

How to make a phylogenetic tree from the ARLEQUIN output.

Output in Arlequin:

NEXUS notation for MST

```
#NEXUS
begin trees; [NEXUS Treefile section generated by Arlequin]
tree W09 MST = [&U] (((((W09-46a:0, W09-50a:1.00000):0,
((((((((((((((((W09-47b:0, W09-56b:1.00000):0, W09-47a:0.00000):0, W09-
56a:0.00000):0, W09-55c:0.00000):0, W09-55b:0.00000):0, W09-
55a:0.00000):0, W09-53b:0.00000):0, W09-53a:0.00000):0, W09-
52b:0.00000):0, W09-52a:0.00000):0, W09-51c:0.00000):0, W09-
51b:0.00000):0, W09-51a:0.00000):0, W09-50c:0.00000):0, W09-
46c:0.00000):0, W09-46b:0.00000):1.00000):0, W09-56c:0.00000):0, W09-
50b:0.00000):0, W09-47c:0.00000);
end;
```

Take the lines marked in yellow, copy them and paste into a spread sheet (e.g. MSWord).

Save as textfile (e.g. test.txt)

Download TreeView from

<http://taxonomy.zoology.gla.ac.uk/rod/treeview.html>

and install on your computer.

Open TreeView, and open your Nexusfile (test.txt) in this program. Then follow the description given in the manual.