# **Analysis Report**

```
library(ggplot2)
# Read discharged_patients.txt into a dataframe

df <- read.table("discharged_patients.txt", header = TRUE, sep = ",", stringsAsFactors = FALE

total_rows <- nrow(df)
total_icu_admit <- sum(tolower(df$icuAdmit) == 'true')

total_ward_admit <- total_rows - total_icu_admit
df$los <- df$dischargeTime - df$admitTime</pre>
```

**Total Patients: 1829** 

ICU admit %: 0.1509021

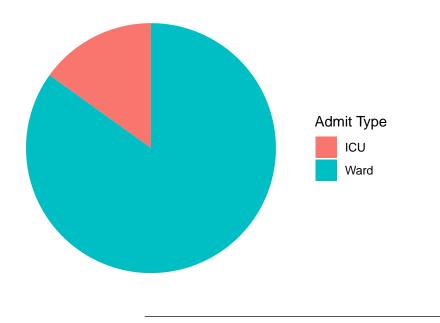
#### Icu-vs-Ward-Admits

```
# Pie chart of ICU vs Ward admits

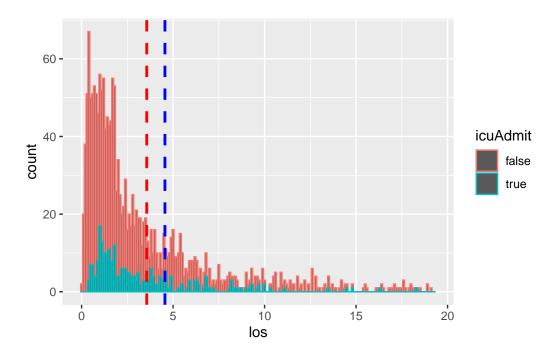
admit_counts <- data.frame(
    type = c("ICU", "Ward"),
    count = c(total_icu_admit, total_ward_admit)
)

ggplot(admit_counts, aes(x = "", y = count, fill = type)) +
    geom_bar(stat = "identity", width = 1) +
    coord_polar(theta = "y") +
    labs(title = "ICU vs Ward Admits", x = NULL, y = NULL, fill = "Admit Type") +
    theme_void()</pre>
```

# ICU vs Ward Admits



## LOS



## summary(icu\_patients\$los)

Min. 1st Qu. Median Mean 3rd Qu. Max. 0.3629 1.2131 2.1605 3.5607 4.3038 33.8704

## summary(ward\_patients\$los)

Min. 1st Qu. Median Mean 3rd Qu. Max. 0.02787 0.93577 1.96473 4.55582 4.95492 92.46698

```
\label{total_ward_discharge} $$ - sum(tolower(df$dischargeLocation) == 'ward')$$ total_icu_discharge <- sum(tolower(df$dischargeLocation) == 'icu')$$ total_transfers_ward <- sum(df$transferTime != -1 & tolower(df$admitLocation) == 'icu')$$ total_transfers_icu <- sum(df$transferTime != -1 & tolower(df$admitLocation) == 'ward')$$
```

## **ADT Summary Stats**

key	value
Total Admissions	1829
Total to Ward	1553
Total to ICU	276
% icu/total	0.1509021
total discharges from ward	1681
total dicharges from icu	148
total transfers to ward	128
total transfers to icu	0
fraction of admissions with transfer to WARD	0.0699836
fraction of admissions with transfer to ICU	0
Average LOS	4.4056553
average LOS for ICU-admits	3.5606909
average length of stay on ICU (transfer or otherwise)	$\operatorname{tbd}$

```
library(dplyr)
df2 <- read.table("visit_data.txt", header = TRUE, sep = ",", stringsAsFactors = FALSE)</pre>
df2$visitDay <- floor(df2$visitTime)</pre>
nvisits <- nrow(df2)</pre>
# Filter for nurse visits
df_nurse_visits <- df2[df2$hcwType == 'NURSE', ]</pre>
df_nurses <- distinct(df_nurse_visits, hcwType, hcwId)</pre>
nurse_count <- nrow(df_nurses)</pre>
# Filter for other HCW types
df_doctor_visits <- df2[df2$hcwType == 'DOCTOR', ]</pre>
df_doctors <- distinct(df_doctor_visits, hcwType, hcwId)</pre>
doctor_count <- nrow(df_doctors)</pre>
                  <- df2[df2$hcwType == 'PT', ]
df_pt_visits
df_pts <- distinct(df_pt_visits, hcwType, hcwId)</pre>
pt_count <- nrow(df_pts)</pre>
df_ot_visits
                  <- df2[df2$hcwType == 'OT', ]
df_ots <- distinct(df_ot_visits, hcwType, hcwId)</pre>
```

```
ot_count <- nrow(df_ots)

df_rt_visits <- df2[df2$hcwType == 'RT', ]

df_rts <- distinct(df_rt_visits, hcwType, hcwId)

rt_count <- nrow(df_rts)</pre>
```

## Total patient visits by hcw type

HCW Type	Total visits (365d)	mean/day
NURSE (26)	357	0.0376185
DOCTOR (18)	312606	47.5808219
OT (9)	765	0.2328767
PT (9)	1019	0.3101979
RT (9)	978	0.2977169

## Average daily visits per patient per hcw per shift

## Important

Future Analysis to be completed:

- average daily visits per patient per hcw per shift
  - total and broken out by how type
- average distinct patients visited per shift
  - broken out by hcw type
- average time between visits per HCW by type
  - verify (intravisit time + duration of visit from Granular Model)
    - \* doctors mean = gamma(0.52, 90.7) + gamma(5.5, 1.2)
    - \* nurses mean = gamma(0.54, 55.7) + gamma(5.5, 1.2)
    - \* therapists mean =  $\operatorname{gamma}(0.52, 61.7) + \operatorname{gamma}(3.0, 1.8)$
- average number of HCW visits per day/patient (total/{type}) val
- average time between visist by HCW per patient
- · average distinct how visits per day by patient

#### **Disease**

• TBD

## punchlist

- Fix the gamma() + gamma() how visit intra-event times
- Fix the HCW assignment procedure for nurse and doctor

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