

## ➤ INSTALLING

- 1) **Download** the latest zip available for your operating system
- 2) **Copy** all the files in the zip to any location
- 3) Install the free software **Mono** .NET framework with GTK# support. You can download the framework for GNU/Linux, Mac OS X or Windows from here:

<http://www.go-mono.com/mono-downloads/download.html>

If you use Ubuntu you can simply install the package "mono-runtime" instead.

## ➤ EXECUTING

- 1) Specific launchers are provided for the different platforms:
  - a) **GNU/Linux**: just click on **PAnalyzer.exe** or execute **PAnalyzer.sh** from a console, which will automatically install the mono runtime if not detected
  - b) **Mac OS X**: just click on the **PAnalyzer** application
  - c) **Windows**: just click on **PAnalyzer.bat**
- 2) Click **Open** and select one or more (multirun sample) protein files from the open dialog. You can download some samples from here:  
<https://code.google.com/p/ehu-bio/downloads/list>
- 3) Click **Execute** and select between the preferences (if applicable).
- 4) **Browse** the results in the Summary, Peptides and Proteins tabs.
- 5) **Save** the protein detection list to a file. A \*.html and a \*.csv output files are always generated and a \*.mzid output file is only generated if the data came from another \*.mzid input file.

## ➤ DOCUMENTATION

See file **SEPROT2011-EHU.pdf**.