

BIOST 544: Homework 1

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End of report. Code appendix begins on the next page.

Code Appendix

```
# setup options
knitr::opts_chunk$set(echo = FALSE, message = FALSE)
options(knitr.kable.NA = '-')
labs = knitr::all_labels()
labs = labs[!labs %in% c("setup", "llm_appendix", "allcode")]
# clear environment
rm(list=ls())

# load relevant packages
# library(dplyr)      # data frame manipulation
# library(knitr)      # table formatting
# library(gtsummary)  # "table 1" summary
# library(tidyverse)

# load data
aml <- read.csv("../data/leukemia_data.csv")
```

End of document.