Comparison of TAF and icesTAF

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

1 Files

```r
compare.files(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)`