

1. Package issue tracking

1.1 Package issues

- `haven::read_xpt()` Some attributes are dropped after using `haven::write_xpt()` `haven::read_xpt()`, e.g., type, length. to do: further check
- `xportr::xportr_length()` `NA_character_` is considered as length 2 - issue resolved on Dec 14, 2022 to do: to be incorporated in pilot3

1.2 Potential improvement

- `metatools::build_from_derived()` <https://github.com/pharmaverse/metatools/issues/46> Not urgent feature but nice to have - especially when `define.xml` is not in good quality

2. Knowledge sharing

Summary of similarities and differences between packages, to help user identify the best tool that suits the need. Maybe this could go to Bayer SAS2R catalog in the future,

`metatools` vs `xportr`

- `metatools::set_variable_labels()` vs `xportr::xportr_label()`

`xportr` vs `haven`

- `xportr::xportr_write()` vs `haven::write_xpt()`

`diffdf` vs `arsenal`

- `diffdf::diffdf()` vs `arsenal::comparedf()`