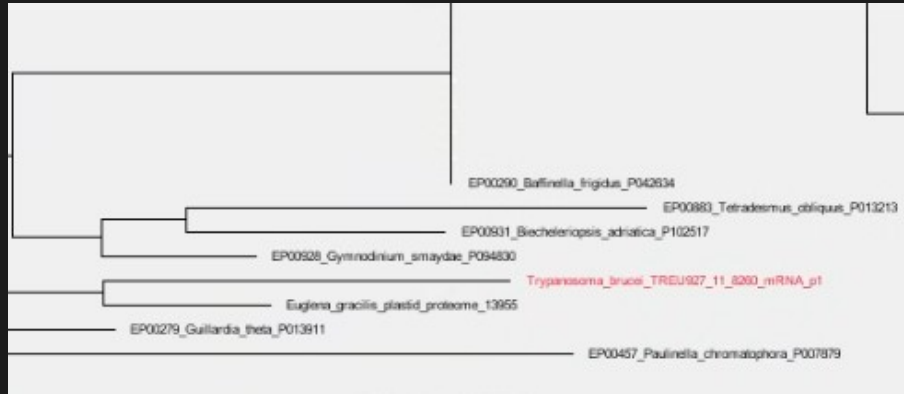


Preparing the input .nexus file

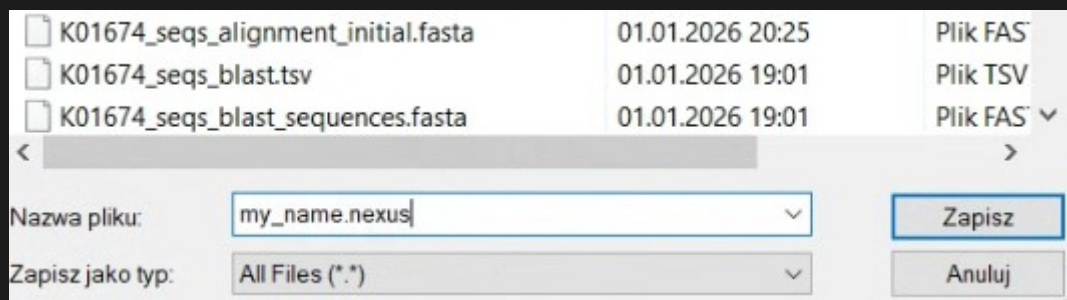
1. Open your tree in FigTree program
2. Mark the taxa to be removed with any colour OTHER than black!



3. Saved file should be in .nexus format, however exporting the tree as nexus file does not include the taxa block with proper colour information. Instead, in left top corner click:

File → Save As...

Choose a name for your file and add .nexus extension to the name!



Save the file. Now it should include the taxa block.

```
awesome-genome-visualisation.txt  chosen probes.txt  K01674.nexus
1  #NEXUS
2  begin:taxa;
3  —>dimensions:ntax=127;
4  —>taxlabels
5  —>Arabidopsis_thaliana_F4HUC4_ATCA5_ARATH
6  —>Arabidopsis_thaliana_F4IHR4_ATCA2_ARATH
7  —>Arabidopsis_thaliana_F4JIK2_ATCA4_ARATH
8  —>Arabidopsis_thaliana_Q04846_ATCA1_ARATH
9  —>Arabidopsis_thaliana_Q8L817_ATCA7_ARATH
10 —>Arabidopsis_thaliana_Q9FM99_ATCA8_ARATH[&!color=#ff0000]
11 —>Arabidopsis_thaliana_Q9FYE3_ATCA3_ARATH
```