

FIRE: THE FIRST-YEAR INNOVATION & RESEARCH EXPERIENCE

SUSTAINABILITY ANALYTICS

How to Create a GitHub-Flavored Markdown Webpage with Quarto

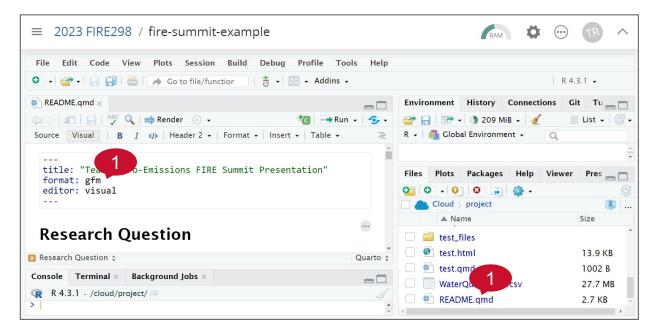
Step 1: Clone GitHub Template into Posit Cloud

- Find the repository
 fire-summit-team-name in GitHub
- 2. Copy the repository's HTTPS
- Clone the repository into the 2023
 FIRE298 Workspace in posit.cloud

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	fire-summit-whistle-blowers (Public)
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Step 2: Open Markdown File and Start Editing

- 1. Click on README.qmd. Note that the format of this document is gfm.
- 2. Start filling out the document



Step 3: Upload files that you would like to showcase

 Upload data and images that you would like to include. Make sure that each file does not exceed 100 MB (the smaller the better).

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Tip 1: Showcase only iteration in one loop

{r} Ξ. #for (i in 1:length(f)) { # print(f[i]) #r<-rast(paste0("G:/Shared drives/2023 FIRE-SA/FALL OUTPUT/Team Microplastic</pre> /NASA output data/",f[i])) r<-rast("201001.nc4") rp<-project(r, crs(cmd))</pre> Evap tavg va<-extract(rp[[1]], cva, fun="mean", na.rm=TRUE)</pre> Evap tavg md<-extract(rp[[1]], cmd, fun="mean", na.rm=TRUE)</pre> Qs tavg va<-extract(rp[[9]], cva, fun="mean", na.rm=TRUE) Qs tavg md<-extract(rp[[9]],cmd,fun="mean",na.rm=TRUE)</pre> Rainf f tavg va<-extract(rp[[12]],cva,fun="mean",na.rm=TRUE)</pre> Rainf f tavg md<-extract(rp[[12]], cmd, fun="mean", na.rm=TRUE)</pre> Tair_f_tavg_va<-extract(rp[[19]], cva, fun="mean", na.rm=TRUE)</pre> Tair f tavg md<-extract(rp[[19]], cmd, fun="mean", na.rm=TRUE)</pre> va<-data.frame(NAME10=cva\$NAME10,</pre> NAMELSAD10=cva\$NAMELSAD10, Evap tavg=Evap tavg va\$Evap tavg, Qs_tavg=Qs_tavg_va\$Qs_tavg, Rainf f tavg=Rainf f tavg va\$Rainf f tavg, Tair f tavg=Tair f tavg va\$Tair f tavg) #Tair f tavg=Tair f tavg va\$Tair f tavg, #file=f[i])

Tip 2: Show an example of a dataframe with head()

An example of control variables from January 2010 in Maryland is shown below. head(md) NAMELSAD10 Evap tavg Qs tavg Rainf f tavg NAME10 Worcester Worcester County 1.402706e-05 6.656126e-06 3.811770e-05 Harford Harford County 1.438178e-05 3.877095e-06 3.511265e-05 2 3 Montgomery Montgomery County 1.093989e-05 2.293109e-06 2.710311e-05 4 Washington Washington County 1.659733e-05 5.653316e-06 3.779316e-05 5 Carroll County 1.466791e-05 1.895020e-06 3.017357e-05 6 St. Mary's St. Mary's County 1.558496e-05 6.449214e-06 4.059291e-05

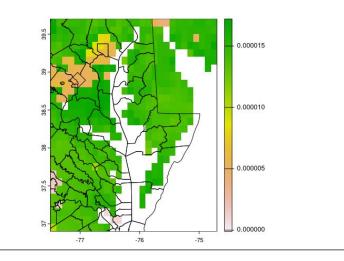
Tip 3: Show a map if you can

Average evapotranspiration levels from January 2010 is shown below.

cmd<-vect("Shapefiles/tl_2020_24_county10.shp")
cva<-vect("Shapefiles/tl_2020_51_county10.shp")</pre>

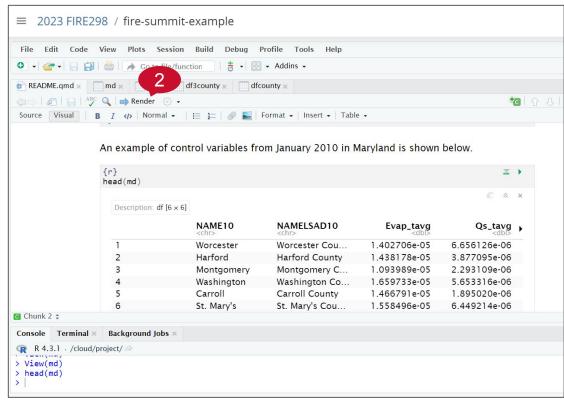
r<-rast("201001.nc4")
rp<-project(r[[1]], crs(cmd))</pre>

plot(rp)
plot(cmd, add=TRUE)
plot(cva, add=TRUE)



Step 4: Render the file

- After everyone in the team has finished their part. Combine everything into the README.qmd file.
- 2. Render the file.



Step 5: Push the file to GitHub to publish the page

- 1. Check each file that you have updated and Click Commit.
- 2. Type in a Commit message and Click Commit in the pop-up window.
- 3. Click Push.
- 4. Type in your GitHub username and Token.

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Step 6: Check your work

- Go to your repository on GitHub and make sure that everything is published.
- You can modify the page as many times as you like by updating README.qmd, rendering it, and pushing it to GitHub.

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