# Installing Simulation Software

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- NOTE 1: This document gives instructions on installing software. There is a second document called testInst.Rmd that you can use to test the installed software.
- NOTE 2: Before working through this installation document, you must copy, or "clone" the class GitHub repository to your computer. The simplest way to do this is to create an RStudio project linked to the git repository https://github.com/SFUStatgen/SFUStat840. See the instructions on the class Canvas page on Computing: https://canvas.sfu.ca/courses/47316/pages/computing for instructions on getting started with git and version control in RStudio.

## Overview

We will work with simulation software called msprime and SLiM. msprime simulates ancestries of chromosomes according to a backward-in-time model called the coalescent that we will study in class. msprime is a Python program that we will interface with through R, using the R package reticulate. SLiM simulates genomic data forwards-in-time using a Wright-Fisher reproductive model that we will also study in class. SLiM is a C program that can be run from either the command line or a graphical user interface. Both msprime and SLiM return a data structure called a succinct tree sequence (STS) that represents ancestral relationships amongst chromosomes. We will use the Python package tskit to work with STSs. We will use the Python package pyslim to import SLiM STSs into Python for processing by tskit. The majority of this document contains instructions for installing Python and the packages msprime, tskit and pyslim. At the end is a short section describing how to install the C program SLiM. The C program SLiM needs to be installed separately from the Python packages, with an installer program that is downloaded from the developer's website.

## Python and Python packages

We will be using the R package **reticulate** to interface between R and Python. Specifically, we will use **reticulate** to install the Python packages **msprime**, **tskit** and **pyslim**. The instructions in this section will lead you through the installation of Python, **reticulate**, **msprime**, **tskit** and **pyslim**.

#### Install Python and conda

Conda is a package management system used by the Python community. Python and conda are installed simultaneously by a program called miniconda, which you can find at https://docs.conda.io/en/latest/miniconda.html

Choose the installer for Python version 3.7 that is appropriate for your operating system and follow the instructions to install. [Note for Mac users: You can choose either a "bash installer" or a ".pkg installer". The .pkg installer may be more familiar to most Mac users. Download the .pkg file to your computer, double-click to start the installer and you will be guided through the installation.]

• NB: The installation of conda changes the configuration of the shell used in your terminal windows. You need to open a *new* terminal after the installation to be able to use conda. From a new terminal, type which python and record the path to the python executable that was installed by miniconda (you will need it in the next step). On my Mac the path is /Users/jgraham/miniconda3/bin/python.

#### Install reticulate

This is an R package and can be installed from CRAN in the usual way. After installing, use library() to load the package and the function use\_python() to set the path to your Python executable, as in the following R code chunk:

```
library(reticulate)
use_python("~/miniconda3/bin/python")
```

## Installing msprime and tskit into a Python "environment".

Python environments are like RStudio projects in that they are meant to encapsulate all code and data for a specific project. However, they take the encapsulation idea a step further than R and also include any Python package that you need for the project. In the following, I create an environment called Stat840 with the conda\_create() function and install msprime and tskit into it with conda\_install(). Since msprime depends on tskit (in much the same way as R packages can depend on each other), tskit will automatically install when we install msprime.

```
envname <- "Stat840"
conda_create(envname)
conda_install(envname, "msprime")</pre>
```

## Installing pyslim

Unfortunately pyslim is not available on the conda-forge website that hosts conda-installable packages. To get around this, I have built and uploaded our own conda-installable version of pyslim to the class GitHub site. To simplify the installation of pyslim from with R, I've also written a function conda\_install\_local(). Both the package and R function are in the class GitHub repository (https://github.com/SFUStatgen/SFUStat840). Before proceeding, you must have a copy of the repository on your computer. In what follows, replace /path/to/SFUStat840 with the path to the copy of the class repository on your computer.

```
pkgname <- "pyslim"
dir <- "/path/to/SFUStat840/PythonPackages/"
source(paste0(dir,"conda_install_local.R"))
conda_install_local(envname,pkgname,dir)</pre>
```

#### Using the Stat840 Python environment in future R sessions

To use the Stat840 Python environment and all the package you have installed into it, include the following code chunk at the top of your RMarkdown documents.

```
library(reticulate)
use_condaenv("Stat840")
```

## Installing SLiM

SLiM is a C program available from https://messerlab.org/slim/

Unfortunately the program is only available for Mac OS X and Linux users. In fact, our group has only been successful installing SLiM on Mac OS X. If you are a Mac user, download and run the OS X installer from the above website. This should install both a command-line version of SLiM (e.g., in /usr/local/bin/slim) and a graphical-user-interface (GUI) version of SLiM (e.g., SLiMgui in your Applications folder).

If you do not have access to a Mac, you can get an account on a Mac Pro housed in Brad McNeney's office. You will have to make an appointment with Brad to log in from the console the first time you use it. After the initial login from the console in his office, you will be able to log in remotely. You will not be able to use the SLiM GUI when you log in remotely.