

Palatnik eda

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```
df <- read.csv('~/Desktop/capstone/ob_gyn_mother_baby.csv')
```

data dictionary:

patient_num: *integer* Masked MRN, the i2b2 patient number

mom_admit_dttm_shifted: *date* date of the mother's delivery admission

mom_disch_ddtm_shifted: *date* date of the mother's delivery discharge

mom_los: *integer* mom's length of stay in days for the delivery encounter

mom_age_at_del: *integer* mom's age at delivery in years

mom_marital_status: *factor 4 levels – Divorced, Married, Single, Other* mom's current marital status

mom_race: *factor 3 levels – White or Caucasian, Black or African American, Other* mom's current race

mom_ethnicity: *factor 3 levels – Hispanic, Non-Hispanic, Other* mom's current ethnicity

mom_ht_ft: *numeric* mom's height (feet counterpart)

mom_ht_in: *numeric* mom's height (inches counterpart)

mom_wt_oz: *integer* mom's weight in ounces

mom_bmi: *numeric* mom's body mass index

preeclampsia: *factor 2 levels – Yes, No* indicates presence of a preeclampsia diagnosis on the delivery encounter

pregest_dm: *factor 2 levels – Yes, No* indicates presence of a pregestational diabetes diagnosis on the delivery encounter

gestestational_dm: *factor 2 levels – Yes, No* indicates presence of a gestational diabetes diagnosis on the delivery encounter

placental_abruption: *factor 2 levels – Yes, No* indicates presence of a placental abruption diabetes diagnosis on the delivery encounter

maternal_dvt: *factor 2 levels – Yes, No* indicates presence of a maternal deep vein thrombosis diabetes diagnosis on the delivery encounter

baby_patient_num: *integer* Masked MRN, the i2b2 patient number

baby_yob_shifted: *integer* baby's year of birth

baby_dob_shifted: *date* baby's date of birth

baby_dod_shifted: *date* baby's date of death

baby_admit_dttm_shifted: *date* date of the baby's birth admission

baby_disch_dttm_shifted: *date* date of the baby's birth discharge

baby_birth_ht_in: *numeric* baby's birth height in inches

baby_birth_wt_oz: *numeric* baby's birth weight in ounces

delivery_method: *factor 4 levels* – *C-Section, Operative Vaginal, Vaginal, Other* name of the delivery method used for the birth

baby_sex: *factor 2 levels* – *Male, Female* baby's current sex

baby_race: *factor 3 levels* – *White or Caucasian, Black or African American, Other* baby's current race

baby_ethnicity: *factor 3 levels* – *Hispanic, Non-Hispanic, Other* baby's current ethnicity

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df$mom_admit_dttm_shifted <-  
  gsub(" .*", "", df$mom_admit_dttm_shifted)  
df$mom_disch_dttm_shifted <-  
  gsub(" .*", "", df$mom_disch_dttm_shifted)  
df$baby_dob_shifted <- gsub(" .*", "", df$baby_dob_shifted)  
df$baby_dod_shifted <- gsub(" .*", "", df$baby_dod_shifted)  
df$baby_admit_dttm_shifted <-  
  gsub(" .*", "", df$baby_admit_dttm_shifted)  
df$baby_disch_dttm_shifted <-  
  gsub(" .*", "", df$baby_disch_dttm_shifted)  
  
df_mutated <- df %>%  
  add_column(mom_ht_ft = NA) %>%  
  add_column(mom_ht_in = NA) %>%  
  select(  
    -delivery_encounter_num,  
    -del_dept_id,  
    -mom_patient_num,  
    -del_dept_name,  
    -mom_disch_disp,  
    -del_stage1_hrs,  
    -del_stage2_hrs,  
    -del_stage3_hrs,  
    -bloodloss_del,  
    -bloodloss_total,  
    -birth_encounter_num,  
    -baby_del_dept,  
    -baby_los,  
    -baby_disch_disp,  
    -gest_age_wks,  
    -apgar1,  
    -apgar5,  
    -apgar10,  
    -anesth_conc,  
    -del_meth_id,  
    -ob_gravidity,  
    -ob_parity,  
    -ob_multiple_births  
  ) %>%  
  mutate(  
    mom_ethnicity = case_when(  
      mom_ethnicity == 'Refused' ~ 'Other',
```

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mom_ethnicity == 'Hispanic' ~ 'Hispanic',
mom_ethnicity == 'Unknown' ~ 'Other',
mom_ethnicity == 'NI' ~ 'Other',
mom_ethnicity == 'Non Hispanic' ~ 'NH'
)
) %>%
mutate(
  mom_race = case_when(
    mom_race == 'Unknown' ~ 'Other',
    mom_race == 'Asian' ~ 'Other',
    mom_race == 'Patient Refused' ~ 'Other',
    mom_race == 'Multiracial' ~ 'Other',
    mom_race == 'Native Hawaiian or Other Pacific Islander' ~ 'Other',
    mom_race == 'American Indian or Alaska Native' ~ 'Other',
    mom_race == 'White or Caucasian' ~ 'White or Caucasian',
    mom_race == 'Black or African American' ~ 'Black or African American',
    mom_race == 'Other' ~ 'Other',
  )
) %>%
mutate(
  baby_ethnicity = case_when(
    baby_ethnicity == 'Refused' ~ 'Other',
    baby_ethnicity == 'Unknown' ~ 'Other',
    baby_ethnicity == 'Hispanic' ~ 'Hispanic',
    baby_ethnicity == 'NI' ~ 'Other',
    baby_ethnicity == 'Non Hispanic' ~ 'NH'
  )
) %>%
mutate(
  baby_race = case_when(
    baby_race == 'Unknown' ~ 'Other',
    baby_race == 'Asian' ~ 'Other',
    baby_race == 'Patient Refused' ~ 'Other',
    baby_race == 'Multiracial' ~ 'Other',
    baby_race == 'Native Hawaiian or Other Pacific Islander' ~ 'Other',
    baby_race == 'American Indian or Alaska Native' ~ 'Other',
    baby_race == 'White or Caucasian' ~ 'White or Caucasian',
    baby_race == 'Black or African American' ~ 'Black or African American',
    baby_race == 'Other' ~ 'Other',
  )
) %>%
mutate(
  mom_marital_status = case_when(
    mom_marital_status == 'Legally Separated' ~ 'Divorced',
    mom_marital_status == 'Divorced' ~ 'Divorced',
    mom_marital_status == 'Patient Refused' ~ 'Other',
    mom_marital_status == 'Unknown' ~ 'Other',
    mom_marital_status == 'Widowed' ~ 'Other',
    mom_marital_status == 'Significant Other' ~ 'Other',
    mom_marital_status == 'Married' ~ 'Married',
    mom_marital_status == 'Single' ~ 'Single'
  )
) %>%

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mutate(mom_marital_status = as.factor(mom_marital_status)) %>%
mutate(mom_race = as.factor(mom_race)) %>%
mutate(mom_ethnicity = as.factor(mom_ethnicity)) %>%
mutate(preeclampsia = as.factor(preeclampsia)) %>%
mutate(pregest_dm = as.factor(pregest_dm)) %>%
mutate(gestational_dm = as.factor(gestational_dm)) %>%
mutate(placental_abruption = as.factor(placental_abruption)) %>%
mutate(maternal_dvt = as.factor(maternal_dvt)) %>%
mutate(baby_birth_wt_gms = as.numeric(baby_birth_wt_gms)) %>%
mutate(
  delivery_method = case_when(
    delivery_method == 'C-Section' ~ 'C-Section',
    delivery_method == 'C-Section w/ BTL' ~ 'C-Section',
    delivery_method == 'C-Section, Classical' ~ 'C-Section',
    delivery_method == 'C-Section, Hysterectomy' ~ 'C-Section',
    delivery_method == 'C-Section, Low Transverse' ~ 'C-Section',
    delivery_method == 'C-Section, Low Vertical' ~ 'C-Section',
    delivery_method == 'C-Section, Unspecified' ~ 'C-Section',
    delivery_method == 'Elective Abortion' ~ 'Other',
    delivery_method == 'Forceps Vaginal Delivery' ~ 'Operative Vaginal',
    delivery_method == 'Miscarriage' ~ 'Other',
    delivery_method == 'NST' ~ 'Other',
    delivery_method == 'Other/Procedure' ~ 'Other',
    delivery_method == 'Spontaneous Abortion' ~ 'Other',
    delivery_method == 'TAB' ~ 'Other',
    delivery_method == 'Vacuum Vaginal Delivery' ~ 'Operative Vaginal',
    delivery_method == 'Vaginal, Breech' ~ 'Vaginal',
    delivery_method == 'Vaginal, Breech Extraction' ~ 'Vaginal',
    delivery_method == 'Vaginal, Forceps' ~ 'Operative Vaginal',
    delivery_method == 'Vaginal, Forceps Delivery' ~ 'Operative Vaginal',
    delivery_method == 'Vaginal, Spontaneous' ~ 'Vaginal',
    delivery_method == 'Vaginal, Spontaneous Breech' ~ 'Vaginal',
    delivery_method == 'Vaginal, Vacuum (Extractor)' ~ 'Operative Vaginal',
    delivery_method == 'Vaginal, Vacuum Delivery' ~ 'Operative Vaginal',
    delivery_method == 'VBAC, Forceps Delivery' ~ 'Operative Vaginal',
    delivery_method == 'VBAC, Spontaneous' ~ 'Vaginal',
    delivery_method == 'VBAC, Vacuum Delivery' ~ 'Operative Vaginal',
  )
) %>%
mutate(delivery_method = as.factor(delivery_method)) %>%
mutate(
  baby_sex = case_when(
    baby_sex == 'Unknown' ~ '',
    baby_sex == 'Male' ~ 'Male',
    baby_sex == 'Female' ~ 'Female'
  )
) %>%
separate(mom_ht_ftin,
  into = c('mom_ht_ft', 'mom_ht_in'),
  sep = "'") %>%
mutate(baby_sex = as.factor(baby_sex)) %>%
mutate(baby_race = as.factor(baby_race)) %>%
mutate(baby_ethnicity = as.factor(baby_ethnicity)) %>%

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mutate(baby_admit_dttm_shifted = as.Date(baby_admit_dttm_shifted, "%m/%d/%y")) %>%
mutate(baby_disch_dttm_shifted = as.Date(baby_disch_dttm_shifted, "%m/%d/%y")) %>%
mutate(baby_dob_shifted = as.Date(baby_dob_shifted, "%m/%d/%y")) %>%
mutate(baby_dod_shifted = as.Date(baby_dod_shifted, "%m/%d/%y")) %>%
mutate(mom_admit_dttm_shifted = as.Date(mom_admit_dttm_shifted, "%m/%d/%y")) %>%
mutate(mom_disch_dttm_shifted = as.Date(mom_disch_dttm_shifted, "%m/%d/%y"))

df_mutated$mom_ht_in <- gsub('\\" , "", df_mutated$mom_ht_in)

df_mutated <- df_mutated %>%
  mutate(mom_ht_ft = as.numeric(mom_ht_ft)) %>%
  mutate(mom_ht_in = as.numeric(mom_ht_in))

df_mutated[df_mutated == ""] <- NA

str(df_mutated, vec.len = 2, strict.width = 'cut')

## 'data.frame': 32499 obs. of 30 variables:
## $ patient_num : int 29732601 19678083 32945117 23004068 28381336 ..
## $ mom_admit_dttm_shifted : Date, format: "2016-10-30" "2019-11-12" ...
## $ mom_disch_dttm_shifted : Date, format: "2016-11-02" "2019-11-14" ...
## $ mom_los : int 3 2 2 2 2 ...
## $ mom_age_at_del : int 34 42 18 33 24 ...
## $ mom_marital_status : Factor w/ 4 levels "Divorced","Married",...: 2 2 4 ..
## $ mom_race : Factor w/ 3 levels "Black or African American",...: ...
## $ mom_ethnicity : Factor w/ 3 levels "Hispanic","NH",...: 2 2 1 2 2 ...
## $ mom_ht_ft : num 5 5 5 5 5 ...
## $ mom_ht_in : num 0 1 3 2 7 ...
## $ mom_wt_oz : int 3616 2832 2144 2608 2864 ...
## $ mom_bmi : num 44.1 33.4 ...
## $ preeclampsia : Factor w/ 2 levels "No","Yes": 1 1 1 1 1 ...
## $ pregest_dm : Factor w/ 2 levels "No","Yes": 1 1 1 1 1 ...
## $ gestational_dm : Factor w/ 2 levels "No","Yes": 1 1 1 1 1 ...
## $ placental_abruption : Factor w/ 2 levels "No","Yes": 1 1 1 1 1 ...
## $ maternal_dvt : Factor w/ 2 levels "No","Yes": 1 1 1 1 1 ...
## $ baby_patient_num : int 28775049 29913097 26379596 18846057 41430988 ..
## $ baby_yob_shifted : int 2016 2019 2020 2020 2018 ...
## $ baby_dob_shifted : Date, format: "2016-10-30" "2019-11-12" ...
## $ baby_dod_shifted : Date, format: NA NA ...
## $ baby_admit_dttm_shifted : Date, format: "2016-10-30" "2019-11-12" ...
## $ baby_disch_dttm_shifted : Date, format: "2016-11-02" "2019-11-14" ...
## $ baby_birth_ht_in : num 19 19 20 19 21 ...
## $ baby_birth_wt_oz : num 132 128 ...
## $ baby_birth_wt_gms : num 3730 3640 ...
## $ delivery_method : Factor w/ 4 levels "C-Section","Operative Vaginal"...
## $ baby_sex : Factor w/ 3 levels "", "Female", "Male": 2 2 2 2 3 ...
## $ baby_race : Factor w/ 3 levels "Black or African American",...: ...
## $ baby_ethnicity : Factor w/ 3 levels "Hispanic","NH",...: 2 2 1 2 2 ...

save(df_mutated, file = "~/Desktop/capstone/Palatnik_2_22.Rdata")

```