Instructions for downloading and installing FastQC.

1) Download the FastQC software package from their website --
   https://www.bioinformatics.babraham.ac.uk/projects/download.html#fastqc

   ## FastQC A quality control application for high throughput sequence data

   - README
   - Installation and setup instructions
   - Release Notes Please read these before using the program.
   - FastQC v0.11.7 (Win/Linux zip file) ⇐ Click Here
   - FastQC v0.11.7 (Mac DMG image)
   - Source Code for the latest FastQC release

2) Upload the fastqc_(version).zip file to your super computing system by dragging and
   dropping it as done before for BBDuk and Trimmomatic

3) Now follow the commands in the screenshot below to decompress the .zip file. This
   creates a folder called "FastQC" in your directory

```
bash-4.2$ ls  //Show files of software directory
bbmap   BBMap_37.90.tar.gz   fastqc_v0.11.7.zip
bash-4.2$ unzip fastqc_v0.11.7.zip //decompress file
Archive:  fastqc_v0.11.7.zip
   creating: FastQC/
  inflating: FastQC/cisd-jhdf5.jar
   creating: FastQC/Configuration/
```