# Workflow disease profiles

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| **Step** | **Notes** |
| RUN-list-header.r |  |
| RUN-list.r |  |
| 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  Result: saved cleaned files in “data/FilesDownload” |  |
| 1. 1impact\_MA.r   Depends on:  Functions.R  Metanalysis\_functions.R |  |
| 1. 1impact.Rmd   Depends on:  Functions.R |  |
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