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0.0.1 genNavbar

genNavbar generates a navigation bar for a web page. The html file created is generally written to the project's docs/Rmd/include directory. However, if this function is used to create a navbar for a Github user web page, the html file should be stored in a sensible location inside the user pages repository, e.g., leonawicz.github.io/assets.

The common navigation bar html is included at the beginning of the body of the html for each web page in the project's website. menu is a vector of names for each dropdown menu. submenus is a list of vectors of menu options corresponding to each menu. files is a similar list of vectors. Each element is either an html file for a web page to be associated with the submenu link, "header" to indicate the corresponding name in submenus is only a group label and not a link to a web page, or "divider" to indicate placement of a bar for separating groups in a dropdown menu.

theme can be either united (default) or cyborg. Both are from Bootswatch. The function must apply some internal differentiation in the construction of the html navigation bar between themes. This is currently the only rpm function which attempts to handle multiple Bootswatch themes with different CSS tags.

```
m <- menu[x]
   s <- submenus[[x]]
        y <- paste0("<li>'a href=\"", f, "\">", m, "</a>\n")
        y <- paste0("<li class=\"dropdown\">\n
            paste(sapply(1:length(s), fillSubmenu, name = s, file = f, theme = theme),
if (include.home)
   home <- paste0("<li>'a href=\"", home.url, "\">Home</a>\n
                                                                              ") else home <- ""
x <- paste0("<div class=\"navbar navbar-default navbar-fixed-top\">\n <div class=\"",
                                                           <button type=\"button\" class=\"".</pre>
                                                                                  <span class=\"icon-b</pre>
                                                                              <div class=\"",
   paste(sapply(1:length(menu), fillMenu, menu = menu, submenus = submenus,
                                                                    <a class=\"btn btn-primary\" href</pre>
                                                                                           </div>
sink(htmlfile)
sink()
```

0.0.2 genOutyaml

genOutyaml generates the _out.yaml file for yaml front-matter common to all html files in the project website. The file should be written to the project's docs/Rmd directory. lib specifies the library directory for any associated files. yaml includes for external html common to all project web pages in the site can also be specified with header, before_body, and after_body. These can be specified by file basename only (no path) and the function assumes these files are in the docs/Rmd/libs. It is recommended. See the project repo [gh-pages](https://github.com/leonawicz/ProjectManagement/tree/gh-pages "gh-pages") branch for an example.

```
genOutyaml <- function(file, theme = "cosmo", highlight = "zenburn", lib = NULL,
   header = NULL, before_body = NULL, after_body = NULL) {
   output.yaml <- paste0("html_document:\n self_contained: false\n theme: ",
        theme, "\n highlight: ", highlight, "\n mathjax: null\n toc_depth: 2\n")
   if (!is.null(lib))
        output.yaml <- paste0(output.yaml, " lib_dir: ", lib, "\n")
   output.yaml <- paste0(output.yaml, " includes:\n")
   if (!is.null(header))</pre>
```