Hands-On Example: Creating a Final Report with Quarto

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Objective

The goal of this section is to use the complete CDISC-compliant SDTM and ADaM datasets to **create a final report using Quarto**. This report will include tables, listings, and figures (TLFs), which are commonly required for regulatory submissions.

Key Points for Presentation

- Overview: Use the complete CDISC-compliant SDTM and ADaM datasets to create a final report using Quarto.
- Required Libraries: {tidyverse} for data manipulation and figures, {gt} and {gtsummary} for listings and tables, and {quarto} for generating the final reproducible report.
- Input Data: Imputed SDTM and ADaM Datasets.
- Data Understanding: Understand how to create TLFs using imputed datasets to ensure compliance with regulatory standards.
- Report Creation Process:
 - Generate tables and listings using {gt} and {gtsummary}.
 - Create figures using {ggplot2}.
 - Compile the final report using {quarto}.

Step-by-Step Report Creation with Quarto

Step 1: Load Required Libraries

```
if (!requireNamespace(c("tidyverse", "gt", "gtsummary", "quarto"), quietly = TRUE)) {
   install.packages(c("tidyverse", "gt", "gtsummary", "quarto"))
}
# Load necessary libraries
library(tidyverse) # Tidyverse for data manipulation and figures
-- Attaching core tidyverse packages ----- tidyverse 2.0.0 --

      v dplyr
      1.1.4
      v readr
      2.1.5

      v forcats
      1.0.0
      v stringr
      1.5.1

      v ggplot2
      3.5.1
      v tibble
      3.2.1

      v lubridate
      1.9.3
      v tidyr
      1.3.1

            1.0.2
v purrr
-- Conflicts ----- tidyverse_conflicts() --
x dplyr::filter() masks stats::filter()
x dplyr::lag() masks stats::lag()
i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become
library(gt)
                # GT for creating high-quality tables
library(gtsummary) # Gtsummary for creating clinical trial summary tables
library(quarto)  # Quarto for generating the final report
# Loading the dataset in the final document
```

Explanation: We start by loading the {tidyverse} package for data manipulation and visualization, {gt} and {gtsummary} for creating tables and listings, and {quarto} for generating the final report. We also load in our previously generated SDTM and ADaM datasets.

sdtm_dm_imputed <- readRDS("sdtm_dm_imputed.rds") # Loads our SDTM data
adam_dm_imputed <- readRDS("adam_dm_imputed.rds") # Loads our ADaM data</pre>

Step 2: Generate Summary Tables

```
# Generate a summary table for demographic characteristics using gtsummary
summary_table <- sdtm_dm_imputed %>%
    # Select relevant columns for summarization: AGE, SEX, ETHNIC, and RACE
    select(AGE, SEX, ETHNIC, RACE) %>%

# Create a summary table with grouping by SEX
    tbl_summary(
```

```
by = SEX, # Group data by SEX to summarize separately for each group
statistic = list(all_continuous() ~ "{mean} ({sd})") # Define how to summarize continuo"
) %>%

# Bold the labels for better visualization in the output table
bold_labels()
```

12 observations missing `SEX` have been removed. To include these observations, use `forcats

```
# View the summary table summary_table
```

Table printed with `knitr::kable()`, not {gt}. Learn why at https://www.danieldsjoberg.com/gtsummary/articles/rmarkdown.html
To suppress this message, include `message = FALSE` in code chunk header.

Characteristic	F, N = 61	M, N = 77
AGE	49 (15)	46 (17)
ETHNIC		
HISPANIC OR LATINO	14 (23%)	30 (39%)
NOT HISPANIC OR LATINO	29 (48%)	5~(6.5%)
NOT REPORTED	6 (9.8%)	21~(27%)
UNKNOWN	12 (20%)	21~(27%)
RACE		
AMERICAN INDIAN OR ALASKA NATIVE	7 (11%)	0 (0%)
ASIAN	7 (11%)	0 (0%)
BLACK OR AFRICAN AMERICAN	0 (0%)	15 (19%)
NATIVE HAWAIIAN OR OTHER PACIFIC	7 (11%)	0 (0%)
ISLANDER		
NOT REPORTED	8 (13%)	26 (34%)
OTHER	8 (13%)	14 (18%)
UNKNOWN	19 (31%)	7 (9.1%)
WHITE	5 (8.2%)	15 (19%)

Explanation: Here, we use {gtsummary} to create a summary table for demographic characteristics, grouped by SEX. The tbl_summary() function is used to summarize the AGE, ETHNIC, and RACE variables, with continuous variables reported as mean and standard deviation.

Step 3: Generate High-Quality Tables for Listings

```
# Generate a high-quality listing using gt
listing_table <- sdtm_dm_imputed %>%
    select(STUDYID, USUBJID, AGE, SEX, ETHNIC, RACE) %>% # Select relevant columns for listing gt() # Create a high-quality table using gt
# View the listing table
listing_table
```

STUDYID	USUBJID	AGE	SEX	ETHNIC	RACE
CDASH_DEMO_01	CDASH_DEMO_01-SUB001	53	F	NOT HISPANIC OR LATINO	NOT
CDASH_DEMO_01	CDASH_DEMO_01-SUB002	60	F	HISPANIC OR LATINO	NATI
CDASH_DEMO_01	CDASH_DEMO_01-SUB003	32	\mathbf{F}	NOT HISPANIC OR LATINO	UNKI
CDASH_DEMO_01	CDASH_DEMO_01-SUB004	49	F	UNKNOWN	AMEI
CDASH_DEMO_01	CDASH_DEMO_01-SUB005	24	\mathbf{M}	HISPANIC OR LATINO	NOT
CDASH_DEMO_01	CDASH_DEMO_01-SUB006	24	\mathbf{M}	HISPANIC OR LATINO	NOT
CDASH_DEMO_01	CDASH_DEMO_01-SUB007	58	\mathbf{F}	NOT HISPANIC OR LATINO	OTHE
CDASH_DEMO_01	CDASH_DEMO_01-SUB008	40	\mathbf{M}	NOT REPORTED	BLAC
CDASH_DEMO_01	CDASH_DEMO_01-SUB009	44	\mathbf{M}	NOT REPORTED	NOT
CDASH_DEMO_01	CDASH_DEMO_01-SUB010	77	\mathbf{M}	HISPANIC OR LATINO	WHIT
CDASH_DEMO_01	CDASH_DEMO_01-SUB011	70	\mathbf{M}	UNKNOWN	OTHE
CDASH_DEMO_01	CDASH_DEMO_01-SUB012	24	\mathbf{F}	NOT HISPANIC OR LATINO	UNKI
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CDASH_DEMO_01	CDASH_DEMO_01-SUB014	44	\mathbf{M}	UNKNOWN	UNKI
CDASH_DEMO_01	CDASH_DEMO_01-SUB015	53	\mathbf{M}	NOT HISPANIC OR LATINO	NOT
CDASH_DEMO_01	CDASH_DEMO_01-SUB016	51	\mathbf{F}	HISPANIC OR LATINO	ASIAl
CDASH_DEMO_01	CDASH_DEMO_01-SUB017	80	\mathbf{F}	NOT REPORTED	UNKI
CDASH_DEMO_01	CDASH_DEMO_01-SUB018	24	\mathbf{M}	NOT REPORTED	WHIT
CDASH_DEMO_01	CDASH_DEMO_01-SUB019	42	\mathbf{F}	UNKNOWN	WHIT
CDASH_DEMO_01	CDASH_DEMO_01-SUB020	55	\mathbf{M}	UNKNOWN	OTHE
CDASH_DEMO_01	CDASH_DEMO_01-SUB001	53	\mathbf{F}	NOT HISPANIC OR LATINO	NOT
CDASH_DEMO_01	CDASH_DEMO_01-SUB002	60	\mathbf{F}	HISPANIC OR LATINO	NATI
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CDASH_DEMO_01	CDASH_DEMO_01-SUB004	49	\mathbf{F}	UNKNOWN	AMEI
CDASH_DEMO_01	CDASH_DEMO_01-SUB005	24	\mathbf{M}	HISPANIC OR LATINO	NOT
CDASH_DEMO_01	CDASH_DEMO_01-SUB006	26	\mathbf{M}	HISPANIC OR LATINO	NOT
CDASH_DEMO_01	CDASH_DEMO_01-SUB007	58	\mathbf{F}	NOT HISPANIC OR LATINO	OTHE
CDASH_DEMO_01	CDASH_DEMO_01-SUB008	40	M	NOT REPORTED	BLAC
CDASH_DEMO_01	CDASH_DEMO_01-SUB009	44	M	NOT REPORTED	NOT

CDASH_DEMO_01	CDASH_DEMO_01-SUB010	53	M	HISPANIC OR LATINO	WHIT
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	CDASH_DEMO_01	CDASH_DEMO_01-SUB010	77	M	HISPANIC OR LATINO	WHI
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Explanation: In this step, we use {gt} to create a listing of the demographic data. The gt() function helps generate publication-quality tables that can be included in the final report.

Step 4: Create Figures Using ggplot2

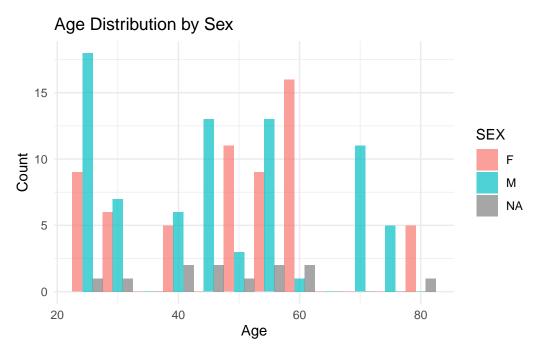
```
# Generate a histogram of AGE by SEX
age_histogram <- sdtm_dm_imputed %>%
  # Use the pipe operator to feed data into ggplot function for visualization
ggplot(aes(x = AGE, fill = SEX)) +

# Create a histogram with 'geom_histogram()'
  # 'aes()' defines the variables for the x-axis and fill color (by SEX)
geom_histogram(binwidth = 5, alpha = 0.7, position = "dodge") +

# Add descriptive labels for the title, x-axis, and y-axis of the plot
labs(title = "Age Distribution by Sex", x = "Age", y = "Count") +

# Apply 'theme_minimal()' for a cleaner, minimal visual style
theme_minimal()

# View the histogram
age_histogram
```



Explanation: We use {ggplot2} from {tidyverse} to create a histogram of AGE by SEX. This figure helps visualize the distribution of age across different sexes in the dataset.

Step 5: Compile the Final Report Using Quarto - hit RENDER

Explanation: Finally, we use {quarto} to compile the report. The quarto_render() function takes the input Quarto markdown file (final_report.qmd) and generates the output in HTML format, incorporating the SDTM and ADaM datasets.

Important CDISC Concepts for Creating Regulatory-Compliant Reports

- Standardization: The use of CDISC standards ensures that datasets are consistent across different clinical trials, which is essential for regulatory review.
- **Traceability**: All derived variables, tables, and figures must be traceable to their source data, ensuring transparency.
- **Metadata Documentation**: Proper documentation of variables and methods used in TLF generation is crucial for regulatory submissions.
- Regulatory Requirements: TLFs are a key part of clinical trial reporting, and adhering to ADaM standards ensures that these outputs meet regulatory guidelines.

Summary of Report Creation

- Tables and Listings: Created summary tables and listings using {gt} and {gtsummary} to meet regulatory submission requirements.
- Figures: Visualized key demographic data using {ggplot2}.
- Quarto Report: Compiled all tables, listings, and figures into a final report using {quarto}.
- Compliance with CDISC Standards: Ensured that all outputs were compliant with CDISC standards, providing traceability and proper documentation.

This hands-on section helps participants understand how to create a final clinical trial report that includes TLFs, emphasizing compliance with CDISC standards for regulatory submissions.