

Sipros

Proteomics data analysis software

Installation instructions

Install and compile

Window

Double click

```
Sipros-ProRata_V3.0_Setup.exe
```

Linux

```
tar -zxzf Sipros-ProRata_V3.0_Linux.tar.gz  
  
cd Sipros-ProRata_V3.0_Linux/src
```

if Sipros will be used on a desktop, you just need Openmp feature. Please execute

```
make
```

You will get binary file Sipros_openmp

If you need the MPI/OpenMP feature of Sipros, please execute

```
make -f MPIMakefile
```

You will get binary file Sipros_openmp_mpi

In this wiki, we will use Sipros to refer to the binary file. Please note that the current makefile uses g++ by default. If you want to switch to other compilers or add flags, please update the corresponding makefile.
