

Download the newick file

The screenshot shows a phylogenetic tree viewer interface. The tree is displayed on the left, with nodes labeled with accession numbers and taxonomic information. The 'Tools' menu is open on the right, showing options for downloading the tree in various formats. The 'Download' option is selected, and the 'Newick file' option is highlighted in the submenu.

Tree labels (from top to bottom):

- ◉ OQ963804.1 |Calidris alba|USA|NY|2022-02-21
- ◉ OQ958820.1 ||USA|NY|2022-04-04
- ◉ OQ958812.1 ||USA|NY|2022-04-04
- ◉ OQ958804.1 ||USA|NY|2022-04-04
- ◉ OQ961564.1 |Bubo virginianus|USA|NY|2022-04
- ◉ OQ961068.1 |Corvus ossifragus|USA|NY|2022-04-04
- ◉ OQ959564.1 |Haliaeetus leucocephalus|USA|NY|
- ◉ OQ963300.1 |Phasianus colchicus|USA|NY|2022-
- ◉ OQ961548.1 |Bubo virginianus|USA|NY|2022-04-
- ◉ OQ961556.1 |Bubo virginianus|USA|NY|2022-04-07
- ◉ OQ962772.1 |Phasianinae|USA|NY|2022-03-22

Tools menu options:

- Download (selected)
- Layout
- Sort
- Zoom behavior
- Clear selection
- Clear subtree
- Clear rerooting
- Expand all
- Edit labels

Download options (from top to bottom):

- ASN text file
- ASN binary file
- Newick file (highlighted)
- NEXUS file
- PDF file

Create an FNA file

HyPhy uses a custom data format (Fasta + Newick or FNA) to make things easier

Simply open your Fasta file and paste the Newick string right below

```
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03-23:0.000000):0.000000, OQ962516.1_|USA|NY|2022-03-21:0.000000):0.000000, OQ960876.1_|Accipiter_cooperii|USA|NY|2022-03-
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