



Identification of Orthologous Genes and Pan-Genomics

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Analysing pan-genomes

Make sure there are not error messages after running each of the following commands!

Run the following command to install the required software for this tutorial:

```
sudo apt install gnuplot5 evince
```

Obtain the data for the Pan-genome exercises from Moodle, look for the file `genomes.pep.tar.gz`, download it to your computer, create a folder called `pangenomics` in your home, and move it there.

Download the software BPGA from <https://sourceforge.net/projects/bpgatool/>, unpack it, change permission so that the files `muscle` and `BPGA-Version-1.3` have execution permission for all users

```
chmod a+x muscle BPGA-Version-1.3
export PATH=~/.Downloads/BPGA-1.3-linux-x86_64-0-0-0/BPGA-
Version-1.3/bin:$PATH
```

Download the software USEARCH from <https://www.drive5.com/usearch/>, make sure to rename the executable to `usearch`, and move it the same folder where you have the sequence files (`~/pangenomics`), make sure the `usearch` executable has execution rights. Copy also the `usearch` program inside the folder: `~/Downloads/BPGA-1.3-linux-x86_64-0-0-0/BPGA-Version-1.3/bin`, move to that folder and run the `BPGA-1.3` executable to generate the database files.

Move yourself to the folder `~/pangenomics`, run the executable `BPGA-1.3` and follow the on-screen instructions, inspect the output files and discuss with your instructor and classmates.

Identification of Orthologous genes

Follow the tutorial: <http://lab.dessimoz.org/tutorials/omapractical/>