sa4ss

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Getting Started

Want to create an assessment document using sa4ss?

Load up the package available on github:

```
remotes::install_github("nwfsc-assess/sa4ss")
library(sa4ss)
```

Now that the package is installed in R, let's create a document.

Step 1

Set the working directory. The document template will be created in the defined location.

```
setwd("C:/Assessments/2021/copper_rockfish_2021/write_up/wa")
```

Step 2

Call the draft function to create a general document with species, authors, and title created:

The above example will automatically populates a document for copper rockfish with the above authors and their associated affiliation. **sa4ss* has a list of common authors and their affiliations already set-up. In order for authors to be found in the available list and their affiliation the names need to be entered exactly as expected here: https://github.com/nwfsc-assess/sa4ss/blob/master/data-raw/authors.csv.

Step 3

Create an Rdata object based upon the base model. The base model can be located anywhere on your machine.

```
sa4ss::read_model(mod_loc = "C:/Assessments/2021/copper_rockfish_2021/models/base")
```

This will create an Rdata object called 00mod.Rdata in your working directory.

Step 4

Render the pdf document based on the draft template created:

```
bookdown::render_book("00a.Rmd", clean = FALSE, output_dir = getwd())
```

This will create build the pdf document which will be save directly in the working directory. However, one may prefer to have the pdf saved in a specific folder. One default approach is the call the render_book function without specifying the output_dir which will put the pdf document in a separate folder in the working directory.

```
bookdown::render_book("00a.Rmd", clean = FALSE)
```

Editing the Document

A number of documents in the specified directory should not be available. These documents correspond to different sections required by the Groundfish Terms of Reference.

When editing make sure to render the document on a regular basis by using either of the above calls to:

```
bookdown::render_book("00a.Rmd", clean = FALSE, output_dir = getwd())
```

This will allow users to identify errors quickly. If the document rendered correctly the user should see that "Output created: main.pdf" followed by miscallaneous other warnings in the R terminal.

Returning to the Document

If you are returning to edit your document at a later point, you will not need to redo steps 1-3 above. To begin working on your document again you should:

- 1. Open an new R terminal.
- 2. Load of the package library(sa4ss).
- 3. Set the working directory to where your document is located.
- 4. Open Rmd files and start editing your document. To render the book at any point in this process you just need to run the following call:

```
bookdown::render_book("00a.Rmd", clean = FALSE)
```

Error in Rendering

If you see an error and the document fails to render you will want to:

- 1. Open the _main.tex file, read the error message in the R terminal, and navigate to that line in the main.tex file.
- 2. Correct the error in the Rmd files.
- 3. Delete the main.Rmd file in the folder.
- 4. Rerun the render line of code:

```
bookdown::render_book("00a.Rmd", clean = FALSE, output_dir = getwd())
```

Updating the Base Model

If the base model changes, the user will need to rerun the **read_model** function to load in the output from the new model run and locate and/or create figures and tables

```
sa4ss::read_model(mod_loc = "C:/Assessments/2021/copper_rockfish_2021/models/new_base")
```

This will create a new model output file which will replace the old file and be used to update the document to the new model the next time the document is rendered.

Tips & Tricks

Share Text Across Documents

If you are creating an area-specific assessment document for species you may want to use the same text across multiple documents. This can be easily done by creating separate folder for sections of text that will be shared across documents. Text should be included in a file with an Rmd extension and can be called using a code chunk with the **knit_child** command in the **knitr** package:

```
other_loc = file.path('folder', 'child_example.Rmd')
res <- knitr::knit_child(other_loc, quiet = TRUE)
cat(res, sep = '\n')
#>
#>
#> Here is the text in the child document.
```

Add Figures from r4ss

There are a number of ways to add figures to the document, either in the executive summary or figures sections. The first approach is to directly specify where the document is located, add a figure caption and alternative text caption (for accessibility), and specific a figure label which allows the user to reference the figure in the document. In the below example the figure (data-plot.png) is located in the same directory as the Rmd files being rendered.

```
# This is the text that is placed in the document:
![Summary of data sources used in the base model.\label{fig:data-plot}](data-plot.png)
{width=100% height=100% alt="Summary of data sources used in the base model"}
```

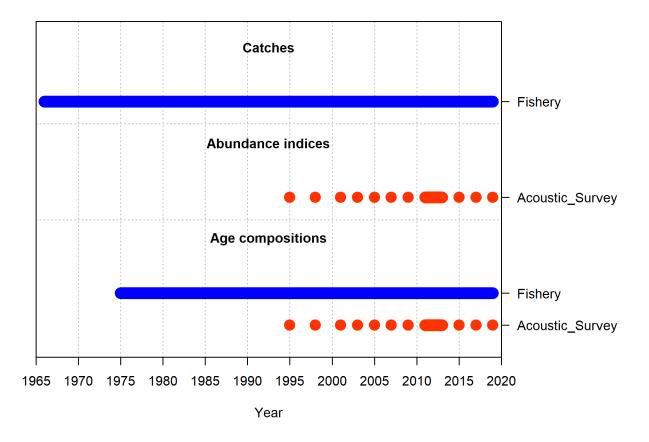


Figure 1: Summary of data sources used in the base model.

Alternatively, the figure you want to include may be located in a folder outside the folder that you are rendering the document within. To do this easily a function has been added to the **sa4ss** package that allows users to do this.

```
# The add_figure function has the user specific the figure and its location to be used.
add_figure(
filein = file.path(mod_loc, "plots", "data_plot.png"),
caption = "Summary of data sources used in the base model",
label = 'data-plot')
```

The above function, **add_figure**, contained in the **sa4ss** package will set the alternative text equal to the caption unless specifically specified by the user.

Both examples above allow the user to reference the figure in the text using {r, eval = FALSE} \ref{fig:data-plot} which will look like this: Figure 1.