

# Data Preprocessing EDA and Feature Engineering

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## Section 1: Data Understanding

### Objectives:

- Understand the structure and purpose of key MIMIC-III tables.
- Identify relationships between tables to facilitate feature extraction.
- Review variable definitions and units of measurement.

### Steps:

#### 1.1 Overview of all Tables:

- Focus on the following pre-processed tables in 'mimic\_data' folder: - `admissions.csv` : Admission details, including admission and discharge times. - `antibiotics.csv` : Antibiotic usage data. - `bloodculture.csv` : Results of blood culture tests. - `gcs_hourly.csv` : Glasgow Coma Score records. - `icd9_diag.csv` : ICD-9 diagnostic codes for patient conditions. - `icustays.csv` : ICU stay details (e.g., admission, discharge times). - `labs_hourly.csv` : Hourly laboratory results. - `output_hourly.csv` : Fluid output data. - `patients.csv` : Demographics and mortality information. - `pt_icu_outcome.csv` : Patient outcomes (e.g., mortality) per ICU stay. - `pt_stay_hr.csv` : Hourly records of ICU stays. - `pt_weight.csv` : Patient weight records. - `pv_mechvent.csv` : Mechanical ventilation data. - `transfers.csv` : Information on patient transfers within the hospital. - `vasopressors.csv` : Administration of vasopressors. - `vitals_hourly.csv` : Hourly vital sign measurements.
- Use `data.table` for efficient loading of large datasets.

#### 1.2 Relationships Between Tables:

- Key relationships include:
  - `subject_id` : Links `patients`, `admissions`, and `icustays`.
  - `hadm_id` : Links `admissions` and `icustays`.
  - `icustay_id` : Links ICU-specific data (e.g., `vitals_hourly`, `labs_hourly`, etc.).
  - Other tables (e.g., `antibiotics`, `bloodculture`) use these IDs to connect to patient-specific data.

#### 1.3 Initial Summarisation:

- Explore each table:
  - Number of rows and columns.
  - Key variables and their data type.
  - Missing data percentages.

```
##
## ### Summary of Table: admissions ###
## Number of rows: 58976
## Number of columns: 19
## Column names and data types:
##          Data_Type Missing_Count Missing_Pct.
## row_id          integer           0         0.00
## subject_id       integer           0         0.00
## hadm_id          integer           0         0.00
## admittime        POSIXct           0         0.00
## dischtime        POSIXct           0         0.00
## deathtime        POSIXct        53122        90.07
## admission_type   character          0         0.00
## admission_location character          0         0.00
## discharge_location character          0         0.00
## insurance        character          0         0.00
## language         character          0         0.00
## religion         character          0         0.00
## marital_status   character          0         0.00
## ethnicity        character          0         0.00
## edregtime        POSIXct        28099        47.64
## edouttime        POSIXct        28099        47.64
## diagnosis        character          0         0.00
## hospital_expire_flag integer          0         0.00
## has_chartevents_data integer          0         0.00
```

```
##
## ---
```

```
## ### Summary of Table: antibiotics ###
## Number of rows: 164927
## Number of columns: 16
## Column names and data types:
##          Data_Type Missing_Count Missing_Pct.
## icustay_id      integer          74         0.04
## starttime       POSIXct          24         0.01
## endtime         POSIXct           7         0.00
## amount          numeric           7         0.00
## amountuom       character          0         0.00
## rate            logical        164927       100.00
## rateuom         logical        164927       100.00
## ordercategoryname character          0         0.00
## patientweight   numeric          31         0.02
## totalamount     integer        8315         5.04
## totalamountuom  character          0         0.00
## statusdescription character          0         0.00
## label           character          0         0.00
## abbreviation    character          0         0.00
## antibiotic      integer           0         0.00
## dbsource        character          0         0.00
```

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##
## ---
##
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```
## ### Summary of Table: bloodculture ###
## Number of rows: 632506
## Number of columns: 10
```

```

## Column names and data types:
##           Data_Type Missing_Count Missing_Pct.
## hadm_id      integer           0         0.00
## icustay_id    integer       156798        24.79
## dy            integer       156798        24.79
## hr            integer       187726        29.68
## charttime     POSIXct         41791         6.61
## chartdate     POSIXct          0         0.00
## org_name      character        0         0.00
## positiveculture integer        0         0.00
## ab_name       character        0         0.00
## antibioticresistance character  0         0.00
##
## ---
##
## ### Summary of Table: gcs_hourly ###
## Number of rows: 1515342
## Number of columns: 7
## Column names and data types:
##           Data_Type Missing_Count Missing_Pct.
## icustay_id    integer           0         0.00
## hr            integer           0         0.00
## gcs           integer           0         0.00
## gcseyes       integer       1444         0.10
## gcsmotor      integer       3768         0.25
## gcsverbal     integer       3568         0.24
## endotrachflag integer           0         0.00
##
## ---
##
## ### Summary of Table: icd9_diag ###
## Number of rows: 651047
## Number of columns: 7
## Column names and data types:
##           Data_Type Missing_Count Missing_Pct.
## row_id        integer           0         0.00
## subject_id     integer           0         0.00
## hadm_id        integer           0         0.00
## seq_num        integer         47         0.01
## icd9_code      character        0         0.00
## short_title    character        0         0.00
## long_title     character        0         0.00
##
## ---
##
## ### Summary of Table: icustays ###
## Number of rows: 61532
## Number of columns: 12
## Column names and data types:
##           Data_Type Missing_Count Missing_Pct.
## row_id        integer           0         0.00
## subject_id     integer           0         0.00
## hadm_id        integer           0         0.00
## icustay_id     integer           0         0.00
## dbsource       character        0         0.00
## first_careunit character        0         0.00

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```

## last_careunit character      0      0.00
## first_wardid integer        0      0.00
## last_wardid integer        0      0.00
## intime POSIXct             0      0.00
## outtime POSIXct            10      0.02
## los numeric                10      0.02
##
## ---
##
## ### Summary of Table: labs_hourly ###
## Number of rows: 928195
## Number of columns: 22
## Column names and data types:
##           Data_Type Missing_Count Missing_Pct.
## icustay_id integer             0      0.00
## hr integer                     0      0.00
## neutrophil numeric           853757    91.98
## creactiveprotein numeric      926422    99.81
## whitebloodcell numeric       576380    62.10
## partialpressureo2 numeric     473825    51.05
## bicarbonate numeric          534726    57.61
## lactate numeric             763646    82.27
## troponin numeric            886778    95.54
## bloodureanitrogen numeric     549233    59.17
## creatinine numeric          547907    59.03
## alaninetransaminase numeric    836577    90.13
## aspartatetransaminase numeric  836642    90.14
## hemoglobin numeric          514069    55.38
## intnormalisedratio numeric    675513    72.78
## platelets numeric           559952    60.33
## albumin numeric            866292    93.33
## chloride numeric           496072    53.44
## glucose numeric            401175    43.22
## sodium numeric            524290    56.48
## bilirubin numeric          816141    87.93
## hematocrit numeric         487396    52.51
##
## ---
##
## ### Summary of Table: output_hourly ###
## Number of rows: 3325543
## Number of columns: 3
## Column names and data types:
##           Data_Type Missing_Count Missing_Pct.
## icustay_id integer             0      0.00
## hr integer                     0      0.00
## urineoutput numeric           11920     0.36
##
## ---
##
## ### Summary of Table: patients ###
## Number of rows: 46520
## Number of columns: 8
## Column names and data types:
##           Data_Type Missing_Count Missing_Pct.
## row_id integer                 0      0.00

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```

## subject_id      integer           0          0.00
## gender          character          0          0.00
## dob             POSIXct           0          0.00
## dod             POSIXct          30761        66.12
## dod_hosp        POSIXct          36546        78.56
## dod_ssn         POSIXct          33142        71.24
## expire_flag     integer           0          0.00
##
## ---
##
## #### Summary of Table: pt_icu_outcome ####
## Number of rows: 61533
## Number of columns: 17
## Column names and data types:
##
##          Data_Type Missing_Count Missing_Pct.
## row_id      integer           0          0.00
## subject_id   integer           0          0.00
## dob          POSIXct           0          0.00
## hadm_id      integer           0          0.00
## admittance   POSIXct          12348         20.07
## dischtime     POSIXct          12348         20.07
## icustay_id    integer           0          0.00
## age_years     numeric           0          0.00
## intime       POSIXct           0          0.00
## outtime      POSIXct           10          0.02
## los          numeric           10          0.02
## hosp_deathtime POSIXct          59256         96.30
## icu_expire_flag integer           0          0.00
## hospital_expire_flag integer          12348         20.07
## dod          POSIXct          37341         60.68
## expire_flag   integer           0          0.00
## ttd_days      integer          37341         60.68
##
## ---

```

```
##
## ### Summary of Table: pt_stay_hr ###
## Number of rows: 3687586
## Number of columns: 9
## Column names and data types:
##          Data_Type Missing_Count Missing_Pct.
## icustay_id    integer             0          0.00
## hadm_id       integer             0          0.00
## subject_id    integer             0          0.00
## intime        POSIXct             0          0.00
## outtime       POSIXct             0          0.00
## starttime     POSIXct             0          0.00
## endtime       POSIXct             0          0.00
## hr            integer             0          0.00
## dy            integer            1398          0.04
##
## ---
##
## ### Summary of Table: pt_weight ###
## Number of rows: 396241
## Number of columns: 11
## Column names and data types:
##          Data_Type Missing_Count Missing_Pct.
## icustay_id    integer             0          0.00
## dy            integer             0          0.00
## starttime     POSIXct             0          0.00
## endtime       POSIXct             0          0.00
## admissionweight numeric          331567        83.68
## dailyweight   numeric          241485        60.94
## previousweight numeric          312334        78.82
## echoweight    numeric          262244        66.18
## avg_weight_naive numeric          20263         5.11
## min_weight    numeric          20263         5.11
## max_weight    numeric          20263         5.11
##
## ---
```

```
##
## ### Summary of Table: pv_mechvent ###
## Number of rows: 694958
## Number of columns: 21
## Column names and data types:
##          Data_Type Missing_Count Missing_Pct.
## icustay_id      integer           0         0.00
## charttime       POSIXct          76         0.01
## starttime       POSIXct           0         0.00
## endtime         POSIXct           0         0.00
## duration_hours  numeric           0         0.00
## ventnum         integer           0         0.00
## minutevolume    numeric        233141        33.55
## settidalvolume  numeric        461276        66.37
## obstidalvolume  numeric        314290        45.22
## sponttidalvolume numeric        439527        63.25
## setpeep         numeric         21446         3.09
## totalpeep       numeric        669702        96.37
## pressurehighprv integer        694415        99.92
## pressurelowprv  numeric        694428        99.92
## timehighprv     numeric        694419        99.92
## timelowprv      numeric        694423        99.92
## meanairwaypressure numeric      236821        34.08
## peakinsppressure numeric      322901        46.46
## neginspforce    numeric        693945        99.85
## insptime        numeric        572792        82.42
## plateaupressure numeric      564736        81.26
```

```
##
## ---
```

```
##
## ### Summary of Table: transfers ###
## Number of rows: 261897
## Number of columns: 13
## Column names and data types:
##          Data_Type Missing_Count Missing_Pct.
## row_id      integer           0         0.00
## subject_id   integer           0         0.00
## hadm_id      integer           0         0.00
## icustay_id   integer        174176        66.51
## dbsource     character         0         0.00
## eventtype    character         0         0.00
## prev_careunit character         0         0.00
## curr_careunit character         0         0.00
## prev_wardid  integer        58933        22.50
## curr_wardid  integer        58943        22.51
## intime       POSIXct          24         0.01
## outtime      POSIXct        58976        22.52
## los          numeric        58976        22.52
```

```
##
## ---
```

```
##
## ### Summary of Table: vasopressors ###
## Number of rows: 314964
## Number of columns: 11
## Column names and data types:
```

```
##          Data_Type Missing_Count Missing_Pct.
## icustay_id      integer           834         0.26
## starttime      POSIXct             0         0.00
## endtime        POSIXct        231185        73.40
## norepinephrine_rate  numeric        231436        73.48
## norepinephrine_amount  numeric        237721        75.48
## epinephrine_rate    numeric        259713        82.46
## epinephrine_amount  numeric        278082        88.29
## dopamine_rate      numeric        181993        57.78
## dopamine_amount    numeric        227051        72.09
## dobutamine_rate    numeric        272450        86.50
## dobutamine_amount  numeric        286648        91.01
##
## ---
##
## ### Summary of Table: vitals_hourly ###
## Number of rows: 7292362
## Number of columns: 11
## Column names and data types:
##          Data_Type Missing_Count Missing_Pct.
## icustay_id      integer             0         0.00
## hr              integer             0         0.00
## spo2            numeric        1972385        27.05
## fio2            numeric        6341965        86.97
## temperature     numeric        5726438        78.53
## resprate        numeric        2550173        34.97
## heartrate       numeric         872012        11.96
## sysbp           numeric        2647890        36.31
## diasbp          numeric        2648668        36.32
## glucose         numeric        6144343        84.26
## meanarterialpressure  numeric        2631818        36.09
##
## ---
```

```
##
## All CSV tables have been successfully loaded and summarized!
```

```
## Loaded tables:
## admissions, antibiotics, bloodculture, gcs_hourly, icd9_diag, icustays, labs_hourly, output
## _hourly, patients, pt_icu_outcome, pt_stay_hr, pt_weight, pv_mechvent, transfers, vasopressors,
## vitals_hourly
```

## 1. admissions

- **Rows:** 58,976 | **Columns:** 19
- **Key Missingness:**
  - `deathtime` : 90.07% missing. Relevant for mortality analysis but likely reflects non-deceased patients.
  - Minimal missingness for core variables like `admittime` , `dischtime` , and demographic details.
- **Observation:** High-quality foundational data with minimal issues, apart from `deathtime` .

## 2. antibiotics

- **Rows:** 164,927 | **Columns:** 16
- **Key Missingness:**



- `rate` and `rateuom` : Both 100% missing, suggesting these variables can be dropped.
- `totalamount` : 5.04% missing.
- **Observation:** Useful for understanding antibiotic administration, though some variables appear irrelevant.

### 3. bloodculture

- **Rows:** 632,506 | **Columns:** 10
- **Key Missingness:**
  - `icustay_id` : 24.79% missing, significant for ICU-related analyses.
  - `hr` : 29.68% missing.
- **Observation:** Moderate missingness for key ICU identifiers may limit linking with other tables.

### 4. gcs\_hourly

- **Rows:** 1,515,342 | **Columns:** 7
- **Key Missingness:**
  - `gcseyes`, `gcsmotor`, and `gcsverbal` : <0.3% missing, indicating good data quality.
- **Observation:** Reliable source for Glasgow Coma Scale (GCS) data with low missingness.

### 5. icd9\_diag

- **Rows:** 651,047 | **Columns:** 7
- **Key Missingness:**
  - Minimal issues, with <0.01% missing in `seq_num`.
- **Observation:** High-quality diagnosis data, ready for analysis.

### 6. icustays

- **Rows:** 61,532 | **Columns:** 12
- **Key Missingness:**
  - `outtime` and `los` : Both 0.02% missing.
- **Observation:** Reliable ICU stay details with minimal issues.

### 7. labs\_hourly

- **Rows:** 928,195 | **Columns:** 22
- **Key Missingness:**
  - Many variables exceed 90% missingness, including `creactiveprotein` (99.81%) and `alaninetransaminase` (90.13%).
  - Core variables like `neutrophil` (91.98%) also have high missingness.
- **Observation:** Key lab data but requires careful selection and imputation due to widespread missingness.

### 8. output\_hourly

- **Rows:** 3,325,543 | **Columns:** 3
- **Key Missingness:**
  - `urineoutput` : 0.36% missing.
- **Observation:** High-quality output data with negligible issues.

### 9. patients

- **Rows:** 46,520 | **Columns:** 8
- **Key Missingness:**
  - Mortality-related fields ( `dod_hosp`, `dod_ssn` ) have >70% missingness.
  - Demographic fields like `gender` and `dob` are complete.
- **Observation:** Core patient demographics are robust, but mortality data requires handling.

## 10. pt\_icu\_outcome

- **Rows:** 61,533 | **Columns:** 17
- **Key Missingness:**
  - Critical fields like `hosp_deathtime` (96.30%) and `dod` (60.68%) have very high missingness.
- **Observation:** ICU outcomes are incomplete for most patients.

## 11. pt\_stay\_hr

- **Rows:** 3,687,586 | **Columns:** 9
- **Key Missingness:**
  - Minimal, with `dy` missing 0.04%.
- **Observation:** Comprehensive hourly stay data with excellent quality.

## 12. pt\_weight

- **Rows:** 396,241 | **Columns:** 11
- **Key Missingness:**
  - Most weight-related fields exceed 60% missingness, e.g., `admissionweight` (83.68%) .
- **Observation:** Data quality for weight variables is poor, limiting analysis.

## 13. pv\_mechvent

- **Rows:** 694,958 | **Columns:** 21
- **Key Missingness:**
  - Ventilation parameters like `pressurehighaprv` and `timelowaprv` exceed 99% missingness.
  - `minutevolume` : 33.55% missing.
- **Observation:** Highly sparse data, with only a few useful variables.

## 14. transfers

- **Rows:** 261,897 | **Columns:** 13
- **Key Missingness:**
  - `icustay_id` : 66.51% missing.
  - Ward identifiers ( `prev_wardid` and `curr_wardid` ) have ~22.5% missingness.
- **Observation:** Transfer details are partially incomplete, limiting their utility.

## 15. vasopressors

- **Rows:** 314,964 | **Columns:** 11
- **Key Missingness:**
  - Missingness ranges from 57.78% ( `dopamine_rate` ) to 91.01% ( `dobutamine_amount` ).
- **Observation:** Sparse data for vasopressor administration, with limited reliable variables.

## 16. vitals\_hourly

- **Rows:** 7,292,362 | **Columns:** 11
- **Key Missingness:**
  - Vital signs like `fio2` and `temperature` exceed 75% missingness.
  - Core variables like `heartrate` and `spo2` are ~10-30% missing.
- **Observation:** Rich time-series data but requires significant preprocessing.

# 1.4 Key Observations from Data:

## 1.4.1 High Missingness in Time-Series Data ( `vitals_hourly` and `labs_hourly` ):

- Many variables in `labs_hourly` and `vitals_hourly` exceed **70% missingness**.
- Some variables ( `creactiveprotein` , `alaninetransaminase` ) in `labs_hourly` have almost **complete missingness**, making them unsuitable for imputation or analysis.

### 1.4.2 Time Discrepancies in `hr` Across Tables:

- `vitals_hourly` : `hr` starts at 1 (post-ICU admission) and increments hourly.
- `labs_hourly` : `hr` includes negative values for pre-ICU measurements.
- Different intervals or irregular sampling times make direct alignment across tables challenging.

### 1.4.3 Key Tables for the First 24 Hours:

- `pt_stay_hr` : Provides a comprehensive hourly structure for ICU stays and can act as a unifying table for `hr` alignment.
- `vitals_hourly` and `labs_hourly` : Crucial for predictive modeling but need proper filtering for the first 24 hours.

### 1.4.4 Predictive Modeling Needs:

- Accurate prediction of mortality requires reliable features extracted from the **first 24 hours**.
- Time-sensitive modeling approaches (e.g., LSTMs, GRUs) need continuous time-series data, while tree-based models (e.g., XGBoost) can use aggregated features.

## 1.5 Consideration for following analysis

Ensure high-quality data is used while minimizing the impact of missingness on analyses. ##### 1.4.1 **Prioritize Tables with Low Missingness:** - `admissions`, `patients`, `icustays`, and `gcs_hourly` are the most reliable tables for initial analysis.

### 1.4.2 Handle High Missingness Strategically:

- For tables like `labs_hourly` and `vitals_hourly`, consider using imputation, variable selection, or excluding highly sparse variables.

### 1.4.3 Focus on Core Time-Series Data:

- `vitals_hourly` and `output_hourly` provide crucial insights into patient conditions, despite moderate missingness.

### 1.4.4 Exclude Variables with >90% Missingness:

- Tables like `vasopressors` and `pv_mechvent` have several variables with near-complete missingness, which may not have value.

## Section 2: Data Preprocessing

### Objectives:

- Prepare the dataset for analysis by filtering, merging, and handling missing data.
- Ensure consistency and completeness in the preprocessed data.

### Define the study population:

- Focus on ICU patients aged between **18 and 89 years**, Aligns with the MIMIC-III age shifting policy for HIPAA compliance and avoids pediatric and super-elderly populations.
- Retain only the **first ICU admission** per patient to ensure independence of observations and avoids over representation of specific patients.
- Ensure the ICU stay duration is **≥ 24 hours** for providing sufficient data for meaningful feature extraction.

- Add **Weekend/Weekday Flag**, which directly supports Aim 2, investigating mortality association with admission timing.

## Validate inclusion and exclusion criteria

- Exclude records missing critical demographic variables like gender, age, or ICU admission/discharge times.
- Align with project aims to predict mortality using data from the **first 24 hours of ICU stay** (for predictors) but not restrict mortality outcomes to the same timeframe.

## Steps:

### 2.1 Merge Patients, Admissions, and ICU Stays

- **Objective:** Combine demographic, admission, and ICU stay data into a cohesive dataset for initial filtering.
- **Why:** `patients`, `admissions`, and `icustays` tables provide core demographic and hospitalization data that form the backbone of our analysis.
- **How:**
  - Merge `patients` and `admissions` using the key `subject_id` to align patient demographics with their hospital admissions.
  - Merge the resulting dataset with `icustays` using the keys `subject_id` and `hadm_id` to include ICU-specific stay details.

### 2.2 Retain the First ICU Admission per Patient

- **Objective:** Ensure that each patient contributes only their first ICU admission to the analysis.
- **Why:** Retaining only the first ICU admission avoids over-representation of patients with multiple ICU stays and ensures independence of observations.
- **How:**
  - Sort the data by `subject_id` and `admittime`.
  - Use `.SD[1]` to retain the first ICU stay for each `subject_id`.

### 2.3 Filter by Age ( $18 \leq \text{Age} \leq 89$ )

- **Objective:** Focus on adult patients while excluding pediatric and super-elderly populations.
- **Why:** The MIMIC-III dataset masks ages above 89 due to HIPAA compliance, making exact age unknown for these patients.
- **How:**
  - Calculate patient age at admission as the difference between `admittime` and `dob`.
  - Retain records where age is between 18 and 89.

### 2.4 Filter ICU Stays Lasting $\geq 24$ Hours

- **Objective:** Exclude ICU stays shorter than 24 hours to ensure sufficient data for analysis.
- **Why:** Short ICU stays may not provide enough information for meaningful predictive modeling.
- **How:**
  - The 'los' variable in the `icustays` table already represents the length of stay in days.
  - Convert it to hours (`los_hours = los * 24`) for consistency.
  - Retain records where `los_hours` is 24 or more.

### 2.5 Add Weekend Admission Flag

- **Objective:** Identify ICU admissions occurring on weekends to address Aim 2 of the project.
- **Why:**
  - Investigate whether weekend ICU admissions are associated with higher mortality rates.

- Weekend admissions could differ in outcomes due to variations in staffing, resource availability, or other factors.
- **How:**
  - Add a new column `intime_weekdays` to display the day of the week (e.g., “Monday”, “Saturday”).
  - Use this column to create a boolean flag `is_weekend_admission`, which is set to `TRUE` for admissions occurring on “Saturday” or “Sunday”.
  - Save the updated dataset to include these new columns for downstream analysis.

## 2.6 Save Intermediate Filtered Data

- **Objective:** Save the filtered dataset for reproducibility and debugging purposes.
- **Why:** Provides a checkpoint to avoid repeating prior filtering steps if further processing needs adjustments.
- **How:**
  - Save the filtered dataset ( `filtered_data` ) as an RDS file using `saveRDS` .

## 2.7 Merge Time-Series Data into `pt_stay_hr`

- **Objective:** Combine hourly clinical measurements into the base time-series structure of `pt_stay_hr` .
- **Why:** Hourly data from `vitals_hourly` , `labs_hourly` , `gcs_hourly` , and `output_hourly` provide critical features for predictive modeling.
- **How:**
  - Sequentially left join `vitals_hourly` , `labs_hourly` , `gcs_hourly` , and `output_hourly` to `pt_stay_hr` using `icustay_id` and `hr` .

## 2.8 Filter Time-Series Data to First 24 Hours

- **Objective:** Retain only the data corresponding to the first 24 hours of ICU stay.
- **Why:** Aligns with the project requirement to use the first 24 hours of ICU data for prediction while not limiting outcomes to the same timeframe.
- **How:**
  - Filter records where the `hr` column is less than or equal to 24.

## 2.9 Save Processed Time-Series Data

- **Objective:** Save the merged and filtered time-series data for further analysis.
- **Why:** Provides a checkpoint for reproducibility and supports efficient debugging.
- **How:**
  - Save the processed time-series dataset as an RDS file.

## 2.10 Merge Filtered Time-Series Data with `filtered_data`

- **Objective:** Combine the filtered demographic and admission data with time-series data for the first 24 hours.
- **Why:** Integrates all relevant information into a single dataset for subsequent analysis and model building.
- **How:**
  - Left join the time-series data with `filtered_data` using `icustay_id` .

## 2.11 Save Final Master Dataset

- **Objective:** Save the fully preprocessed dataset for predictive modeling and hypothesis testing.
- **Why:** Ensures the final dataset is ready for downstream tasks and avoids repetition of preprocessing steps.
- **How:**
  - Save the final dataset ( `master_data` ) as an RDS file using `saveRDS` .

```
##      hadm_id      icustay_id      MDROs
## Min.   :100001   Min.     :    -1   Mode :logical
## 1st Qu.:125063   1st Qu.:    -1   FALSE:67737
## Median :149996   Median :228138   TRUE :318
## Mean   :149982   Mean    :174167
## 3rd Qu.:174895   3rd Qu.:264167
## Max.   :199999   Max.    :299998
```

```
## No duplicates exist for hadm_id.
```

```
## [1] "C"
```

```
## Final filtered data saved with 36522 rows and 40 columns.
```

```
## Filtered Time-Series data saved with 568037 rows and 44 columns.
```

```
## Updated Filtering and Merging Steps Completed.
```

```
## Filtered Dataset Rows: 36522
```

```
## Master Dataset Rows: 375552
```

```
## Master Dataset Columns: 83
```

## Section 3: Exploratory Data Analysis (EDA)

### Objectives:

- Understand the structure and relationships in the filtered data.
- Identify trends, distributions, and potential outliers.
- Evaluate key predictors and their correlations with the target variable (mortality).

### Steps:

#### 3.1 Basic Descriptive Statistics:

- **Objective:** Summarize the dataset to understand its structure and identify potential issues.
- **Why:**
  - Ensure numerical and categorical variables are within expected ranges.
  - Identify missing values that may need handling during modeling.
- **How:**
  - Calculate summary statistics for numerical variables (mean, median, standard deviation, min, max).
  - Tabulate categorical variables (frequency and proportions).
  - Summarize missing values for all variables to identify those requiring imputation or exclusion.
  - Stratify statistics by mortality ( `EXPIRE_FLAG` ) to detect differences between survivors and non-survivors.

### 3.2 Target Variable Analysis:

- **Objective:** Understand the distribution of the target variable and its relationship with key features.
- **Why:**
  - Explore the prevalence of mortality ( `EXPIRE_FLAG` ).
  - Analyze survival times for additional insights.
- **How:**
  - Visualize the distribution of mortality ( `EXPIRE_FLAG` ) as proportions or counts.
  - Use histograms and bar plots to compare mortality trends across age groups, gender, and ICU types.
  - Explore survival times using Kaplan-Meier curves or other survival analysis techniques.

### 3.3 Key Predictor Exploration:

- **Objective:** Investigate the distribution and predictive power of key clinical variables.
- **Why:**
  - Determine whether predictors show significant differences across mortality outcomes.
  - Identify potential predictive patterns or outliers in vital signs and lab results.
- **How:**
  - Use boxplots and density plots to visualize distributions of vital signs and lab values.
  - Focus on predictors with lower percentages of missing values to ensure robust analysis.
  - Stratify by `EXPIRE_FLAG` to compare trends between survivors and non-survivors.

### 3.4 Correlation Analysis:

- **Objective:**
  - Focus on numerical variables in `master_data`.
  - Address potential multicollinearity by identifying highly correlated variables ( $> 0.8$  or  $< -0.8$ ).
- **Why:**
  - Identify groups of correlated variables to avoid redundancy in modeling.
  - Highlight potential key predictors.
- **How:**
  - Compute a correlation matrix for numerical variables using complete cases.
  - Visualize correlations using a heatmap with hierarchical clustering to reveal relationships.

### 3.5 Demographics and ICU Characteristics:

- **Objective:** Explore the relationships between patient demographics, ICU characteristics, and mortality outcomes.
- **Why:**
  - Assess the impact of variables such as age, gender, and ICU type on mortality.
  - Investigate potential differences in outcomes between weekend and weekday admissions.
- **How:**
  - Analyze mortality rates across demographic groups (age, gender, ethnicity).
  - Visualize age distribution and compare across survival groups.
  - Visualize the distribution of ICU types ( `first_careunit` ) and their association with mortality.
  - Assess the impact of weekend ( `is_weekend_admission` ) vs. weekday admissions.
  - Perform t-tests for continuous variables (e.g., age).
  - Use chi-squared tests for categorical variables (e.g., gender, ICU types).

## Insights of step 3.1 results

```
##
## ### Summary of Table: final_filtered_data ###
## Number of rows: 36522
## Number of columns: 40
##
## Column names and data types:
##           Data_Type Missing_Count Missing_Pct.
## hadm_id          integer           0         0.00
## icustay_id        integer           0         0.00
## subject_id        integer           0         0.00
## row_id            integer           0         0.00
## dob.x             POSIXct           0         0.00
## admittime.x       POSIXct        5860        16.05
## disctime.x        POSIXct        5860        16.05
## age_years         numeric           0         0.00
## intime            POSIXct           0         0.00
## outtime           POSIXct           2         0.01
## los               numeric           2         0.01
## hosp_deathtime    POSIXct       35249        96.51
## icu_expire_flag   integer           0         0.00
## hospital_expire_flag.x integer     5860        16.05
## dod               POSIXct       22343        61.18
## expire_flag.x     integer           0         0.00
## ttd_days          integer       22343        61.18
## first_careunit     character         0         0.00
## last_careunit      character         0         0.00
## first_wardid       integer           0         0.00
## last_wardid        integer           0         0.00
## insurance          character         0         0.00
## language           character         0         0.00
## religion           character         0         0.00
## marital_status     character         0         0.00
## ethnicity          character         0         0.00
## admission_type     character         0         0.00
## admission_location  character         0         0.00
## hospital_expire_flag.y integer         0         0.00
## admittime.y        POSIXct           0         0.00
## disctime.y         POSIXct           0         0.00
## deathtime          POSIXct       32570       89.18
## icd9_code          character         0         0.00
## intime_weekdays   character         0         0.00
## is_weekend_admission logical         0         0.00
## gender             character         0         0.00
## dob.y             POSIXct           0         0.00
## expire_flag.y      integer           0         0.00
## avg_weight_naive   numeric        2312         6.33
## MDROs             logical        6859        18.78
##
## ---
```



```

##
## ### Summary of Table: master_data ###
## Number of rows: 375552
## Number of columns: 83
##
## Column names and data types:
##
##          Data_Type Missing_Count Missing_Pct.
## icustay_id      integer           0         0.00
## hadm_id.x        integer           0         0.00
## subject_id.x     integer           0         0.00
## row_id           integer           0         0.00
## dob.x            POSIXct           0         0.00
## admittance.x     POSIXct        57789        15.39
## dischtime.x      POSIXct        57789        15.39
## age_years        numeric           0         0.00
## intime.x          POSIXct           0         0.00
## outtime.x        POSIXct           2         0.00
## los              numeric           2         0.00
## hosp_deathtime   POSIXct       362480        96.52
## icu_expire_flag  integer           0         0.00
## hospital_expire_flag.x integer      57789        15.39
## dod              POSIXct       231193        61.56
## expire_flag.x    integer           0         0.00
## ttd_days         integer      231193        61.56
## first_careunit   character          0         0.00
## last_careunit    character          0         0.00
## first_wardid     integer           0         0.00
## last_wardid      integer           0         0.00
## insurance        character          0         0.00
## language         character          0         0.00
## religion          character          0         0.00
## marital_status   character          0         0.00
## ethnicity        character          0         0.00
## admission_type    character          0         0.00
## admission_location character          0         0.00
## hospital_expire_flag.y integer          0         0.00
## admittance.y     POSIXct           0         0.00
## dischtime.y      POSIXct           0         0.00
## deathtime        POSIXct       336408        89.58
## icd9_code         character          0         0.00
## intime_weekdays character          0         0.00
## is_weekend_admission logical          0         0.00
## gender           character          0         0.00
## dob.y            POSIXct           0         0.00
## expire_flag.y     integer           0         0.00
## avg_weight_naive  numeric       21751         5.79
## MDR0s            logical       68089        18.13
## hr               integer       21246         5.66
## hadm_id.y        integer       21246         5.66
## subject_id.y     integer       21246         5.66
## intime.y         POSIXct       21246         5.66
## outtime.y        POSIXct       21246         5.66
## starttime        POSIXct       21246         5.66
## endtime          POSIXct       21246         5.66
## dy              integer       21582         5.75

```

## spo2	numeric	53170	14.16
## fio2	numeric	356903	95.03
## temperature	numeric	256856	68.39
## resprate	numeric	60883	16.21
## heartrate	numeric	55848	14.87
## sysbp	numeric	65010	17.31
## diasbp	numeric	65063	17.32
## glucose.x	numeric	286346	76.25
## meanarterialpressure	numeric	64372	17.14
## neutrophil	numeric	372230	99.12
## creactiveprotein	numeric	375428	99.97
## whitebloodcell	numeric	351211	93.52
## partialpressureo2	numeric	338361	90.10
## bicarbonate	numeric	349620	93.09
## lactate	numeric	360710	96.05
## troponin	numeric	370982	98.78
## bloodureanitrogen	numeric	349457	93.05
## creatinine	numeric	349354	93.02
## alaninetransaminase	numeric	369411	98.36
## aspartatetransaminase	numeric	369409	98.36
## hemoglobin	numeric	345076	91.89
## intnormalisedratio	numeric	355970	94.79
## platelets	numeric	349277	93.00
## albumin	numeric	371438	98.90
## chloride	numeric	346295	92.21
## glucose.y	numeric	333377	88.77
## sodium	numeric	349594	93.09
## bilirubin	numeric	369437	98.37
## hematocrit	numeric	341138	90.84
## gcs	integer	260984	69.49
## gcseyes	integer	261112	69.53
## gcsmotor	integer	261276	69.57
## gcsverbal	integer	261288	69.57
## endotrachflag	integer	260984	69.49
## urineoutput	numeric	151427	40.32
##			
## ---			

## Summary of final\_filtered\_data

### 1. Overall Dataset Shape:

- **Rows:** 36,522
- **Columns:** 40

### 2. Key Observations:

- Variables like `dod` and `ttd_days` have a significant percentage of missing values (61.18%).
- `deathtime` and `hosp_deathtime` has 89.18% and 96.51% missing values, indicating most records lack time-of-death information.
- Most categorical fields have no missing values.
- All numeric values ( `los` , `age` , `icu_los_hours` ) are complete and ready for analysis.

## Summary of master\_data

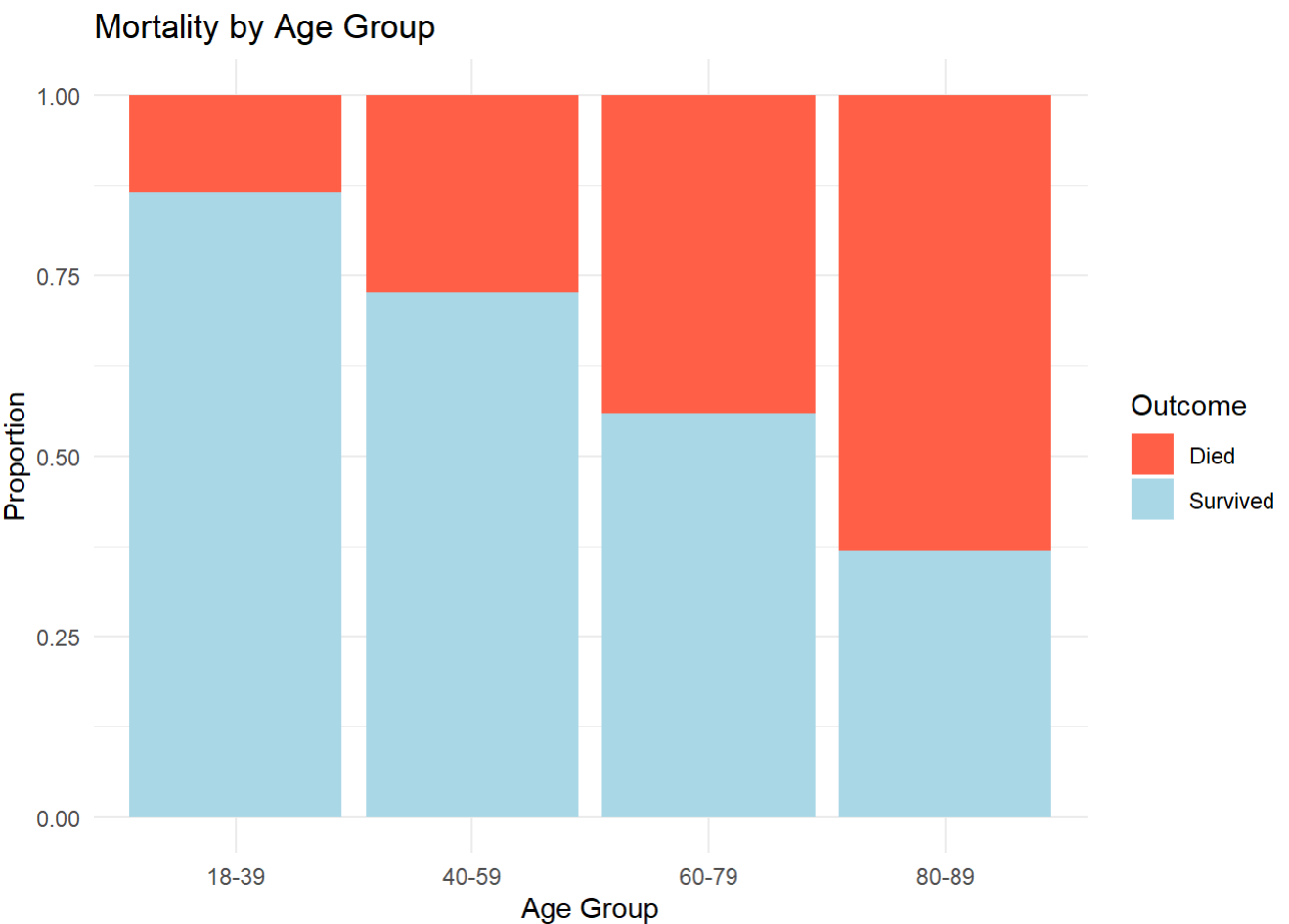
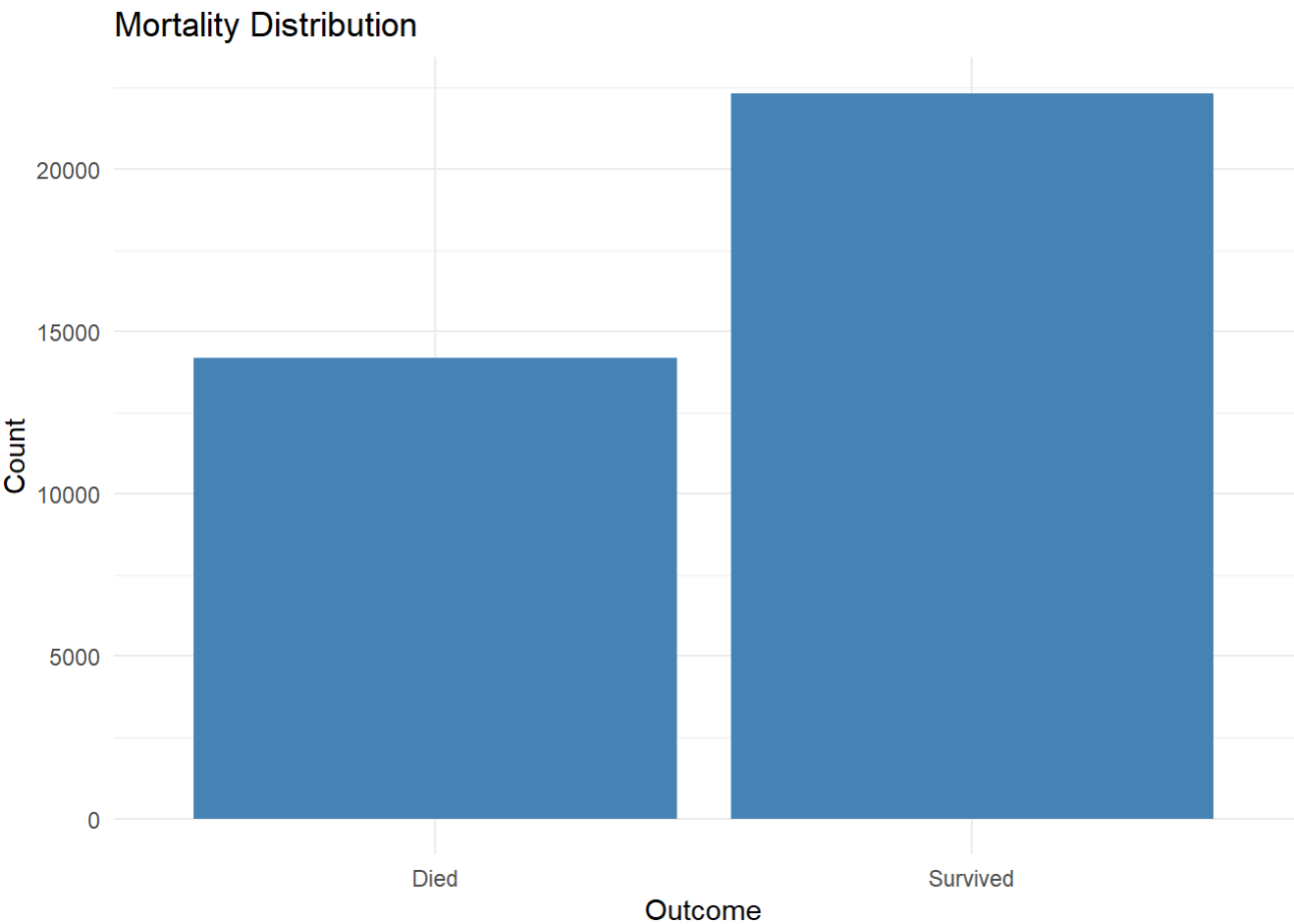
### 1. Overall Dataset Shape:

- **Rows:** 375,552
- **Columns:** 83

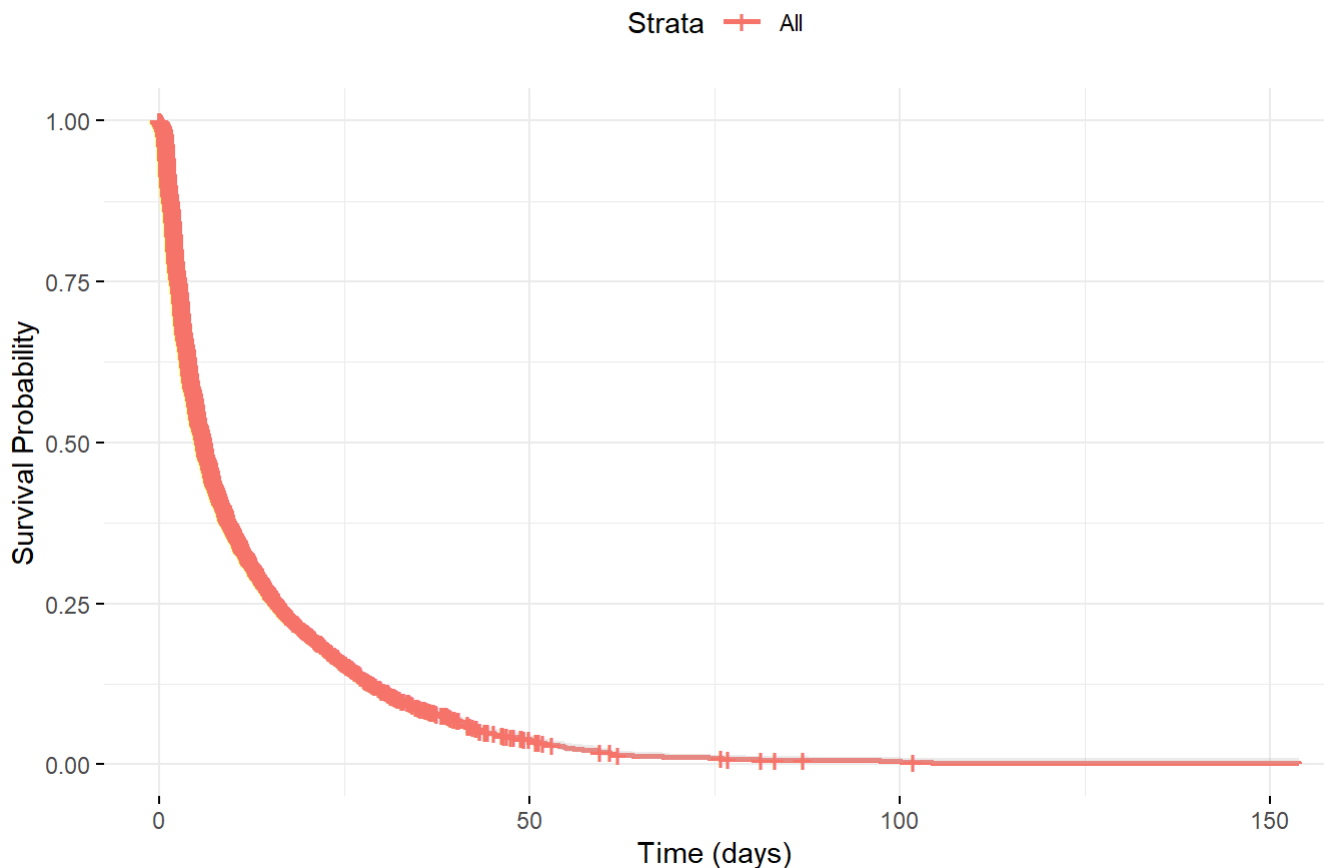
### 2. Key Observations:

- High missingness in many clinical variables ( `creactiveprotein` , `alaninetransaminase` , **etc.**), with some exceeding 90%.
- Critical time-series variables ( `spo2` , `temperature` , **etc.**) also show significant missingness, requiring imputation or exclusion strategies.
- Demographic and admission-related variables ( `gender` , `ethnicity` , `admittime` ) are fully populated, which is good for initial analysis.

# Observation and insights of step 3.2 results



## Survival Analysis



## Mortality Distribution

- **Observation:**
  - A larger proportion of ICU patients survived, as shown by the taller bar labeled “Survived.”
  - A smaller proportion of the patients did not survive (“Died”).
- **Insight:**
  - The dataset is imbalanced, with a majority of the patients surviving. This imbalance could influence predictive modeling, requiring techniques like balancing the dataset or using metrics robust to class imbalance (e.g., F1 score, AUC).

## Mortality by Age Group

- **Observation:**
  - The mortality rate increases with age.
  - In the age group 18–39, the proportion of patients who died is minimal compared to those who survived.
  - In the 80–89 age group, a significant proportion of patients did not survive, nearly matching or exceeding the survivors.
- **Insight:**
  - Age is a critical factor influencing ICU outcomes, with older patients at a much higher risk of mortality.
  - Predictive models should incorporate age as a key feature, potentially treating it as a non-linear variable to capture this trend.

## Survival Analysis

- **Observation:**
  - The survival probability drops steeply during the initial days of ICU stay and gradually levels off as time progresses.
  - The steep decline indicates that the first few days in the ICU are critical for patient survival.

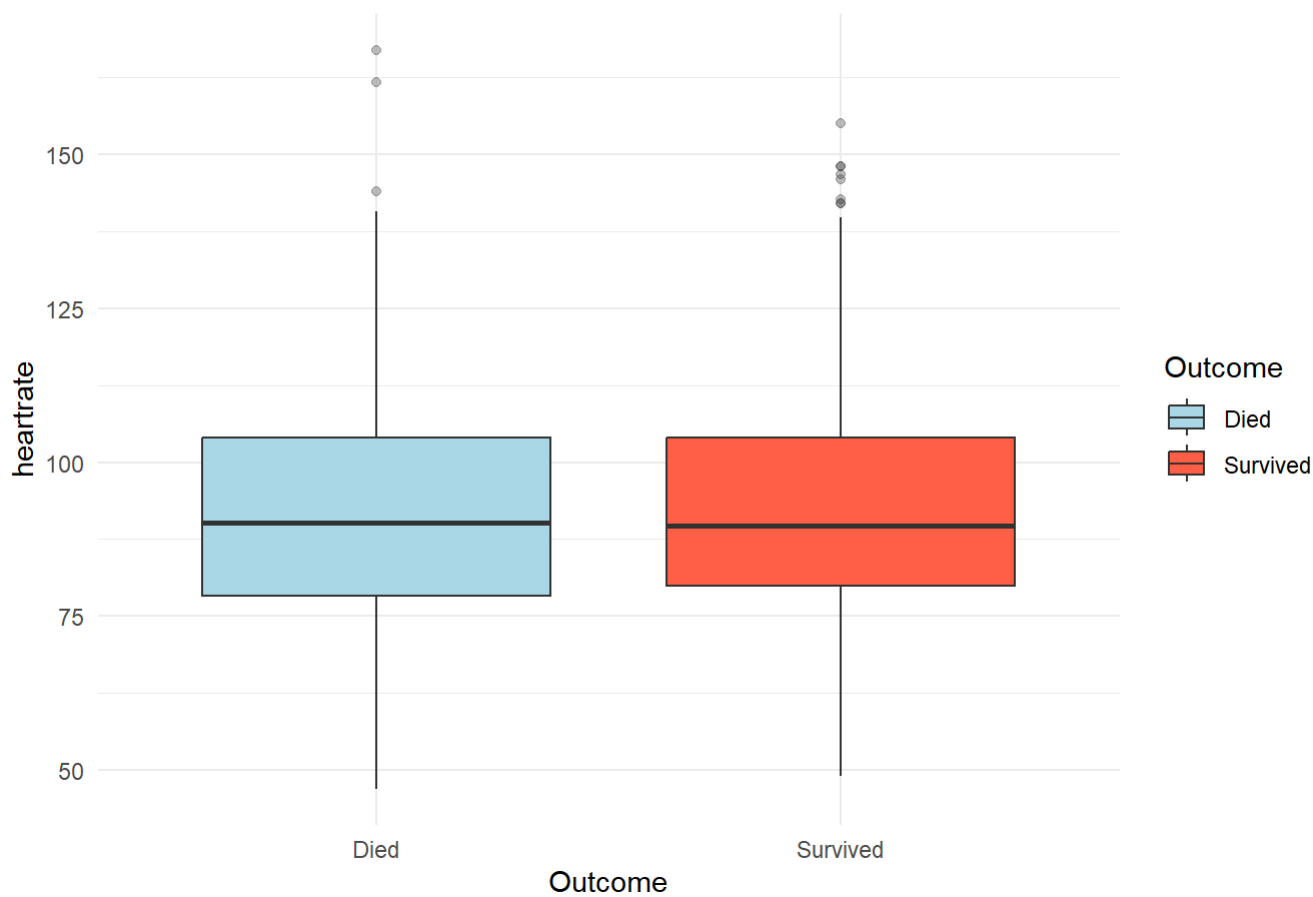
- **Insight:**

- This suggests that immediate and intensive care during the initial period is crucial for improving survival rates.
- The leveling off of survival probability after a certain point may indicate a higher likelihood of recovery or stabilization for longer-staying patients.
- Survival analysis supports the hypothesis that time-dependent features and early intervention are vital for predicting mortality.

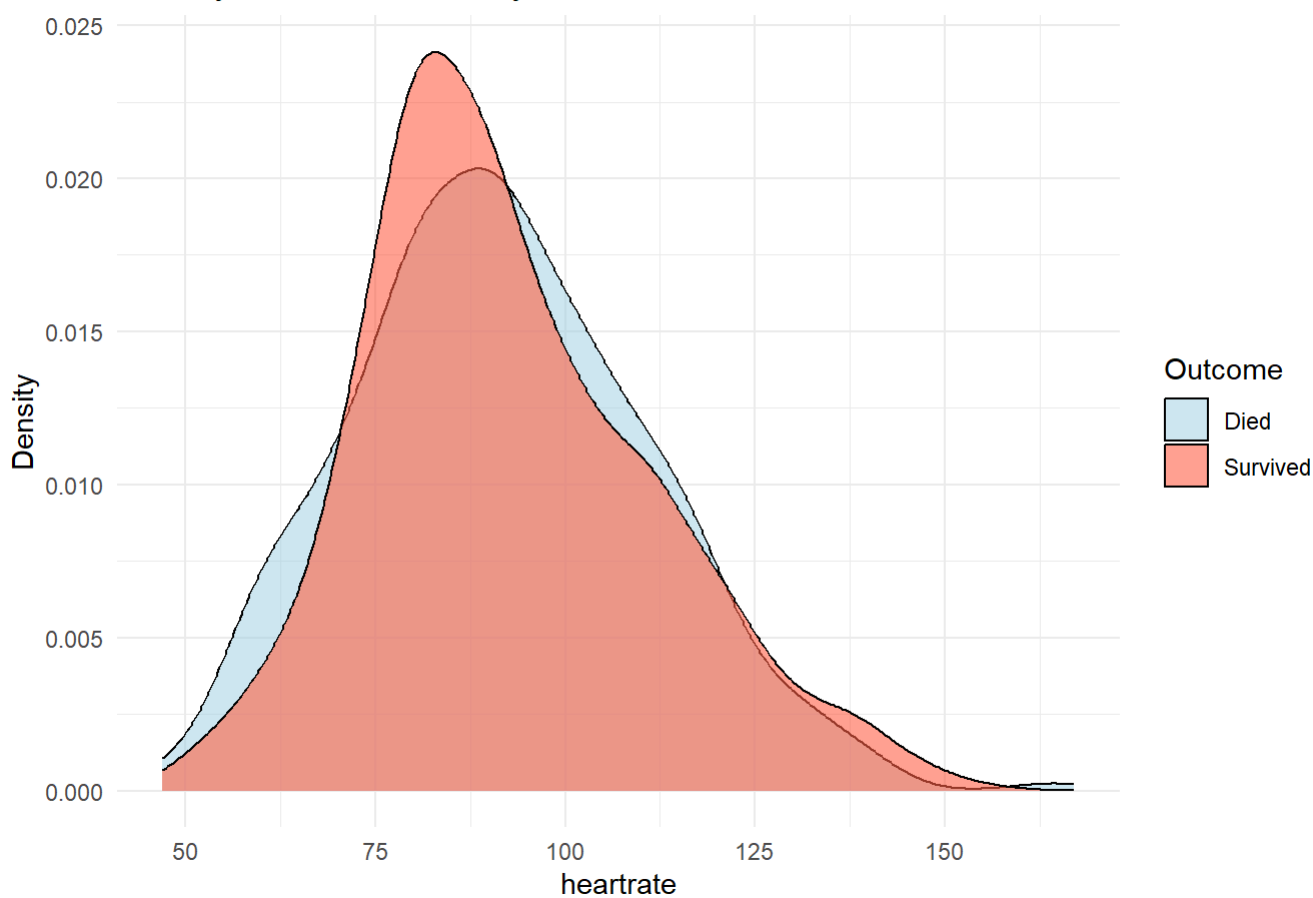
## Insights of step 3.3 results

```
##  
## --- Visualizations for heartrate ---
```

Boxplot of heartrate by Outcome

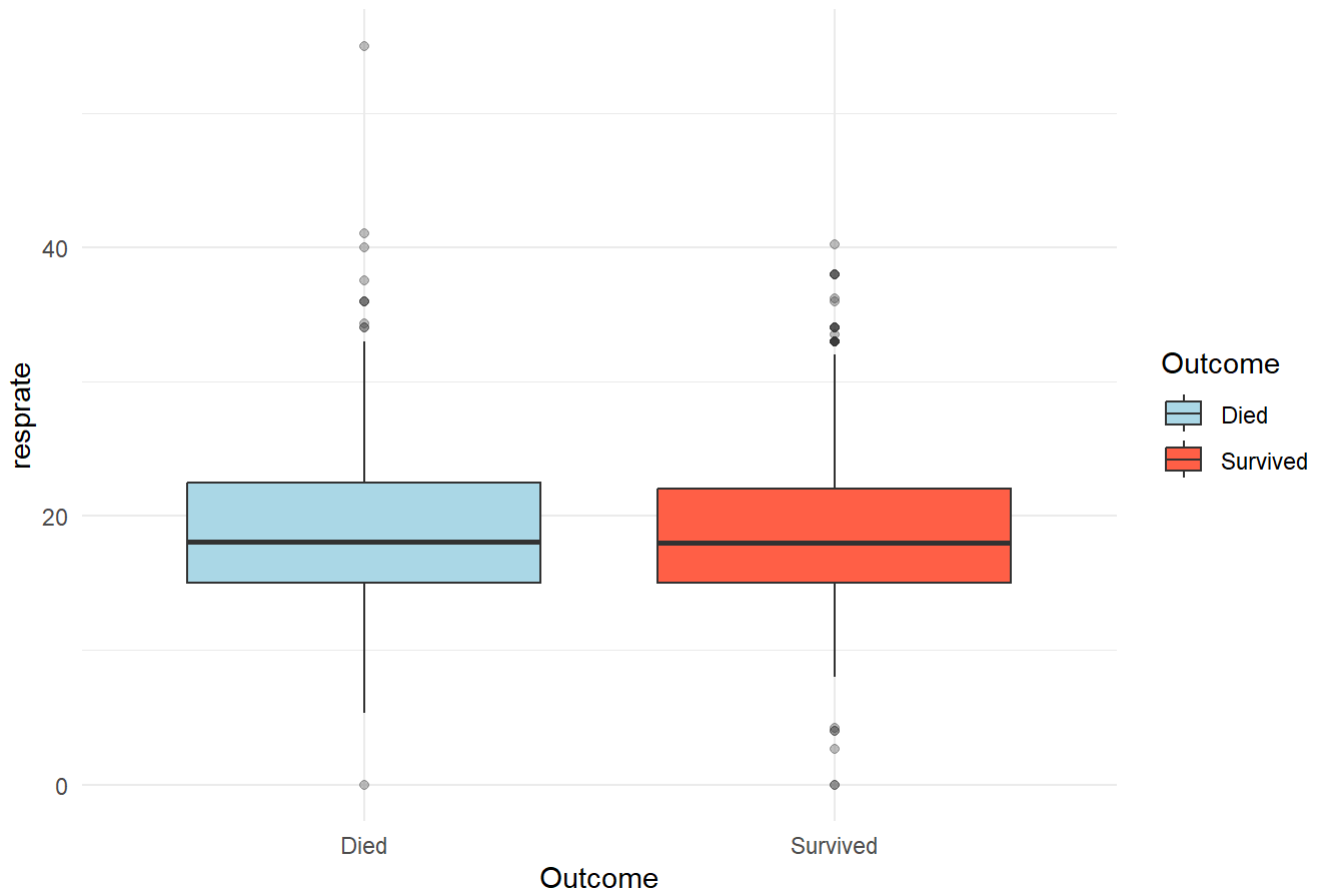


Density Plot of heartrate by Outcome

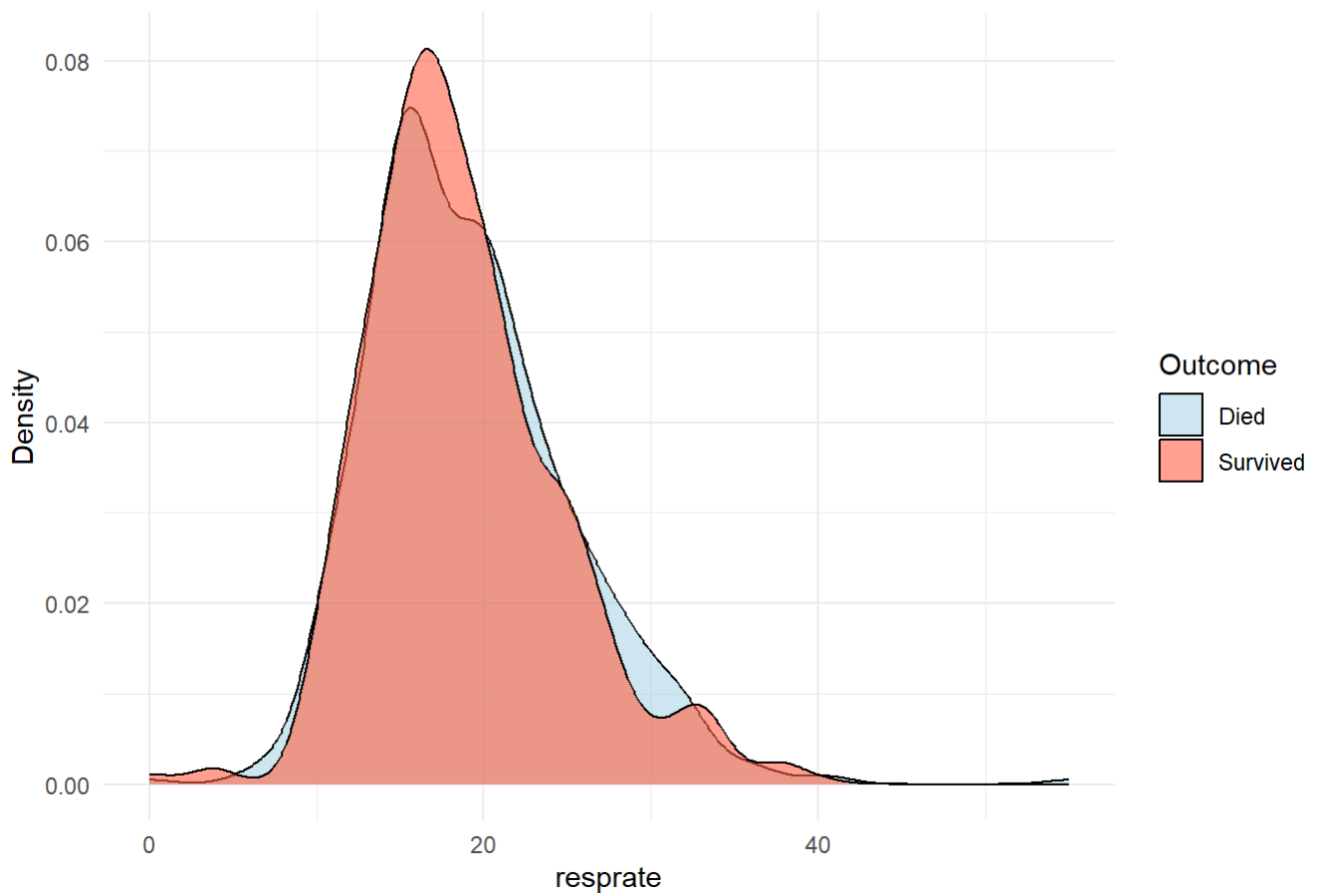


```
##  
## --- Visualizations for resprate ---
```

Boxplot of resprate by Outcome



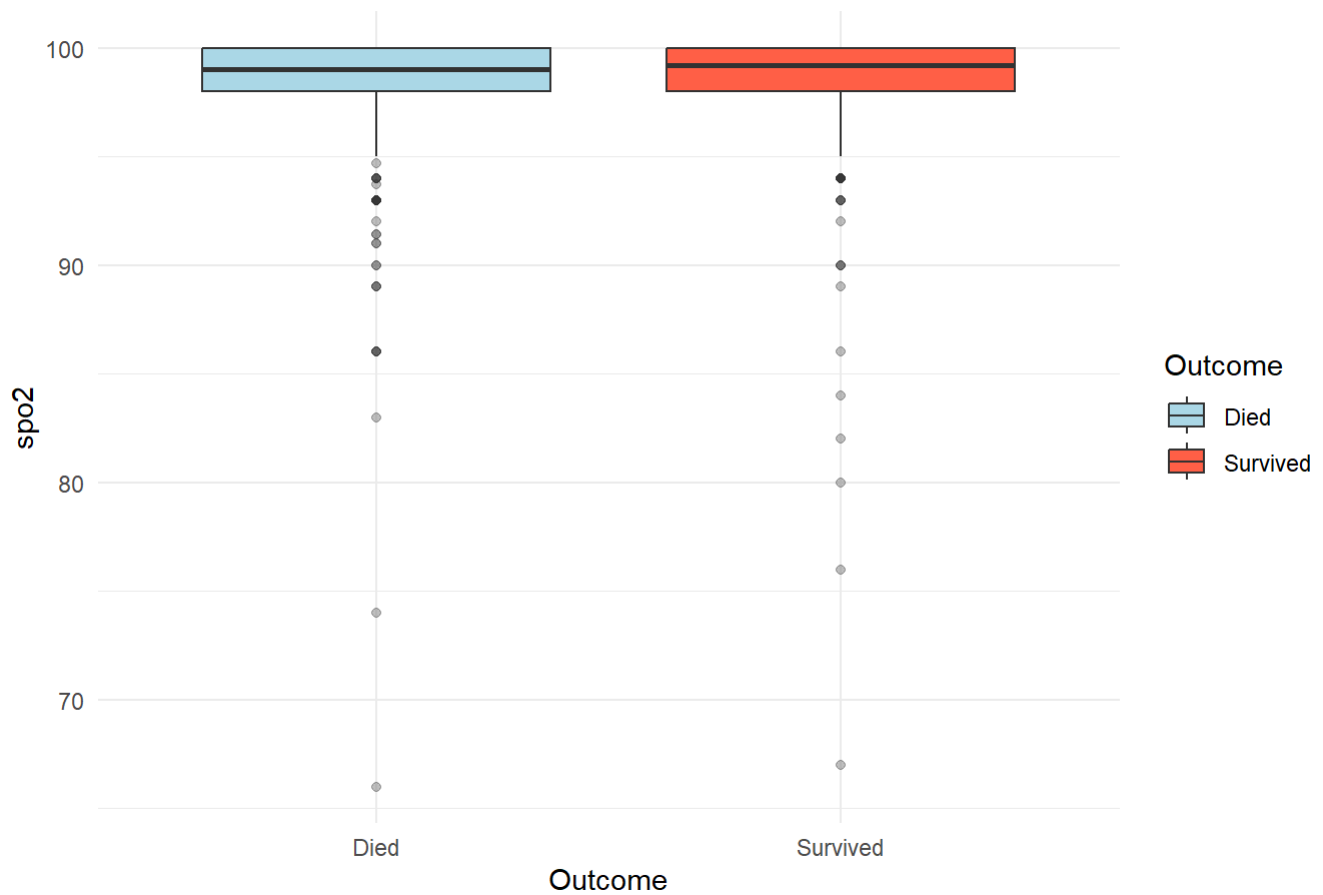
Density Plot of resprate by Outcome



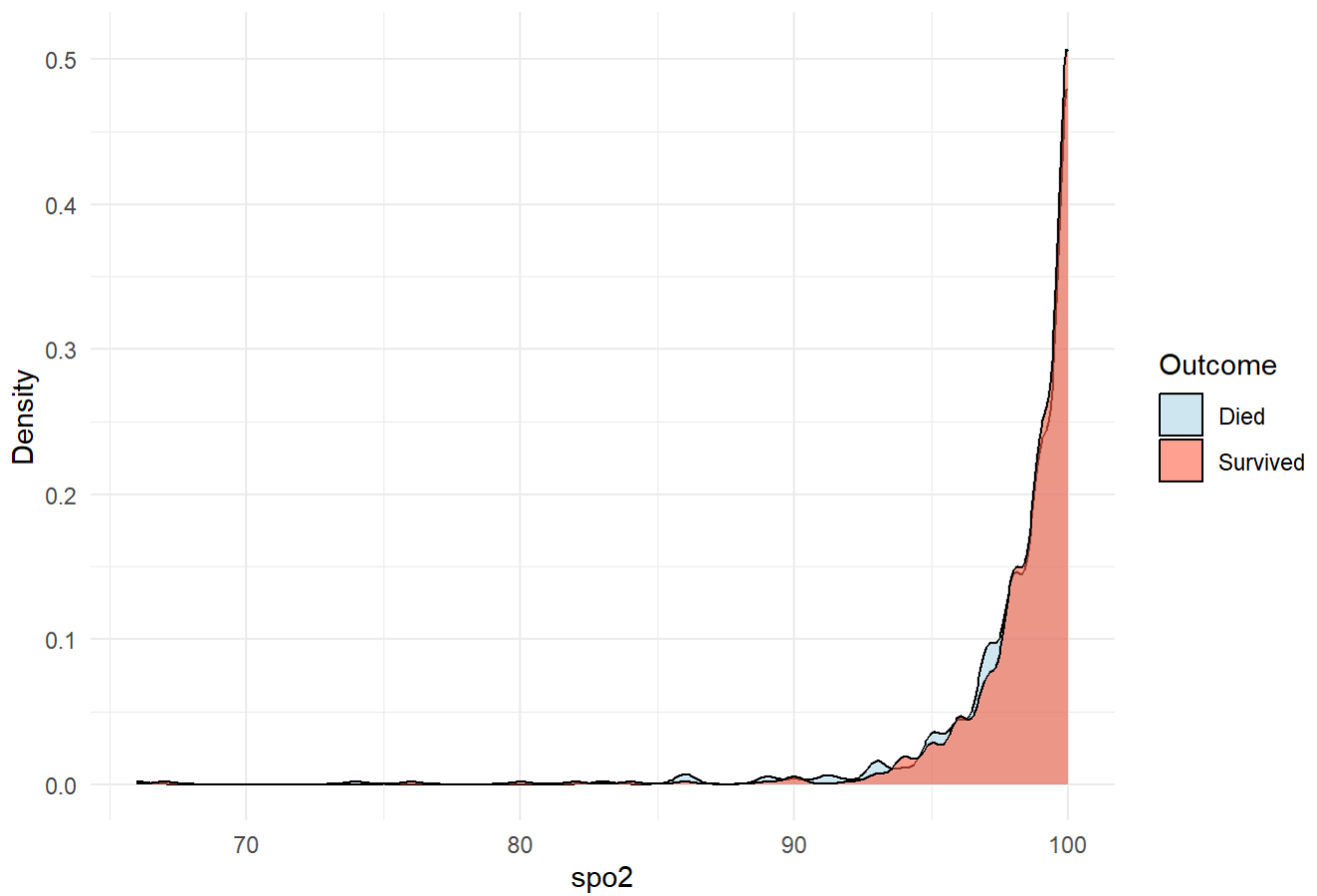
```
##  
## --- Visualizations for spo2 ---
```



Boxplot of spo2 by Outcome

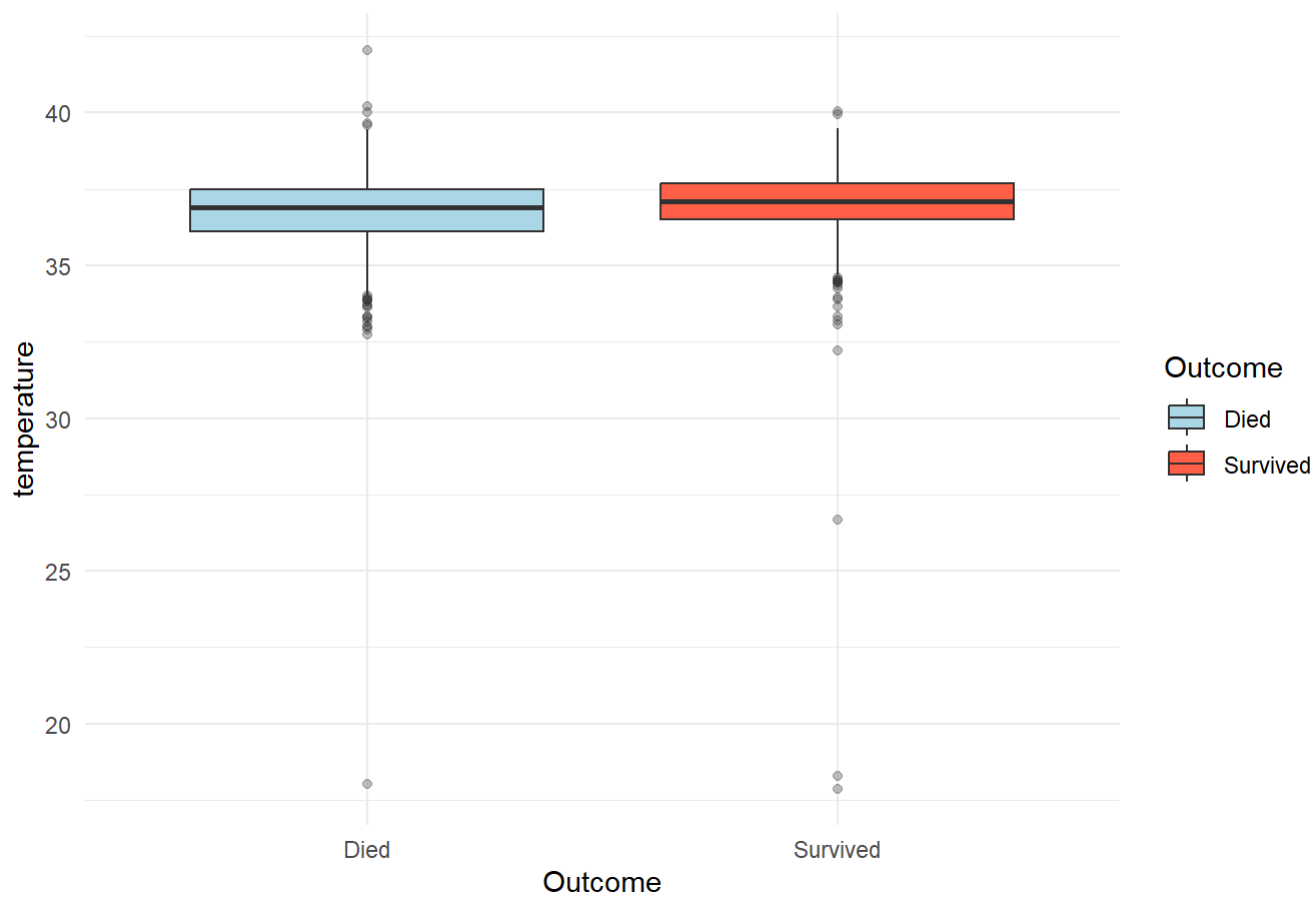


Density Plot of spo2 by Outcome

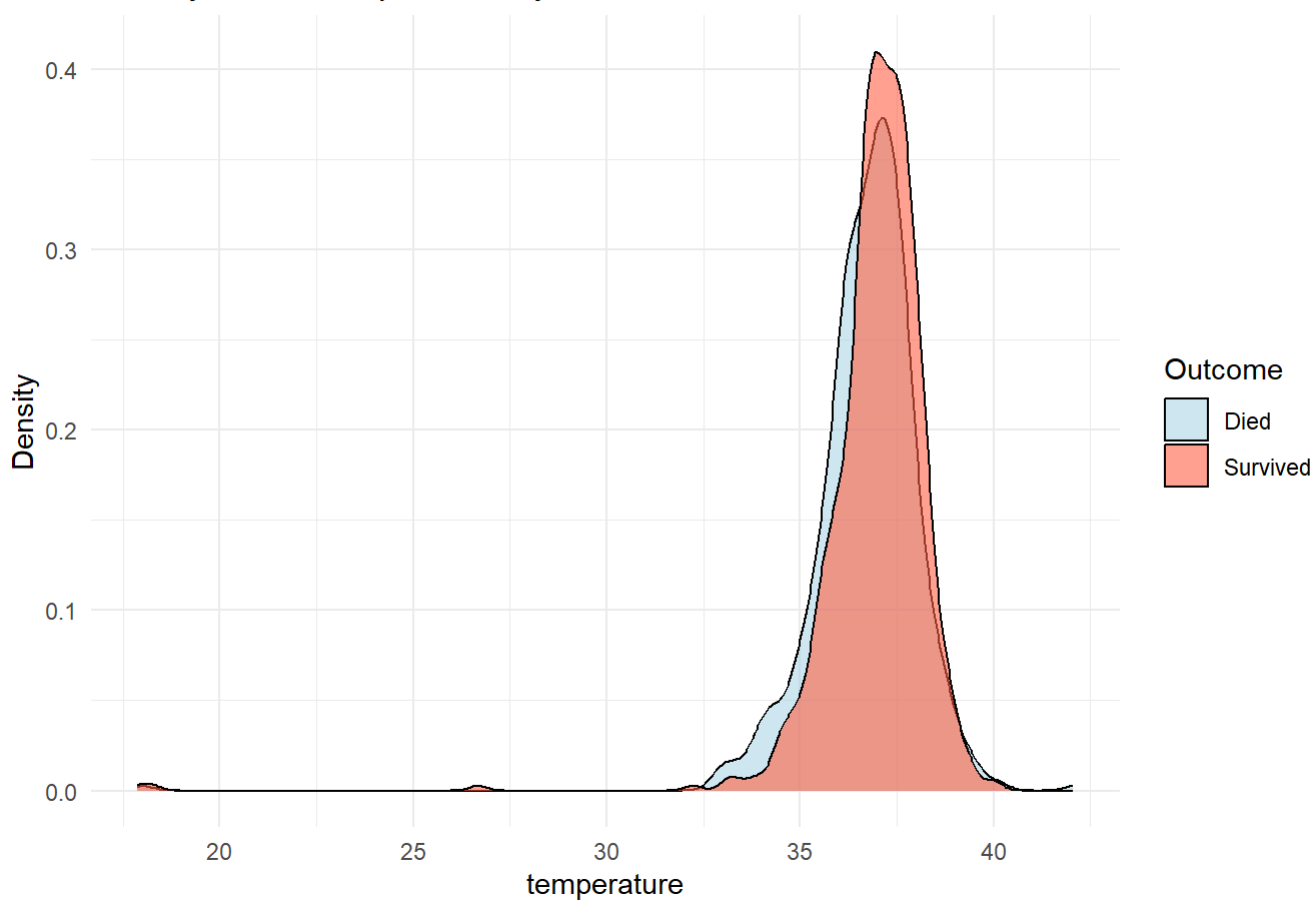


```
##  
## --- Visualizations for temperature ---
```

Boxplot of temperature by Outcome

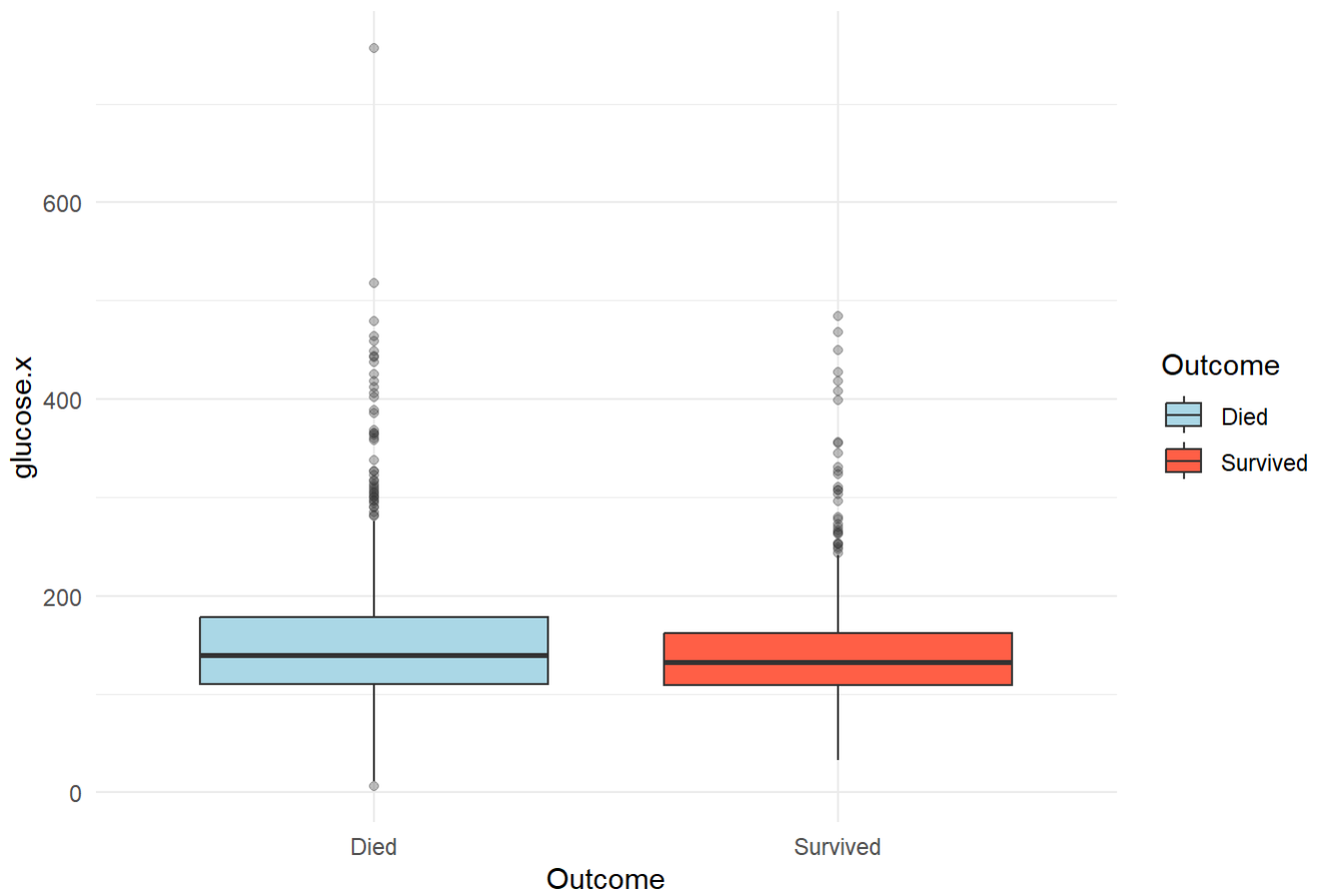


Density Plot of temperature by Outcome

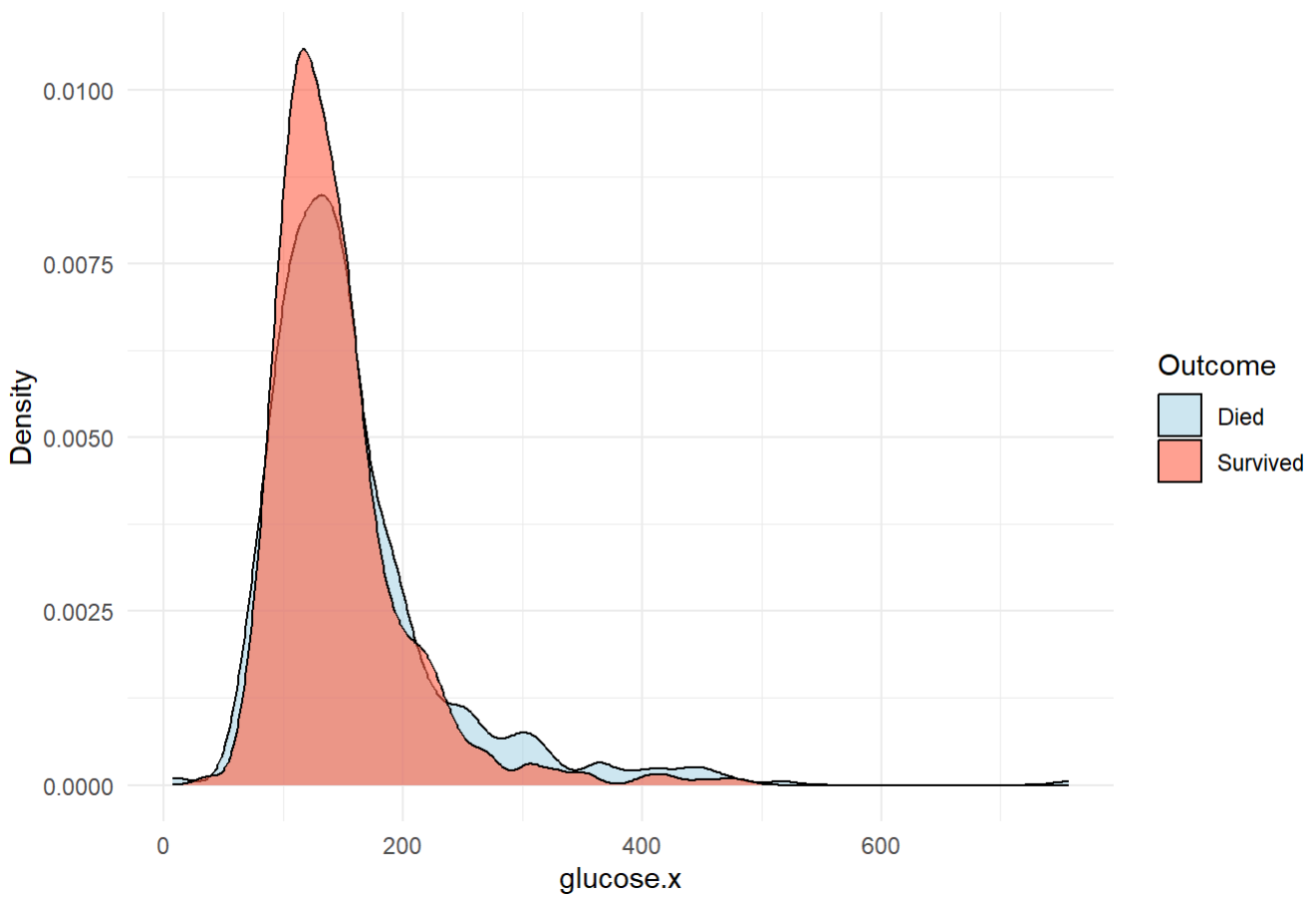


```
##  
## --- Visualizations for glucose.x ---
```

Boxplot of glucose.x by Outcome

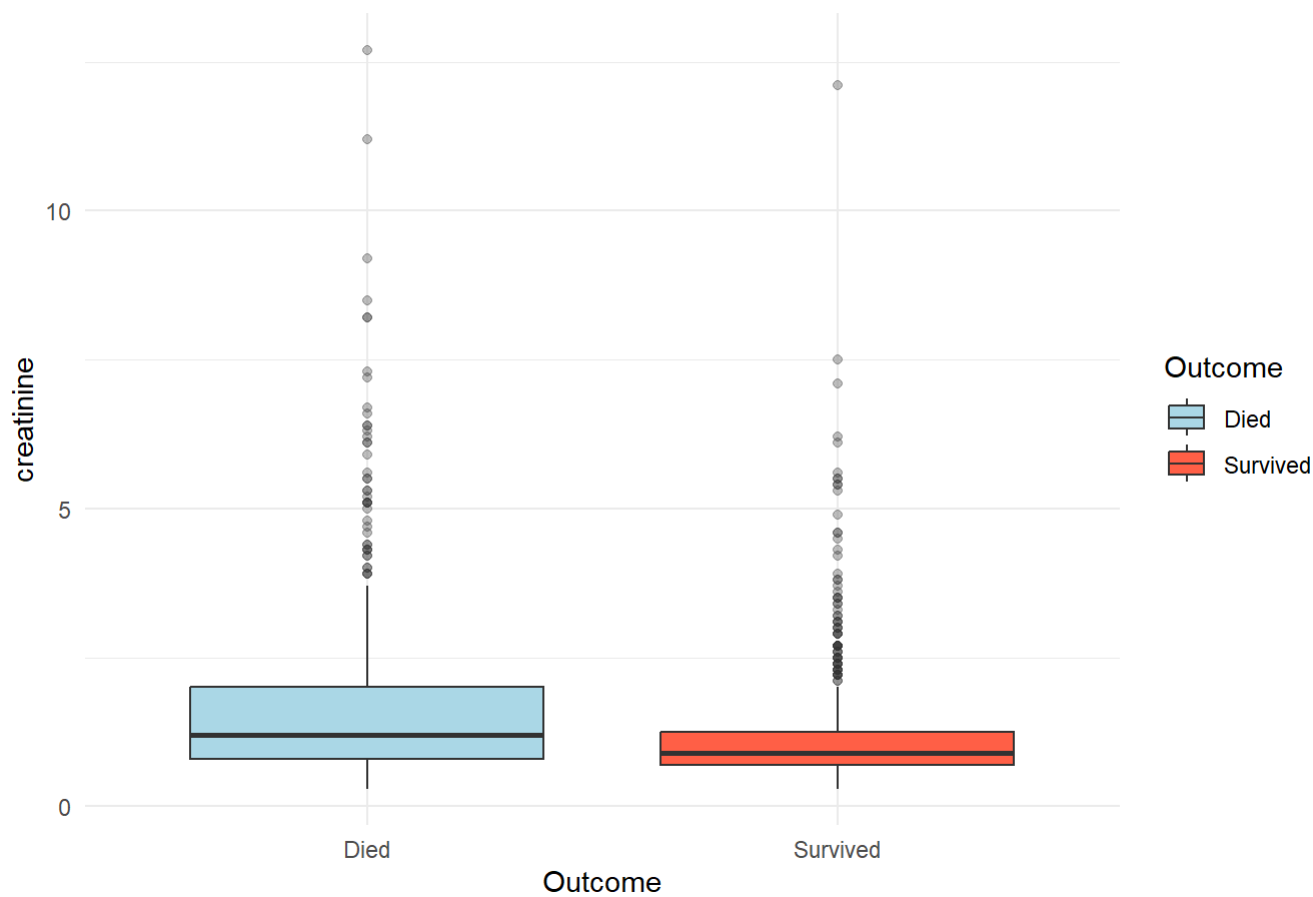


Density Plot of glucose.x by Outcome

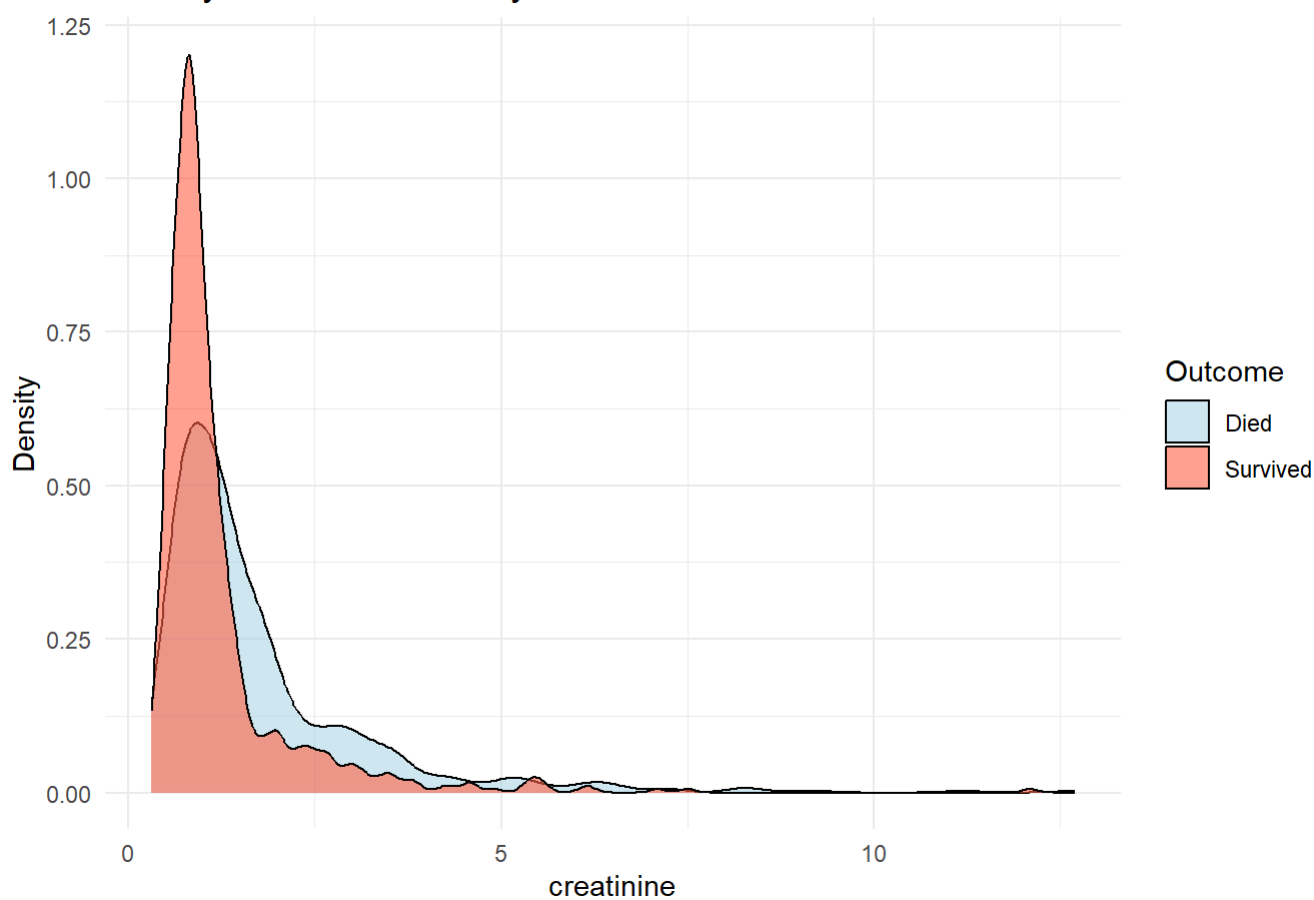


```
##  
## --- Visualizations for creatinine ---
```

Boxplot of creatinine by Outcome

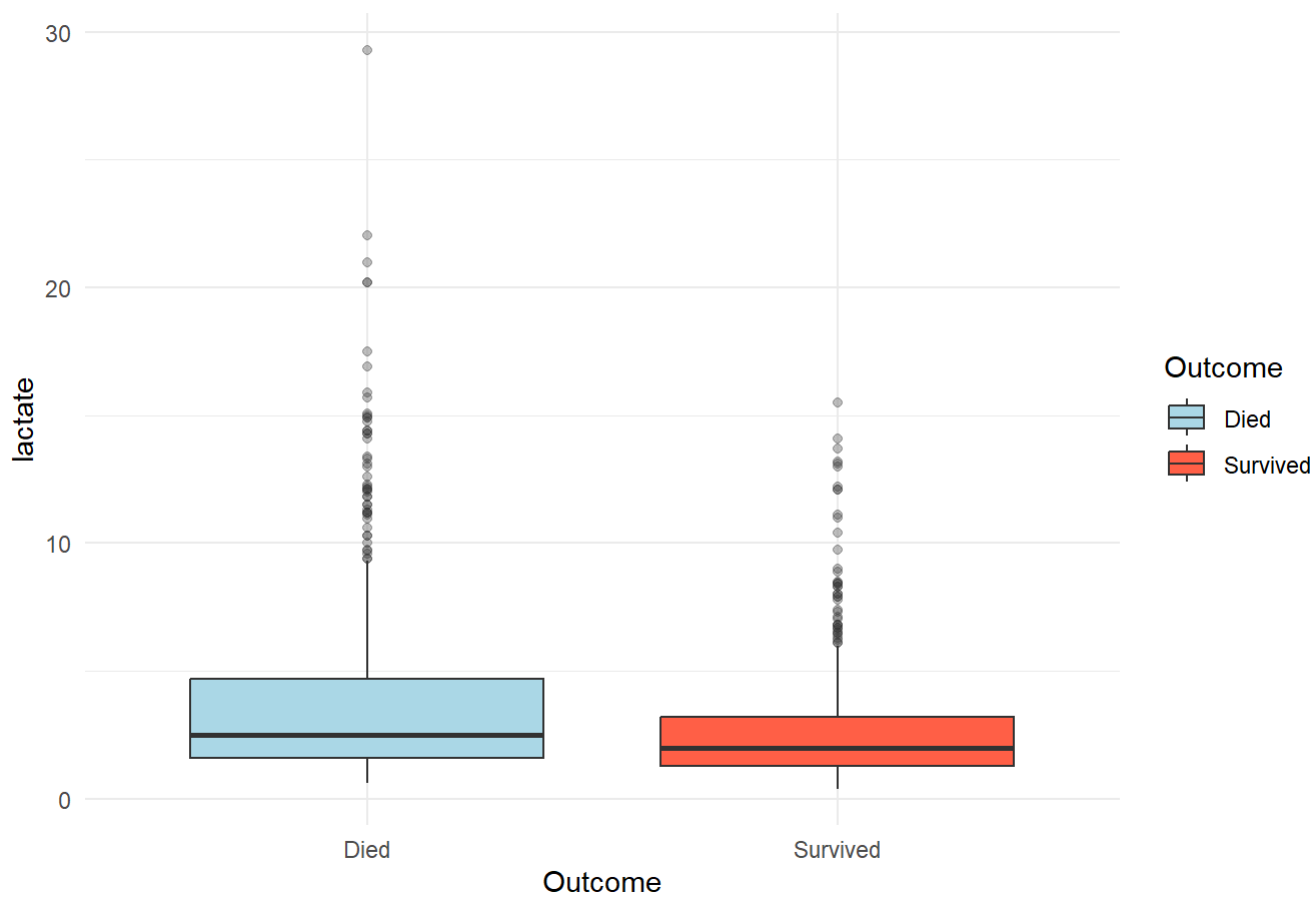


Density Plot of creatinine by Outcome

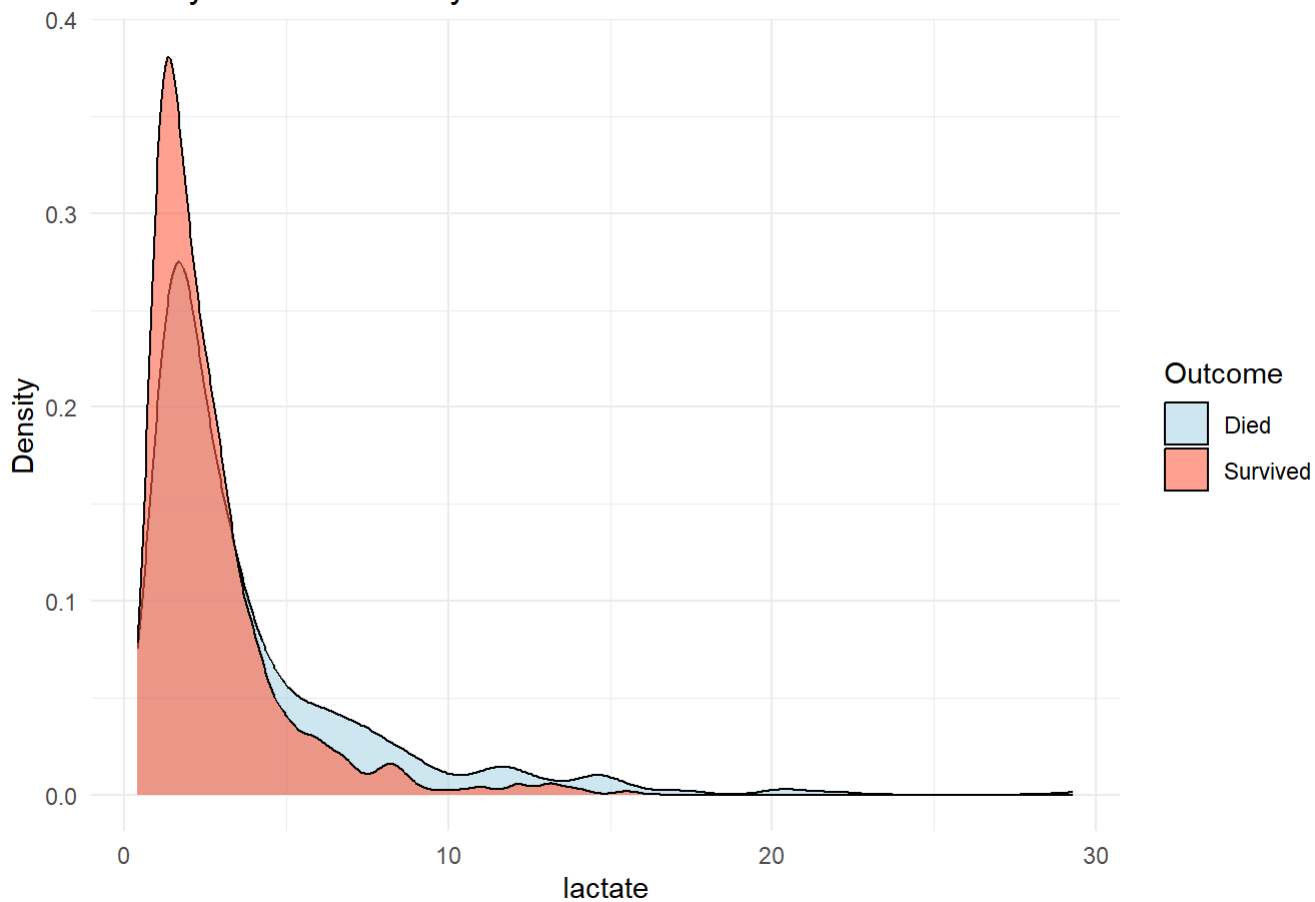


```
##  
## --- Visualizations for lactate ---
```

Boxplot of lactate by Outcome

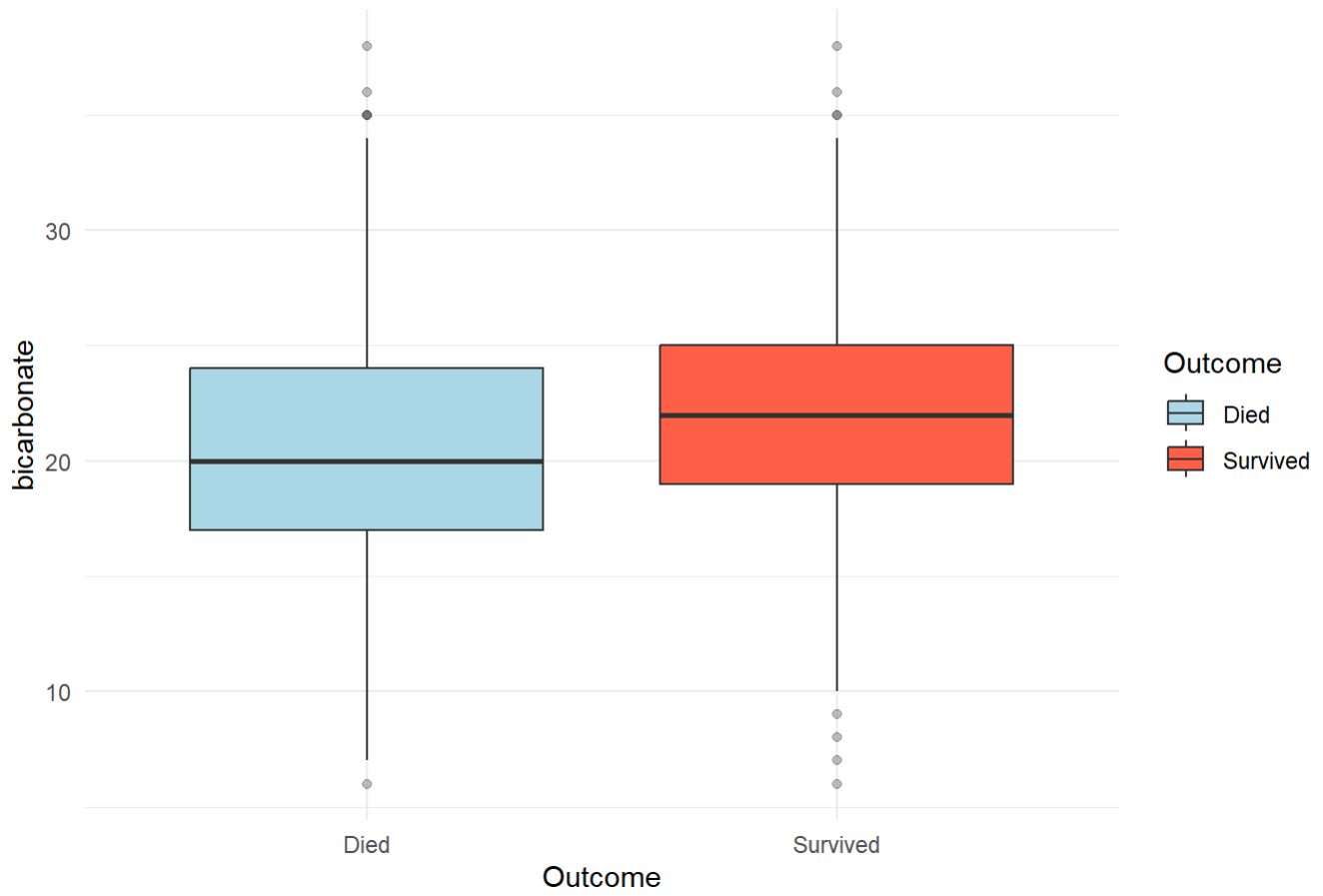


Density Plot of lactate by Outcome

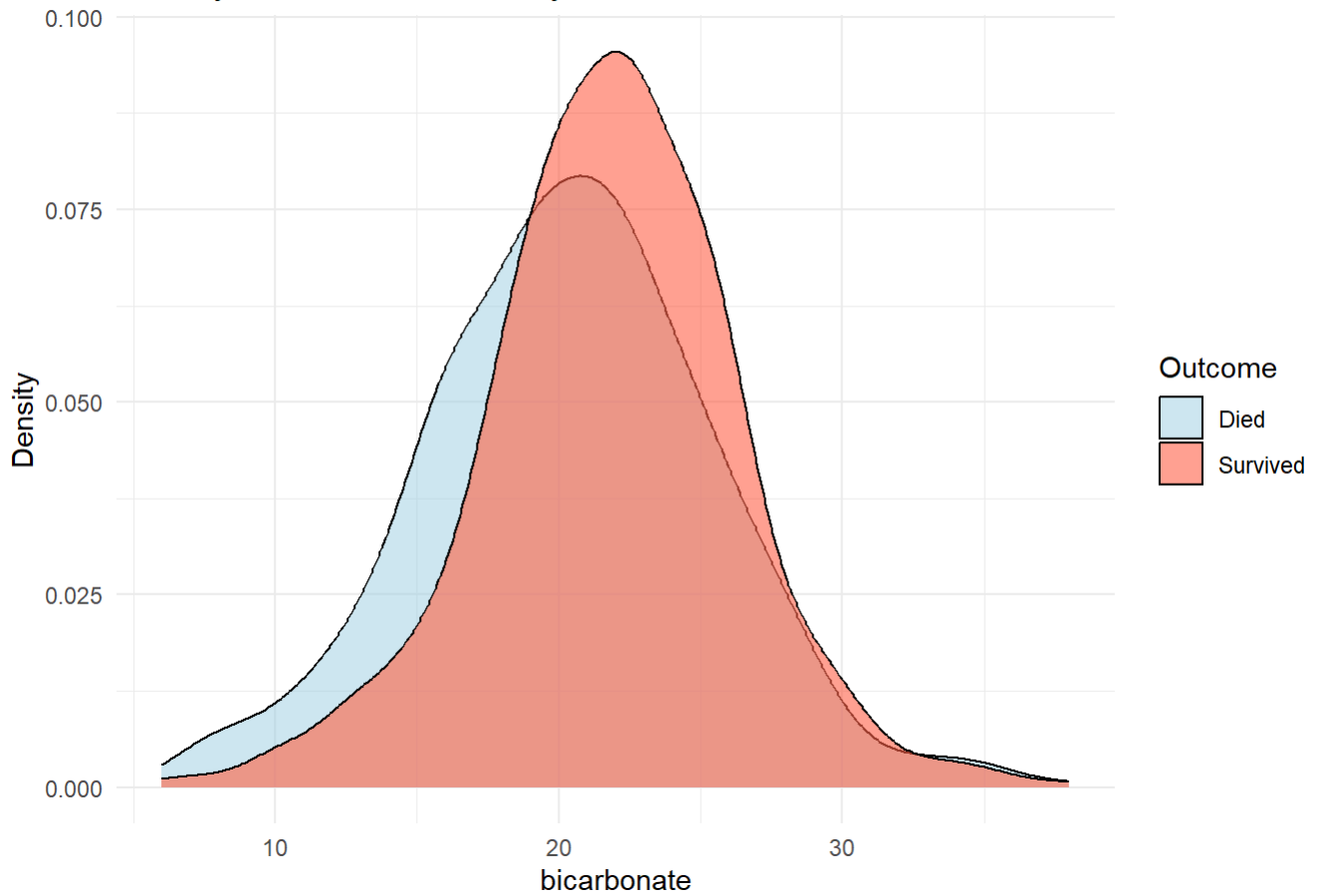


##  
## --- Visualizations for bicarbonate ---

Boxplot of bicarbonate by Outcome



Density Plot of bicarbonate by Outcome



##		Variable	P_Value	Mean_Survived	Mean_Died
## mean in group 01		resprate	0.000000e+00	18.387521	19.325925
## mean in group 05		creatinine	1.166878e-134	1.182105	1.660277
## mean in group 04		glucose.x	8.100044e-126	139.132221	149.799808
## mean in group 06		lactate	1.881095e-114	2.477553	3.483804
## mean in group 0		heartrate	9.081887e-83	84.640268	85.901786
## mean in group 07		bicarbonate	1.564196e-27	23.507164	22.833644
## mean in group 03		temperature	8.359552e-14	37.038217	36.903344
## mean in group 02		spo2	3.004859e-01	102.239292	97.155565

##		Variable	P_Value
## 2		first_careunit	0.000000e+00
## 3		intime_weekdays	1.183170e-142
## 1		gender	3.479749e-83
## 4		is_weekend_admission	1.573501e-69

## Statistical Test Results:

- Variables like `resprate`, `creatinine`, `glucose.x`, and `lactate` show strong statistical significance (p-values close to 0), indicating a clear difference between survivors and non-survivors.
- Less significant variables such as `spo2` (p-value ~0.30) may not contribute significantly to outcome prediction.

## Categorical Variable Analysis:

- Categorical predictors like `first_careunit` and `intime_weekdays` exhibit highly significant associations with mortality (p-values ~0), suggesting these are strong predictors.
- The variable `is_weekend_admission` shows weaker significance but is still worth considering due to its contextual relevance.

```
##
## Call:
## glm(formula = expire_flag.x ~ heartrate + resprate + spo2 + temperature +
##       glucose.x + creatinine + lactate + bicarbonate + gender +
##       first_careunit + is_weekend_admission, family = "binomial",
##       data = master_data)
##
## Coefficients:
##               Estimate Std. Error z value Pr(>|z|)
## (Intercept)      2.774997   3.001960   0.924  0.35528
## heartrate        -0.007743   0.003437  -2.253  0.02427 *
## resprate         -0.013222   0.011494  -1.150  0.24998
## spo2             -0.013253   0.023265  -0.570  0.56891
## temperature     -0.024385   0.043306  -0.563  0.57338
## glucose.x        -0.000257   0.001016  -0.253  0.80027
## creatinine        0.303653   0.062077   4.892 1.00e-06 ***
## lactate          0.152457   0.028754   5.302 1.15e-07 ***
## bicarbonate      -0.009335   0.014587  -0.640  0.52220
## genderM          -0.130711   0.124169  -1.053  0.29249
## first_careunitCSRU -0.664066   0.257079  -2.583  0.00979 **
## first_careunitMICU -0.022545   0.245116  -0.092  0.92672
## first_careunitSICU  0.164567   0.256174   0.642  0.52061
## first_careunitTSICU -0.519692   0.254471  -2.042  0.04113 *
## is_weekend_admissionTRUE -0.018291   0.146443  -0.125  0.90060
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##    Null deviance: 1758.7  on 1268  degrees of freedom
## Residual deviance: 1617.6  on 1254  degrees of freedom
##   (因为不存在, 374283个观察量被删除了)
## AIC: 1647.6
##
## Number of Fisher Scoring iterations: 4
```

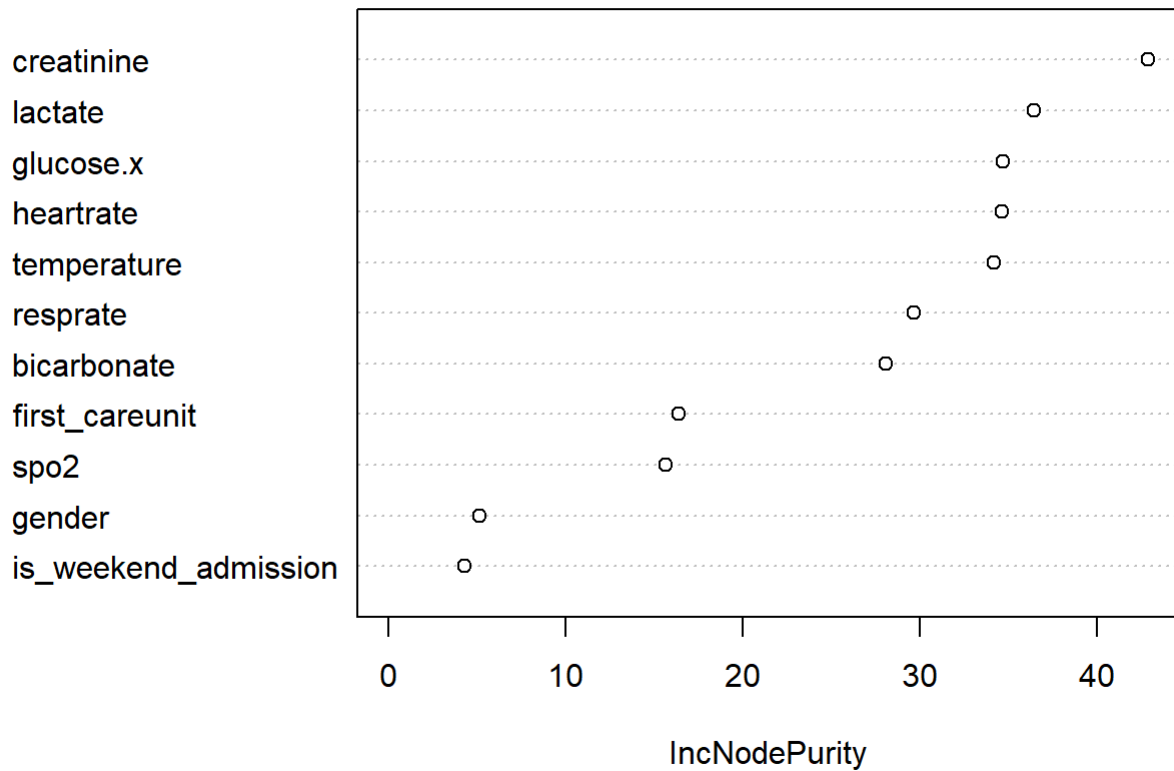
## Logistic Regression:

- Predictors like creatinine, lactate, and certain ICU units ( first\_careunitCSRU and first\_careunitTSICU ) are statistically significant with strong effects.
- Variables such as gender and is\_weekend\_admission are not significant, indicating limited predictive value for mortality.

```
##               IncNodePurity
## heartrate      34.650075
## resprate       29.705450
## spo2           15.685075
## temperature    34.217176
## glucose.x      34.698600
## creatinine     42.932079
## lactate        36.439048
## bicarbonate    28.124834
## gender         5.158534
## first_careunit 16.383881
## is_weekend_admission 4.293866
```



## Random Forest Feature Importance



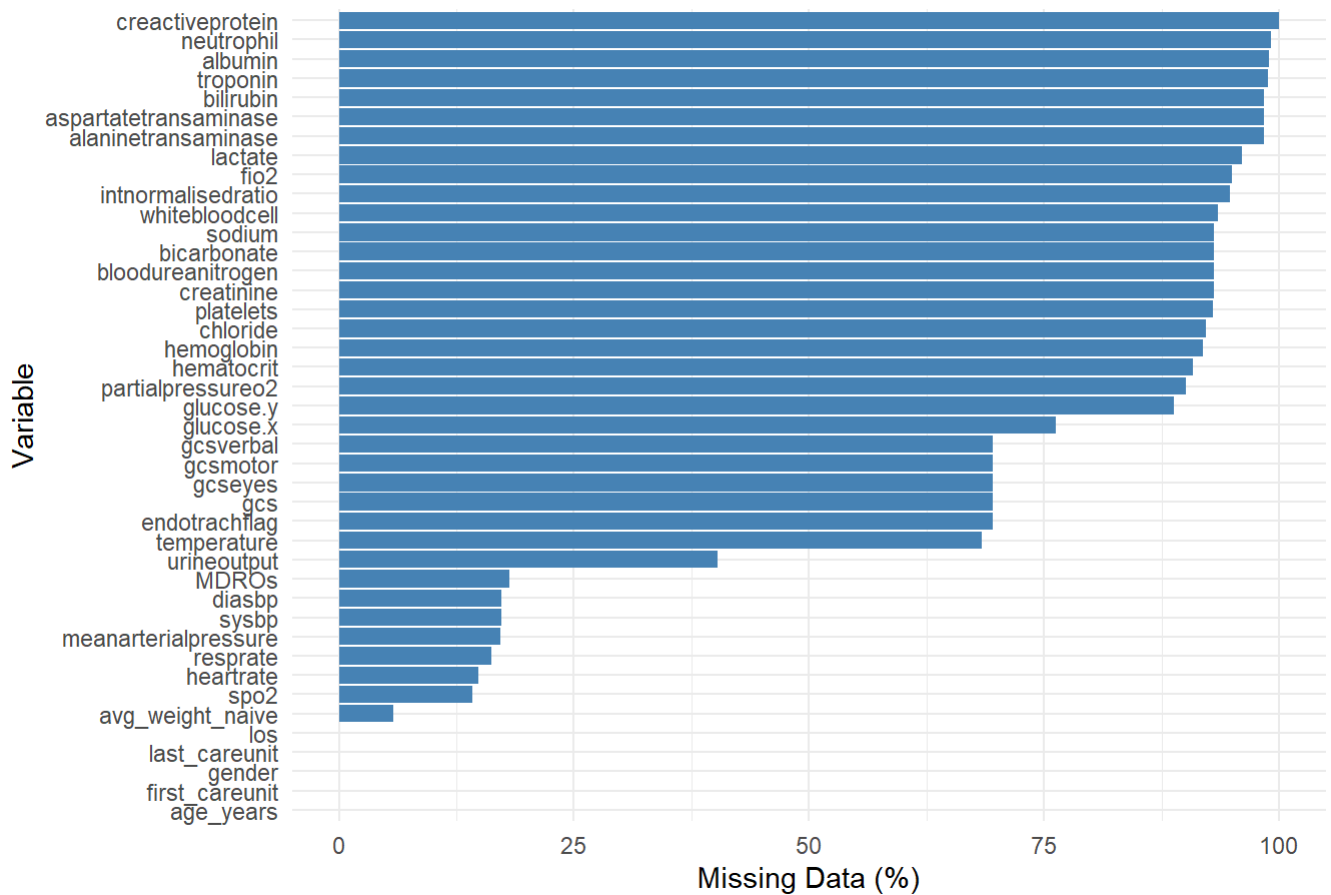
##### **Random Forest Feature Importance:** - Top predictors include `creatinine`, `lactate`, `glucose.x`, and `temperature`, aligning with both t-test and logistic regression results. - Categorical variables like `first_careunit` also play a significant role, reaffirming their importance.

## Observation and insights of step 3.4 results

```
## ### Missing Data Summary for Clinically Important Variables ###
```

##	Variable	Missing_Count	Missing_Pct
## los	los	2	0.00
## age_years	age_years	0	0.00
## avg_weight_naive	avg_weight_naive	21751	5.79
## spo2	spo2	53170	14.16
## fio2	fio2	356903	95.03
## temperature	temperature	256856	68.39
## resprate	resprate	60883	16.21
## heartrate	heartrate	55848	14.87
## sysbp	sysbp	65010	17.31
## diasbp	diasbp	65063	17.32
## glucose.x	glucose.x	286346	76.25
## meanarterialpressure	meanarterialpressure	64372	17.14
## neutrophil	neutrophil	372230	99.12
## creactiveprotein	creactiveprotein	375428	99.97
## whitebloodcell	whitebloodcell	351211	93.52
## partialpressureo2	partialpressureo2	338361	90.10
## bicarbonate	bicarbonate	349620	93.09
## lactate	lactate	360710	96.05
## troponin	troponin	370982	98.78
## bloodureanitrogen	bloodureanitrogen	349457	93.05
## creatinine	creatinine	349354	93.02
## alaninetransaminase	alaninetransaminase	369411	98.36
## aspartatetransaminase	aspartatetransaminase	369409	98.36
## hemoglobin	hemoglobin	345076	91.89
## intnormalisedratio	intnormalisedratio	355970	94.79
## platelets	platelets	349277	93.00
## albumin	albumin	371438	98.90
## chloride	chloride	346295	92.21
## glucose.y	glucose.y	333377	88.77
## sodium	sodium	349594	93.09
## bilirubin	bilirubin	369437	98.37
## hematocrit	hematocrit	341138	90.84
## urineoutput	urineoutput	151427	40.32
## gcs	gcs	260984	69.49
## gcseyes	gcseyes	261112	69.53
## gcsmotor	gcsmotor	261276	69.57
## gcsverbal	gcsverbal	261288	69.57
## MDROs	MDROs	68089	18.13
## endotrachflag	endotrachflag	260984	69.49
## first_careunit	first_careunit	0	0.00
## last_careunit	last_careunit	0	0.00
## gender	gender	0	0.00

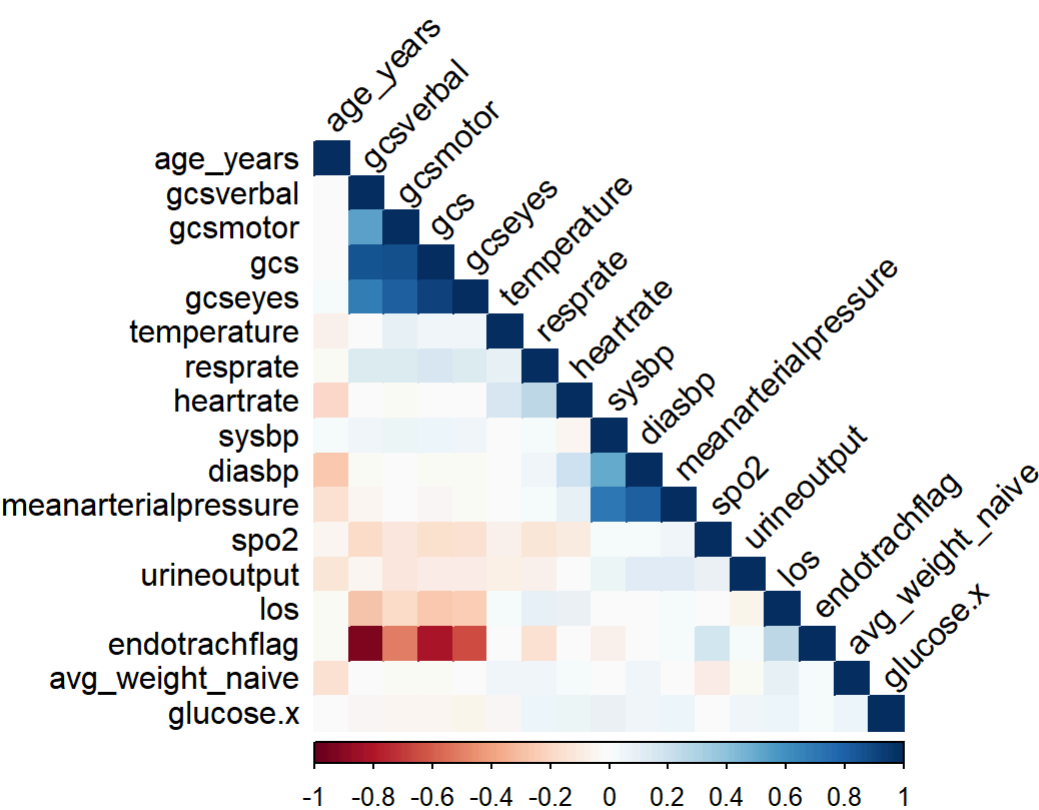
## Missing Data Percentage by Variable



## Missing Data Analysis

- Missing data percentages are clearly calculated and visualized.
- Variables with significant missingness, such as `creactiveprotein` (99.97%) and `neutrophil` (99.12%), highlight potential candidates for exclusion or imputation.

## Correlation Heatmap for Numeric Variables



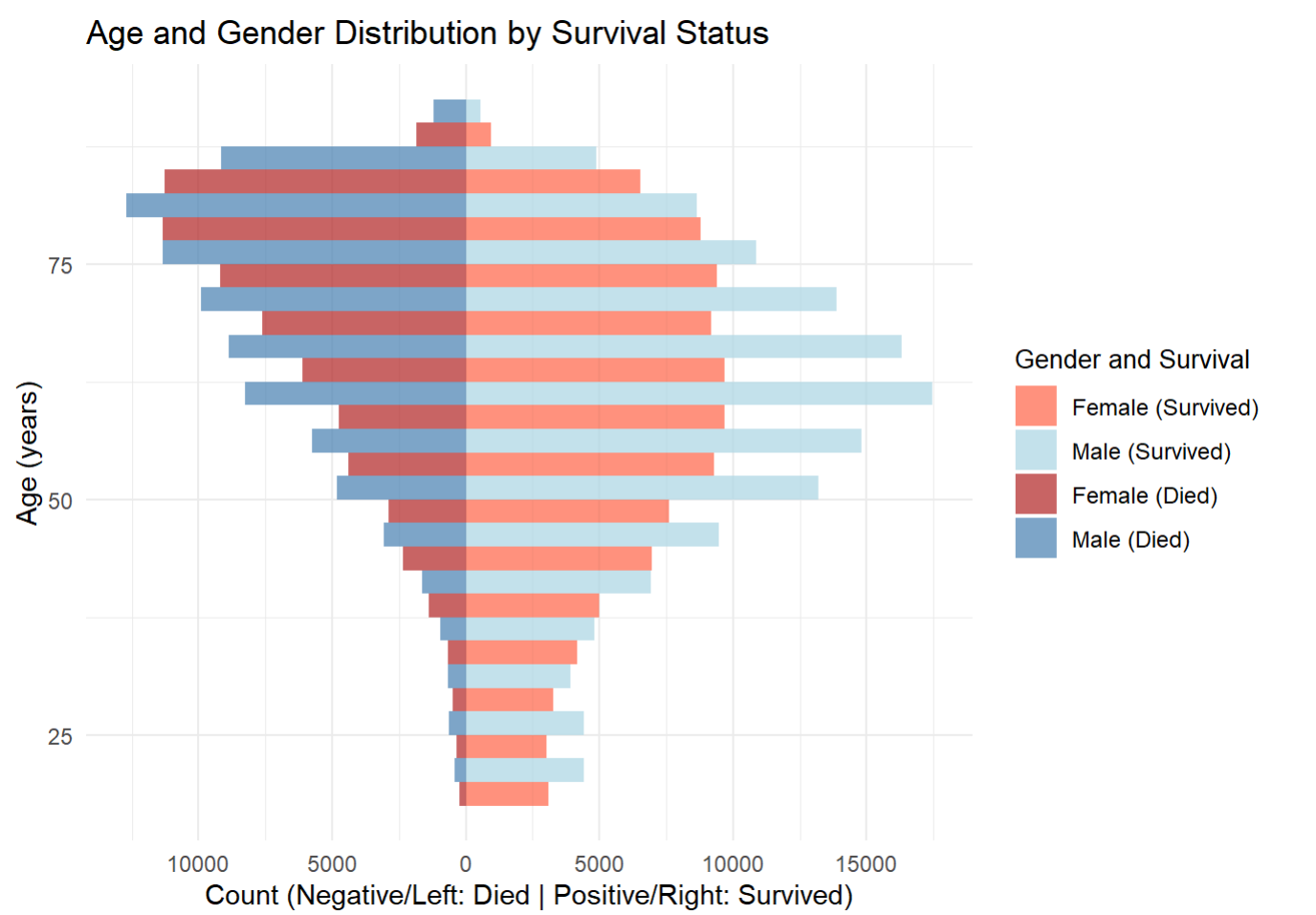
#### **Correlation Analysis** - The correlation heatmap effectively visualizes relationships among numeric variables. - Significant correlations, such as `gcs` and its subcomponents (`gcseyes`, `gcsmotor`, `gcsverbal`) and blood pressure (`sysbp`, `diasbp`, `meanarterialpressure`), are expected due to their clinical relationships. - Variables like `gcs` and its subcomponents might exhibit multicollinearity. Consider removing highly correlated variables before regression or model training. - The negative correlation of `gcsverbal` and `endotrachflag` makes sense as intubation often impairs verbal response.

```
##
## Highly Correlated Numeric Variable Pairs:
##      Var1              Var2 Correlation
## 1   diasbp meanarterialpressure  0.8194698
## 2     gcs              gcseyes  0.9188379
## 3     gcs              gcsmotor  0.8780389
## 4   gcseyes              gcsmotor  0.8150518
## 5     gcs              gcsverbal  0.8593701
## 6     gcs      endotrachflag -0.8035365
## 7 gcsverbal      endotrachflag -0.9378105

## Data filtering and correlation analysis completed.
```

If highly correlated numeric variable pairs found, use Principal Component Analysis (PCA) or select one representative variable from each correlated group to avoid redundancy in predictive modeling.

# Insights of step 3.5 results



```
##
## T-Test for Age by Mortality Status:
```

```
##
## Welch Two Sample t-test
##
## data:  age_years by expire_flag.x
## t = -207.58, df = 345654, p-value < 2.2e-16
## alternative hypothesis: true difference in means between group 0 and group 1 is not equal to 0
## 95 percent confidence interval:
##  -10.71817 -10.51766
## sample estimates:
## mean in group 0 mean in group 1
##      57.94767      68.56558
```

```
##
## Gender Distribution by Mortality:
```

```
##
##      0      1
## F  96636 64974
## M 134557 79385
```

```
##  
## Chi-Squared Test for Gender by Mortality:
```

```
##  
## Pearson's Chi-squared test with Yates' continuity correction  
##  
## data: gender_table  
## X-squared = 373.36, df = 1, p-value < 2.2e-16
```

## Age and gender Distribution:

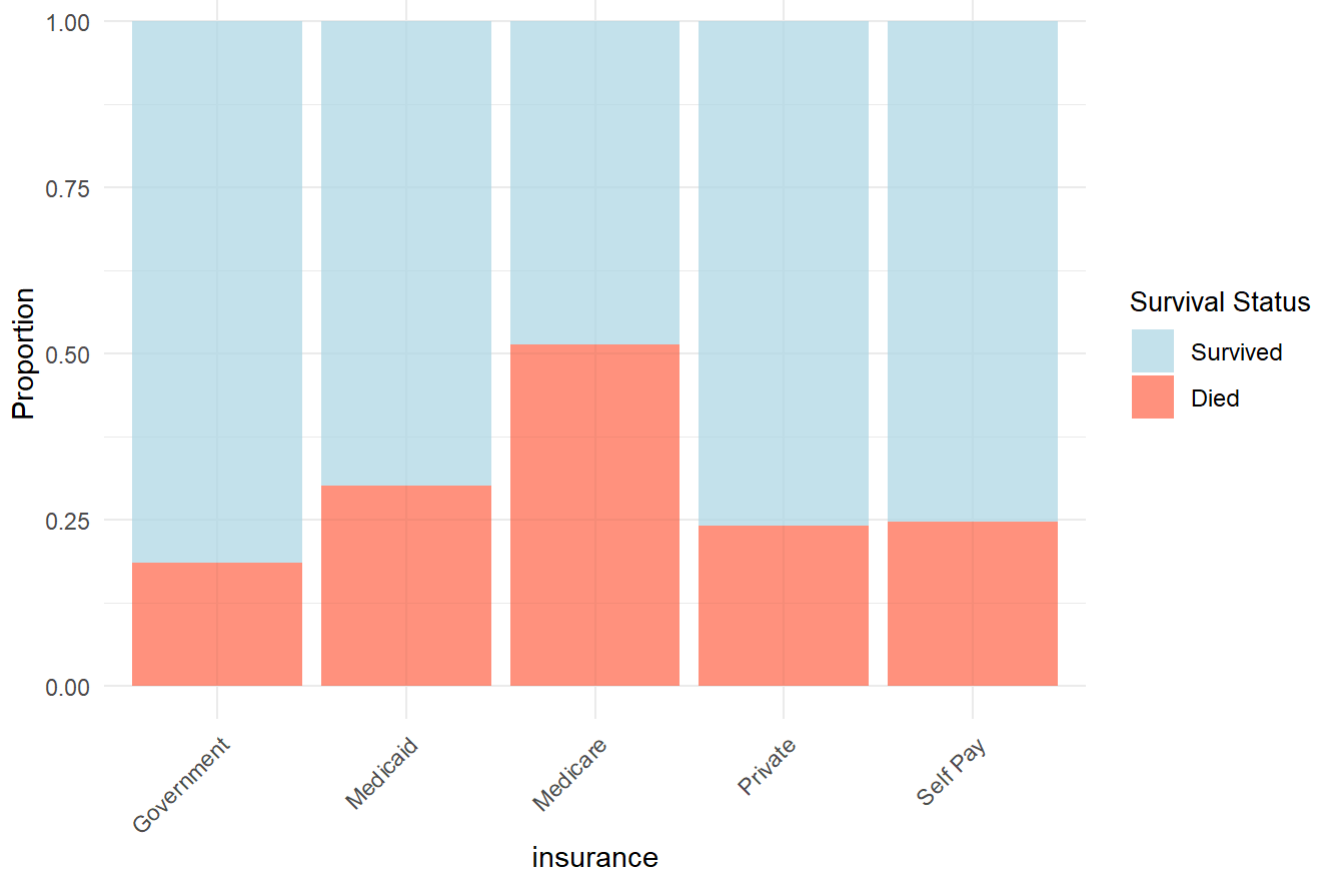
### 1. Visualization:

- Mortality is higher among older age groups for both genders, with non-survivor bars dominating at higher ages.
- Males have a slightly higher proportion of survivors in the younger age groups compared to females.
- The overlap in the middle age range indicates similar mortality rates between genders for these age categories.

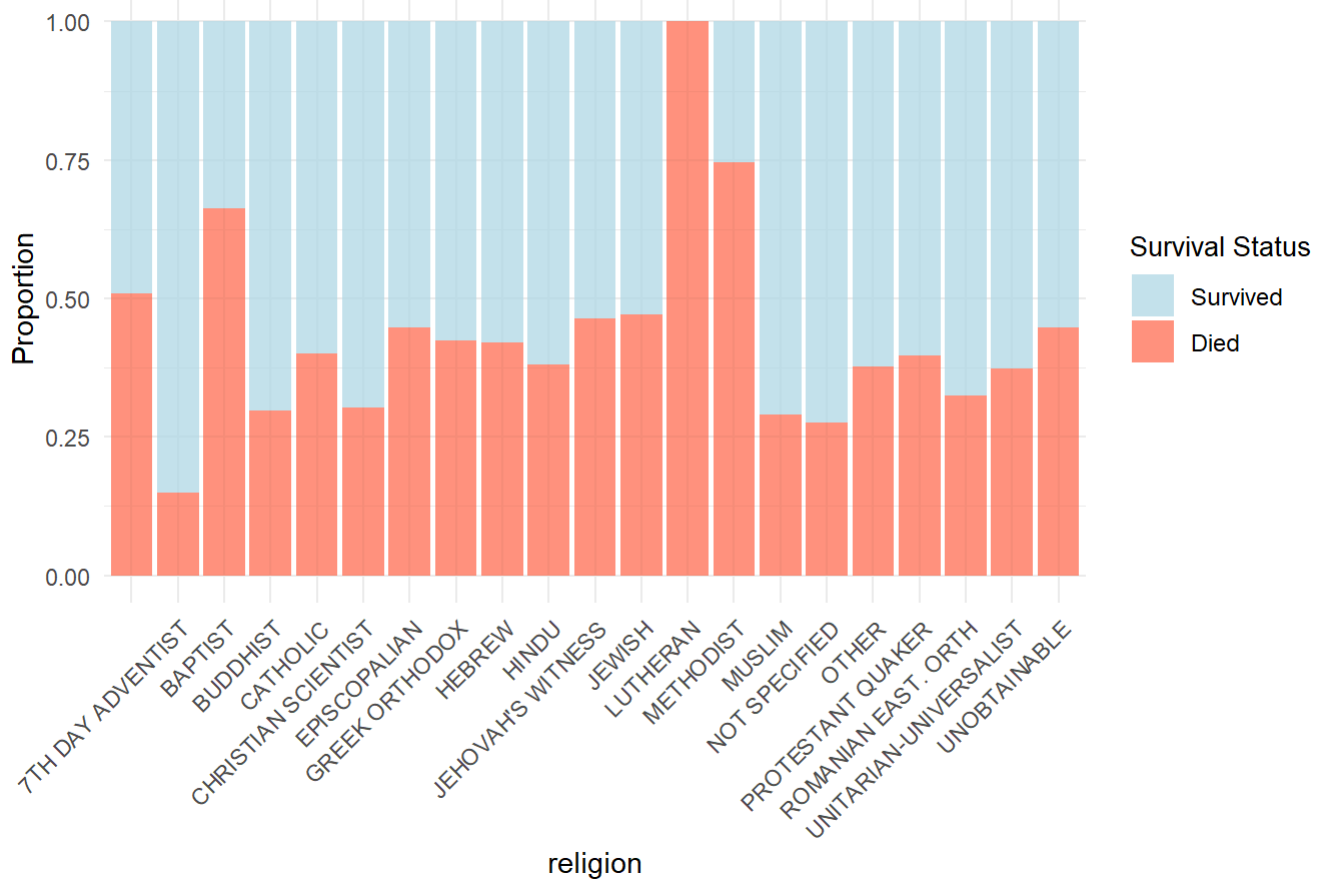
### 2. Statistical Test:

- The Welch Two Sample t-test reveals a statistically significant difference in the mean age of survivors (57.95 years) and non-survivors (68.57 years), with a p-value < 2.2e-16. This highlights that age is a critical factor associated with mortality.
- The mortality is slightly higher among males (59.02% of non-survivors) compared to females (40.98% of non-survivors).
- The Chi-squared test confirms a significant association between gender and mortality (p-value < 2.2e-16). However, the effect size would need further exploration to assess its clinical relevance.

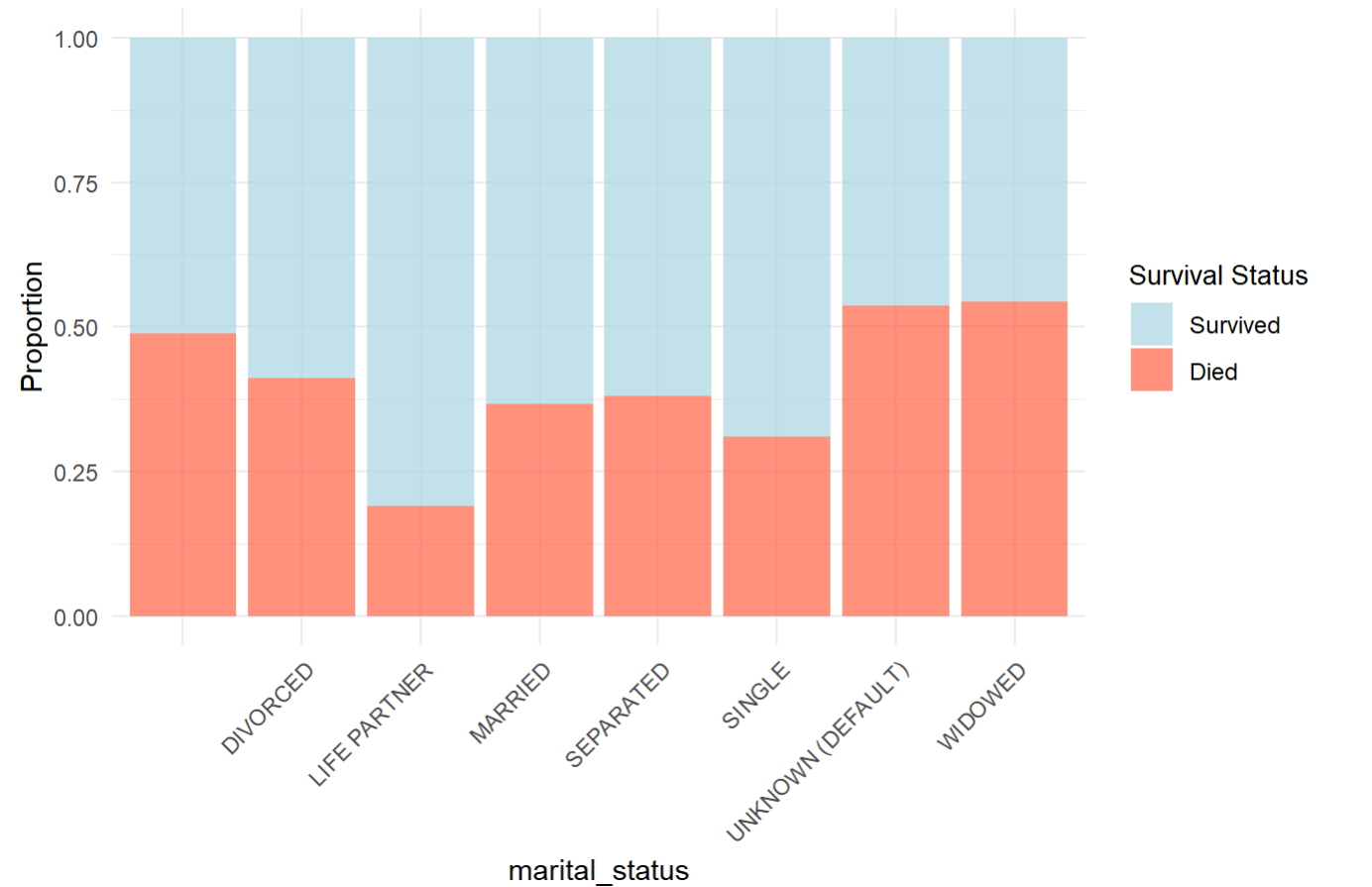
### Distribution of insurance by Mortality Status



### Distribution of religion by Mortality Status



Distribution of marital\_status by Mortality Status



## Language Distribution:



##	Category	Count	Percentage
##	<char>	<int>	<num>
## 1:	ENGL	203603	5.421433e+01
## 2:		140543	3.742305e+01
## 3:	SPAN	8601	2.290229e+00
## 4:	PTUN	4700	1.251491e+00
## 5:	RUSS	4154	1.106105e+00
## 6:	CANT	2666	7.098884e-01
## 7:	PORT	2638	7.024327e-01
## 8:	CAPE	1690	4.500043e-01
## 9:	MAND	1041	2.771920e-01
## 10:	HAIT	1027	2.734641e-01
## 11:	VIET	755	2.010374e-01
## 12:	ITAL	665	1.770727e-01
## 13:	GREE	520	1.384628e-01
## 14:	ARAB	303	8.068124e-02
## 15:	AMER	269	7.162790e-02
## 16:	PERS	267	7.109535e-02
## 17:	HIND	224	5.964554e-02
## 18:	CAMB	184	4.899455e-02
## 19:	POLI	157	4.180513e-02
## 20:	KORE	155	4.127258e-02
## 21:	*BEN	122	3.248551e-02
## 22:	ETHI	121	3.221924e-02
## 23:	FREN	104	2.769257e-02
## 24:	ALBA	98	2.609492e-02
## 25:	LAOT	98	2.609492e-02
## 26:	THAI	77	2.050315e-02
## 27:	*ARM	76	2.023688e-02
## 28:	*GUJ	50	1.331374e-02
## 29:	JAPA	49	1.304746e-02
## 30:	*BUL	49	1.304746e-02
## 31:	SOMA	28	7.455692e-03
## 32:	*URD	27	7.189417e-03
## 33:	*DUT	25	6.656868e-03
## 34:	TURK	25	6.656868e-03
## 35:	*FAR	24	6.390593e-03
## 36:	*TEL	24	6.390593e-03
## 37:	*NEP	24	6.390593e-03
## 38:	TAGA	24	6.390593e-03
## 39:	*TOI	24	6.390593e-03
## 40:	*KHM	24	6.390593e-03
## 41:	*PUN	24	6.390593e-03
## 42:	** T	24	6.390593e-03
## 43:	*MAN	24	6.390593e-03
## 44:	*PHI	24	6.390593e-03
## 45:	* BE	24	6.390593e-03
## 46:	*TOY	24	6.390593e-03
## 47:	*YOR	24	6.390593e-03
## 48:	*YID	24	6.390593e-03
## 49:	*ARA	24	6.390593e-03
## 50:	BENG	24	6.390593e-03
## 51:	* FU	24	6.390593e-03
## 52:	*IBO	5	1.331374e-03
## 53:	*CDI	4	1.065099e-03

## 54:	*HUN	2	5.325494e-04
## 55:	*BUR	2	5.325494e-04
## 56:	URDU	2	5.325494e-04
## 57:	**TO	2	5.325494e-04
## 58:	*AMH	2	5.325494e-04
## 59:	*LEB	2	5.325494e-04
## 60:	*PER	1	2.662747e-04
## 61:	**SH	1	2.662747e-04
## 62:	*SPA	1	2.662747e-04
## 63:	*FIL	1	2.662747e-04
## 64:	*BOS	1	2.662747e-04
## 65:	*ROM	1	2.662747e-04
## 66:	*MOR	1	2.662747e-04
## 67:	SERB	1	2.662747e-04
## 68:	*CAN	1	2.662747e-04
## 69:	*DEA	1	2.662747e-04
## 70:	*FUL	1	2.662747e-04
## 71:	*TAM	1	2.662747e-04
##	Category	Count	Percentage

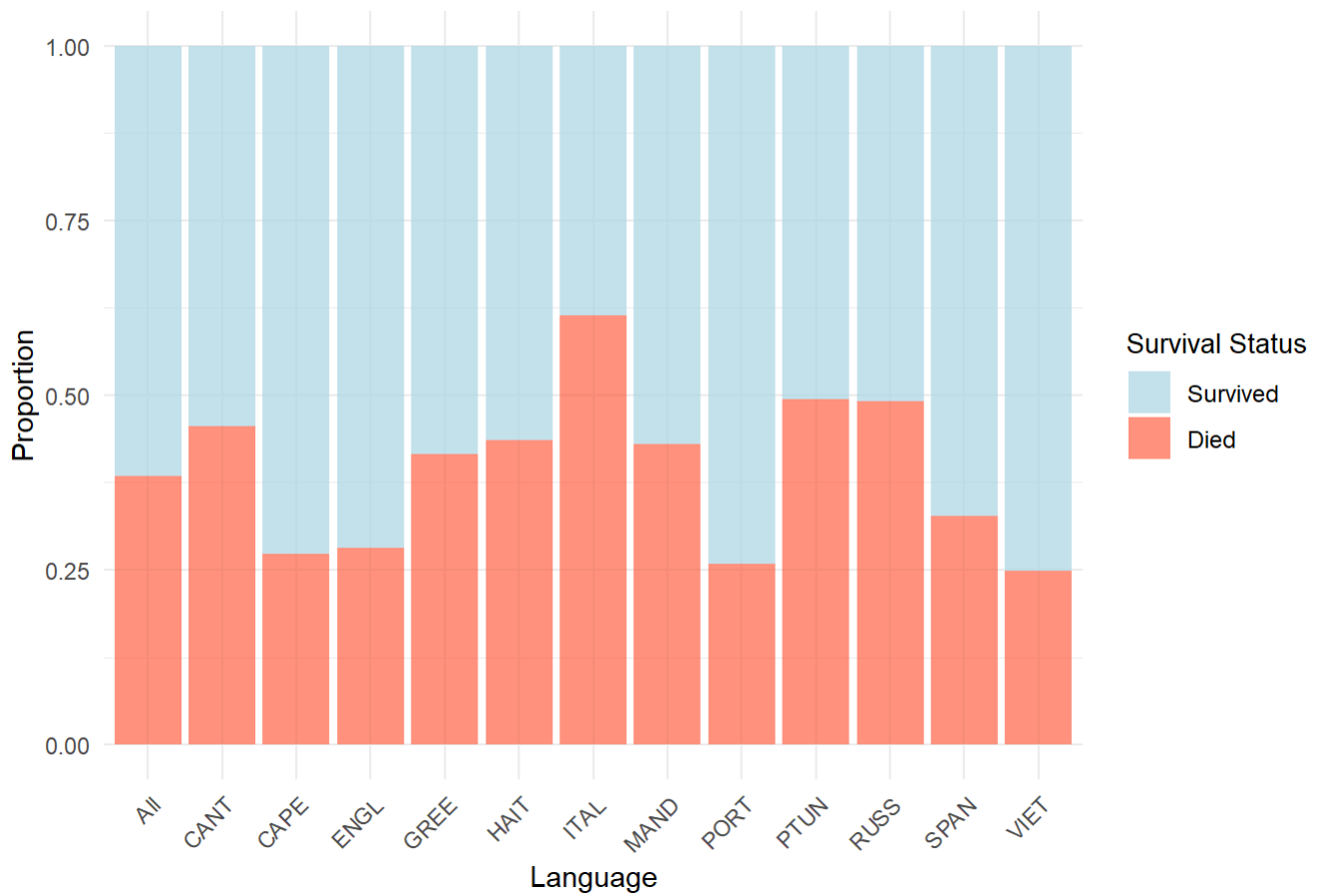
##  
## Ethnicity Distribution:

##	Category	Count
##	<char>	<int>
## 1:	WHITE	261734
## 2:	UNKNOWN/NOT SPECIFIED	37445
## 3:	BLACK/AFRICAN AMERICAN	27384
## 4:	HISPANIC OR LATINO	9905
## 5:	OTHER	9199
## 6:	UNABLE TO OBTAIN	7514
## 7:	ASIAN	5655
## 8:	PATIENT DECLINED TO ANSWER	4142
## 9:	ASIAN - CHINESE	1685
## 10:	HISPANIC/LATINO - PUERTO RICAN	1578
## 11:	BLACK/CAPE VERDEAN	1313
## 12:	MULTI RACE ETHNICITY	837
## 13:	WHITE - RUSSIAN	813
## 14:	HISPANIC/LATINO - DOMINICAN	664
## 15:	BLACK/HAITIAN	660
## 16:	WHITE - OTHER EUROPEAN	634
## 17:	WHITE - BRAZILIAN	520
## 18:	ASIAN - ASIAN INDIAN	419
## 19:	ASIAN - VIETNAMESE	374
## 20:	PORTUGUESE	331
## 21:	BLACK/AFRICAN	301
## 22:	MIDDLE EASTERN	300
## 23:	HISPANIC/LATINO - GUATEMALAN	281
## 24:	WHITE - EASTERN EUROPEAN	204
## 25:	HISPANIC/LATINO - CUBAN	181
## 26:	ASIAN - OTHER	148
## 27:	ASIAN - FILIPINO	128
## 28:	AMERICAN INDIAN/ALASKA NATIVE	127
## 29:	HISPANIC/LATINO - SALVADORAN	127
## 30:	ASIAN - CAMBODIAN	125
## 31:	HISPANIC/LATINO - MEXICAN	124
## 32:	HISPANIC/LATINO - COLOMBIAN	123
## 33:	CARIBBEAN ISLAND	117
## 34:	ASIAN - KOREAN	102
## 35:	NATIVE HAWAIIAN OR OTHER PACIFIC ISLANDER	80
## 36:	HISPANIC/LATINO - CENTRAL AMERICAN (OTHER)	76
## 37:	SOUTH AMERICAN	74
## 38:	ASIAN - JAPANESE	51
## 39:	ASIAN - THAI	26
## 40:	HISPANIC/LATINO - HONDURAN	26
## 41:	AMERICAN INDIAN/ALASKA NATIVE FEDERALLY RECOGNIZED TRIBE	25
##	Category	Count

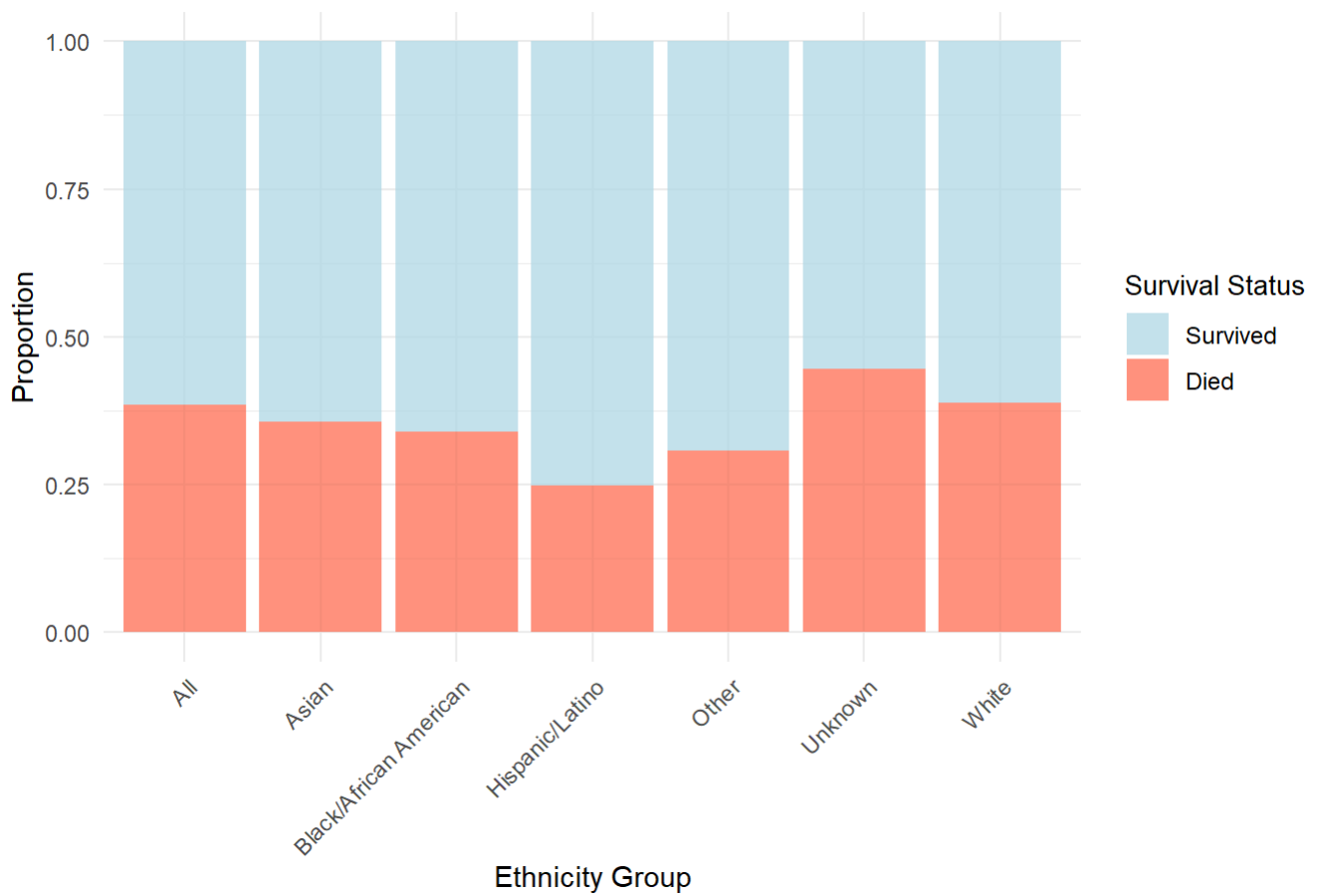
##	Percentage
##	<num>
## 1:	69.693145024
## 2:	9.970656527
## 3:	7.291666667
## 4:	2.637451005
## 5:	2.449461060
## 6:	2.000788173
## 7:	1.505783487
## 8:	1.102909850
## 9:	0.448672887

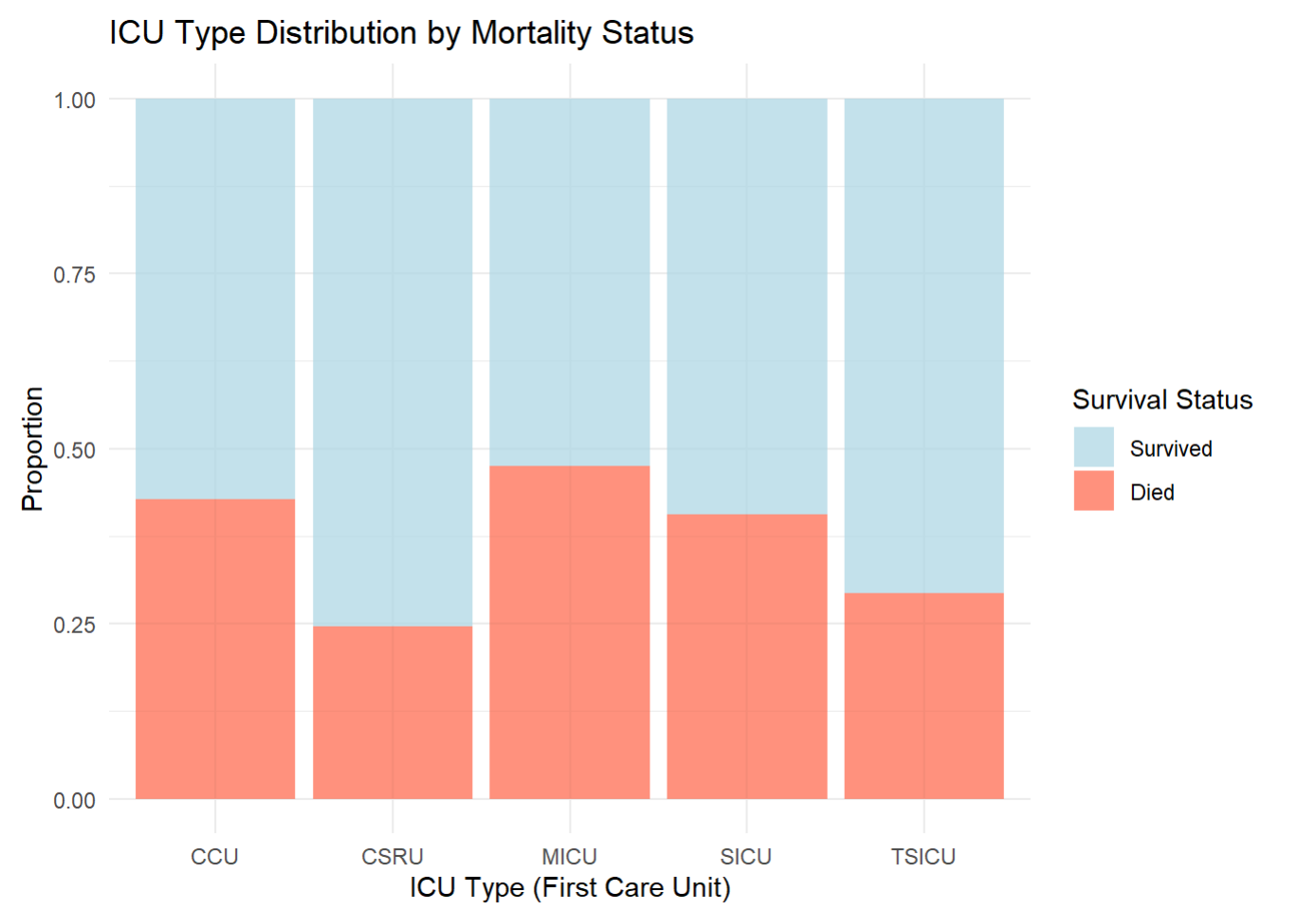
##	10:	0.420181493
##	11:	0.349618695
##	12:	0.222871933
##	13:	0.216481339
##	14:	0.176806408
##	15:	0.175741309
##	16:	0.168818166
##	17:	0.138462849
##	18:	0.111569104
##	19:	0.099586742
##	20:	0.088136929
##	21:	0.080148688
##	22:	0.079882413
##	23:	0.074823194
##	24:	0.054320041
##	25:	0.048195723
##	26:	0.039408657
##	27:	0.034083163
##	28:	0.033816888
##	29:	0.033816888
##	30:	0.033284339
##	31:	0.033018064
##	32:	0.032751789
##	33:	0.031154141
##	34:	0.027160020
##	35:	0.021301977
##	36:	0.020236878
##	37:	0.019704329
##	38:	0.013580010
##	39:	0.006923142
##	40:	0.006923142
##	41:	0.006656868
##		Percentage

Distribution of Language by Mortality Status (With Overall >500)



Distribution of Ethnicity by Mortality Status (Filtered and Grouped)





```
##  
## ICU Type Distribution by Mortality:
```

```
##  
##           0      1  
## CCU   31674 23622  
## CSRU   58908 19283  
## MICU   67127 60734  
## SICU   38033 25961  
## TSICU  35451 14759
```

```
##  
## Chi-Squared Test for ICU Type by Mortality:
```

```
##  
## Pearson's Chi-squared test  
##  
## data:  icu_table  
## X-squared = 12995, df = 4, p-value < 2.2e-16
```

## ICU Characteristics:

### 1. Visualization Insights:

- **CCU (Coronary Care Unit):** Patients in CCU have a moderate mortality rate, reflecting the focused care for cardiac conditions.

- **CSRU (Cardiac Surgery Recovery Unit):** This unit has one of the lowest mortality rates, likely due to the controlled recovery environment post-surgery.
- **MICU (Medical Intensive Care Unit):** The highest mortality rate is observed here, which is expected given its focus on managing critical medical conditions.
- **SICU (Surgical Intensive Care Unit):** Mortality rates are moderate, possibly related to complex post-surgical care cases.
- **TSICU (Trauma/Surgical Intensive Care Unit):** The lowest mortality rate suggests effective care for trauma/surgical emergencies.

## 2. Statistical Analysis:

- The chi-squared test for ICU type by mortality yields a **highly significant result** (p-value < 2.2e-16), indicating a strong association between ICU type and mortality status.

## 3. Clinical Interpretation:

- The results suggest that ICU type is a critical factor influencing patient outcomes.
- MICU patients, likely being the most critically or complexity ill, exhibit a substantially higher mortality rate.
- Further analysis could explore patient characteristics (e.g., age, comorbidities) within each ICU type to better understand these differences.

Weekend vs. Weekday Admissions by Mortality Status



```
##
## Weekend Admission Distribution by Mortality:
```

```
##
##           0           1
##  FALSE 185322 112254
##   TRUE   45871  32105
```

```
##  
## Chi-Squared Test for Weekend Admission by Mortality:
```

```
##  
## Pearson's Chi-squared test with Yates' continuity correction  
##  
## data: weekend_table  
## X-squared = 310.65, df = 1, p-value < 2.2e-16
```

## Weekend vs. Weekday Admissions:

### 1. Visualization Insights:

- **Weekday Admissions (FALSE):** Higher total admissions compared to weekends, with a slightly lower proportion of mortality.
- **Weekend Admissions (TRUE):** Lower total admissions, with a slightly higher mortality proportion compared to weekdays.

### 2. Statistical Analysis:

- The chi-squared test (p-value < 2.2e-16) confirms a statistically significant association between weekend admissions and mortality. This suggests a potential difference in outcomes based on the day of admission.

### 3. Clinical Interpretation:

- The slightly higher mortality proportion for weekend admissions may reflect differences in resource availability, staffing, or severity of cases during weekends. Further investigation into staffing levels, patient profiles, and care processes during weekends is recommended.
- Hospitals could consider optimizing weekend staffing and resources to ensure consistent care quality throughout the week.

---

## Section 4: Feature Engineering

### Objectives:

- Create meaningful features to enhance the predictive power of the dataset.
- Transform raw time-series data into aggregated features suitable for machine learning models.

### Steps:

#### 1. Aggregate Time-Series Data:

- Identify key vitals/lab variables and define clinical thresholds for abnormal values.
- Generate flags for abnormal values (e.g., heart rate > 120 bpm, lactate > 4 mmol/L).

#### 2. Feature Transformation:

- Standardization: Apply z-score normalization to continuous variables (e.g., age, vitals, labs).
- Categorical Encoding: Convert categorical variables (e.g., ICU type, gender) to one-hot encoding.

#### 3. Interaction and Derived Features:

- Interaction Terms: Create interaction terms for meaningful combinations (e.g., age × ICU type, age × lactate).

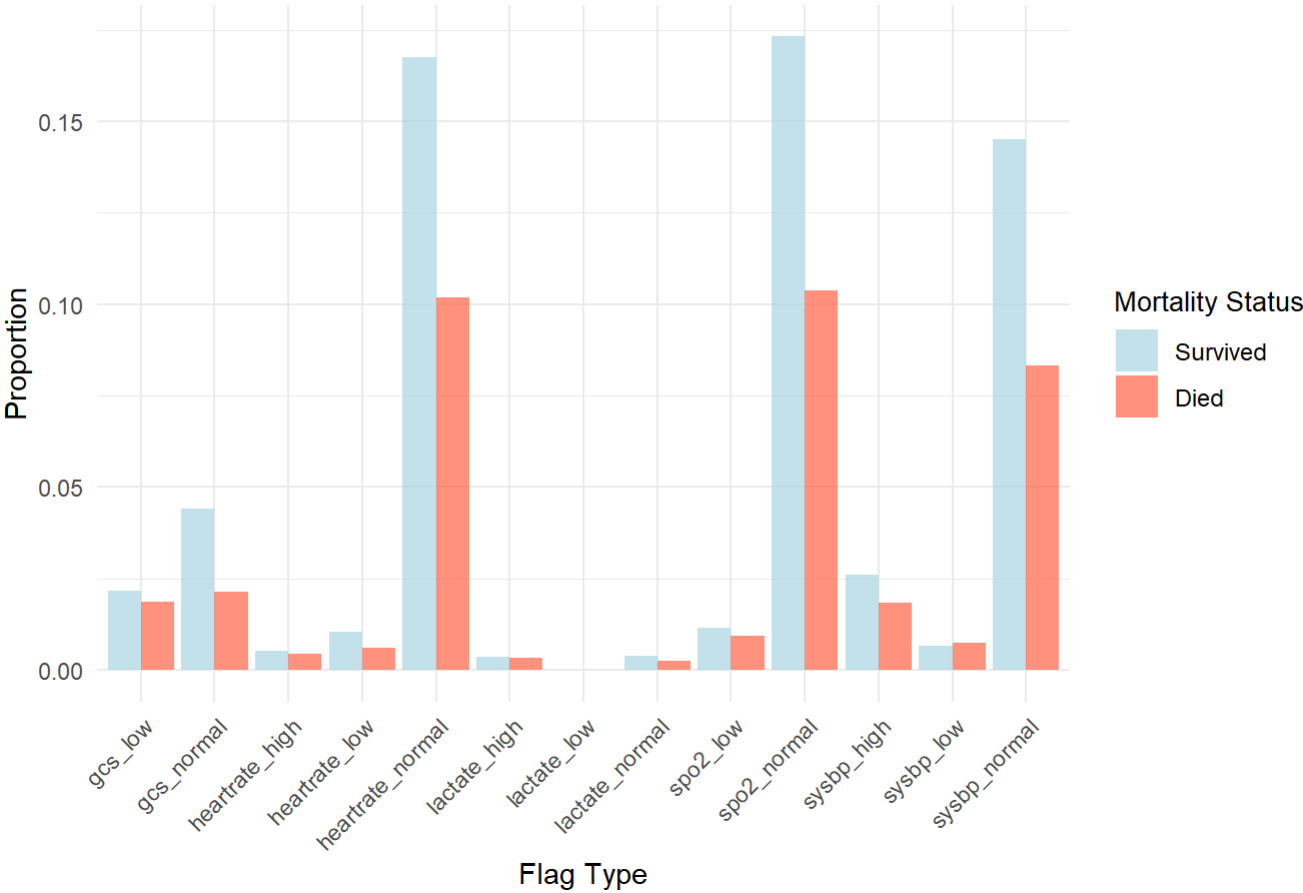


- Calculate clinically relevant ratios, such as:
  - Systolic/diastolic blood pressure (sysbp/diasbp).
  - BUN/creatinine ratio.
- Cumulative Measures: Add aggregate features like total urine output in the first 24 hours.

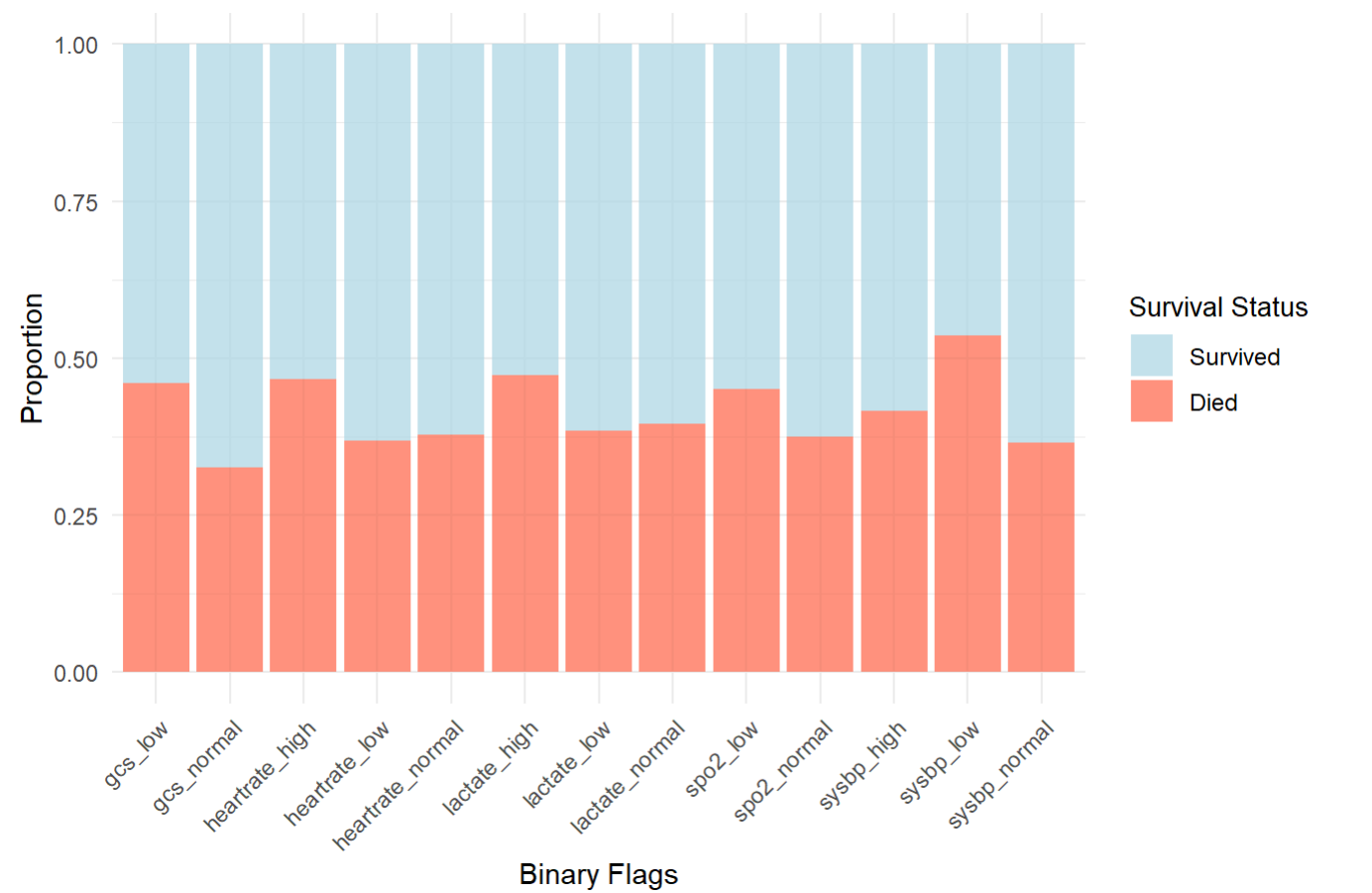
```
## [1] "Summary of Normal and Abnormal Flags:"
```

```
##  heartrate_normal lactate_normal spo2_normal sysbp_normal gcs_normal
## 1          291424          7170      299699      246987      70942
##  heartrate_high lactate_high sysbp_high heartrate_low lactate_low spo2_low
## 1          10503          7659      48259      17777          13      22683
##  sysbp_low gcs_low
## 1       15296   43626
```

Distribution of Normal and Abnormal Flags by Mortality Status

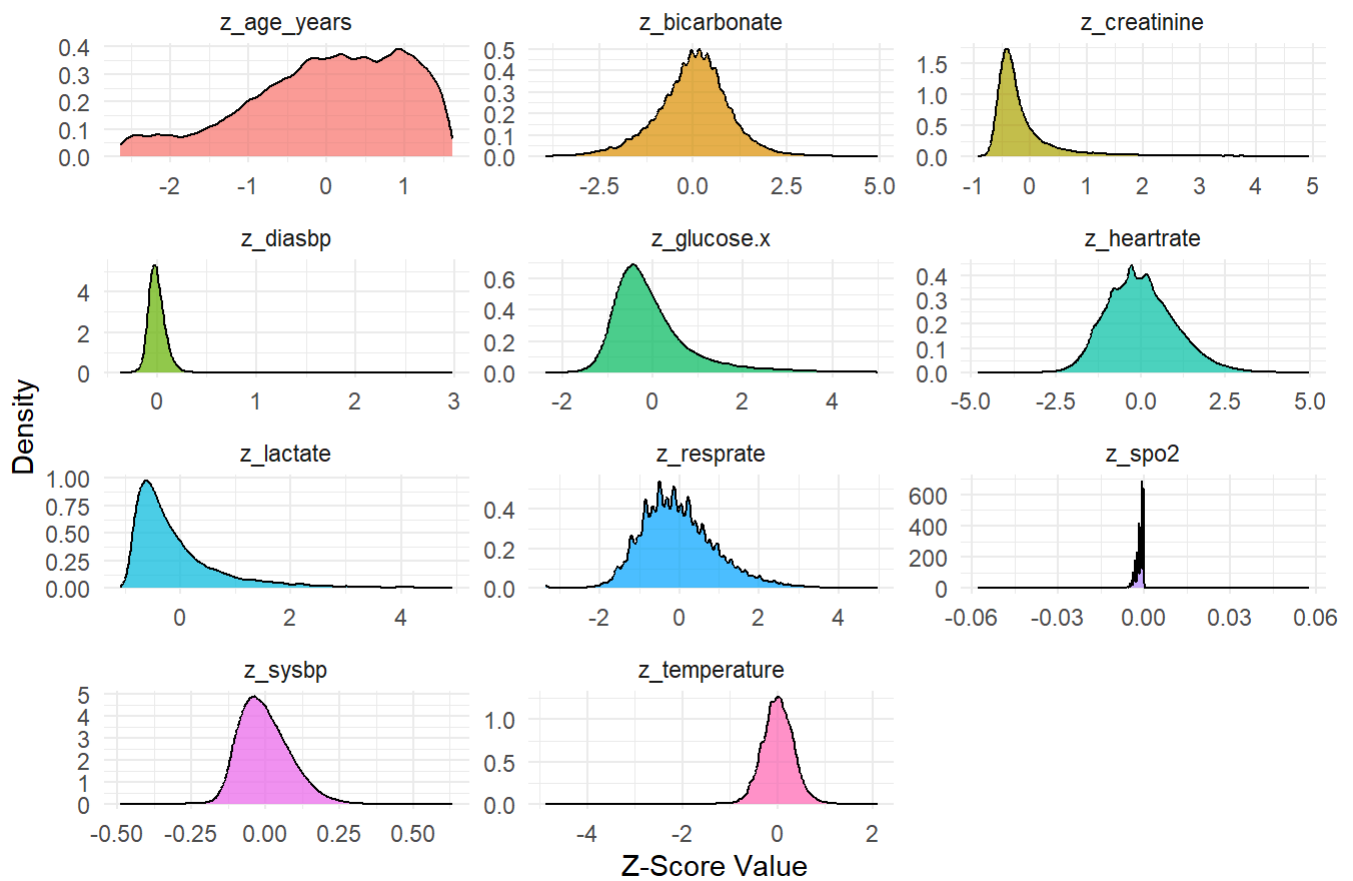


Distribution of Normal and Abnormal Flags by Mortality Status

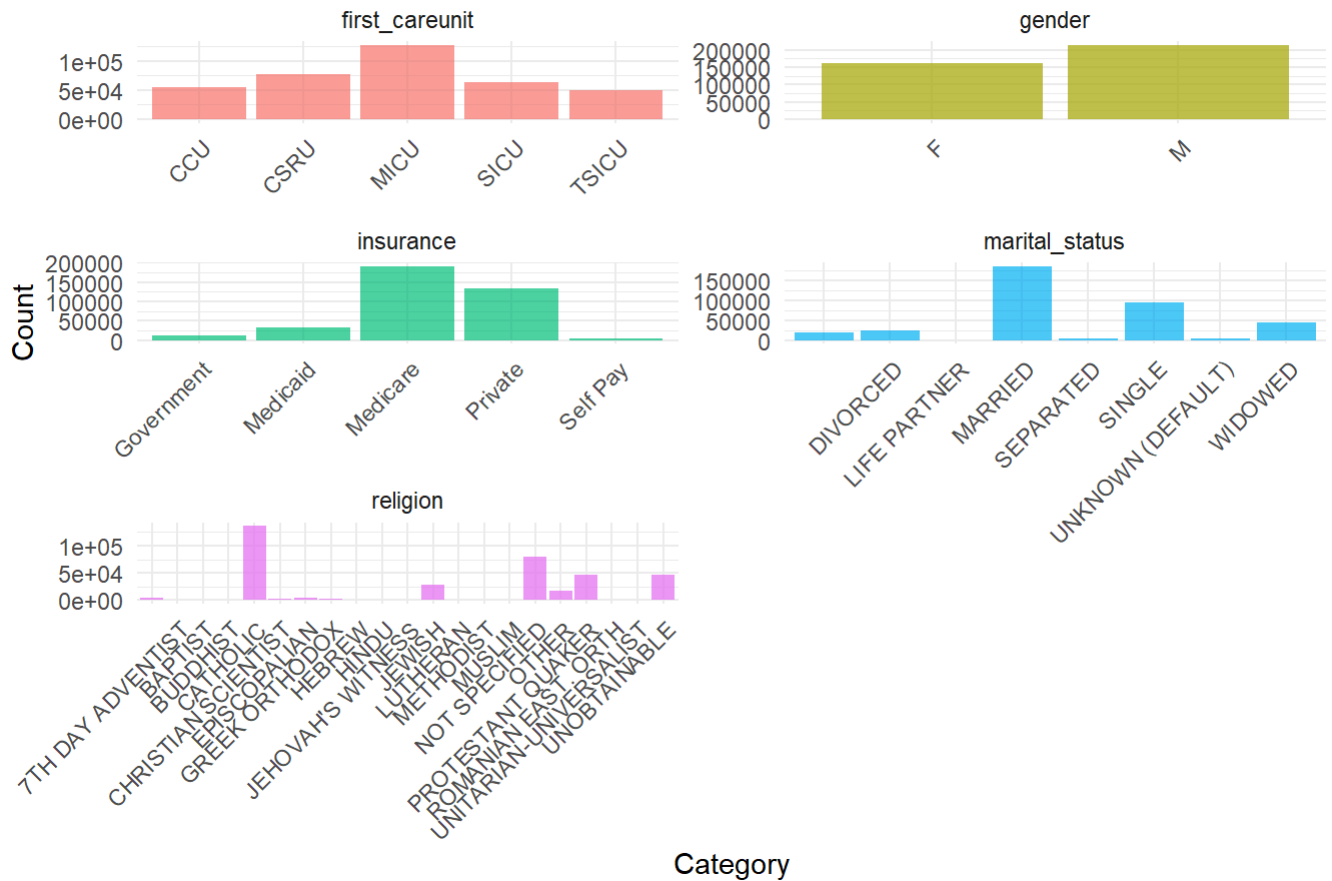


```
## Feature transformation completed: Standardized continuous variables and one-hot encoded categorical variables.
```

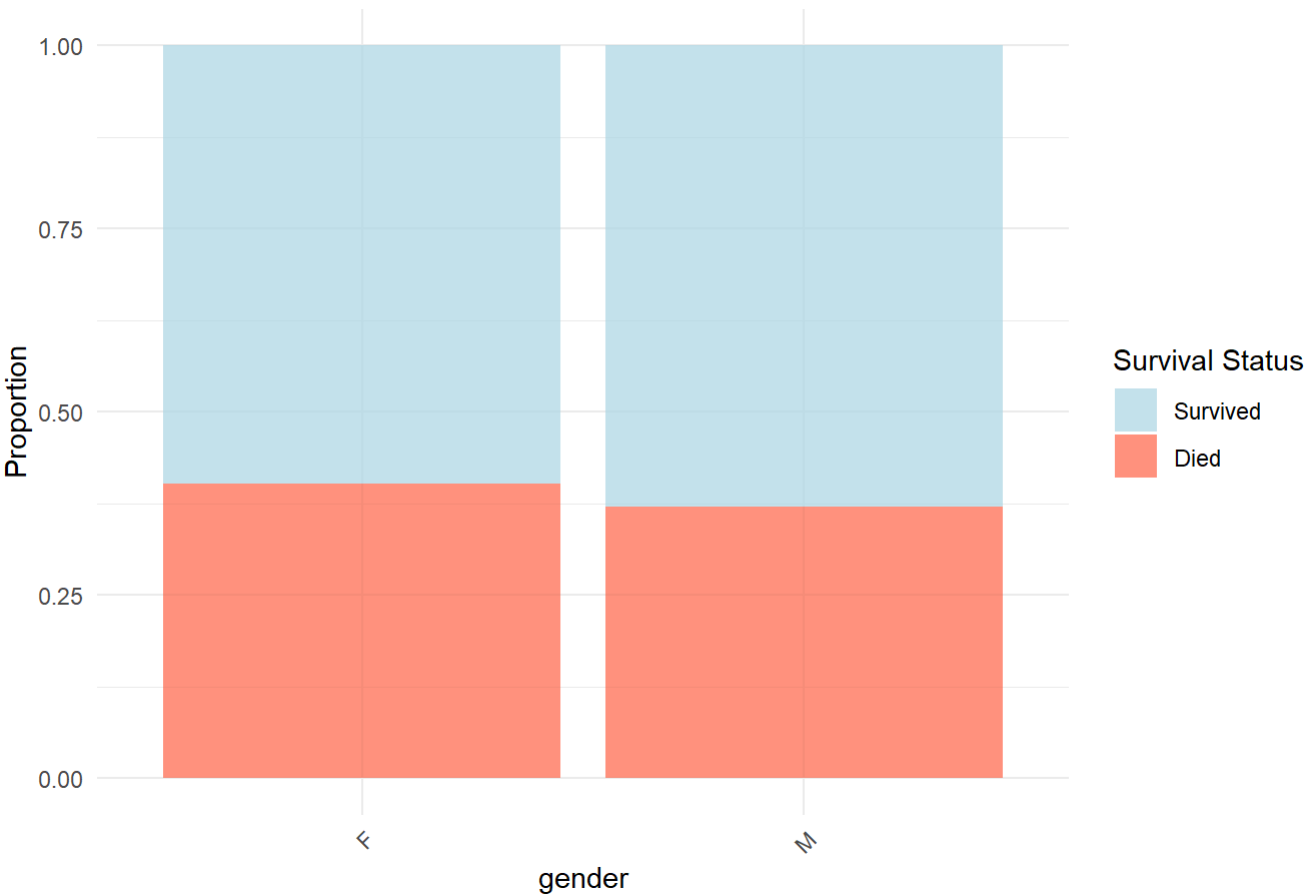
## Distribution of Standardized Variables



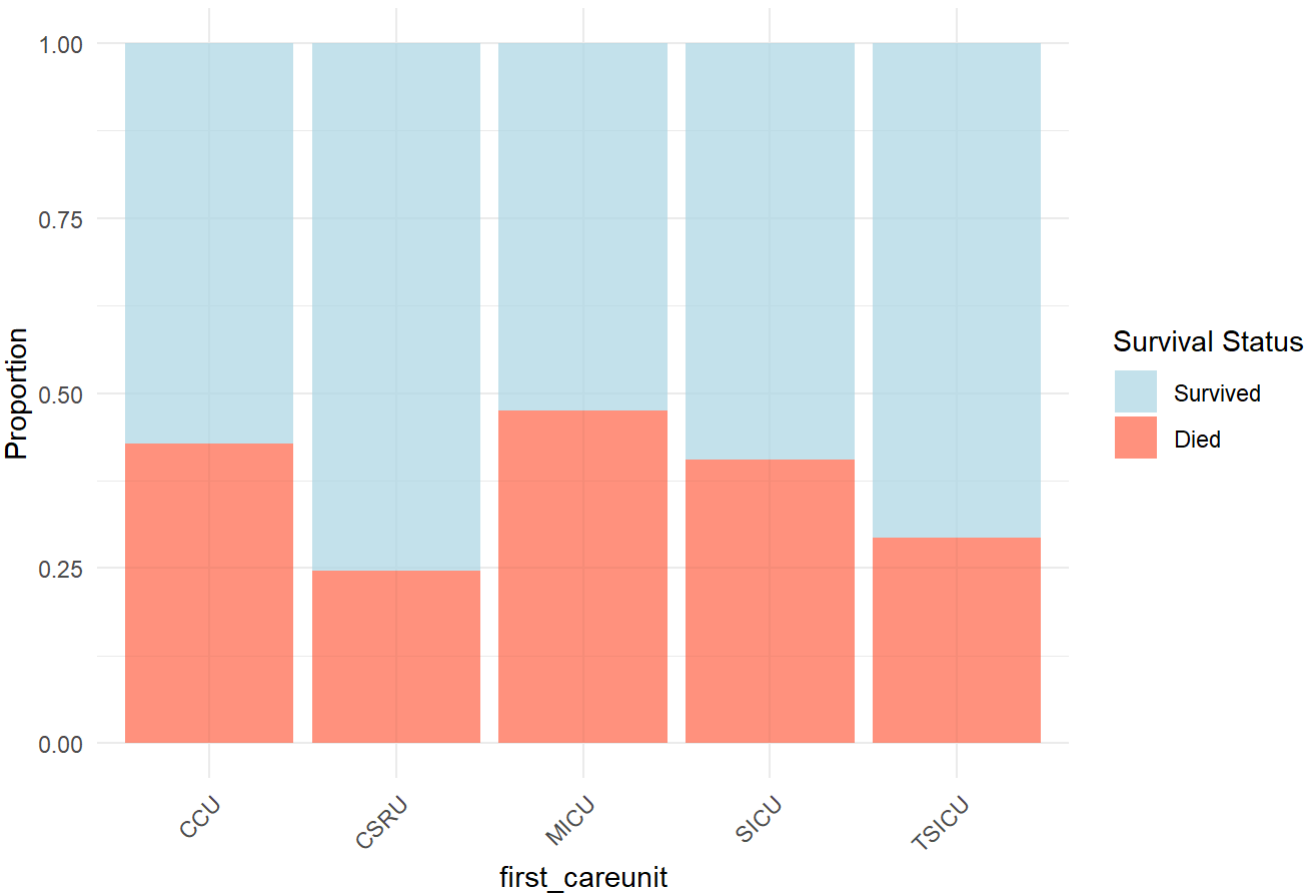
## Counts of One-Hot Encoded Categorical Variables



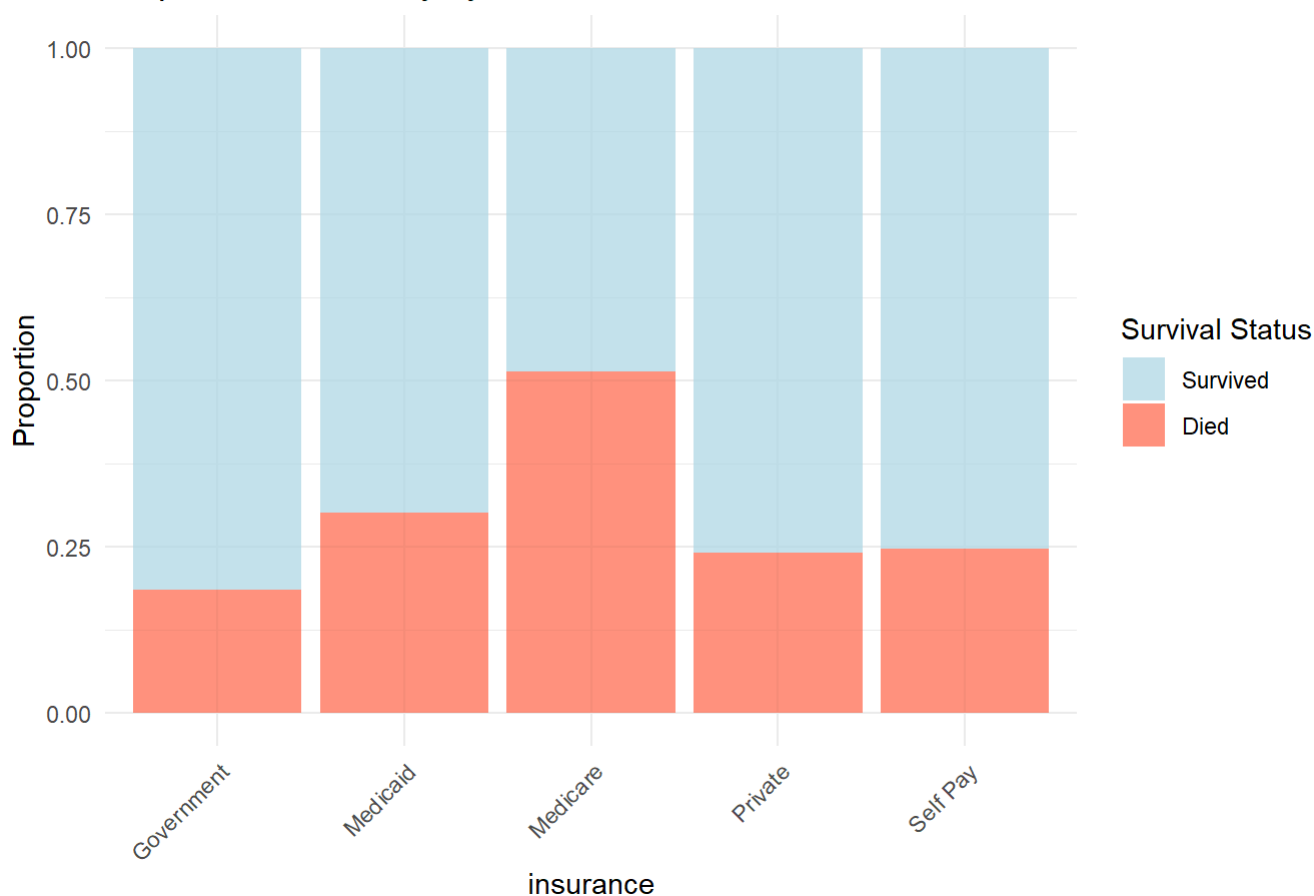
Proportion of Mortality by gender



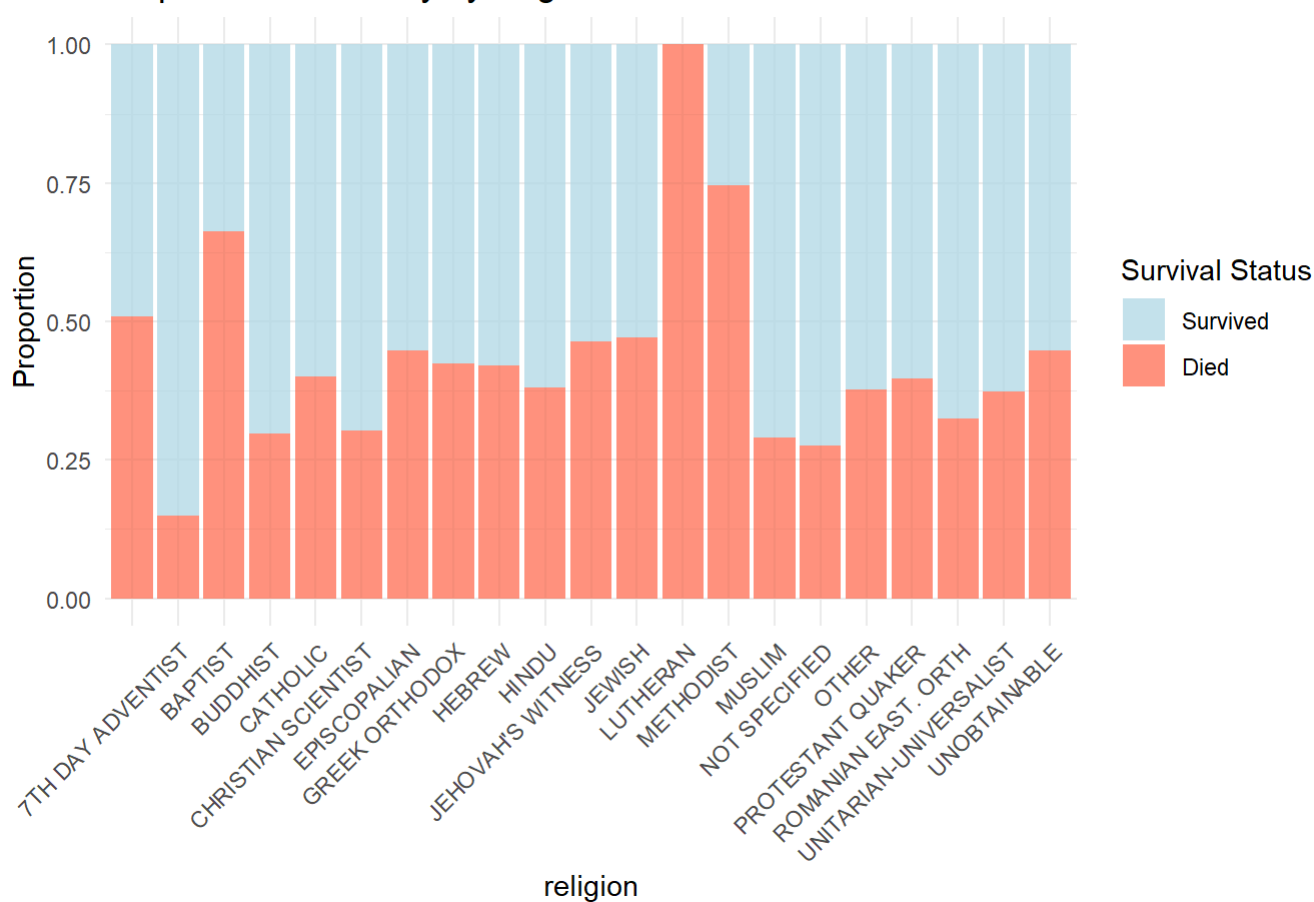
Proportion of Mortality by first\_careunit



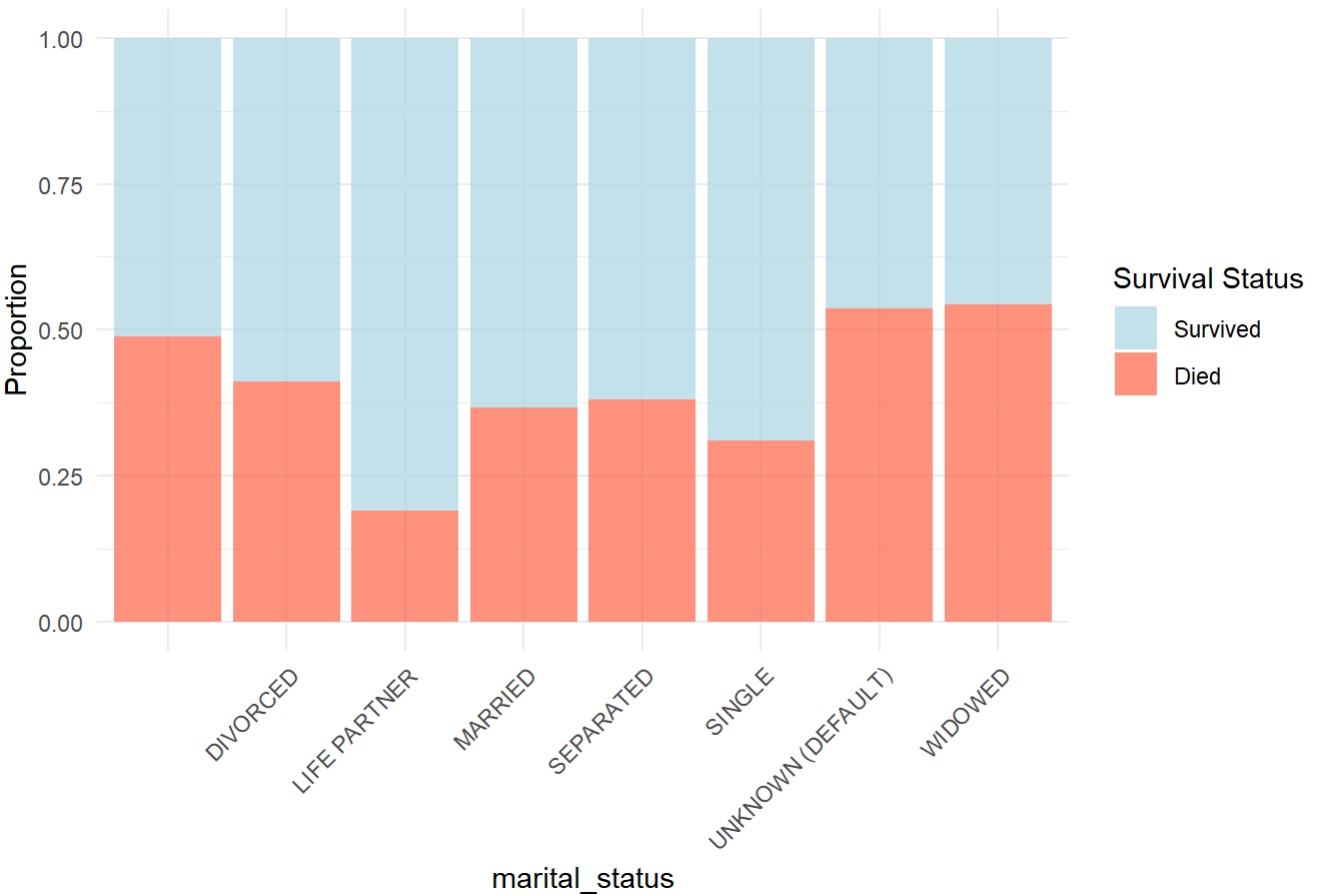
Proportion of Mortality by insurance



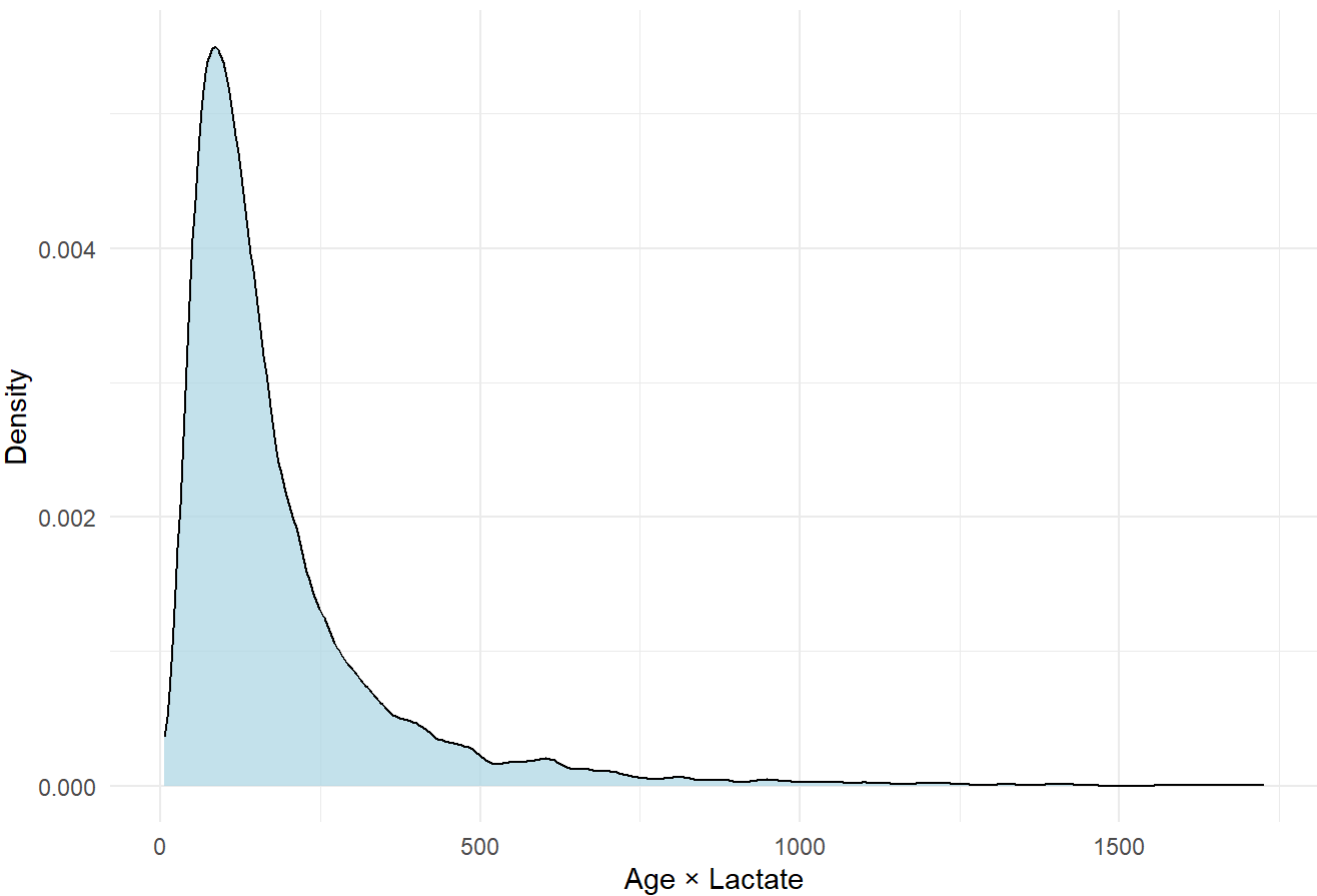
Proportion of Mortality by religion



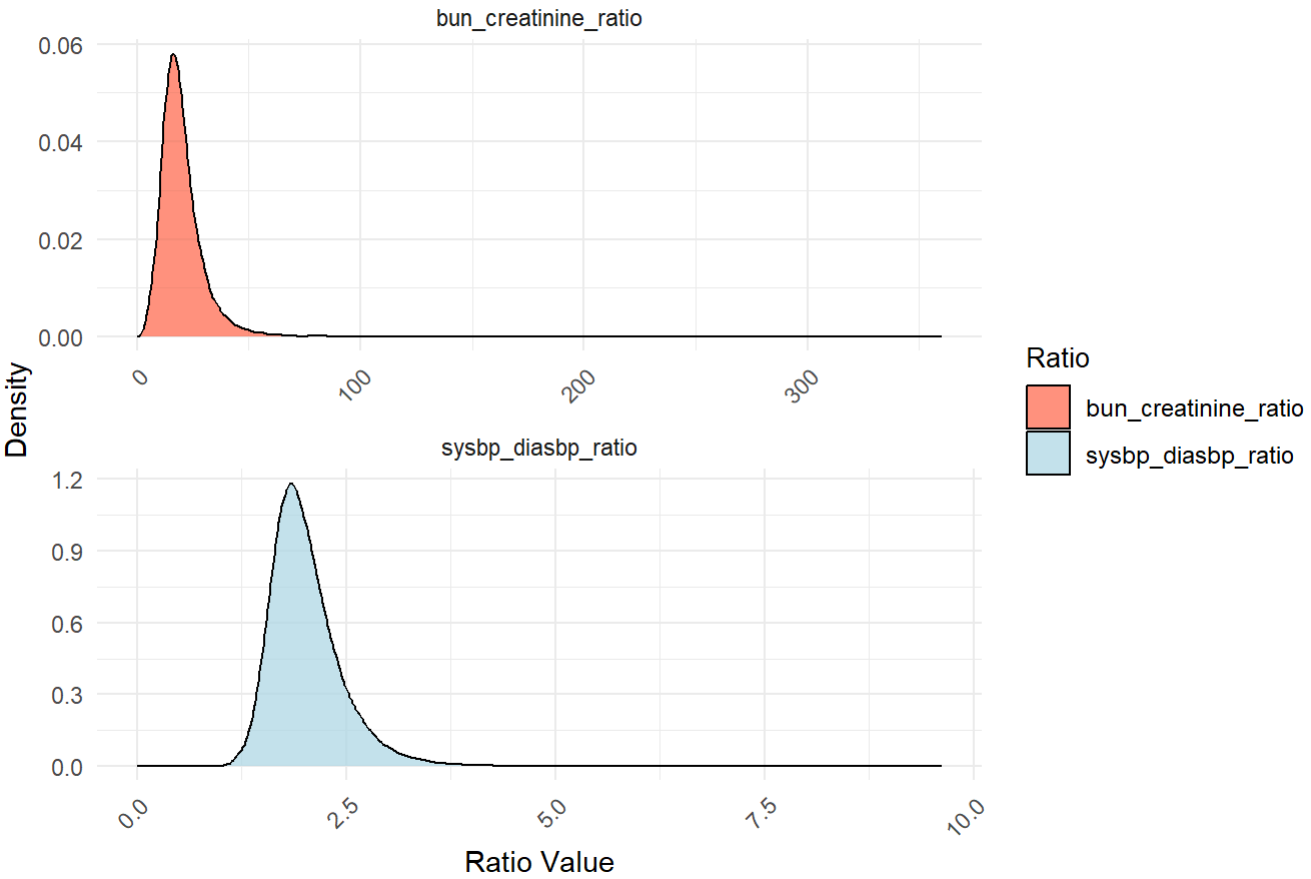
Proportion of Mortality by marital\_status



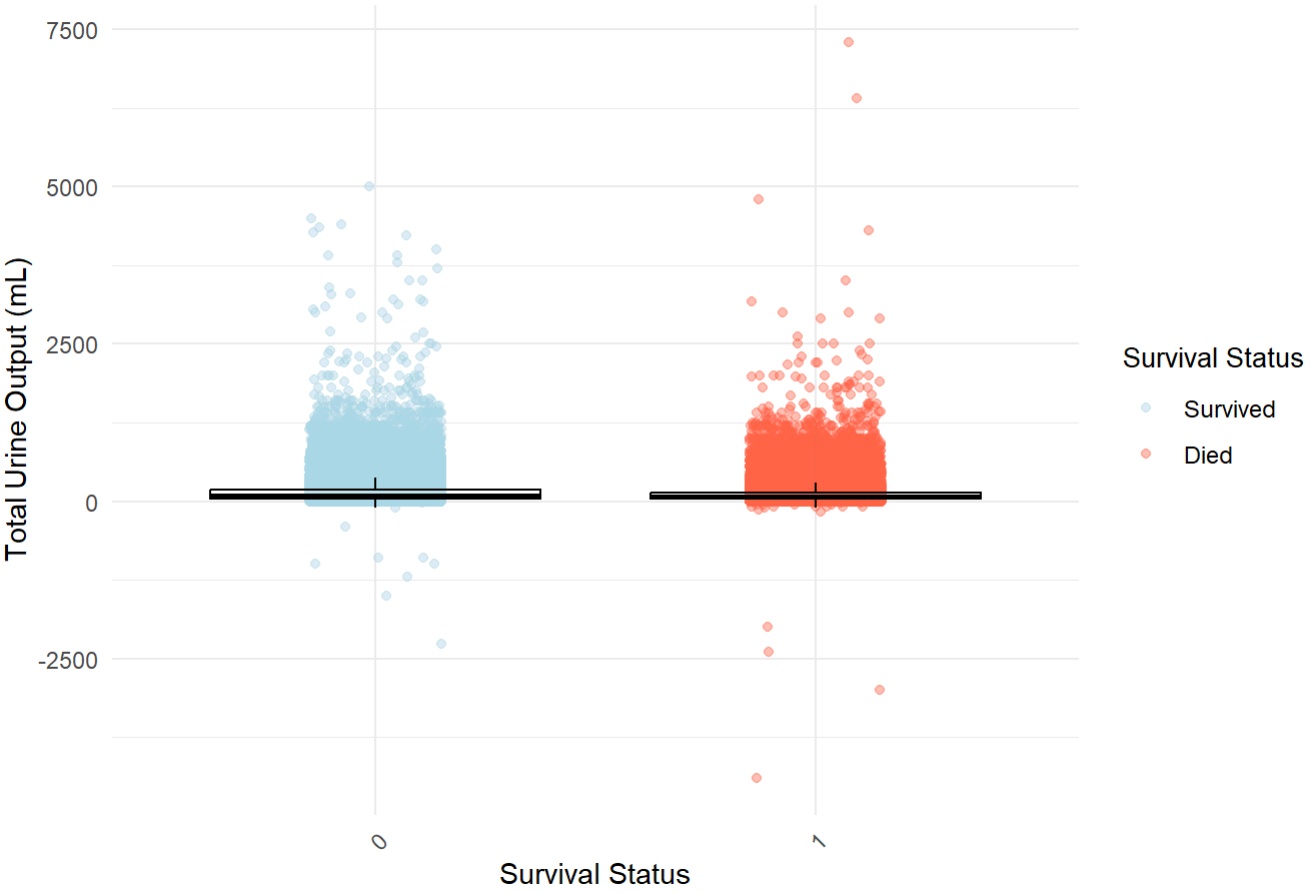
Distribution of Age  $\times$  Lactate Interaction Term



Distribution of Clinically Relevant Ratios (Filtered)



Scatter Plot of Total Urine Output by Mortality Status



## Section 5: Save All Processed Datasets to CSV Format