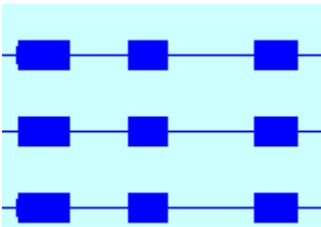




- The font increased appropriately.

- ☐ mac
- ☐ linux
- ☐ windows

Uncheck the **Arrow** option in the **Show Strand** box.



- The barbed wire disappeared from the intron sections of gene models.

- ☐ mac
- ☐ linux
- ☐ windows

Check the **+/-** option in the **Show Strand** box to combine the plus and minus strands.

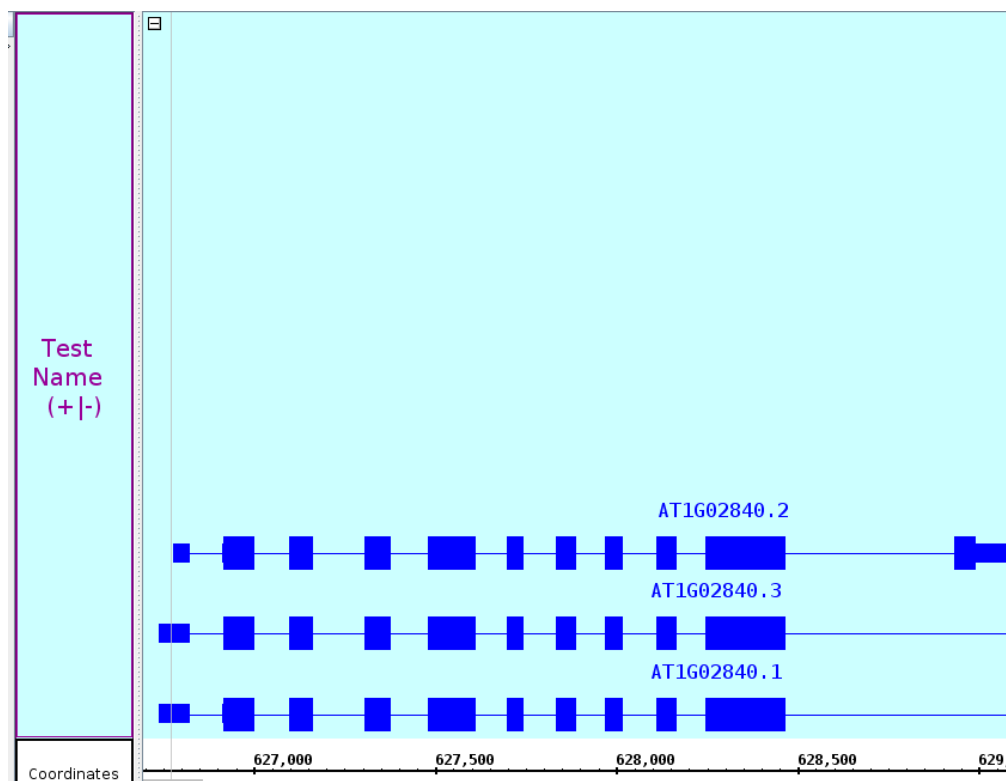
**Show Strand**

☐ Arrow

☒ +/- 

☐ Color

+  - 



- The two strands are now one.

- ☐ mac
- ☐ linux
- ☐ windows

Check the **Color** option in the **Show Strand** box.

**Show Strand**

☐ Arrow

☐ +/-

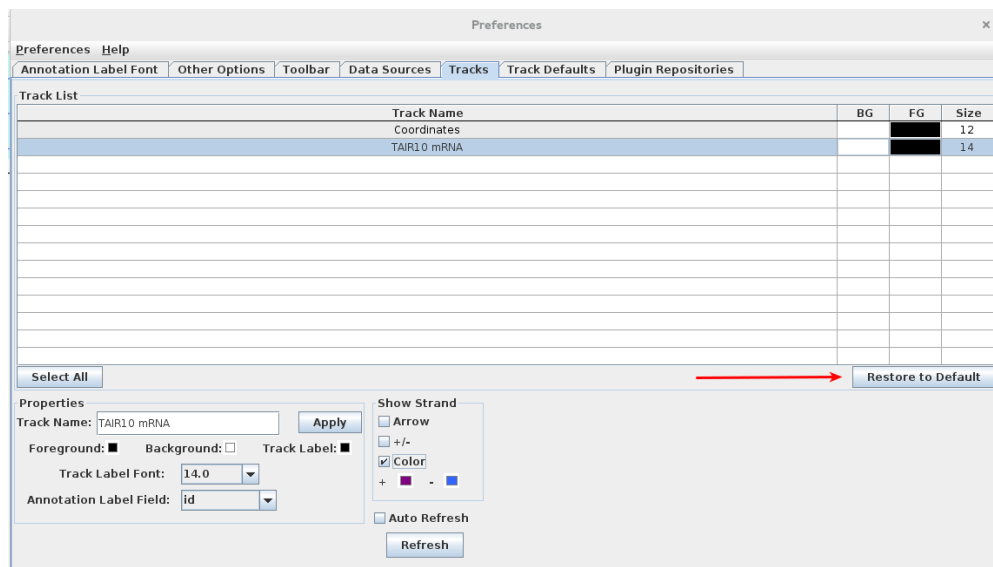
☒ Color 

+  - 



- ☐ mac
- ☐ linux
- ☐ windows

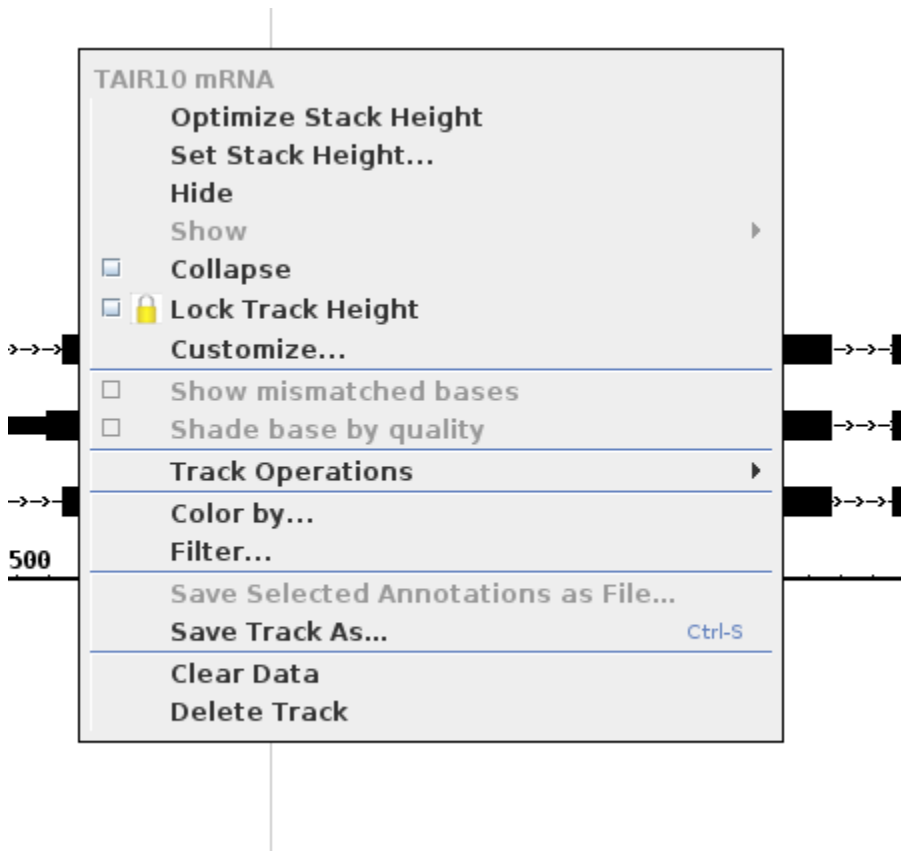
Click **Restore to Default** in the **Tracks** tab in **Preferences**.



- ☐ mac
- ☐ linux
- ☐ windows

## Adding and Removing Tracks

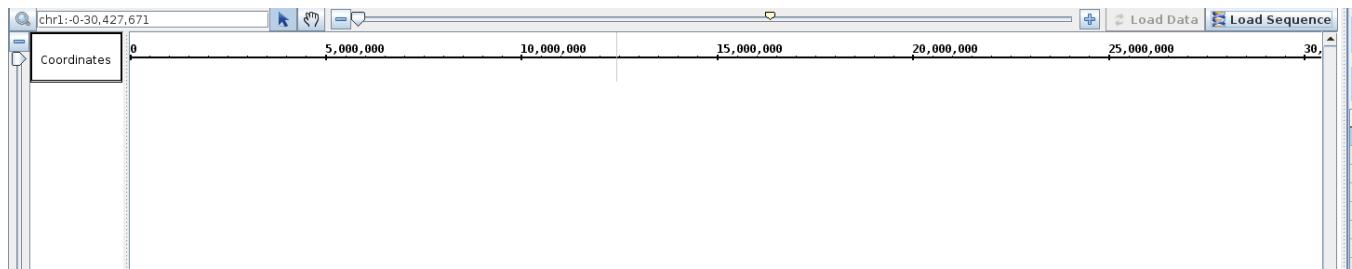
Right click on the white space surrounding the gene models.



- The context menu looks like the one pictured above.

- ☐ mac
- ☐ linux
- ☐ windows

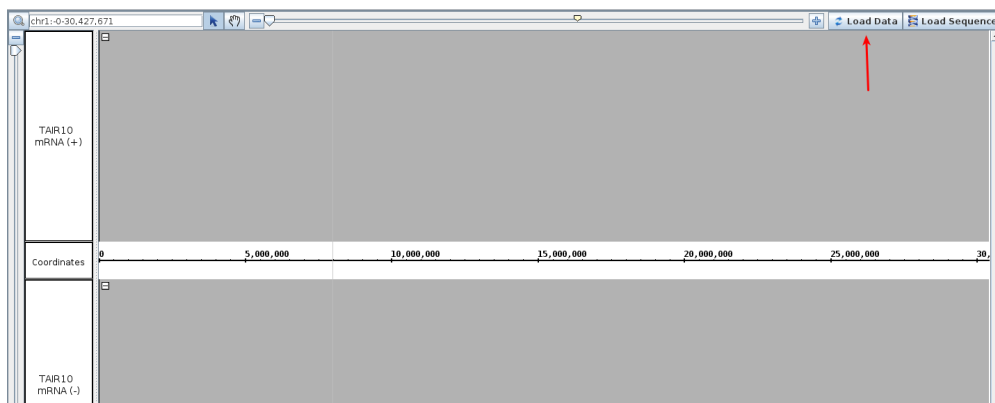
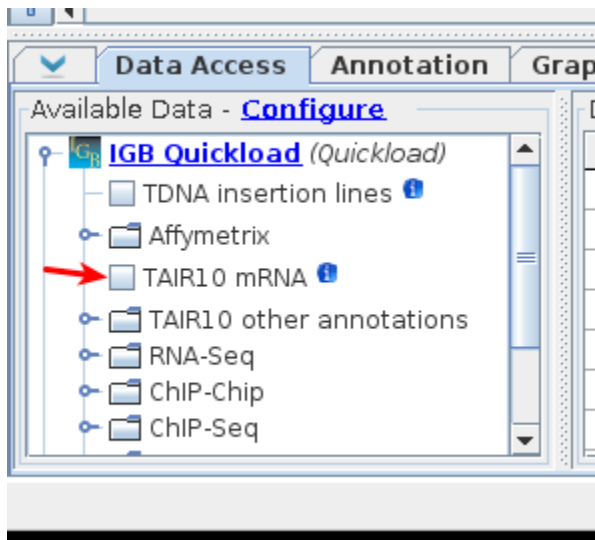
Select **Delete Track**.



- The track disappeared.

- ☐ mac
- ☐ linux
- ☐ windows

Add the **TAIR10 mRNA** track back in via the **Available Data** section.



- The track reappeared with no data loaded.

- ☐ mac
- ☐ linux
- ☐ windows