

Comparison of TAF and icesTAF

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Preamble

```
library(diff)
tools.prod <- "~/git/ices/tools-prod"
```

1 R directory

```
compare.rdir(tools.prod)
```

```
## $TAF
## [1] "TAF-internal.R" "TAF-package.R"
##
## $icesTAF
## [1] "draft.data.script.R" "icesTAF-internal.R" "icesTAF-package.R"
## [4] "taf.roxygenise.R"
```

Check out these files:

1. draft.data.script.R
2. get.stockassessment.results.R
3. get.stockassessments.R
4. taf.roxygenise.R
5. upload.fay.R

2 Namespace

```
compare.namespace(tools.prod)
```

```
## $TAF
## character(0)
##
## $icesTAF
## [1] "S3method(roxy_tag_parse,roxy_tag_tafAccess)"
## [2] "S3method(roxy_tag_parse,roxy_tag_tafKey)"
## [3] "S3method(roxy_tag_parse,roxy_tag_tafOriginator)"
## [4] "S3method(roxy_tag_parse,roxy_tag_tafPeriod)"
## [5] "S3method(roxy_tag_parse,roxy_tag_tafSource)"
## [6] "S3method(roxy_tag_parse,roxy_tag_tafYear)"
## [7] "export(draft.data.script)"
## [8] "export(taf.roxygenise)"
## [9] "importFrom(glue,glue)"
## [10] "importFrom(jsonlite,parse_json)"
## [11] "importFrom(purrr,flatten)"
## [12] "importFrom(roxygen2,roxy_tag_parse)"
```

Check out these entries:

1. `export(draft.data.script)`
2. `export(get.stockassessment.results)`
3. `export(get.stockassessments)`
4. `export(taf.roxygenise)`
 - `S3method(roxy_tag_parse,roxy_tag_tafAccess)`
 - `S3method(roxy_tag_parse,roxy_tag_tafKey)`
 - `S3method(roxy_tag_parse,roxy_tag_tafOriginator)`
 - `S3method(roxy_tag_parse,roxy_tag_tafPeriod)`
 - `S3method(roxy_tag_parse,roxy_tag_tafSource)`
 - `S3method(roxy_tag_parse,roxy_tag_tafYear)`
5. `export(upload.fay)`

3 Environment

```
compare.environment()
```

```
## $TAF
## character(0)
##
## $icesTAF
## [1] "draft.data.script"
## [2] "list2bibtex"
## [3] "loadpkg"
## [4] "roclet_output.roclet_taf"
## [5] "roclet_process.roclet_taf"
## [6] "roxy_tag_parse.roxy_tag_tafAccess"
## [7] "roxy_tag_parse.roxy_tag_tafKey"
## [8] "roxy_tag_parse.roxy_tag_tafOriginator"
## [9] "roxy_tag_parse.roxy_tag_tafPeriod"
## [10] "roxy_tag_parse.roxy_tag_tafSource"
## [11] "roxy_tag_parse.roxy_tag_tafYear"
## [12] "taf_roclet"
## [13] "taf.roxygenise"
```

Check out these objects:

1. draft.data.script
2. taf.roxygenise
 - list2bibtex
 - loadpkg
 - roclet_output.roclet_taf
 - roclet_process.roclet_taf
 - roxy_tag_parse.roxy_tag_tafAccess
 - roxy_tag_parse.roxy_tag_tafKey
 - roxy_tag_parse.roxy_tag_tafOriginator
 - roxy_tag_parse.roxy_tag_tafPeriod
 - roxy_tag_parse.roxy_tag_tafSource
 - roxy_tag_parse.roxy_tag_tafYear
 - taf_roclet