

# Introduction to Data Cleaning with OpenRefine

Alexander Botzki and Herwig Van Marck  
2015-11



**BITS** VIB  
BIOINFORMATICS TRAINING  
AND SERVICES

# What is OpenRefine?

A free, open source, power tool for working with messy data

Formerly known as Google Refine

Being rebranded as **OpenRefine** (<http://openrefine.org>)

VIB-Bits plugins to facilitate usage  
(<http://www.bits.vib.be/software>)

# How does OpenRefine compare with other tools?

- Compared to **spreadsheets**
  - Basic unit of interaction is column (versus cells)
  - Pro: easier to import data, explore, manipulate and export again
- Compared to **scripting**
  - Pro: you see your data, while it is being transformed
  - Con: for medium size data sets
- Compared to **databases**
  - Pro: you see your data, while it is being queried
  - Con: for simple data structures

# What are typical use cases for OpenRefine?

- Explore unknown/new data files
- Manipulate/clean data to prepare for other tools
  - E.g. GraphPad Prism
- Get data from web services
- Use as a workflow tool
- Create dashboards

# How to install OpenRefine

Installation instructions can be found on

<https://github.com/OpenRefine/OpenRefine/wiki/Installation-Instructions>

(version 2.6 beta 1 is preferable)

Platforms:

- Windows
- Mac OSX
- Linux

# How to run OpenRefine

- Windows

Run the .exe file in the installation folder

- Mac OSX

Double click the OpenRefine app in the Applications folder

- Linux

Start ./refine in the installation folder

# How to shut down OpenRefine

- Windows

Press **Ctrl-C** in the OpenRefine Command windows

- Mac OSX

Invoke the **Quit** command on the OpenRefine app

- Linux

Press **Ctrl-C** in the shell

# User Interface overview

## Home screen

Create Project

Import Project

Open Project

Delete projects

Rename projects



# User Interface overview

## Project

Data table

Column menu

Side bar

Facet / Filter

Undo / Redo

Home button

# Input data demo

## Configuration screen

Preview area

Options area

Parsing format

Parsing options

Ignore first lines

Parse as column header

# Exercise 1: importing files

a) Import the file `qPcr results.txt`

# Exercise 1: importing files

a) Import the file `qPcr results.txt`

Hints:

- Use `ignore first ... line(s)` at beginning of file
- Check if correct separator is used

# Explore data using facets demo

## Text facet

- Sorting by name/count

- Facet counts in tab separated format

## Querying

- Select choice

- Inverted query

- Resetting facet

- Facet counts reflect query

# Explore data using facets demo

## Numeric facet

Query using slider

## Timeline facet

Histogram

Query using slider

Query is reflected in histogram

# Explore data using facets demo

## Custom text facet

Create selectable items using expression

e.g. `value.toLowerCase().contains("good")`

Query by selecting item

# Explore data using facets demo

Data and history is always saved, but **query is not!**

Current query can be saved

Using the [Permalink](#) link and bookmarking the page

Facet box size can not be saved



## Exercise 2: using facets

Use the file `syst-nocallsCG69.bed` to determine

- a) the number of no-call regions that are larger than 1040 bases long in chromosome 21
- b) the length of the longest region in chromosome 1

## Exercise 2: using facets

Use the file `syst-nocallsCG69.bed` to determine

- a) the number of no-call regions that are larger than 1040 bases long in chromosome 21
- b) the length of the longest region in chromosome 1

Hints:

- The 5th column contains the length of a region
- Use a custom facet with '`>`' in the expression
- Use sort to determine the longest region

# Cleaning data demo

## Cluster facet choices

Try different keying functions

Merge clusters of similar values

[Browse this cluster](#) link opens cluster in a new window

## Exercise 3a: cleaning data

Clean the file `qPcr results.txt` you loaded earlier.

## Exercise 3a: cleaning data

Clean the file `qPcr results.txt` you loaded earlier.

Hints:

- Check columns using facets
- Use the clustering tool in facets

# Manipulate data demo

Use 'Line based text files' for complex data files

Edit column>Split into several columns...

- Regular expressions for separator

E.g. '+'

- Split into ... columns at most

Edit cells>Blank down on index column (1st)

Use custom text facet to check if separator is used

E.g. `value.contains("|")`

# Expression window details

Preview tab

Help tab

Expressions can also use chained form

e.g. `value.contains("|")` instead of `contains(value,"|")`

History tab

Reuse recently used expressions

Starred tab

Expressions that were starred in the history tab

## Exercise 3b: cleaning data

Clean the numerical column in file `qPcr results.txt` you loaded earlier.



## Exercise 3b: cleaning data

Clean the numerical column in file `qPcr results.txt` you loaded earlier.

Hints:

- Use numeric facet to explore column
- Use the `replaceChars(...)` command and the `toNumber()` command in transform column

# Extended course material

For extended course material

Go to [www.bits.vib.be](http://www.bits.vib.be)

Click on [Training](#)

Click on [Previous trainings](#)

Click on [Data Manipulation with OpenRefine](#)

and also

Click on [Custom trainings](#)

See [OpenRefine](#)