

Import Tab

The screenshot shows the 'Import Tab' interface. At the top, there are tabs for 'Samples', 'Import', and 'Citation'. The 'Import' tab is active. On the left, a 'Files' sidebar lists a directory structure including Applications, Library, Network, System, Users, Volumes, bin, cores, dev, etc, home, net, opt, private, sbin, tmp, usr, and var. A blue arrow points from the 'var' folder to instruction 1. The main panel has a title 'Import' and two warning messages: 'Add your taxonomy file first, then .tsv, .csv or .txt of second dataset (2 files remaining)' and 'Choose a name for your import'. Below these is a text input field 'Name your import' with a '6 Characters left' indicator. A blue arrow points from this field to instruction 3. Under 'Import settings', there are three radio button options: 'Everything' (selected), 'Binned scaffolds', and 'Unbinned scaffolds'. A blue arrow points from the 'Everything' option to instruction 4. At the bottom of the panel is an 'Import' button with a download icon. A blue arrow points from this button to instruction 5. In the top right corner, there is a 'Database' link with a green icon.

1. navigate to the uBin input files

2. select first the taxonomy file and second the SCGS.csv file by double-clicking them

5. Click import and start curating

3. Name the sample

4. Choose the import option

- Everything = both binned and unbinned scaffolds get imported

- Binned scaffolds = only binned scaffolds get imported

Unbinned scaffolds = only unbinned scaffolds get imported