

Laboratory 2: installing software

The software installed on your computer by default is capable of doing many wonderful things, but there will come a time when you will need something a bit more specialized. This laboratory exercise is designed to expose you to the most common mechanisms of software installation available in Ubuntu.

Tasks

- (1) Install the font 'Hack'. This font is specifically designed for programmers and helps one easily spot errors. For example, '0' and 'o' have a very different appearance in Hack. It is the font used to typeset commands in these laboratory exercises.
 - (a) Make sure your computer is connected to the internet (e.g. open a web page to check for connectivity or type `ping -c 5 9.9.9.9` in the terminal and confirm that there is 0% packet loss).
 - (b) In the terminal, type `sudo apt update` to update the stored list of packages available for your computer (the package cache). Type your password when prompted. Answer question (1).
 - (c) Type `sudo apt upgrade` in the terminal to upgrade to the latest versions of each package and then type `sudo apt autoremove` to remove any old versions (when installing software always remember these steps as skipping them can cause problems). Type your password, if prompted, and agree to continue when prompted.
 - (d) Type `sudo apt install fonts-hack` to install Hack. Type your password, if prompted, and agree to continue when prompted.
 - (e) Change the default font in the terminal to Hack by selecting 'Preferences...' from the 'File' menu and setting 'Font' to be Hack. Test by opening a new terminal window.
- (2) Install the program 'Visual Studio Code'. It is a very nice text editor designed specifically for programming. There are a number of built-in features and optional extensions to help programmers enter code faster and identify potential errors before the code is run. Visual Studio Code is available, for free, from Microsoft.
 - (a) In the terminal, type `wget -q -O - https://packages.microsoft.com/keys/microsoft.asc | gpg --dearmor > packages.microsoft.gpg` to download and process the Microsoft GNU Privacy Guard code signing key.
 - (b) Type `sudo install -o root -g root -m 644 packages.microsoft.gpg /usr/share/keyrings/` in the terminal to install the key. If prompted, type your password.
 - (c) In the terminal, type `echo 'deb [arch=amd64 signed-by=/usr/share/keyrings/packages.microsoft.gpg] https://packages.microsoft.com/repos/vscode stable main' | sudo tee /etc/apt/sources.list.d/vscode.list > /dev/null` to add the Visual Studio Code repository to your system. If prompted, type your password.
 - (d) Update the package cache by typing `sudo apt update` in the terminal. If prompted, type your password. Answer question (2).
 - (e) Finally, type `sudo apt install code` in the terminal to install the program. Type your password, if prompted, and agree to continue when prompted.

- (f) Open Visual Studio Code by typing `code &` in the terminal (the '&' allows you to continue working in the terminal without opening another window). You may wish to add Visual Studio Code to the quick launcher for easy access.
- (3) Install a Perl script for local use.
- (a) Download the 'B.pl' (Little 2010) using the command line by typing:
`wget http://www.nybg.org/files/scientists/dlittle/B-1.2.tar.gz` in the terminal.
 - (b) Unarchive and decompress the files by typing `tar xvzf B-1.2.tar.gz` in the terminal. Answer question (3).
 - (c) Create your own 'scripts' folder by typing `mkdir scripts` in the terminal.
 - (d) Move the script to the 'scripts' folder by typing `mv B.pl scripts/` in the terminal.
 - (e) Delete the extra files by typing `rm B-1.2.tar.gz ._B.pl README.txt ._README.txt gpl.txt ._example.CAF example.spf ._example.spf` in the terminal.
 - (f) Determine the current contents of your \$PATH variable by typing `echo $PATH` in the terminal.
 - (g) Read over the invisible file '.bashrc' using a simple text editor by typing `nano .bashrc` in the terminal. This script is read by bash every time a new (non-login) shell is started, so one can set 'preferences' by including actions in the file. Exit nano by pressing `ctrl` and `x` at the same time.
 - (h) Add scripts to your PATH by typing `echo 'export PATH=$PATH:$HOME/scripts' >> .bashrc` in the terminal. Answer question (4).
 - (i) Exit the terminal and reopen it (this loads the newly edited version of '.bashrc').
 - (j) Determine the current contents of your \$PATH variable by typing `echo $PATH` in the terminal. It should have changed from step (f). If your scripts directory is not appended to the end, something is wrong.
 - (k) Type `B.pl` in the terminal to see the script instructions.
 - (l) Type `B.pl -i example.CAF` in the terminal to run the script. Answer question (5).
- (4) Install the ABySS assembler (<https://github.com/bcgsc/abyss/>; Jackman et al. 2017).
- (a) To find out if ABySS is available as a package, type `apt-cache search abyss` in the terminal.
 - (b) To determine which version of ABySS is available, type `apt-cache show abyss` in the terminal. Answer question (6).
 - (c) To compile ABySS from source, you must install automake, make, a compiler, a linker, and standard libraries. Type `sudo apt install build-essential automake` in the terminal to install the basics. Type your password, if prompted, and agree to continue when prompted.
 - (d) Download the current release of ABySS by typing
`wget https://github.com/bcgsc/abyss/archive/refs/tags/2.3.5.tar.gz` in the terminal.
 - (e) Expand the archive by typing `tar xvzf 2.3.5.tar.gz` in the terminal.
 - (f) Change to the ABySS directory by typing `cd abyss-2.3.5` in the terminal.
 - (g) Generate a config file for your system by typing `./autogen.sh` in the terminal.

- (h) Create a MakeFile for your system by typing `./configure --with-mpi` in the terminal. Answer question (7).
 - (i) Install the boost C++ library by typing `sudo apt install libboost-all-dev` in the terminal. Type your password, if prompted, and agree to continue when prompted.
 - (j) Rerun the configure script by typing `./configure --with-mpi` in the terminal. Answer question (8).
 - (k) Install the Google sparse hash library by typing `sudo apt install libsparsehash-dev` in the terminal. Type your password, if prompted, and agree to continue when prompted.
 - (l) Rerun the configure script by typing `./configure --with-mpi` in the terminal.
 - (m) Attempt to build ABySS by typing `make -j$(nproc) CFLAGS='-march=native -O2'` in the terminal. Answer question (9).
 - (n) Test the build by typing `make check` in the terminal. Answer question (10).
 - (o) Install ABySS by typing `sudo make install` in the terminal. If prompted, type your password.
- (5) Install a Docker image of Goalign (<https://github.com/evolbioinfo/goalign>; Lemoine & Gas-cuel 2021).
- (a) Install the Docker code signing key by typing `wget -q -O - https://download.docker.com/linux/ubuntu/gpg | gpg --dearmor > docker.gpg` to download and process the Docker GNU Privacy Guard code signing key.
 - (b) Type `sudo install -o root -g root -m 644 docker.gpg /usr/share/keyrings/` in the terminal to install the key. If prompted, type your password.
 - (c) Add the Docker repository by typing `echo "deb [arch=$(dpkg --print-architecture) signed-by=/etc/apt/keyrings/docker.gpg] https://download.docker.com/linux/ubuntu $(lsb_release -cs) stable" | sudo tee /etc/apt/sources.list.d/docker.list > /dev/null` in the terminal.
 - (d) Update the package lists and install Docker by typing `sudo apt update && sudo apt install docker-ce docker-ce-cli containerd.io` in the terminal. Agree to continue when prompted.
 - (e) Add yourself to the 'docker' group (that does not exist yet) by typing `sudo usermod -aG docker ${USER}` in the terminal.
 - (f) Create the 'docker' group by typing `newgrp docker` in the terminal. You can test that the group is working correctly by typing `id -nG` in the terminal—you should see 'docker' as one of the groups listed.
 - (g) Type `docker run --rm pegi3s/goalign goalign help` to install the Goalign Docker image and then display help information. Answer question (11).

Questions (<https://forms.gle/fHdqspviX4wheaCS7>)

- (1) For task (1)(b), which urls are providing package lists?
- (2) For task (2)(d), which urls were added by task (2)(c)?
- (3) For task (3)(b), what does each of the tar options do?

- (4) For task (3)(h):
 - (a) What does each part of the command do?
 - (b) What is '\$HOME'?
 - (c) Is the value of '\$HOME' the same for everyone?
 - (d) What could you use in place of '\$HOME' in the .bashrc file?
- (5) For task (3)(l), what is the barcode sequence quality for the example file?
- (6) For task (4)(b), what version of ABySS is available from the package repository and what is the current release version?
- (7) For task (4)(h):
 - (a) Was the configure script able to create a MakeFile?
 - (b) What dependencies were missing?
 - (c) How else could you have determined what was required?
- (8) For task (4)(j), are all of the dependencies satisfied now? Explain.
- (9) For task (4)(m):
 - (a) What does 'nproc' do?
 - (b) Why is it surrounded by parentheses and preceded by a dollar sign?
 - (c) Would it behave any differently if it were surrounded by grave accents?
- (10) For task (4)(n), how can you tell if the build is good or not?
- (11) For task (5)(g), which version of Galign did you install?

Literature cited

- Jackman, S. D., B. P. Vandervalk, H. Mohamadi, J. Chu, S. Yeo, S. A. Hammond, G. Jahesh, H. Khan, L. Coombe, R. L. Warren & I. Birol.** 2017. ABySS 2.0: resource-efficient assembly of large genomes using a Bloom filter. *Genome Research* 27: 768–777.
- Lemoine, F. & O. Gascuel.** 2021. Gotree/Goalign: toolkit and Go API to facilitate the development of phylogenetic workflows. *NAR Genomics and Bioinformatics* 3.
- Little, D. P.** 2010. A unified index of sequence quality and contig overlap for DNA barcoding. *Bioinformatics* 26: 2780–2781.

Due at the start of class February 7.