



# DIA Biostatistics Industry and Regulator Forum

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#BioStats22



# Achieving Regulatory Approval Using R

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FDA/CDER/OB

DIA

# Disclaimer

This presentation reflects the views of the presenter and should not be construed to represent Food and Drug Administration views or policies

# Statistical Software Clarifying Statement

FDA does not require use of any specific software for statistical analyses, and statistical software is not explicitly discussed in Title 21 of the Code of Federal Regulations [e.g., in 21CFR part 11]. However, the software package(s) used for statistical analyses should be fully documented in the submission, including version and build identification.

# Outline

## ► Overview of Prograf RWE sNDA

- Clinical/Regulatory Background
- Summary of Study
- Scientific Registry of Transplant Recipients
- Standard Analytic Files

## ► Pre-/Post-Submission Discussion

## ► Using R-Markdown for Regulatory Submission

- Raw Data Format and Analysis Data Creation
- Examples of R-Markdown Practices and Output
- Lessons Learned from Regulatory Perspective

# Overview of Prograf RWE Application

## ► Clinical/Regulatory Background

- Prograf® (Tacrolimus) was indicated for the prophylaxis of organ rejection in adult and pediatric patients receiving allogeneic lung transplant in combination with other immunosuppressants
- Prograf was approved for prophylaxis of organ rejection in patients receiving liver transplants in 1994 (later for kidney & heart), based on RCT Evidence
- RCTs for lung was not submitted to FDA, but drug has been used widely in clinical care; sponsor (Astellas) submitted supplemental New Drug Application to FDA on Dec 15, 2020
- Approval for preventing rejection/death in lung transplant granted Jul 16, 2021

# Overview of Prograf RWE Application

## ► Summary of Study

- **Study Design:** Non-interventional (observational) treatment arm, compared to historical controls
- **Primary Endpoint:** A composite endpoint of graft failure (GF) or death (due to any cause) within one year (365 days) after transplant
- **Data Source:** Scientific Registry of Transplant Recipients (SRTR) data on all lung transplants in US during 1999–2017
- **Study Population:** Adult and pediatric patients in tacrolimus immediate release (TAC IR) in combination with mycophenolate mofetil (MMF) or azathioprine (AZA)

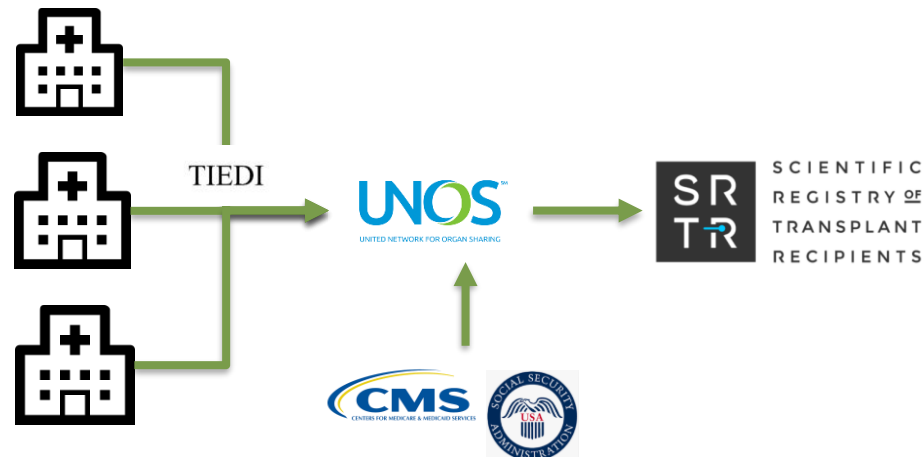


<https://www.fda.gov/drugs/news-events-human-drugs/fda-approves-new-use-transplant-drug-based-real-world-evidence>

# Overview of Prograf RWE Application

## ► Scientific Registry of Transplant Recipients (SRTR)

- The SRTR is a national transplant registry and made available under a data use agreement to external researchers and includes outcomes for all transplant recipients, candidates, recipients and donors in the United States from Oct 1987 onward
- For Prograf RWE application, the sponsor submitted the SRTR standard analytic files (SAFs) that contain data on all lung transplant candidates, recipients, and donors in the United States between 1999 and 2017





# Overview of Prograf RWE Application

## Public Data Dictionary of SRTR Standard Analytic Files

ABOUT SRTR ▾ ABOUT THE DATA ▾ REPORTS ▾ TOOLS ▾ NEWS & MEDIA ▾ REQUESTING SRTR DATA ▾ FAQs ▾					
<a href="#">◀ Home</a>   SAF Data Dictionary					
SRTR 2112 Public SAFs Data Dictionary					
Data Sets		Data File: DONOR_DECEASED			
All Deceased Donor Information					
<a href="#">File Linking Diagram</a>					
General		Variable	Type	Length	Format
DONOR_DECEASED		DONOR_ID	num	8	
DONOR_DISPOSITION		DON_A1	num	8	<a href="#">ALOCUS</a>
DONOR_LIVE		DON_A2	num	8	<a href="#">ALOCUS</a>
DON_LIV_FOL		DON_ABNORM_CONGEN	char	1	
FOL_IMMUNO		DON_ABNORM_LVH	char	1	
HIST_OPO_TXC		DON_ABNORM_VALVES	char	1	
IMMUNO		DON_ABO	char	3	<a href="#">SABO</a>
INSTITUTION		DON_AGE	num	8	
MALIG		DON_AGE_IN_MONTHS	num	8	
REC_HISTO		DON_ALLOC_ECD_CVA	num	3	
REC_HISTO_XMAT		DON_ALLOC_ECD_HYPERTEN	num	3	
TREATMENT		DON_ALLOC_ECD_SERUM_CREAT	num	8	
Heart & Lungs		DON_ALLOC_ECD_SERUM_CREAT_DT	num	8	MMDDYY
Kidney & Pancreas		DON_ALLOC_REMAIN_LI_SEG	char	1	
Liver & Intestines		DON_ANTI_CMV	char	2	<a href="#">SSRLSTT</a>
		DON_ANTI_CONVULS	char	1	
		DON_ANTI_HBC	char	2	<a href="#">SSRLSTT</a>
		DON_ANTI_HCV	char	2	<a href="#">SSRLSTT</a>
		DON_ANTI_HIV	char	2	<a href="#">SSRLSTT</a>
		DON_ANTI_HTLV	char	2	<a href="#">SSRLSTT</a>
		DON_ANTI_HYPERTEN	char	1	
		DON_ARGININE	char	1	
		DON_B1	num	8	<a href="#">BLOCUS</a>
		DON_B2	num	8	<a href="#">BLOCUS</a>
		DON_BIOGV_DCN	num	8	<a href="#">BLOCUS</a>
		DON_BIOGV_DCN	num	8	<a href="#">BLOCUS</a>

20680 subjects with single observation, 453 variables included in one final dataset

# Pre-submission Discussions

	Typical NDA Submission	Prograf RWE Submission
<b>Study Report</b>	<b>Y</b>	<b>Y</b>
<b>Define Document</b>	<b>Y</b>	<b>Y</b>
<b>FDA ESG</b>	<b>Y</b>	<b>Y</b>
<b>Standardized Data</b>	<b>Y</b>	<b>N</b>
<b>SAS XPORT</b>	<b>V5</b>	<b>V8</b>
<b>Programming Language</b>	<b>SAS</b>	<b>R</b>

*ESG: Electronic Submission Gateway*

# Pre-submission Discussions

- ▶ Type C meeting on Feb 2020: Asked whether hybrid of SAS and R (employing packages such as “survival”, the “tidyverse” suite and “rmarkdown”) is acceptable
- ▶ Type B meeting on Aug 2020: Asked whether the programs submitted in .Rmd and html formats will be acceptable for review
  - Agreed with the program submission using R markdown. Asked sponsor to generate the final format into both html and pdf formats. Also, specify alternative packages if applicable.
  - Asked sponsor to submit a sample data (200 subjects) with relevant programs and document.
- ▶ Sample data/code submission and testing on Nov 2020: FDA assessed sample data with relevant R-markdown before sNDA submission on Dec 2020

# Post-submission Discussions

- ▶ Information Request on Jan 2021: Experienced difficulties installing company-owned R package; Not compatible with FDA (R 3.6.1 and R 4.0.3) and does not disclose a GitHub allowing for manual download
  - Submit the GitHub address
  - Clarify the required R version
  - Provide a reference manual (static pdf vignette)
- ▶ Information Request on Apr 2021: Experienced difficulties using sponsor suggested LaTeX editor
  - Submitted Rmd programs in a LaTeX-free environment

# Using R-Markdown for Regulatory Submission

## ► Raw Data Format of Standard Analytic Files (SAFs)

- SRTR provides data only in SAS format (“.xpt/sas7bdat” extension)
- SAS 9.4 (SAS/STAT 15.1) was used to read SAF .xpt/sas7bdat files
- Sponsor provided programming code and analysis data in R-Markdown (.Rmd)
- Formatted SAS variables were converted into corresponding CSV variables that were composed of both the unformatted and formatted values
- Imported the CSV files into R to create dataset of interest
- R-Markdown provided both the R code as well as comments and intermittent results that create transparent documentation of the process



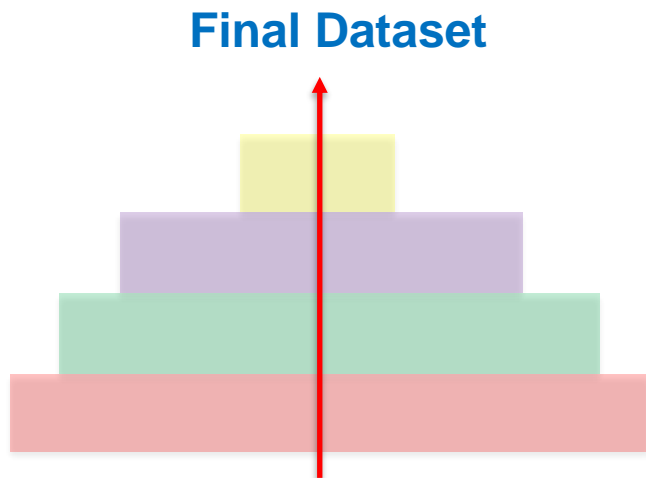
# Using R-Markdown for Regulatory Submission

## ► Analysis Dataset Description

- Analysis datasets support all protocol- and statistical analysis plan-specified objectives
  - tx\_data (Lung Transplant Analysis Data): A dataset with one record per patient containing all variables used in the tables and figures
  - Efficacy, safety, baseline characteristics
  - No PK/PD
- No ADaM conformance checks were performed
- Although the define file has the “look and feel” of define.xml v2.0, the underlying xml is not compliant with the standard

# Using R-Markdown for Regulatory Submission

## Steps for Data Management

[illegible]

# Using R-Markdown for Regulatory Submission

## Snapshots of final data management stage

Load Data

Set up Functions

Derive Variables

Output

### Lung Transplant Derive Post Transplant Outcome Variables

This document shows the derivation of the post transplant outcome variables.

Code ▾

### Load Data

```
#set up

library(dplyr)
library(tidyr)
library(CDMSDataAnalysisTools)
library(lubridate)
library(knitr)

library(ggplot2)
library(survival)
library(survminer)
library(kableExtra)

colorblind <- c(
  "#E69F00",
  "#56B4E9",
  "#009E73",
  "#F0E442",
  "#0072B2",
  "#D5E080")

DATADIR1 <- file.path("./csv_datasets")

rx_data <- load_all_analysis_datasets(dataset_directory = DATADIR1, include,
  set_names = c("rx_tx", "Followup"))

load(paste0(DATADIR1, "/derive_post_tx_variables.RData"))
```

Hide

- Load Data
- Set up Functions
- Derive Variables
- Output

```
_3yr, ltf_end_date_3yr, na.rm=TRUE))

.dgt_bx_act_ltf = ifelse(rx_year %in% 1999:2005, NA, death_gf ~ bx ~ ar_created ~ ltf)
.dgt_bx_act_ltf_all = ifelse(rx_year %in% 1999:2005, NA, death_gf_all ~ bx_all ~ ar_created_all
) ~ ltf_all)
.dgt_bx_act_ltf_3yr = ifelse(rx_year %in% 1999:2005, NA, death_gf_3yr ~ bx_3yr ~ ar_created_3yr
) ~ ltf_3yr)
.dgt_bx_act_ltf_2yr = ifelse(rx_year %in% c(1999:2005, 2017), NA, death_gf_2yr ~ bx_2yr ~ ar_c
reated_2yr ~ ltf_2yr)
.dgt_bx_act_ltf_1yr = ifelse(rx_year %in% c(1999:2005, 2016:2017), NA, death_gf_1yr ~ bx_1yr
) ~ ar_created_1yr ~ ltf_1yr)

.dgt_bx_act_ltf_dt = as_data(pmin(death_gf_dt, bx_dt, ar_created_dt, ltf_dt, na.rm=TRUE))
.dgt_bx_act_ltf_end_date_all = as_data(pmin(death_gf_end_date_all, bx_end_date_all, ar_created
_end_date_all, ltf_end_date_all, na.rm=TRUE))
.dgt_bx_act_ltf_end_date_3yr = as_data(pmin(death_gf_end_date_3yr, bx_end_date_3yr, ar_created
_end_date_3yr, ltf_end_date_3yr, na.rm=TRUE))
.dgt_bx_act_ltf_end_date_2yr = as_data(pmin(death_gf_end_date_2yr, bx_end_date_2yr, ar_created
_end_date_2yr, ltf_end_date_2yr, na.rm=TRUE))
.dgt_bx_act_ltf_end_date_1yr = as_data(pmin(death_gf_end_date_1yr, bx_end_date_1yr, ar_created
_end_date_1yr, ltf_end_date_1yr, na.rm=TRUE))

}
```

### Output

```
tx_data<- tx_data%>%
  select(-TRR_ID, -TFL_LAFUDATE, -TFL_GRAFT_DT, -PERS_RET_X)

save(tx_data, file=paste0(DATADIR1, "/derive_ltf_variables.RData"))
```

Hide



# Using R-Markdown for Regulatory Submission

## HTML Output of R-Markdown

Overview

Study Population

Non-Cumulative Exclusions

Cumulative Exclusions

Immunosuppressive Regimen at Discharge

Baseline Characteristics of Lung Transplant Recipients

Trends by Transplant Era

- Exclusion Criteria: See: [exclusion\\_criteria.html](#)
- Derive Drug Variables: See: [derive\\_drug\\_variables.html](#)
  - Derive Other Variables: See: [derive\\_variables.html](#)

### Study Population

There were a total of 29772 lung transplants between 1999 and 2017; after exclusions 26080 lung transplant recipients remained in the study cohort.

### Non-Cumulative Exclusions

Study Population Exclusions by Age: Non-cumulative

Grouping	Characteristic	Total	Adult	Pediatric
		N(%)	N(%)	N(%)
All		29772	28817	955
Received any previous transplant, lung or otherwise		1240 (4.2%)	1174 (4.1%)	66 (6.9%)
Multi-organ transplant		134 (0.5%)	120 (0.4%)	14 (1.5%)
Living donor transplant		245 (0.8%)	152 (0.5%)	93 (9.7%)
Missing discharge date		248 (0.8%)	236 (0.8%)	12 (1.3%)
Discharged after 1 year		34 (0.1%)	34 (0.1%)	0
Missing maintenance immunosuppression information at discharge		975 (3.3%)	941 (3.3%)	34 (3.6%)
Death, graft failure, or retransplant before discharge		1837 (6.2%)	1767 (6.1%)	70 (7.3%)
Incorrect Death Date		1 (0.0%)	1 (0.0%)	0
Incorrect Retransplant Date		1 (0.0%)	1 (0.0%)	0
Any Exclusion		3692 (12.4%)	3462 (12.0%)	230 (24.1%)
Included		26080 (87.6%)	25355 (88.0%)	725 (75.9%)

Note:

A patient may be excluded for more than one reason. Non-cumulative data report counts and percent among all patients. The sum of all non-cumulative exclusions exceeds the total number of excluded patients.

\* Program: './Create\_cohort\_files/NoLatex\_Demographics\_Report.Rmd'

#### Overview

Data Availability at Annually Collected Outcomes

One, Two, and Three Year Outcomes By Era

One Year Death or Graft Failure By Subgroup

Adult: Cumulative Incidence

Adult: Proportional Hazard

Pediatrics: Cumulative Incidence

Cumulative Incidence By Month

Trends Over Time

Efficacy

Safety

### One Year Death or Graft Failure By Subgroup

#### Adult: Cumulative Incidence

Kaplan-Meier Estimate of Cumulative Incidence Percent (95% CI) of Death or Graft Failure at 1y Post-transplant by Subgroup: Adults

Grouping		At Risk	Events	% (95% CI)
Age at Transplant	18-34 years	2884	225	18.3% (15.0, 20.2)
Age at Transplant	35-49 years	3786	272	16.0% (14.5, 20.3)
Age at Transplant	50-64 years	13257	1214	11.7% (10.0, 15.3)
Age at Transplant	65+ years	3508	633	11.8% (11.0, 12.7)
Recipient Sex	Female	10629	920	12.7% (7.2, 17.8)
Recipient Sex	Male	14526	1424	13.2% (11.7, 17.4)
Lung Transplant Procedure	Double	16190	1297	13.9% (11.2, 19.2)
Lung Transplant Procedure	Single	9165	1047	12.0% (11.1, 12.8)
Transplant Era	1999-2005	6284	725	16.8% (7.9, 24.8)
Transplant Era	2006-2009	3024	519	11.7% (9.3, 14.0)
Transplant Era	2010-2017	14047	1190	12.3% (10.3, 17.9)
Diagnosis Group	A - COPD	9529	896	17.0% (15.8, 26.1)
Diagnosis Group	B - Pulmonary hypertension	966	80	23.8% (10.0, 46.8)
Diagnosis Group	C - Cystic fibrosis	3196	247	15.0% (10.0, 28.8)
Diagnosis Group	D - Pulmonary fibrosis	11621	1154	10.3% (9.7, 10.8)
Diagnosis Group	Unknown	41	7	17.1% (4.7, 27.8)
BMI	<18.5	2323	243	10.6% (9.4, 11.9)
BMI	18.5-<25	10228	844	13.0% (10.5, 19.9)
BMI	25-<30	8889	857	13.8% (7.9, 19.4)
BMI	30+	3646	360	10.0% (9.4, 11.6)
BMI	Unknown	269	40	15.3% (10.8, 19.5)

# Using R-Markdown for Regulatory Submission

## ► Lesson Learned from Regulatory Perspective

- RWD sources are disparate and non-uniform
- For this submission, RWD structure was complex and had no prior knowledge or experience
- R-Markdown provided transparent data management and analysis processes for review
- Easy for reproducing and navigating results and supported interdisciplinary collaboration
- Challenge existed when using R-Markdown throughout the review
  - Provide all necessary documents to aid review process
  - Submit all relevant analytic packages
  - Complex features are unnecessary and may not be compatible with FDA environment
  - Make it analyzable/executable: simple coding practices are acceptable



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# Appendix: Submission Issues

## ► Data Submission with SAS XPORT (.xpt) version 8

	Version 5†	Version 8
Variable Name	8 chars	32 chars
Variable Label	40 chars	256 chars
Variable Length	200 chars	32,767 chars

†FDA Study Data Technical Conformance Guide (version 4.5.1, July 2020)

- Data Submission through File Transfer Protocol (FTP)
- Sending a link to sponsor's database instead of using Electronic Submission Gateway Code Submission using R Markdown
- No SAS codes for data management and analysis

# Appendix: Submission Issues

## ► Data Submission with SAS XPORT (.xpt) version 8

- Comment: V8 is beneficial for lifting limits of characters. However, it will further increase file size, concerns on no native mechanism for audit trails, referencing source data. Although, it is advertised as nonproprietary, not sure whether it is vendor neutral.
- Conclusion: Although FDA cannot require v8, accepting it is different. We decided to accept v8 by reviewing a sample data.

## ► Data Submission through File Transfer Protocol (FTP)

- Comment: According to eCTD binding guidance† (section III M), the sponsor must use the FDA electronic submission gateway (ESG) for all submissions that are 10 gigabytes or smaller
- Conclusion: FDA does not accept regulatory submissions with File Transfer Protocol (FTP). The FDA ESG enables the secure submission of regulatory information for review and is our preferred method of transmission

†Providing Regulatory Submissions in Electronic Format — Certain Human Pharmaceutical Product Applications and Related Submissions Using the eCTD Specifications (Feb 2020).

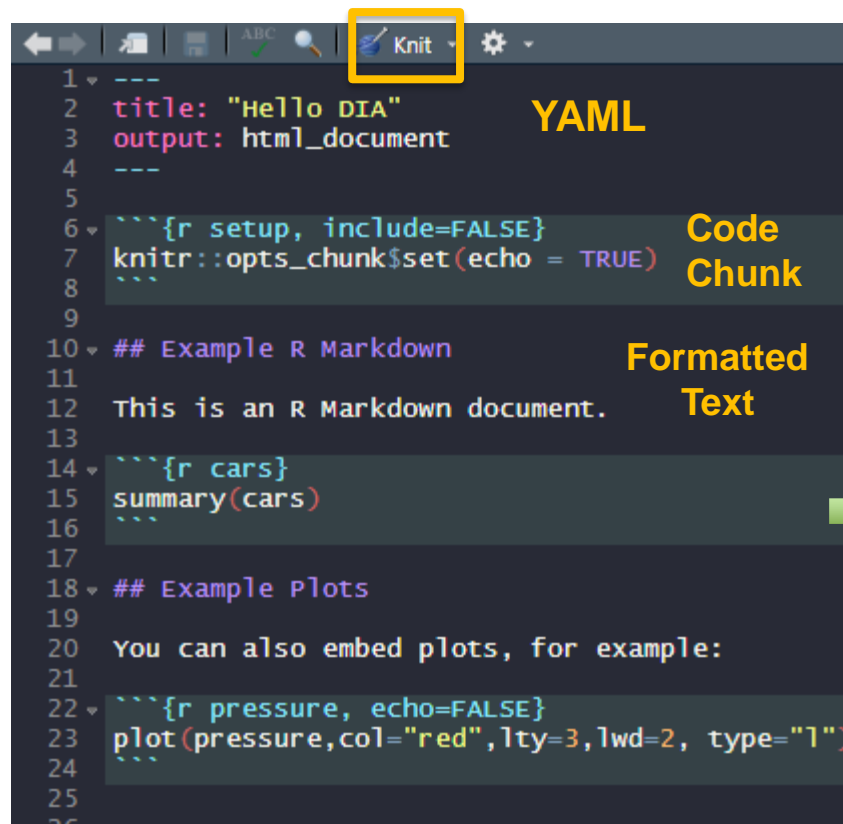
## ► Code Submission using R Markdown

- Conclusion: FDA agree with the program submission using R markdown. If your programs use R packages, specify alternative packages if applicable.

# Using R-Markdown for Regulatory Submission

## R Markdown

- Provides a unified framework for combining code, results, and comments
- Reproducible and shareable
- Comes pre-installed with RStudio



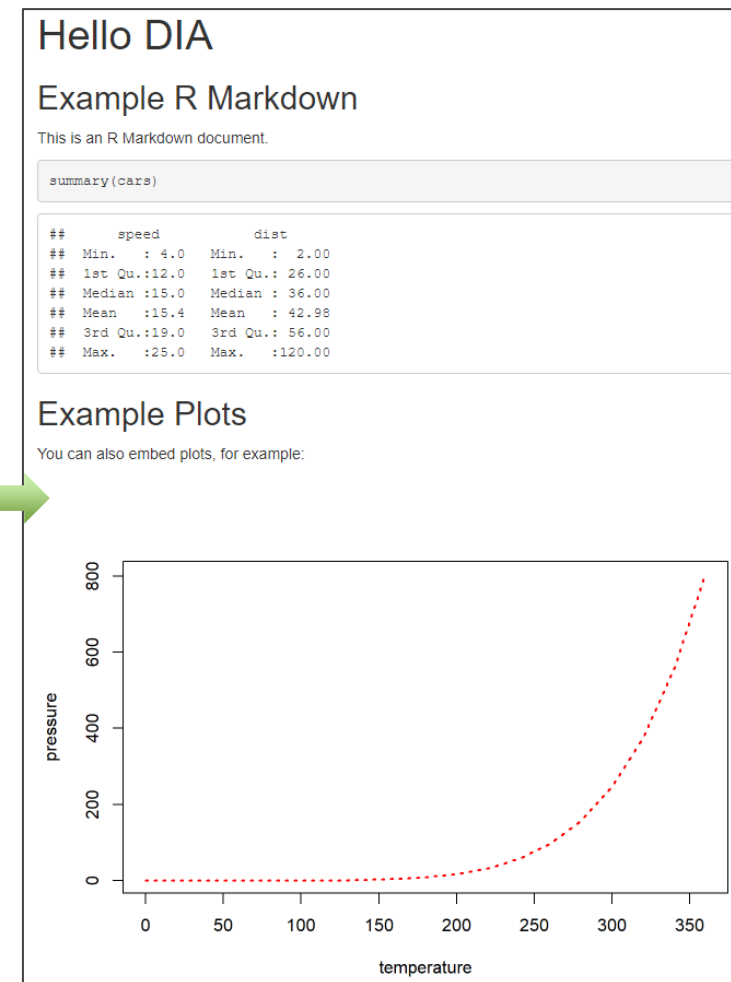
```
1 ---
2 title: "Hello DIA"
3 output: html_document
4 ---
5
6 {r setup, include=FALSE}
7 knitr::opts_chunk$set(echo = TRUE)
8
9
10 ## Example R Markdown
11
12 This is an R Markdown document.
13
14 {r cars}
15 summary(cars)
16
17
18 ## Example Plots
19
20 You can also embed plots, for example:
21
22 {r pressure, echo=FALSE}
23 plot(pressure,col="red",lty=3,lwd=2, type="l")
24
25
26
```

YAML

Code Chunk

Formatted Text

\*. Rmd



\*. html