

Biostat 203B Homework 4

Due Mar 9 @ 11:59PM

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Display machine information:

```
sessionInfo()
```

```
R version 4.4.2 (2024-10-31)
Platform: x86_64-pc-linux-gnu
Running under: Ubuntu 24.04.1 LTS
```

```
Matrix products: default
BLAS: /usr/lib/x86_64-linux-gnu/blas/libblas.so.3.12.0
LAPACK: /usr/lib/x86_64-linux-gnu/lapack/liblapack.so.3.12.0
```

```
locale:
 [1] LC_CTYPE=C.UTF-8      LC_NUMERIC=C           LC_TIME=C.UTF-8
 [4] LC_COLLATE=C.UTF-8    LC_MONETARY=C.UTF-8    LC_MESSAGES=C.UTF-8
 [7] LC_PAPER=C.UTF-8      LC_NAME=C              LC_ADDRESS=C
[10] LC_TELEPHONE=C        LC_MEASUREMENT=C.UTF-8 LC_IDENTIFICATION=C
```

```
time zone: America/Los_Angeles
tzcode source: system (glibc)
```

```
attached base packages:
[1] stats      graphics  grDevices  utils      datasets  methods    base
```

```
loaded via a namespace (and not attached):
 [1] compiler_4.4.2    fastmap_1.2.0      cli_3.6.4          tools_4.4.2
 [5] htmltools_0.5.8.1 rstudioapi_0.17.1  yaml_2.3.10        rmarkdown_2.29
 [9] knitr_1.49         jsonlite_1.9.0     xfun_0.50          digest_0.6.37
[13] rlang_1.1.5       evaluate_1.0.3
```

Display my machine memory.

```
memuse::Sys.meminfo()
```

Totalram: 7.686 GiB

Freeram: 6.247 GiB

Load database libraries and the tidyverse frontend:

```
library(bigrquery)
library(dbplyr)
library(DBI)
library(gt)
library(gtsummary)
library(tidyverse)
```

```
-- 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  1.9.4      v tidyr      1.3.1
v purrr      1.0.4
```

```
-- Conflicts ----- tidyverse_conflicts() --
x dplyr::filter() masks stats::filter()
x dplyr::ident()   masks dbplyr::ident()
x dplyr::lag()     masks stats::lag()
x dplyr::sql()     masks dbplyr::sql()
```

i Use the conflicted package (<<http://conflicted.r-lib.org/>>) to force all conflicts to become

```
library(forcats)
```

```
library(stringr)
```

0.1 Q1. Compile the ICU cohort in HW3 from the Google BigQuery database

Below is an outline of steps. In this homework, we exclusively work with the BigQuery database and should not use any MIMIC data files stored on our local computer. Transform data as much as possible in BigQuery database and collect() the tibble **only at the end of Q1.7**.

0.1.1 Q1.1 Connect to BigQuery

Authenticate with BigQuery using the service account token. Please place the service account token (shared via BruinLearn) in the working directory (same folder as your qmd file). Do **not** ever add this token to your Git repository. If you do so, you will lose 50 points.

```
# path to the service account token
satoken <- "biostat-203b-2025-winter-4e58ec6e5579.json"
# BigQuery authentication using service account
bq_auth(path = satoken)
```

Connect to BigQuery database `mimiciv_3_1` in GCP (Google Cloud Platform), using the project billing account `biostat-203b-2025-winter`.

```
# connect to the BigQuery database `biostat-203b-2025-mimiciv_3_1`
con_bq <- dbConnect(
  bigrquery::bigquery(),
  project = "biostat-203b-2025-winter",
  dataset = "mimiciv_3_1",
  billing = "biostat-203b-2025-winter"
)
con_bq
```

```
<BigQueryConnection>
  Dataset: biostat-203b-2025-winter.mimiciv_3_1
  Billing: biostat-203b-2025-winter
```

List all tables in the `mimiciv_3_1` database.

```
dbListTables(con_bq)
```

[1] "admissions"	"caregiver"	"chartevents"
[4] "d_hcpcs"	"d_icd_diagnoses"	"d_icd_procedures"
[7] "d_items"	"d_labitems"	"datetimeevents"
[10] "diagnoses_icd"	"drgcodes"	"emar"
[13] "emar_detail"	"hpcsevents"	"icustays"
[16] "ingredientevents"	"inputevents"	"labevents"
[19] "microbiologyevents"	"omr"	"outputevents"
[22] "patients"	"pharmacy"	"poe"
[25] "poe_detail"	"prescriptions"	"procedureevents"
[28] "procedures_icd"	"provider"	"services"
[31] "transfers"		

0.1.2 Q1.2 icustays data

Connect to the `icustays` table.

```
# full ICU stays table
icustays_tble <- tbl(con_bq, "icustays") |>
  arrange(subject_id, hadm_id, stay_id) |>
  # show_query() |>
  print(width = Inf)
```

```
# Source:      SQL [?? x 8]
```

```

# Database:    BigQueryConnection
# Ordered by: subject_id, hadm_id, stay_id
  subject_id  hadm_id  stay_id first_careunit
      <int>    <int>    <int> <chr>
1    10000032  29079034  39553978 Medical Intensive Care Unit (MICU)
2    10000690  25860671  37081114 Medical Intensive Care Unit (MICU)
3    10000980  26913865  39765666 Medical Intensive Care Unit (MICU)
4    10001217  24597018  37067082 Surgical Intensive Care Unit (SICU)
5    10001217  27703517  34592300 Surgical Intensive Care Unit (SICU)
6    10001725  25563031  31205490 Medical/Surgical Intensive Care Unit (MICU/SICU)
7    10001843  26133978  39698942 Medical/Surgical Intensive Care Unit (MICU/SICU)
8    10001884  26184834  37510196 Medical Intensive Care Unit (MICU)
9    10002013  23581541  39060235 Cardiac Vascular Intensive Care Unit (CVICU)
10   10002114  27793700  34672098 Coronary Care Unit (CCU)
      last_careunit                                intime
      <chr>                                         <dtm>
1 Medical Intensive Care Unit (MICU)                2180-07-23 14:00:00
2 Medical Intensive Care Unit (MICU)                2150-11-02 19:37:00
3 Medical Intensive Care Unit (MICU)                2189-06-27 08:42:00
4 Surgical Intensive Care Unit (SICU)                2157-11-20 19:18:02
5 Surgical Intensive Care Unit (SICU)                2157-12-19 15:42:24
6 Medical/Surgical Intensive Care Unit (MICU/SICU)  2110-04-11 15:52:22
7 Medical/Surgical Intensive Care Unit (MICU/SICU)  2134-12-05 18:50:03
8 Medical Intensive Care Unit (MICU)                2131-01-11 04:20:05
9 Cardiac Vascular Intensive Care Unit (CVICU)      2160-05-18 10:00:53
10 Coronary Care Unit (CCU)                          2162-02-17 23:30:00
      outtime      los
      <dtm>          <dbl>
1 2180-07-23 23:50:47 0.410
2 2150-11-06 17:03:17 3.89
3 2189-06-27 20:38:27 0.498
4 2157-11-21 22:08:00 1.12
5 2157-12-20 14:27:41 0.948
6 2110-04-12 23:59:56 1.34
7 2134-12-06 14:38:26 0.825
8 2131-01-20 08:27:30 9.17
9 2160-05-19 17:33:33 1.31
10 2162-02-20 21:16:27 2.91
# i more rows

```

0.1.3 Q1.3 admissions data

Connect to the admissions table.

```

# # TODO
admissions_tble <- tbl(con_bq, "admissions") |>

```

```

print(width = Inf)

# Source:   table<`admissions`> [?? x 16]
# Database: BigQueryConnection
  subject_id  hadm_id  admittime              disctime
    <int>      <int> <dtm>                  <dtm>
1  10106244  26713233 2147-05-09 10:34:00 2147-05-12 13:43:00
2  13700703  20448599 2172-09-25 01:01:00 2172-10-03 13:25:00
3  15443666  27961368 2168-12-30 23:30:00 2169-01-05 16:02:00
4  16299919  26977065 2193-05-15 08:37:00 2193-05-17 16:03:00
5  14149715  24191358 2181-10-25 19:37:00 2181-10-29 14:38:00
6  14446098  20543394 2182-04-04 20:11:00 2182-05-07 19:00:00
7  10584718  23485217 2165-02-12 15:41:00 2165-03-06 08:20:00
8  12224488  25909420 2158-10-29 15:59:00 2158-11-01 15:45:00
9  15845632  28189199 2124-10-05 02:44:00 2124-10-12 15:00:00
10 18131667  28337235 2195-11-18 02:58:00 2195-11-27 13:34:00
  deathtime      admission_type  admit_provider_id
    <dtm>          <chr>         <chr>
1 NA              DIRECT EMER.    <NA>
2 NA              OBSERVATION ADMIT <NA>
3 NA              OBSERVATION ADMIT <NA>
4 NA              OBSERVATION ADMIT <NA>
5 NA              OBSERVATION ADMIT P00230
6 NA              URGENT           P004G6
7 2165-03-06 08:20:00 EW EMER.     P004G6
8 NA              EW EMER.         P004G6
9 NA              EW EMER.         P004G6
10 NA             EW EMER.         P004G6
  admission_location      discharge_location  insurance
    <chr>                  <chr>             <chr>
1 PHYSICIAN REFERRAL      HOME                Private
2 EMERGENCY ROOM          HOME                Private
3 EMERGENCY ROOM          HOME HEALTH CARE    Medicare
4 EMERGENCY ROOM          HOSPICE              Medicare
5 EMERGENCY ROOM          SKILLED NURSING FACILITY Medicare
6 TRANSFER FROM HOSPITAL   SKILLED NURSING FACILITY Medicare
7 TRANSFER FROM SKILLED NURSING FACILITY DIED    Medicare
8 WALK-IN/SELF REFERRAL   HOME                Medicare
9 PHYSICIAN REFERRAL      HOME                Private
10 PHYSICIAN REFERRAL     HOME HEALTH CARE    Medicare
  language marital_status race              edregtime
    <chr>      <chr>        <chr>            <dtm>
1 English  SINGLE         WHITE              NA
2 English  MARRIED        WHITE              2172-09-24 17:38:00
3 English  SINGLE         BLACK/AFRICAN AMERICAN 2168-12-30 11:19:00

```

```

4 English WIDOWED BLACK/AFRICAN AMERICAN 2193-05-15 04:36:00
5 English SINGLE WHITE 2181-10-25 08:48:00
6 English MARRIED WHITE NA
7 English MARRIED WHITE NA
8 English SINGLE WHITE - OTHER EUROPEAN 2158-10-28 20:22:00
9 English MARRIED WHITE 2124-10-04 19:30:00
10 English SINGLE WHITE 2195-11-17 21:04:00
  edouttime      hospital_expire_flag
  <dtm>          <int>
1 NA 0
2 2172-09-25 03:07:00 0
3 2168-12-31 01:22:00 0
4 2193-05-15 14:27:00 0
5 2181-10-26 15:18:00 0
6 NA 0
7 NA 1
8 2158-10-29 18:01:00 0
9 2124-10-05 04:10:00 0
10 2195-11-18 04:51:00 0
# i more rows

```

0.1.4 Q1.4 patients data

Connect to the patients table.

```

# # TODO
patients_tble <- tbl(con_bq, "patients") |>
  print(width = Inf)

```

```

# Source:   table<`patients`> [?? x 6]
# Database: BigQueryConnection
  subject_id gender anchor_age anchor_year anchor_year_group dod
    <int> <chr>      <int>      <int> <chr>      <date>
1   10078138 F          18        2110 2017 - 2019 NA
2   10180372 M          18        2110 2008 - 2010 NA
3   10686175 M          18        2110 2011 - 2013 NA
4   10851602 F          18        2110 2014 - 2016 NA
5   10902424 F          18        2110 2017 - 2019 NA
6   11092326 M          18        2110 2008 - 2010 NA
7   11289691 F          18        2110 2017 - 2019 NA
8   11595073 M          18        2110 2011 - 2013 NA
9   11739764 F          18        2110 2017 - 2019 NA
10  11776346 F          18        2110 2008 - 2010 NA
# i more rows

```

0.1.5 Q1.5 labevents data

Connect to the `labevents` table and retrieve a subset that only contain subjects who appear in `icustays_tble` and the lab items listed in HW3. Only keep the last lab measurements (by `storetime`) before the ICU stay and pivot lab items to become variables/columns. Write all steps in *one* chain of pipes.

```
# Define the desired column order (ensuring subject_id and stay_id appear first)
column_order <- c("subject_id", "stay_id",
                  "bicarbonate", "chloride",
                  "creatinine", "glucose",
                  "potassium", "sodium",
                  "hematocrit", "wbc")

# Load labevents data from BigQuery and filter for relevant lab tests
labevents_tble <- tbl(con_bq, "labevents") |>

# Keep only the selected lab test item IDs
filter(itemid %in% c(50912, 50971, 50983, 50902,
                    50882, 51221, 51301, 50931)) |>

# Select relevant columns for processing
select(subject_id, storetime, itemid, valuenum) |>

# Join with ICU stay data to get stay_id and admission time
inner_join(
  select(icustays_tble, subject_id, stay_id, intime),
  by = "subject_id") |>

# Keep only lab results recorded before ICU admission time
filter(storetime < intime) |>

# Group by subject, stay, and itemid to retain the most recent measurement
group_by(subject_id, stay_id, itemid) |>
slice_max(order_by = storetime) |>
ungroup() |> # Remove grouping for subsequent operations

# Convert itemid numeric codes into meaningful lab test names
mutate(itemid = case_when(
  itemid == 51301 ~ "wbc",
  itemid == 51221 ~ "hematocrit",
  itemid == 50983 ~ "sodium",
  itemid == 50971 ~ "potassium",
  itemid == 50931 ~ "glucose",
  itemid == 50912 ~ "creatinine",
  itemid == 50902 ~ "chloride",
```

```

    itemid == 50882 ~ "bicarbonate"
  )) |>

  # Keep only necessary columns after renaming
  select(subject_id, stay_id, itemid, valuenum) |>

  # Convert from long format to wide format with mean aggregation for duplicates
  pivot_wider(names_from = itemid,
              values_from = valuenum,
              values_fn = mean) |>

  # Ensure the final column order follows the defined structure
  select(all_of(column_order)) |>

  # Arrange rows for better readability
  arrange(subject_id, stay_id) |>

  # Print the result with a wide display to prevent truncation
  print(width = Inf)

```

Warning: ORDER BY is ignored in subqueries without LIMIT
 i Do you need to move arrange() later in the pipeline or use window_order() instead?

Warning: Missing values are always removed in SQL aggregation functions.
 Use `na.rm = TRUE` to silence this warning
 This warning is displayed once every 8 hours.

Warning: ORDER BY is ignored in subqueries without LIMIT
 i Do you need to move arrange() later in the pipeline or use window_order() instead?

```

# Source:      SQL [?? x 10]
# Database:    BigQueryConnection
# Ordered by: subject_id, stay_id
  subject_id  stay_id bicarbonate chloride creatinine glucose potassium sodium
    <int>      <int>      <dbl>    <dbl>      <dbl>    <dbl>      <dbl> <dbl>
1  10000032  39553978         25        95         0.7     102         6.7  126
2  10000690  37081114         26       100          1      85         4.8  137
3  10000980  39765666         21       109         2.3     89         3.9  144
4  10001217  34592300         30       104         0.5     87         4.1  142
5  10001217  37067082         22       108         0.6    112         4.2  142
6  10001725  31205490         NA        98         NA       NA         4.1  139
7  10001843  39698942         28        97         1.3    131         3.9  138
8  10001884  37510196         30        88         1.1    141         4.5  130
9  10002013  39060235         24       102         0.9    288         3.5  137
10 10002114  34672098         18        NA         3.1     95         6.5  125
  hematocrit  wbc
    <dbl> <dbl>

```



```

1      41.1    6.9
2      36.1    7.1
3      27.3    5.3
4      37.4    5.4
5      38.1   15.7
6      NA     NA
7      31.4   10.4
8      39.7   12.2
9      34.9    7.2
10     34.3   16.8
# i more rows

```

```

# Check the number of rows
labevents_tble |>
  tally() |>
  pull(n)

```

```

Warning: ORDER BY is ignored in subqueries without LIMIT
i Do you need to move arrange() later in the pipeline or use window_order() instead?
ORDER BY is ignored in subqueries without LIMIT
i Do you need to move arrange() later in the pipeline or use window_order() instead?

[1] 88086

```

0.1.6 Q1.6 chartevents data

Connect to `chartevents` table and retrieve a subset that only contain subjects who appear in `icustays_tble` and the chart events listed in HW3. Only keep the first chart events (by `storetime`) during ICU stay and pivot chart events to become variables/columns. Write all steps in *one* chain of pipes.

```

# # TODO
# Define the desired column order (ensuring subject_id and stay_id appear first)
column_order <- c("subject_id", "stay_id",
                  "heart_rate",
                  "non-invasive_blood_pressure_systolic",
                  "non-invasive_blood_pressure_diastolic",
                  "temperature_fahrenheit",
                  "respiratory_rate")

# Load chartevents data from BigQuery and filter for relevant vital measurements
chartevents_tble <- tbl(con_bq, "chartevents") |>

# Keep only the selected vital measurements item IDs
filter(itemid %in% c(220045, 220179, 220180,
                    223761, 220210)) |>

# Select relevant columns for processing

```

```

select(subject_id, storetime, itemid, valuenum) |>

# Join with ICU stay data to get stay_id and ICU intime and outtime
inner_join(
  select(icustays_tble, subject_id,
        stay_id, intime,outtime),
  by = "subject_id") |>

# Keep only lab results recorded in ICU
filter(storetime >= intime ) |>
filter(storetime <= outtime ) |>

# Group by subject, stay, and itemid to retain the earliest measurement
group_by(subject_id, stay_id, itemid) |>
slice_min(order_by = storetime) |>
ungroup() |> # Remove grouping for subsequent operations

# Convert itemid numeric codes into meaningful lab test names
mutate(itemid = case_when(
  itemid == 220045 ~ "heart_rate",
  itemid == 220179 ~ "non-invasive_blood_pressure_systolic",
  itemid == 220180 ~ "non-invasive_blood_pressure_diastolic",
  itemid == 223761 ~ "temperature_fahrenheit",
  itemid == 220210 ~ "respiratory_rate"
)) |>

# Keep only necessary columns after renaming
select(subject_id, stay_id, itemid, valuenum) |>

# Convert from long format to wide format with mean aggregation for duplicates
pivot_wider(names_from = itemid,
            values_from = valuenum,
            values_fn = ~ round(mean(.), 1)) |>

# Ensure the final column order follows the defined structure
select(all_of(column_order)) |>

# Arrange rows for better readability
arrange(subject_id, stay_id) |>

# Print the result with a wide display to prevent truncation
print(width = Inf)

```

Warning: ORDER BY is ignored in subqueries without LIMIT

i Do you need to move arrange() later in the pipeline or use window_order() instead?

```

ORDER BY is ignored in subqueries without LIMIT
i Do you need to move arrange() later in the pipeline or use window_order() instead?

# Source:      SQL [?? x 7]
# Database:    BigQueryConnection
# Ordered by:  subject_id, stay_id
  subject_id  stay_id heart_rate `non-invasive_blood_pressure_systolic`
      <int>    <int>      <dbl>                                <dbl>
1    10000032 39553978      91                                  84
2    10000690 37081114      78                                  106
3    10000980 39765666      76                                  154
4    10001217 34592300     79.3                               156
5    10001217 37067082      86                                  151
6    10001725 31205490      86                                  73
7    10001843 39698942    124.                                110
8    10001884 37510196      49                                  174.
9    10002013 39060235      80                                  98.5
10   10002114 34672098    110.                                112
  `non-invasive_blood_pressure_diastolic` temperature_fahrenheit
                                <dbl>                                <dbl>
1                                48                                  98.7
2                                56.5                               97.7
3                                102                                 98
4                                93.3                               97.6
5                                90                                  98.5
6                                56                                  97.7
7                                78                                  97.9
8                                30.5                               98.1
9                                62                                  97.2
10                               80                                  97.9
  respiratory_rate
      <dbl>
1          24
2        24.3
3        23.5
4          14
5          18
6          19
7        16.5
8          13
9          14
10         21
# i more rows

# Check the number of rows
chartevents_tble |>

```

```
tally() |>
pull(n)
```

Warning: ORDER BY is ignored in subqueries without LIMIT
i Do you need to move arrange() later in the pipeline or use window_order() instead?
ORDER BY is ignored in subqueries without LIMIT
i Do you need to move arrange() later in the pipeline or use window_order() instead?
[1] 94364

0.1.7 Q1.7 Put things together

This step is similar to Q7 of HW3. Using *one* chain of pipes |> to perform following data wrangling steps: (i) start with the `icustays_tble`, (ii) merge in admissions and patients tables, (iii) keep adults only (age at ICU intime >= 18), (iv) merge in the `labevents` and `chartevents` tables, (v) collect the tibble, (vi) sort `subject_id`, `hadm_id`, `stay_id` and `print(width = Inf)`.

```
# # TODO
mimic_icu_cohort <- icustays_tble |>

# Merge with admissions and patients tables
left_join(select(admissions_tble, -subject_id), by = "hadm_id") |>

# Merge with patients table
left_join(patients_tble, by = "subject_id") |>

# Keep only adults (age at ICU intime >= 18)
mutate(age_intime = year(intime) - anchor_year + anchor_age) |>
filter(age_intime >= 18) |>

# Merge with labevents and chartevents tables
left_join(labevents_tble, by = c("subject_id", "stay_id")) |>
left_join(chartevents_tble, by = c("subject_id", "stay_id")) |>

# Collect data into memory
collect() |>

# Sort by subject_id, hadm_id, stay_id
arrange(subject_id, hadm_id, stay_id) |>

# Print the full width to ensure readability
print(width = Inf)
```

Warning: ORDER BY is ignored in subqueries without LIMIT
i Do you need to move arrange() later in the pipeline or use window_order() instead?
ORDER BY is ignored in subqueries without LIMIT

```

i Do you need to move arrange() later in the pipeline or use window_order() instead?
ORDER BY is ignored in subqueries without LIMIT
i Do you need to move arrange() later in the pipeline or use window_order() instead?
ORDER BY is ignored in subqueries without LIMIT
i Do you need to move arrange() later in the pipeline or use window_order() instead?
ORDER BY is ignored in subqueries without LIMIT
i Do you need to move arrange() later in the pipeline or use window_order() instead?

# A tibble: 94,458 x 41
  subject_id hadm_id stay_id first_careunit
    <int>      <int>    <int> <chr>
1  10000032 29079034 39553978 Medical Intensive Care Unit (MICU)
2  10000690 25860671 37081114 Medical Intensive Care Unit (MICU)
3  10000980 26913865 39765666 Medical Intensive Care Unit (MICU)
4  10001217 24597018 37067082 Surgical Intensive Care Unit (SICU)
5  10001217 27703517 34592300 Surgical Intensive Care Unit (SICU)
6  10001725 25563031 31205490 Medical/Surgical Intensive Care Unit (MICU/SICU)
7  10001843 26133978 39698942 Medical/Surgical Intensive Care Unit (MICU/SICU)
8  10001884 26184834 37510196 Medical Intensive Care Unit (MICU)
9  10002013 23581541 39060235 Cardiac Vascular Intensive Care Unit (CVICU)
10 10002114 27793700 34672098 Coronary Care Unit (CCU)
  last_careunit      intime
    <chr>            <dtm>
1 Medical Intensive Care Unit (MICU) 2180-07-23 14:00:00
2 Medical Intensive Care Unit (MICU) 2150-11-02 19:37:00
3 Medical Intensive Care Unit (MICU) 2189-06-27 08:42:00
4 Surgical Intensive Care Unit (SICU) 2157-11-20 19:18:02
5 Surgical Intensive Care Unit (SICU) 2157-12-19 15:42:24
6 Medical/Surgical Intensive Care Unit (MICU/SICU) 2110-04-11 15:52:22
7 Medical/Surgical Intensive Care Unit (MICU/SICU) 2134-12-05 18:50:03
8 Medical Intensive Care Unit (MICU) 2131-01-11 04:20:05
9 Cardiac Vascular Intensive Care Unit (CVICU) 2160-05-18 10:00:53
10 Coronary Care Unit (CCU) 2162-02-17 23:30:00
  outtime      los admittance      disctime
    <dtm>      <dbl> <dtm>      <dtm>
1 2180-07-23 23:50:47 0.410 2180-07-23 12:35:00 2180-07-25 17:55:00
2 2150-11-06 17:03:17 3.89 2150-11-02 18:02:00 2150-11-12 13:45:00
3 2189-06-27 20:38:27 0.498 2189-06-27 07:38:00 2189-07-03 03:00:00
4 2157-11-21 22:08:00 1.12 2157-11-18 22:56:00 2157-11-25 18:00:00
5 2157-12-20 14:27:41 0.948 2157-12-18 16:58:00 2157-12-24 14:55:00
6 2110-04-12 23:59:56 1.34 2110-04-11 15:08:00 2110-04-14 15:00:00
7 2134-12-06 14:38:26 0.825 2134-12-05 00:10:00 2134-12-06 12:54:00
8 2131-01-20 08:27:30 9.17 2131-01-07 20:39:00 2131-01-20 05:15:00
9 2160-05-19 17:33:33 1.31 2160-05-18 07:45:00 2160-05-23 13:30:00
10 2162-02-20 21:16:27 2.91 2162-02-17 22:32:00 2162-03-04 15:16:00
  deathtime      admission_type      admit_provider_id

```

	<dtm>	<chr>	<chr>		
1	NA	EW EMER.	P060TX		
2	NA	EW EMER.	P26QQ4		
3	NA	EW EMER.	P060TX		
4	NA	EW EMER.	P3610N		
5	NA	DIRECT EMER.	P2760U		
6	NA	EW EMER.	P32W56		
7	2134-12-06 12:54:00	URGENT	P67ATB		
8	2131-01-20 05:15:00	OBSERVATION ADMIT	P49AFC		
9	NA	SURGICAL SAME DAY ADMISSION	P8286C		
10	NA	OBSERVATION ADMIT	P46834		
	admission_location	discharge_location	insurance	language	marital_status
	<chr>	<chr>	<chr>	<chr>	<chr>
1	EMERGENCY ROOM	HOME	Medicaid	English	WIDOWED
2	EMERGENCY ROOM	REHAB	Medicare	English	WIDOWED
3	EMERGENCY ROOM	HOME HEALTH CARE	Medicare	English	MARRIED
4	EMERGENCY ROOM	HOME HEALTH CARE	Private	Other	MARRIED
5	PHYSICIAN REFERRAL	HOME HEALTH CARE	Private	Other	MARRIED
6	PACU	HOME	Private	English	MARRIED
7	TRANSFER FROM HOSPITAL	DIED	Medicare	English	SINGLE
8	EMERGENCY ROOM	DIED	Medicare	English	MARRIED
9	PHYSICIAN REFERRAL	HOME HEALTH CARE	Medicare	English	SINGLE
10	PHYSICIAN REFERRAL	HOME HEALTH CARE	Medicaid	English	<NA>
	race	edregtime	edouttime		
	<chr>	<dtm>	<dtm>		
1	WHITE	2180-07-23 05:54:00	2180-07-23 14:00:00		
2	WHITE	2150-11-02 11:41:00	2150-11-02 19:37:00		
3	BLACK/AFRICAN AMERICAN	2189-06-27 06:25:00	2189-06-27 08:42:00		
4	WHITE	2157-11-18 17:38:00	2157-11-19 01:24:00		
5	WHITE	NA	NA		
6	WHITE	NA	NA		
7	WHITE	NA	NA		
8	BLACK/AFRICAN AMERICAN	2131-01-07 13:36:00	2131-01-07 22:13:00		
9	OTHER	NA	NA		
10	UNKNOWN	2162-02-17 19:35:00	2162-02-17 23:30:00		
	hospital_expire_flag	gender	anchor_age	anchor_year	anchor_year_group
	<int>	<chr>	<int>	<int>	<chr>
1	0	F	52	2180	2014 - 2016
2	0	F	86	2150	2008 - 2010
3	0	F	73	2186	2008 - 2010
4	0	F	55	2157	2011 - 2013
5	0	F	55	2157	2011 - 2013
6	0	F	46	2110	2011 - 2013
7	1	M	73	2131	2017 - 2019
8	1	F	68	2122	2008 - 2010
9	0	F	53	2156	2008 - 2010

```

10          0 M          56          2162 2020 - 2022
      dod      age_intime bicarbonate chloride creatinine glucose potassium
      <date>      <int>      <dbl>      <dbl>      <dbl>      <dbl>      <dbl>
1 2180-09-09      52          25          95          0.7          102          6.7
2 2152-01-30      86          26          100          1            85          4.8
3 2193-08-26      76          21          109          2.3          89          3.9
4 NA              55          22          108          0.6          112          4.2
5 NA              55          30          104          0.5          87          4.1
6 NA              46          NA          98          NA            NA          4.1
7 2134-12-06      76          28          97          1.3          131          3.9
8 2131-01-20      77          30          88          1.1          141          4.5
9 NA              57          24          102          0.9          288          3.5
10 2162-12-11      56          18          NA          3.1          95          6.5
      sodium hematocrit   wbc heart_rate `non-invasive_blood_pressure_systolic`
      <dbl>      <dbl> <dbl>      <dbl>      <dbl>
1 126          41.1  6.9          91          84
2 137          36.1  7.1          78          106
3 144          27.3  5.3          76          154
4 142          38.1  15.7         86          151
5 142          37.4  5.4         79.3         156
6 139          NA    NA          86          73
7 138          31.4  10.4        124.         110
8 130          39.7  12.2         49          174.
9 137          34.9  7.2          80          98.5
10 125          34.3  16.8        110.         112
      `non-invasive_blood_pressure_diastolic` temperature_fahrenheit
      <dbl>      <dbl>
1 48          98.7
2 56.5         97.7
3 102          98
4 90          98.5
5 93.3         97.6
6 56          97.7
7 78          97.9
8 30.5         98.1
9 62          97.2
10 80          97.9
      respiratory_rate
      <dbl>
1 24
2 24.3
3 23.5
4 18
5 14
6 19
7 16.5

```

```

8           13
9           14
10          21
# i 94,448 more rows

```

0.1.8 Q1.8 Preprocessing

Perform the following preprocessing steps. (i) Lump infrequent levels into “Other” level for `first_careunit`, `last_careunit`, `admission_type`, `admission_location`, and `discharge_location`. (ii) Collapse the levels of `race` into `ASIAN`, `BLACK`, `HISPANIC`, `WHITE`, and `Other`. (iii) Create a new variable `los_long` that is `TRUE` when `los` is greater than or equal to 2 days. (iv) Summarize the data using `tbl_summary()`, stratified by `los_long`. Hint: `fct_lump_n` and `fct_collapse` from the `forcats` package are useful.

Hint: Below is a numerical summary of my tibble after preprocessing:

```

# Process mimic_icu_cohort data
mimic_icu_cohort <- mimic_icu_cohort |>
  mutate(
    first_careunit = fct_lump(first_careunit, n = 4),
    last_careunit = fct_lump(last_careunit, n = 4),
    admission_type = fct_lump(admission_type, n = 4),
    admission_location = fct_lump(admission_location, n = 3),
    discharge_location = fct_lump(discharge_location, n = 4)
  ) |>

# Sort by subject_id, hadm_id, stay_id
arrange(subject_id, hadm_id, stay_id) |>

# Print the full width to ensure readability
print(width = Inf)

```

```

# A tibble: 94,458 x 41
  subject_id hadm_id stay_id first_careunit
    <int>    <int>   <int> <fct>
1  10000032 29079034 39553978 Medical Intensive Care Unit (MICU)
2  10000690 25860671 37081114 Medical Intensive Care Unit (MICU)
3  10000980 26913865 39765666 Medical Intensive Care Unit (MICU)
4  10001217 24597018 37067082 Surgical Intensive Care Unit (SICU)
5  10001217 27703517 34592300 Surgical Intensive Care Unit (SICU)
6  10001725 25563031 31205490 Medical/Surgical Intensive Care Unit (MICU/SICU)
7  10001843 26133978 39698942 Medical/Surgical Intensive Care Unit (MICU/SICU)
8  10001884 26184834 37510196 Medical Intensive Care Unit (MICU)
9  10002013 23581541 39060235 Cardiac Vascular Intensive Care Unit (CVICU)
10 10002114 27793700 34672098 Other

```


last_careunit			intime
<fct>			<dtm>
1 Medical Intensive Care Unit (MICU)			2180-07-23 14:00:00
2 Medical Intensive Care Unit (MICU)			2150-11-02 19:37:00
3 Medical Intensive Care Unit (MICU)			2189-06-27 08:42:00
4 Surgical Intensive Care Unit (SICU)			2157-11-20 19:18:02
5 Surgical Intensive Care Unit (SICU)			2157-12-19 15:42:24
6 Medical/Surgical Intensive Care Unit (MICU/SICU)			2110-04-11 15:52:22
7 Medical/Surgical Intensive Care Unit (MICU/SICU)			2134-12-05 18:50:03
8 Medical Intensive Care Unit (MICU)			2131-01-11 04:20:05
9 Cardiac Vascular Intensive Care Unit (CVICU)			2160-05-18 10:00:53
10 Other			2162-02-17 23:30:00

outtime	los	admittime	dischtime
<dtm>	<dbl>	<dtm>	<dtm>
1 2180-07-23 23:50:47	0.410	2180-07-23 12:35:00	2180-07-25 17:55:00
2 2150-11-06 17:03:17	3.89	2150-11-02 18:02:00	2150-11-12 13:45:00
3 2189-06-27 20:38:27	0.498	2189-06-27 07:38:00	2189-07-03 03:00:00
4 2157-11-21 22:08:00	1.12	2157-11-18 22:56:00	2157-11-25 18:00:00
5 2157-12-20 14:27:41	0.948	2157-12-18 16:58:00	2157-12-24 14:55:00
6 2110-04-12 23:59:56	1.34	2110-04-11 15:08:00	2110-04-14 15:00:00
7 2134-12-06 14:38:26	0.825	2134-12-05 00:10:00	2134-12-06 12:54:00
8 2131-01-20 08:27:30	9.17	2131-01-07 20:39:00	2131-01-20 05:15:00
9 2160-05-19 17:33:33	1.31	2160-05-18 07:45:00	2160-05-23 13:30:00
10 2162-02-20 21:16:27	2.91	2162-02-17 22:32:00	2162-03-04 15:16:00

deathtime	admission_type	admit_provider_id
<dtm>	<fct>	<chr>
1 NA	EW EMER.	P060TX
2 NA	EW EMER.	P26QQ4
3 NA	EW EMER.	P060TX
4 NA	EW EMER.	P3610N
5 NA	Other	P276OU
6 NA	EW EMER.	P32W56
7 2134-12-06 12:54:00	URGENT	P67ATB
8 2131-01-20 05:15:00	OBSERVATION ADMIT	P49AFC
9 NA	SURGICAL SAME DAY ADMISSION	P8286C
10 NA	OBSERVATION ADMIT	P46834

admission_location	discharge_location	insurance	language	marital_status
<fct>	<fct>	<chr>	<chr>	<chr>
1 EMERGENCY ROOM	HOME	Medicaid	English	WIDOWED
2 EMERGENCY ROOM	Other	Medicare	English	WIDOWED
3 EMERGENCY ROOM	HOME HEALTH CARE	Medicare	English	MARRIED
4 EMERGENCY ROOM	HOME HEALTH CARE	Private	Other	MARRIED
5 PHYSICIAN REFERRAL	HOME HEALTH CARE	Private	Other	MARRIED
6 Other	HOME	Private	English	MARRIED
7 TRANSFER FROM HOSPITAL	DIED	Medicare	English	SINGLE
8 EMERGENCY ROOM	DIED	Medicare	English	MARRIED

9	PHYSICIAN REFERRAL	HOME HEALTH CARE	Medicare	English	SINGLE		
10	PHYSICIAN REFERRAL	HOME HEALTH CARE	Medicaid	English	<NA>		
	race	edregtime	edouttime				
	<chr>	<dtm>	<dtm>				
1	WHITE	2180-07-23 05:54:00	2180-07-23 14:00:00				
2	WHITE	2150-11-02 11:41:00	2150-11-02 19:37:00				
3	BLACK/AFRICAN AMERICAN	2189-06-27 06:25:00	2189-06-27 08:42:00				
4	WHITE	2157-11-18 17:38:00	2157-11-19 01:24:00				
5	WHITE	NA	NA				
6	WHITE	NA	NA				
7	WHITE	NA	NA				
8	BLACK/AFRICAN AMERICAN	2131-01-07 13:36:00	2131-01-07 22:13:00				
9	OTHER	NA	NA				
10	UNKNOWN	2162-02-17 19:35:00	2162-02-17 23:30:00				
	hospital_expire_flag	gender	anchor_age	anchor_year	anchor_year_group		
	<int>	<chr>	<int>	<int>	<chr>		
1	0	F	52	2180	2014 - 2016		
2	0	F	86	2150	2008 - 2010		
3	0	F	73	2186	2008 - 2010		
4	0	F	55	2157	2011 - 2013		
5	0	F	55	2157	2011 - 2013		
6	0	F	46	2110	2011 - 2013		
7	1	M	73	2131	2017 - 2019		
8	1	F	68	2122	2008 - 2010		
9	0	F	53	2156	2008 - 2010		
10	0	M	56	2162	2020 - 2022		
	dod	age_intime	bicarbonate	chloride	creatinine	glucose	potassium
	<date>	<int>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>
1	2180-09-09	52	25	95	0.7	102	6.7
2	2152-01-30	86	26	100	1	85	4.8
3	2193-08-26	76	21	109	2.3	89	3.9
4	NA	55	22	108	0.6	112	4.2
5	NA	55	30	104	0.5	87	4.1
6	NA	46	NA	98	NA	NA	4.1
7	2134-12-06	76	28	97	1.3	131	3.9
8	2131-01-20	77	30	88	1.1	141	4.5
9	NA	57	24	102	0.9	288	3.5
10	2162-12-11	56	18	NA	3.1	95	6.5
	sodium	hematocrit	wbc	heart_rate	non-invasive_blood_pressure_systolic`		
	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>		
1	126	41.1	6.9	91	84		
2	137	36.1	7.1	78	106		
3	144	27.3	5.3	76	154		
4	142	38.1	15.7	86	151		
5	142	37.4	5.4	79.3	156		
6	139	NA	NA	86	73		

```

7    138      31.4  10.4    124.      110
8    130      39.7  12.2     49      174.
9    137      34.9   7.2     80      98.5
10   125      34.3  16.8    110.      112

```

```

`non-invasive_blood_pressure_diastolic` temperature_fahrenheit
<dbl> <dbl>
1      48      98.7
2     56.5     97.7
3     102      98
4      90     98.5
5     93.3     97.6
6      56     97.7
7      78     97.9
8     30.5     98.1
9      62     97.2
10     80     97.9

```

```

respiratory_rate
<dbl>
1      24
2     24.3
3     23.5
4      18
5      14
6      19
7     16.5
8      13
9      14
10     21

```

```
# i 94,448 more rows
```

```
# Collapse race categories into ASIAN, BLACK, HISPANIC, WHITE, and Other
```

```
mimic_icu_cohort <- mimic_icu_cohort |>
```

```

mutate(
  race = case_when(
    str_detect(race, regex("^ASIAN", ignore_case = TRUE)) ~ "ASIAN",
    str_detect(race, regex("^BLACK", ignore_case = TRUE)) ~ "BLACK",
    str_detect(race, regex("HISPANIC|LATINO", ignore_case = TRUE)) ~ "HISPANIC",
    str_detect(race, regex("^WHITE", ignore_case = TRUE)) ~ "WHITE",
    TRUE ~ "Other" # Assign everything else to "Other"
  )
)|>

```

```

# Print the full width to ensure readability
print(width = Inf)

```

```
# A tibble: 94,458 x 41
```

```
subject_id hadm_id stay_id first_careunit
```

	<int>	<int>	<int>	<fct>
1	10000032	29079034	39553978	Medical Intensive Care Unit (MICU)
2	10000690	25860671	37081114	Medical Intensive Care Unit (MICU)
3	10000980	26913865	39765666	Medical Intensive Care Unit (MICU)
4	10001217	24597018	37067082	Surgical Intensive Care Unit (SICU)
5	10001217	27703517	34592300	Surgical Intensive Care Unit (SICU)
6	10001725	25563031	31205490	Medical/Surgical Intensive Care Unit (MICU/SICU)
7	10001843	26133978	39698942	Medical/Surgical Intensive Care Unit (MICU/SICU)
8	10001884	26184834	37510196	Medical Intensive Care Unit (MICU)
9	10002013	23581541	39060235	Cardiac Vascular Intensive Care Unit (CVICU)
10	10002114	27793700	34672098	Other

	last_careunit	intime
	<fct>	<dtm>
1	Medical Intensive Care Unit (MICU)	2180-07-23 14:00:00
2	Medical Intensive Care Unit (MICU)	2150-11-02 19:37:00
3	Medical Intensive Care Unit (MICU)	2189-06-27 08:42:00
4	Surgical Intensive Care Unit (SICU)	2157-11-20 19:18:02
5	Surgical Intensive Care Unit (SICU)	2157-12-19 15:42:24
6	Medical/Surgical Intensive Care Unit (MICU/SICU)	2110-04-11 15:52:22
7	Medical/Surgical Intensive Care Unit (MICU/SICU)	2134-12-05 18:50:03
8	Medical Intensive Care Unit (MICU)	2131-01-11 04:20:05
9	Cardiac Vascular Intensive Care Unit (CVICU)	2160-05-18 10:00:53
10	Other	2162-02-17 23:30:00

	outtime	los	admittime	dischtime
	<dtm>	<dbl>	<dtm>	<dtm>
1	2180-07-23 23:50:47	0.410	2180-07-23 12:35:00	2180-07-25 17:55:00
2	2150-11-06 17:03:17	3.89	2150-11-02 18:02:00	2150-11-12 13:45:00
3	2189-06-27 20:38:27	0.498	2189-06-27 07:38:00	2189-07-03 03:00:00
4	2157-11-21 22:08:00	1.12	2157-11-18 22:56:00	2157-11-25 18:00:00
5	2157-12-20 14:27:41	0.948	2157-12-18 16:58:00	2157-12-24 14:55:00
6	2110-04-12 23:59:56	1.34	2110-04-11 15:08:00	2110-04-14 15:00:00
7	2134-12-06 14:38:26	0.825	2134-12-05 00:10:00	2134-12-06 12:54:00
8	2131-01-20 08:27:30	9.17	2131-01-07 20:39:00	2131-01-20 05:15:00
9	2160-05-19 17:33:33	1.31	2160-05-18 07:45:00	2160-05-23 13:30:00
10	2162-02-20 21:16:27	2.91	2162-02-17 22:32:00	2162-03-04 15:16:00

	deathtime	admission_type	admit_provider_id
	<dtm>	<fct>	<chr>
1	NA	EW EMER.	P060TX
2	NA	EW EMER.	P26QQ4
3	NA	EW EMER.	P060TX
4	NA	EW EMER.	P3610N
5	NA	Other	P276OU
6	NA	EW EMER.	P32W56
7	2134-12-06 12:54:00	URGENT	P67ATB
8	2131-01-20 05:15:00	OBSERVATION ADMIT	P49AFC
9	NA	SURGICAL SAME DAY ADMISSION	P8286C

10	NA	OBSERVATION ADMIT		P46834			
	admission_location	discharge_location	insurance	language	marital_status		
	<fct>	<fct>	<chr>	<chr>	<chr>		
1	EMERGENCY ROOM	HOME	Medicaid	English	WIDOWED		
2	EMERGENCY ROOM	Other	Medicare	English	WIDOWED		
3	EMERGENCY ROOM	HOME HEALTH CARE	Medicare	English	MARRIED		
4	EMERGENCY ROOM	HOME HEALTH CARE	Private	Other	MARRIED		
5	PHYSICIAN REFERRAL	HOME HEALTH CARE	Private	Other	MARRIED		
6	Other	HOME	Private	English	MARRIED		
7	TRANSFER FROM HOSPITAL	DIED	Medicare	English	SINGLE		
8	EMERGENCY ROOM	DIED	Medicare	English	MARRIED		
9	PHYSICIAN REFERRAL	HOME HEALTH CARE	Medicare	English	SINGLE		
10	PHYSICIAN REFERRAL	HOME HEALTH CARE	Medicaid	English	<NA>		
	race	edregtime	edouttime	hospital_expire_flag	gender		
	<chr>	<dtm>	<dtm>	<int>	<chr>		
1	WHITE	2180-07-23 05:54:00	2180-07-23 14:00:00		0 F		
2	WHITE	2150-11-02 11:41:00	2150-11-02 19:37:00		0 F		
3	BLACK	2189-06-27 06:25:00	2189-06-27 08:42:00		0 F		
4	WHITE	2157-11-18 17:38:00	2157-11-19 01:24:00		0 F		
5	WHITE	NA			0 F		
6	WHITE	NA			0 F		
7	WHITE	NA			1 M		
8	BLACK	2131-01-07 13:36:00	2131-01-07 22:13:00		1 F		
9	Other	NA			0 F		
10	Other	2162-02-17 19:35:00	2162-02-17 23:30:00		0 M		
	anchor_age	anchor_year	anchor_year_group	dod	age_intime	bicarbonate	
	<int>	<int>	<chr>	<date>	<int>	<dbl>	
1	52	2180	2014 - 2016	2180-09-09	52	25	
2	86	2150	2008 - 2010	2152-01-30	86	26	
3	73	2186	2008 - 2010	2193-08-26	76	21	
4	55	2157	2011 - 2013	NA	55	22	
5	55	2157	2011 - 2013	NA	55	30	
6	46	2110	2011 - 2013	NA	46	NA	
7	73	2131	2017 - 2019	2134-12-06	76	28	
8	68	2122	2008 - 2010	2131-01-20	77	30	
9	53	2156	2008 - 2010	NA	57	24	
10	56	2162	2020 - 2022	2162-12-11	56	18	
	chloride	creatinine	glucose	potassium	sodium	hematocrit	wbc heart_rate
	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>
1	95	0.7	102	6.7	126	41.1	6.9 91
2	100	1	85	4.8	137	36.1	7.1 78
3	109	2.3	89	3.9	144	27.3	5.3 76
4	108	0.6	112	4.2	142	38.1	15.7 86
5	104	0.5	87	4.1	142	37.4	5.4 79.3
6	98	NA	NA	4.1	139	NA	NA 86
7	97	1.3	131	3.9	138	31.4	10.4 124.

8	88	1.1	141	4.5	130	39.7	12.2	49
9	102	0.9	288	3.5	137	34.9	7.2	80
10	NA	3.1	95	6.5	125	34.3	16.8	110.

```

`non-invasive_blood_pressure_systolic`
<dbl>
1      84
2     106
3     154
4     151
5     156
6      73
7     110
8     174.
9     98.5
10    112
`non-invasive_blood_pressure_diastolic` temperature_fahrenheit
<dbl> <dbl>
1      48      98.7
2     56.5     97.7
3     102      98
4      90     98.5
5     93.3     97.6
6      56     97.7
7      78     97.9
8     30.5     98.1
9      62     97.2
10     80     97.9

```

```

respiratory_rate
<dbl>
1      24
2     24.3
3     23.5
4      18
5      14
6      19
7     16.5
8      13
9      14
10     21

```

i 94,448 more rows

```

# Step (iii): Create los_long variable
mimic_icu_cohort <- mimic_icu_cohort |>
  mutate(los_long = los >= 2) |>
  #collect() />
  print(width = Inf)

```

A tibble: 94,458 x 42

	subject_id	hadm_id	stay_id	first_careunit
	<int>	<int>	<int>	<fct>
1	10000032	29079034	39553978	Medical Intensive Care Unit (MICU)
2	10000690	25860671	37081114	Medical Intensive Care Unit (MICU)
3	10000980	26913865	39765666	Medical Intensive Care Unit (MICU)
4	10001217	24597018	37067082	Surgical Intensive Care Unit (SICU)
5	10001217	27703517	34592300	Surgical Intensive Care Unit (SICU)
6	10001725	25563031	31205490	Medical/Surgical Intensive Care Unit (MICU/SICU)
7	10001843	26133978	39698942	Medical/Surgical Intensive Care Unit (MICU/SICU)
8	10001884	26184834	37510196	Medical Intensive Care Unit (MICU)
9	10002013	23581541	39060235	Cardiac Vascular Intensive Care Unit (CVICU)
10	10002114	27793700	34672098	Other

	last_careunit	intime
	<fct>	<dtm>
1	Medical Intensive Care Unit (MICU)	2180-07-23 14:00:00
2	Medical Intensive Care Unit (MICU)	2150-11-02 19:37:00
3	Medical Intensive Care Unit (MICU)	2189-06-27 08:42:00
4	Surgical Intensive Care Unit (SICU)	2157-11-20 19:18:02
5	Surgical Intensive Care Unit (SICU)	2157-12-19 15:42:24
6	Medical/Surgical Intensive Care Unit (MICU/SICU)	2110-04-11 15:52:22
7	Medical/Surgical Intensive Care Unit (MICU/SICU)	2134-12-05 18:50:03
8	Medical Intensive Care Unit (MICU)	2131-01-11 04:20:05
9	Cardiac Vascular Intensive Care Unit (CVICU)	2160-05-18 10:00:53
10	Other	2162-02-17 23:30:00

	outtime	los	admittime	dischtime
	<dtm>	<dbl>	<dtm>	<dtm>
1	2180-07-23 23:50:47	0.410	2180-07-23 12:35:00	2180-07-25 17:55:00
2	2150-11-06 17:03:17	3.89	2150-11-02 18:02:00	2150-11-12 13:45:00
3	2189-06-27 20:38:27	0.498	2189-06-27 07:38:00	2189-07-03 03:00:00
4	2157-11-21 22:08:00	1.12	2157-11-18 22:56:00	2157-11-25 18:00:00
5	2157-12-20 14:27:41	0.948	2157-12-18 16:58:00	2157-12-24 14:55:00
6	2110-04-12 23:59:56	1.34	2110-04-11 15:08:00	2110-04-14 15:00:00
7	2134-12-06 14:38:26	0.825	2134-12-05 00:10:00	2134-12-06 12:54:00
8	2131-01-20 08:27:30	9.17	2131-01-07 20:39:00	2131-01-20 05:15:00
9	2160-05-19 17:33:33	1.31	2160-05-18 07:45:00	2160-05-23 13:30:00
10	2162-02-20 21:16:27	2.91	2162-02-17 22:32:00	2162-03-04 15:16:00

	deathtime	admission_type	admit_provider_id
	<dtm>	<fct>	<chr>
1	NA	EW EMER.	P060TX
2	NA	EW EMER.	P26QQ4
3	NA	EW EMER.	P060TX
4	NA	EW EMER.	P3610N
5	NA	Other	P276OU
6	NA	EW EMER.	P32W56
7	2134-12-06 12:54:00	URGENT	P67ATB

	admission_location	discharge_location	insurance	language	marital_status
	<fct>	<fct>	<chr>	<chr>	<chr>
1	EMERGENCY ROOM	HOME	Medicaid	English	WIDOWED
2	EMERGENCY ROOM	Other	Medicare	English	WIDOWED
3	EMERGENCY ROOM	HOME HEALTH CARE	Medicare	English	MARRIED
4	EMERGENCY ROOM	HOME HEALTH CARE	Private	Other	MARRIED
5	PHYSICIAN REFERRAL	HOME HEALTH CARE	Private	Other	MARRIED
6	Other	HOME	Private	English	MARRIED
7	TRANSFER FROM HOSPITAL	DIED	Medicare	English	SINGLE
8	EMERGENCY ROOM	DIED	Medicare	English	MARRIED
9	PHYSICIAN REFERRAL	HOME HEALTH CARE	Medicare	English	SINGLE
10	PHYSICIAN REFERRAL	HOME HEALTH CARE	Medicaid	English	<NA>

	race	edregtime	edouttime	hospital_expire_flag	gender
	<chr>	<dtm>	<dtm>	<int>	<chr>
1	WHITE	2180-07-23 05:54:00	2180-07-23 14:00:00		0 F
2	WHITE	2150-11-02 11:41:00	2150-11-02 19:37:00		0 F
3	BLACK	2189-06-27 06:25:00	2189-06-27 08:42:00		0 F
4	WHITE	2157-11-18 17:38:00	2157-11-19 01:24:00		0 F
5	WHITE	NA	NA		0 F
6	WHITE	NA	NA		0 F
7	WHITE	NA	NA		1 M
8	BLACK	2131-01-07 13:36:00	2131-01-07 22:13:00		1 F
9	Other	NA	NA		0 F
10	Other	2162-02-17 19:35:00	2162-02-17 23:30:00		0 M

	anchor_age	anchor_year	anchor_year_group	dod	age_intime	bicarbonate
	<int>	<int>	<chr>	<date>	<int>	<dbl>
1	52	2180	2014 - 2016	2180-09-09	52	25
2	86	2150	2008 - 2010	2152-01-30	86	26
3	73	2186	2008 - 2010	2193-08-26	76	21
4	55	2157	2011 - 2013	NA	55	22
5	55	2157	2011 - 2013	NA	55	30
6	46	2110	2011 - 2013	NA	46	NA
7	73	2131	2017 - 2019	2134-12-06	76	28
8	68	2122	2008 - 2010	2131-01-20	77	30
9	53	2156	2008 - 2010	NA	57	24
10	56	2162	2020 - 2022	2162-12-11	56	18

	chloride	creatinine	glucose	potassium	sodium	hematocrit	wbc	heart_rate
	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>
1	95	0.7	102	6.7	126	41.1	6.9	91
2	100	1	85	4.8	137	36.1	7.1	78
3	109	2.3	89	3.9	144	27.3	5.3	76
4	108	0.6	112	4.2	142	38.1	15.7	86
5	104	0.5	87	4.1	142	37.4	5.4	79.3

6	98	NA	NA	4.1	139	NA	NA	86
7	97	1.3	131	3.9	138	31.4	10.4	124.
8	88	1.1	141	4.5	130	39.7	12.2	49
9	102	0.9	288	3.5	137	34.9	7.2	80
10	NA	3.1	95	6.5	125	34.3	16.8	110.

```

`non-invasive_blood_pressure_systolic`
<dbl>
1      84
2     106
3     154
4     151
5     156
6      73
7     110
8     174.
9     98.5
10    112
`non-invasive_blood_pressure_diastolic` temperature_fahrenheit
<dbl>
1      48
2     56.5
3     102
4      90
5     93.3
6      56
7      78
8     30.5
9      62
10     80

```

```

respiratory_rate los_long
<dbl> <lgl>
1      24 FALSE
2     24.3 TRUE
3     23.5 FALSE
4      18 FALSE
5      14 FALSE
6      19 FALSE
7     16.5 FALSE
8      13 TRUE
9      14 FALSE
10     21 TRUE

```

```

# i 94,448 more rows

```

```

mimic_icu_cohort <- mimic_icu_cohort |>
  select(-c("subject_id", "hadm_id", "stay_id",
            "intime", "outtime", "admittime", "dischtime",

```

```

      "deathtime", "admit_provider_id", "edregtime",
      "edouttime", "anchor_age", "anchor_year",
      "anchor_year_group")) |>
tbl_summary(by = "los_long")

```

14 missing rows in the "los_long" column have been removed.
 The following errors were returned during `tbl_summary()`:
 x For variable `dod` (`los_long = FALSE`) and "p75" statistic: * not defined
 for "Date" objects

```

# Now print the summary table output
mimic_icu_cohort

```

0.1.9 Q1.9 Save the final tibble

Save the final tibble to an R data file `mimic_icu_cohort.rds` in the `mimiciv_shiny` folder.

```

# make a directory mimiciv_shiny
if (!dir.exists("mimiciv_shiny")) {
  dir.create("mimiciv_shiny")
}
# save the final tibble
mimic_icu_cohort |>
  write_rds("mimiciv_shiny/mimic_icu_cohort.rds", compress = "gz")

```

Close database connection and clear workspace.

```

if (exists("con_bq")) {
  dbDisconnect(con_bq)
}
rm(list = ls())

```

Although it is not a good practice to add big data files to Git, for grading purpose, please add `mimic_icu_cohort.rds` to your Git repository.

0.2 Q2. Shiny app

Develop a Shiny app for exploring the ICU cohort data created in Q1. The app should reside in the `mimiciv_shiny` folder. The app should contain at least two tabs. One tab provides easy access to the graphical and numerical summaries of variables (demographics, lab measurements, vitals) in the ICU cohort, using the `mimic_icu_cohort.rds` you curated in Q1. The other tab allows user to choose a specific patient in the cohort and display the patient's ADT and ICU stay information as we did in Q1 of HW3, by dynamically retrieving the patient's ADT and ICU stay information from BigQuery database. Again, do **not** ever add the BigQuery token to your Git repository. If you do so, you will lose 50 points.

Characteristic		TRUE N = 46,337 [†]
first_careunit		
	Cardiac Vascular Intensive Care Unit (CVICU)	7,353 (16%)
	Medical Intensive Care Unit (MICU)	9,837 (21%)
	Medical/Surgical Intensive Care Unit (MICU/SICU)	6,667 (14%)
	Surgical Intensive Care Unit (SICU)	6,434 (14%)
	Other	16,046 (35%)
last_careunit		
	Cardiac Vascular Intensive Care Unit (CVICU)	7,353 (16%)
	Medical Intensive Care Unit (MICU)	9,837 (21%)
	Medical/Surgical Intensive Care Unit (MICU/SICU)	6,667 (14%)
	Surgical Intensive Care Unit (SICU)	6,434 (14%)
	Other	16,046 (35%)
los		3.9 (2.7, 6.8)
admission_type		
	EW EMER.	23,012 (50%)
	OBSERVATION ADMIT	7,393 (16%)
	SURGICAL SAME DAY ADMISSION	4,001 (8.6%)
	URGENT	8,691 (19%)
	Other	3,240 (7.0%)
admission_location		
	EMERGENCY ROOM	17,058 (37%)
	PHYSICIAN REFERRAL	11,013 (24%)
	TRANSFER FROM HOSPITAL	13,904 (30%)
	Other	4,362 (9.4%)
discharge_location		
	DIED	6,884 (15%)
	HOME	6,879 (15%)
	HOME HEALTH CARE	10,620 (23%)
	SKILLED NURSING FACILITY	8,785 (19%)
	Other	13,092 (28%)
	Unknown	77
insurance		
	Medicaid	6,768 (15%)
	Medicare	26,330 (58%)
	No charge	5 (<0.1%)
	Other	1,091 (2.4%)
	Private	11,515 (25%)
	Unknown	628
language	27	
	American Sign Language	29 (<0.1%)
	Amharic	14 (<0.1%)
	Arabic	87 (0.2%)
	Armenian	12 (<0.1%)
	Bengali	22 (<0.1%)
	Chinese	550 (1.2%)