

# Epiverse pipeline applications: challenges and lessons learned

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#### **Epiverse pipelines**

- Aim: to contain the steps to conduct outbreak analytic tasks from start to finish in a reliable manner
- Divided in early, middle, and late tasks
- Collected as R markdown templates in the <u>episoap</u> package





Epiverse pipelines roadmap, Andree Valle

#### **Pipeline applications/ Case studies**

- Aim: to showcase the functionality of Epiverse's pipelines to potential users
- R packages applied to real outbreak data
- Tailored to specific end-user needs and interests
  - MVD-Severity and underreporting
  - Cholera-Transmissibility



- 1. Interoperability
  - a. Incompatible packages

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Linelist objects not usable with functions such as `mutate()` or `filter()`

- 1. Interoperability
  - a. Incompatible packages
  - b. Output vs input format

```
\underline{incidence2} \leftrightarrow \underline{cfr}
```

```
# Convert to incidence
MVD_cases_deaths <-
incidence2::incidence(MVD_linelist_cut,c("Onset_week","Death_week")) |>
complete_dates()
# Pivot table
MVD_cases_deaths <- pivot_wider(MVD_cases_deaths, names_from =
count_variable, values_from = count)
# Change column names for function
names(MVD_cases_deaths) <- c("date_index","deaths","cases")
# Reorder for function
MVD_cases_deaths <- MVD_cases_deaths[,c("date_index","cases","deaths")]
# Convert to data frame
MVD_cases_deaths <- as.data.frame(MVD_cases_deaths)</pre>
```

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  - a. Documentation

#### ## Data required

The data required to estimate how the severity of a disease changes over time using the \_cfr\_ package includes:

\* A time-series of cases, hospitalisations or some other proxy for infections over time;

\* A time-series of deaths;

\* A delay distribution, describing the probability an individual will die \$t\$ days after they were initially exposed. Such distributions come from the literature, where studies have typically fit distributions to data describing the process.

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  - b. Non-informative error messages

```
Previous error message of `estimate_static()` function from `cfr`:
Error in data.frame(severity_me = severity_me, severity_lo =
severity_lims[[1]], : arguments imply differing number of rows: 0, 1
```



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Current error message of `estimate\_static()` function from `cfr`: Error in estimate\_static(daily\_cases\_deaths\_missing\_data, epi\_dist = onset\_to\_death\_ebola, : Input data must have sequential dates with none missing or duplicated

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#### 3. Sometimes don't translate to certain real-life scenarios

`<u>cfr</u>` currently uses *days* as input for dates, whereas some data sources provide weeks of onset/death



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- 3. Sometimes don't translate to certain real-life scenarios
- 4. Packages under development

Unstable functions, features that are removed, names changed, etc.  $\rightarrow$  difficult for users to keep track

```
Error in format_output(estimate_static(df_ebola_subset,
correct_for_delays = TRUE, :
    could not find function "format_output"
```

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```
Error in delay_opts(list(mean = onset_to_death_logmean, mean_sd = 0.1, :
    Delay distributions must be of given either using a call to `dist_spec` or one of the
`get_...` functions such as `get_incubation_period`. This behaviour has changed from
previous versions of `EpiNow2` and any code using it may need to be updated. For
examples and more information, see the relevant documentation pages using
`?delay_opts`.
```

#### **Lessons learned - Pipeline applications**

- Testing data pipelines is as relevant as testing the functionality of packages individually
- Ideally there would be at least one person within the team to carry out the testing
- Challenges are also an opportunity to optimise the pipelines
  - E.g.: `cleanepi` to remove duplicated data across `linelist` tags

cleanepi(df, remove.duplicates=T, duplicates.from="tags")

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- Challenges are also an opportunity to optimise the pipelines
- ... and an opportunity for RSEs and RFs for collaborative development
- In the future, this process must be carried out also by users outside the team



## Thank you for your attention!

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