

Scientific Reproducibility in Biology Research

Day 4

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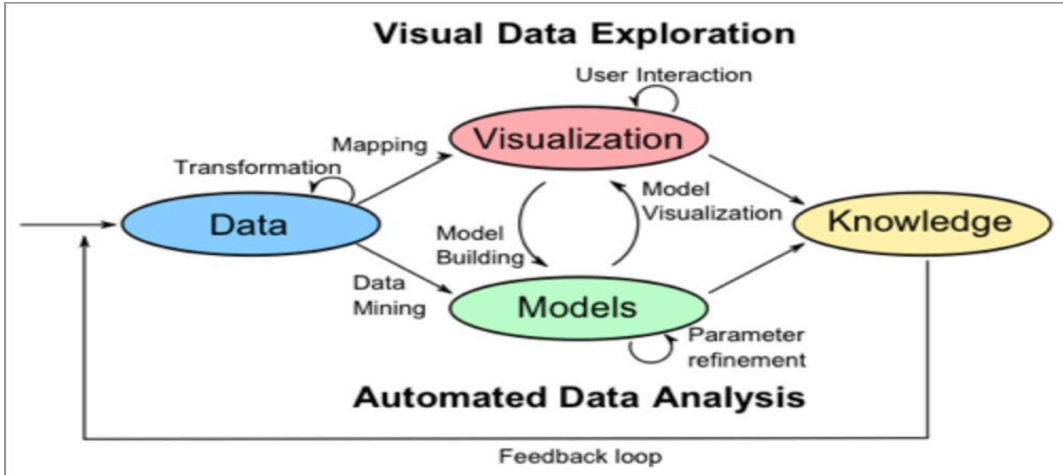
Australian National University

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Review of workshop highlights

- ▶ Making scientific reproducibility a priority
 - ▶ Clear directory/file structures
 - ▶ FAIR data principles
 - ▶ Good coding practices
 - ▶ Automated processes

Review of workshop highlights



Review of workshop highlights

- ▶ create Rmarkdown documents
- ▶ create tidy data for easy visualisation
- ▶ table summaries using gtsummary package
- ▶ data wrangling using tidyverse packages
- ▶ data visualisation using ggplot()

Fun data wrangling tools!

- ▶ `filter()`
- ▶ `select()`
- ▶ `arrange()`
- ▶ `mutate()`
- ▶ `group_by() %>% summarise()`
- ▶ `pivot_longer()` & `pivot_wider()`
- ▶ `fct_recode()`

Fun ggplot() commands!

- ▶ `facet_wrap()`
- ▶ `geom_point()`, `geom_histogram()`, `geom_bar()`, `geom_beeswarm`,
`geom_line()`, `geom_etc()`
- ▶ `geom_smooth()`
- ▶ `scale_y_log10()`
- ▶ `labs()`
- ▶ `scale_colour_manual()`
- ▶ `theme()`

What we'll cover today

- ▶ Review Day 3 assignment
- ▶ Jam session: applying your new skills to real data