# How to use the **ss3sim** package to run simulations in **SS3**

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# 1 Installing the ss3sim R package

The package can be installed and loaded with:

```
# install.packages(c("r4ss", "plyr", "MCMCpack")) # dependencies, if needed
# install.packages("devtools")
# devtools::install_github("ss3sim", username="seananderson")
library(ss3sim)
```

While the package is under active development, it's a good idea to install a new version every day or so using the install\_github command:

```
devtools::install_github("ss3sim", username="seananderson")
```

You can read the help files and access this vignette again with:

```
help(package = "ss3sim")
vignette("ss3sim")
```

## 2 Shortcut to running FISH 600 models

Assuming you've set up SS3 in your path and are familiar with how the package works (as explained below), you can run FISH 600 models as follows. Note that this example runs 1 iteration and 2 bias adjustment runs.

```
run_fish600(iterations = 1, scenarios = "D0-E0-F0-G0-M0-R0-S0-cod",
bias_adjust = TRUE, bias_nsim = 2)
```

Typically we will run 100 iterations and 10 bias adjustment runs.

```
run_fish600(iterations = 1:100, scenarios = "D0-E0-F0-G0-M0-R0-S0-cod",
bias_adjust = TRUE, bias_nsim = 10)
```

You may want to run the iterations in chunks. In that case you'll want to tell run\_ss3sim that you've already run the bias correction by setting the bias\_already\_run argument to TRUE and the bias\_adjust argument to FALSE. When you run the bias correction runs you need to run at least one regular run. Therefore, you might do something like this:

```
run_fish600(iterations = 1, scenarios = "DO-EO-FO-GO-MO-RO-SO-cod",
bias_adjust = TRUE, bias_nsim = 10)
```

```
run_fish600(iterations = 2:100, scenarios = "D0-E0-F0-G0-M0-R0-S0-cod",
bias_adjust = FALSE, bias_already_run = TRUE)
```

Through this strategy you could run the second set (after running bias adjustment) using parallel processing or on separate machines. (Separate machines would require copying the existing folder structure to the second machine first.)

You can run multiple scenarios in sequence by expanding the vector of values passed to scenarios. For example,

```
run_fish600(iterations = 1:100,
scenarios = c("D0-E0-F0-G0-M0-R0-S0-cod", "D0-E0-F0-G0-M1-R0-S0-cod"),
bias_adjust = TRUE, bias_nsim = 10)
```

## 3 Putting SS3 in your path

SS3 must be in your path for the ss3sim package to work. Your "path" is a list of folders that your operating system looks in whenever you type the name of a program on the command line. Having a binary in your path means that your operating system knows where to look for the file regardless of what folder you're working in.

### 3.1 For Unix (Linux and OS X)

To check if SS3 is in your path: open a Terminal window and type which SS3 and hit enter. If you get nothing returned then SS is not in your path. The easiest way to fix this is to move the SS3 binary to a folder that's already in your path. To find existing path folders type echo \$PATH in the terminal and hit enter. Now move the SS3 binary to one of these folders. For example, in a Terminal window type:

#### sudo cp ~/Downloads/SS3 /usr/bin/

You will need to use **sudo** and enter your password after to have permission to move a file to a folder like /usr/bin/.

If you've previously modified your path to add a non-standard location for the SS3 binary, you may need to also tell R about the new path. The path that R sees may not include additional paths that you've added through a configuration file like .bash\_profile. You can add to the path that R sees by including a line like this in your .Rprofile file. (This is an invisible file in your home directory.)

Sys.setenv(PATH=paste(Sys.getenv("PATH"),"/my/non-standard/folder",sep=":"))

#### 3.2 For Windows

To check if SS is in your path: open a DOS prompt and type ss3 -? and hit enter. If you get a line like "ss3 is not recognized . . . " then SS is not in your path. To add it to your path:

- 1. Find the latest version of the ss3.exe binary on your computer
- 2. Record the folder location. E.g. C:/SS3.24o/
- 3. Click on the start menu and type "environment"
- 4. Choose "Edit environment variables for your account" under Control Panel
- 5. Click on PATH if it exists, create it if doesn't exist
- 6. Choose PATH and click edit
- 7. In the "Edit User Variable" window add to the **end** of the "Variable value" section a semicolon and the SS3 folder location you recorded earlier. E.g.; C:/SS3.24o/
- 8. Restart your computer
- 9. Go back to the DOS prompt and try typing ss3 -? and hitting return again.

## 4 Setting up the file structure

This package is set up assuming that you have an established base case operating model and estimation model to work with. Each operating model and estimation model should be in their own folder. The operating model folder should have the files:

```
yourmodel.ctl
yourmodel.dat
ss3.par
starter.ss
forecast.ss
```

The estimation model folder should have:

```
yourmodel.ctl
yourmodel.dat # optional; not used
starter.ss
forecast.ss
```

In both cases, nothing more and nothing less. The names of the .ctl and .dat files are not important. The package functions will rename them after they are copied to appropriate folders. These files should be derived from the .ss\_new files but named as listed above. It's important to use these .ss\_new files so they have consistent formatting. Many of the functions in this package depend on that formatting.

For the purposes of the FISH 600 project, we have unique case identifiers which combine to create unique scenarios. The types of cases are: natural mortality (M), fishing mortality (F), data quality (D), and retrospective (R). And the species are cod (cod), sardine-like (sar), and flatfish (fla). It's important to use these three letter abbreviations for the species since the functions assume that the last three letters represent a species (or some other identifier for a different project).

The different version of each case are identified with numbers. So, for example, the base case scenario for cod is identified as M1-F1-D1-R1-cod. We will have a spreadsheet that describes each of these in plain language.

The function copy\_ss3models creates a folder structure and copies over the operating and estimation models. The folder structure looks like:

```
M1-F1-D1-R1-cod/1/om
M1-F1-D1-R1-cod/1/em
```

```
M1-F1-D1-R1-cod/2/om
M1-F1-D1-R1-cod/2/em
```

If you are using bias adjustment (bias\_adjust = TRUE) then there will be some additional folders. In that case the folders will look like:

```
M1-F1-D1-R1-cod/bias/1/om
M1-F1-D1-R1-cod/bias/1/em
M1-F1-D1-R1-cod/bias/2/om
M1-F1-D1-R1-cod/bias/2/em
...
M1-F1-D1-R1-cod/1/om
M1-F1-D1-R1-cod/1/em
M1-F1-D1-R1-cod/2/om
M1-F1-D1-R1-cod/2/em
```

Note that the operating and estimating model folders have been renamed om and em within each iteration, the operating and estimation models have been checked to make sure they contain the minimal files (as listed above), the filenames have been made all lowercase, the data file has been renamed data.dat, the control files have been renamed om.ctl or em.ctl, and the starter and control files have been adjusted to reflect these new file names.

The functions in this package assume you've set your working directory in R to be the base folder where you will store the scenario folders.

# 5 Creating the input configuration files

You will need to have a folder containing "case" argument definitions. These plain text files are read by get\_caseval and turned into argument lists that are passed to run\_ss3sim. You can create template input files by running create\_argfiles. It reads the various functions and parses the arguments and default values into plain text files. The default settings create these files:

```
1. MO-spp.txt
```

```
3. index0-spp.txt
```

- 4. agecomp0-spp.txt
- 5. lcomp0-spp.txt
- 6. RO-spp.txt
- 7. S0-spp.txt
- 8. GO-spp.txt
- 9. E0-spp.txt

Look in your working directory for the template files. Change the case ID number (defaults to 0) and the species identifier to a three letter identifier. For the FISH 600 project use one of cod, sar, or fla for cod, sardine, or flatfish. An example filename would be M1-sar.txt or lcomp2-fla.txt. The case D1 corresponds to the files index1-spp.txt, agecomp1-spp.txt, and lcomp0-spp.txt. The other case types have single argument files.

The first column in the text files denotes the argument to be passed to a function. The second argument denotes the value to be passed. You can use any simple R syntax. For example: c(1, 2, 4), or seq(1, 100) or 1:100 or matrix(). Character objects don't need to be quoted, but can be if you'd like. However, be careful not to use the delimiter (set up as a semicolon) anywhere else in the file besides to denote columns. You can add comments after any # symbols. Internally, the functions evaluate in R any entries that have no character values (e.g. 1:100) or have an alphanumeric character followed by a (. Anything that is character only or has character mixed with numeric but doesn't have the regular expression "[A-Za-z0-9](" gets turned into a character argument.

Putting that all together, here's what an example F1-cod.txt file might look like:

years; 1:100 years\_alter; NA

fvals; NA

# 6 Running the models

The run\_ss3sim function is a wrapper function. It adjusts the natural mortality, adjusts the fishing mortality, adds recruitment deviations, calls run\_ss3model to run the operating model, samples various survey estimates from the operating model, changes the age composition data as specified, changes the length composition data as specified, copies and renames files as necessary, and calls run\_ss3model again to run the estimation model.

The run\_fish600 function is a higher-level wrapper that deals with parsing the case input files and then passes these arguments on to run\_ss3sim. This is what we will use for the FISH 600 project.

See the PDF version of this vignette for a flow chart illustrating how run\_fish600 and run\_ss3sim work:

vignette("ss3sim")

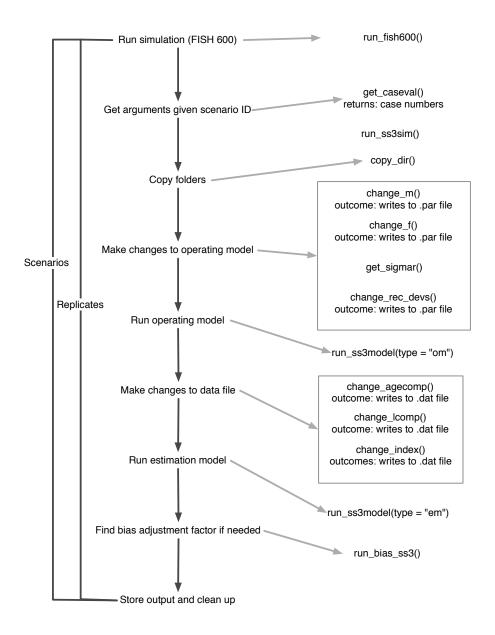


Figure 1: Simulation steps. Higher-level function calls are shown on the right.