

Project Proposal

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Load Packages

```
library(tidyverse)
library(haven)
eprocedures <- read_sas("/home/guest/STA198/team-name-tbd/eprocedures_03ref.sas7bdat")
esituation1 <- read_sas("/home/guest/STA198/team-name-tbd/esituation_10ref.sas7bdat")
esituation2 <- read_sas("/home/guest/STA198/team-name-tbd/esituation_09ref.sas7bdat")
```

Load Data

[to be inserted upon receiving expanded OIT container]

Introduction and Data, including Research Questions

Our data is provided by NEMSIS, the National EMS Information System, and was collected as an effort to help standardize and improve EMS databases.

Every year, over 16 million patients are treated by EMS. In comparison to White patients, Asian and Hispanic patients were less likely to be evaluated for their pain. White and Black patients were relatively equal in pain assessment frequency, but only 13.9% of Black patients were given pain medication while 20.1% of White patients were given pain medication (Soares et al, 2019). While there are lots of limitations to interpreting EMS treatment of patients, previous findings have shown that race directly leads to a difference of treatment in emergency situations. We wish to dive deeper into this to see if other factors influence patients' treatment of care. From the data provided by NEMSIS, we ask, what factors influence somebody's process of emergency care in an EMS call?

To answer our overarching research question, we plan to center our analysis around these questions: Is there a correlation between drug use and observed quality of care? Is there a correlation between race and observed quality of care? Are there biases present in the EMS quality of care sample size? What is the relationship between symptoms and vital sign ranges? Is race correlated to technicians' assessment of GCS?

Variables of interest include Primary Symptom, Other Associated Symptoms, Cause of Injury, Barriers to Patient Care, Alcohol/Drug Use Indicators, Protocols Used, Medication Given, Procedure, Procedure Successful/Response to Procedure, Suspected EMS Work Related Exposure/Injury/Death, Race, Initial Patient Acuity vs Final Patient Acuity, and Vital Signs—Blood Pressure, Pulse Oximetry, Respiratory Rate, O2 Saturation, Blood Glucose, Glasgow Coma Score, and Level of Responsiveness.

Glimpse

```
glimpse(eprocedures)
```

```
## Rows: 1,095
## Columns: 2
## $ eProcedures_03      <chr> "103696004", "103715008", "103720008", "103721007", ~
## $ ProcedureCodeDescr <chr> "Patient referral to specialist (procedure)", "Remo~
glimpse(esituation1)
```

```
## Rows: 3,010
## Columns: 2
## $ eSituation_10        <chr> "A31", "A31.8", "A40.8", "A41", "A41.8", "A41.89", ~
## $ DiagnosisCodeDescr <chr> "Infection due to other mycobacteria", "Other mycob~
glimpse(esituation2)
```

```
## Rows: 3,314
## Columns: 2
## $ eSituation_09        <chr> "A08.4", "A15.9", "A31", "A31.8", "A37", "A40.8", "~
## $ DiagnosisCodeDescr <chr> "Viral gastroenteritis NOS", "Respiratory tuberculo~
```

We have achieved extraction of three of our variables with the help of Phuc Nguyen and will continue to upload the full data to Github when it is separated into unidentifiable random samples, either by means of working with an experienced R user or obtaining a larger container from the OIT.

We will be using ~500 observations with 10-20 variables as personally approved by the NEMSIS research team for uploading to Github. We will be including a substantial credit line and clarifying the nature of the data as NOT population-based and NOT used to represent public health data. The process will include uploading the data onto a Windows computer, bypassing the encryption with the corresponding Bitlocker passcode, uploading the zip file onto a high-capacity R-studio container, reading it from SAS to R, joining the files, narrowing down the sample to 500 random observations using R code, saving that file, and uploading it to Github in place of the original data set to work from. Currently, we have an open help desk request of number RITM4124724 to obtain access to the high-capacity container.

Data Analysis Plan

As a part of our data analysis, a few of the specific predictors we are looking at are race, symptoms, cause of injury, BGL, and Alcohol/Drug Use Indicators. As a result, the outcomes we will be looking at are primarily related to the quality of care, which we will be able to determine from Response Time, GCS, Barriers to Patient Care, and Vital Signs. Furthermore, in order to address our research questions we will be using race, cause of injury, and symptoms as comparison groups.

In addition, we have identified a few vital statistical methods needed for analysis. These include p-value/t-test (null hypothesis), standard deviation for the symptoms/conditions as related to different vital signs, and conditional probability. As a result, we expect to see a correlation between race and quality of care, blood glucose level, alcohol/drug use indicators, symptoms, and vital signs. We also expect that the null hypothesis can be rejected ($P < 0.05$) and a standard distribution of vitals will be related to their associated symptoms. Various conditional probabilities are expected to bolster our expected conclusion as well

Visualizations may include a faceted histogram or boxplot of distributions of blood pressures by body system conditions associated with the vital signs. Categorical visuals will be of use to us because of the nature of many of our variables, so we will make use of creative dodged and filled bar plots to show alcohol and drug use indicators by race. We can also use a tree map to show proportion related to the total body systems associated with different patient IDs.

Calculations include the proportion of work-related exposures, injuries, and deaths out of the total data set.

References:

Soares, W. E., 3rd, Knowles, K. J., 2nd, & Friedmann, P. D. (2019). A Thousand Cuts: Racial and Ethnic

Disparities in Emergency Medicine. Medical care, 57(12), 921–923. <https://doi.org/10.1097/MLR.0000000000001250>

CHANGES FOR RESUBMISSION: We added the data to our project, added more acknowledgements, and did thorough glimpses of our data.