

Download the newick file

The screenshot displays a phylogenetic tree viewer interface. The tree is rooted on the left and branches out to the right. The nodes are labeled with accession numbers, species names, and location/date information. The 'Tools' dropdown menu is open, showing options for downloading the tree in various formats. The 'Newick file' option is highlighted.

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
- Layout
- Sort
- Zoom behavior
- Clear selection
- Clear subtree
- Clear rerooting
- Expand all
- Edit labels

Download options (visible on the right):

- ASN text file
- ASN binary file
- Newick file
- 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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