# Workflow disease profiles

|  |  |  |
| --- | --- | --- |
| Step | Review to add to public repository | Notes |
| ExperimentalInfections (\*) |  |  |
| 1. Download from Distiller as CSV   DATA\_EXTRACT\_STORYMAPS  REF\_extract |  |  |
| 1. Place CSVs (2) on “data/DistillerData\_raw” |  |  |
| 1. Run   \*\_data\_cleaning.R  depends on:  \*\_columnNames.csv  \*\_definitions.R  Result: saved cleaned files in “data/FilesDownload” |  | If we organize the headers properly in Distiller maybe we won’t need the “column names” |
| 1. 1impact\_MA.r |  |  |
| 1. 1impact.Rmd   Depends on:  Functions.R |  |  |
|  |  |  |
|  |  |  |
|  |  |  |

**Test agents**

AHFV

ASFV

BTV

GETV

Linfantum