

P8106 Final Project: Predicting COVID-19 Recovery Time and Identifying Significant Risk Factors

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Contents

Background:	2
Data:	2

Background:

Check the report.

Data:

Description check the report.

```
# Dataset Loading
load("data/recovery.Rdata")

set.seed(3521) # Runze Cui's uni(2183):
# Create a first random sample of 2000 participants:
dat1 <- dat[sample(1:10000, 2000),]

set.seed(3555) # Yuchen Hua's uni(3555)
# Create a second random sample of 2000 participants:
dat2 <- dat[sample(1:10000, 2000),]

# Merged the two datasets and remove repeated observations:
dat <- unique(rbind(dat1, dat2))

# Get rid of the id variable from the merged dataset and do the data cleaning:
dat = dat %>%
  select(-id) %>%
  mutate(gender = as.factor(gender)) %>%
  mutate(race = as.factor(race)) %>%
  mutate(smoking = as.factor(smoking)) %>%
  mutate(hypertension = as.factor(hypertension)) %>%
  mutate(diabetes = as.factor(diabetes)) %>%
  mutate(vaccine = as.factor(vaccine)) %>%
  mutate(severity = as.factor(severity)) %>%
  mutate(study = as.factor(study)) %>%
  na.omit() %>%
  relocate(recovery_time)

head(dat)
```

##	recovery_time	age	gender	race	smoking	height	weight	bmi	hypertension
## 8158	52	61	0	1	1	169.9	87.6	30.4	0
## 3387	24	60	1	1	2	173.4	70.6	23.5	0
## 1709	36	60	1	1	1	178.2	79.9	25.1	0
## 4051	23	70	1	4	0	167.4	77.7	27.7	1
## 954	24	63	1	4	0	175.4	88.7	28.8	1
## 531	36	65	0	1	0	160.4	74.4	28.9	1

##	diabetes	SBP	LDL	vaccine	severity	study
## 8158	0	118	103	0	0	C
## 3387	0	129	101	1	0	B
## 1709	0	130	107	1	0	A
## 4051	0	145	128	1	0	B
## 954	0	131	100	0	0	A
## 531	0	137	153	1	0	A

```

# Separate the data as training and test data:
set.seed(3521)
# Specify rows of training data:
trRows <- createDataPartition(dat$recovery_time, p = 0.7, list = FALSE)

# training data
training <- dat[trRows, ]
## matrix of predictors
x <- model.matrix(recovery_time~.,dat)[trRows,-1]
## vector of response
y <- dat$recovery_time[trRows]

# test data
test <- dat[-trRows, ]
## matrix of predictors
x2 <- model.matrix(recovery_time~.,dat)[-trRows,-1]
## vector of response
y2 <- dat$recovery_time[-trRows]

```