# DAS Project2 Group18

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```
#load the data
  shelter_01 <- read.csv("dataset18.csv")</pre>
  #Checking for missing value
  any_na <- apply(shelter_01, 2, function(x) any(is.na(x)))</pre>
  any_na
    animal_type
                           month
                                              year
                                                       intake_type
                                                                        outcome_type
                           FALSE
                                            FALSE
                                                              FALSE
                                                                               FALSE
          FALSE
    chip_status time_at_shelter
          FALSE
                           FALSE
  total_na <- sum(is.na(shelter_01))</pre>
  total_na
[1] 0
  #Converting a string variable to a factor type and make a summary statistics
  shelter_01$animal_type <- as.factor(shelter_01$animal_type)</pre>
  shelter_01$intake_type <- as.factor(shelter_01$intake_type)</pre>
  shelter_01$outcome_type <- as.factor(shelter_01$outcome_type)</pre>
  shelter_01$chip_status <- as.factor(shelter_01$chip_status)</pre>
  summary(shelter_01)
    animal_type
                      month
                                         year
                                                              intake_type
BIRD
          : 2
                  Min. : 1.000
                                    Min.
                                           :2016
                                                    CONFISCATED
CAT
          :238
                  1st Qu.: 4.000
                                    1st Qu.:2017
                                                    OWNER SURRENDER: 363
```

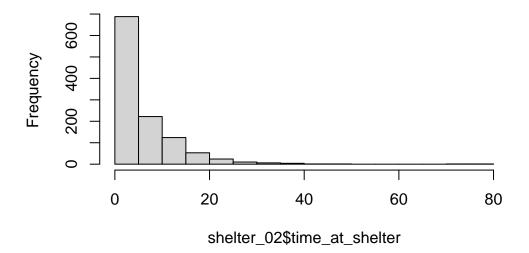
```
DOG
               Median : 7.000
                              Median:2017
                                             STRAY
       :880
                                                       :713
LIVESTOCK: 1
               Mean
                    : 6.574 Mean
                                    :2017
WILDLIFE: 14
               3rd Qu.: 9.000
                               3rd Qu.:2017
               Max.
                     :12.000
                              Max.
                                     :2017
                              chip_status time_at_shelter
          outcome_type
                                          Min. : 0.00
ADOPTION
                :474
                      SCAN CHIP
                                   :214
                      SCAN NO CHIP :860
                                          1st Qu.: 1.00
DIED
                : 14
                      UNABLE TO SCAN: 61
                                          Median: 4.00
EUTHANIZED
                :417
FOSTER
               : 30
                                          Mean : 6.12
                                          3rd Qu.: 9.00
RETURNED TO OWNER:200
                                          Max. :78.00
```

#Converting shelter\_01 to dataframe
shelter\_02 <- as.data.frame(shelter\_01)
summary(shelter\_02)</pre>

animal_type mont		th year		ear	intake_type		
BIRD : 2	Min.	: 1.000	Min.	:201	6 CONFISCATED : 59		
CAT :238	1st Qu.	: 4.000	1st Qu	.:201	7 OWNER SURRENDER:363		
DOG :880	Median	: 7.000	Median	:201	7 STRAY :713		
LIVESTOCK: 1	Mean	: 6.574	Mean	:201	7		
WILDLIFE : 14	3rd Qu.	: 9.000	3rd Qu	.:201	7		
	Max.	:12.000	Max.	:201	7		
outo	ome_type		chip_st	atus	time_at_shelter		
ADOPTION	:474	SCAN CHI	P ::	214	Min. : 0.00		
DIED	: 14	SCAN NO	CHIP :	860	1st Qu.: 1.00		
EUTHANIZED	:417	UNABLE T	O SCAN:	61	Median: 4.00		
FOSTER	: 30				Mean : 6.12		
RETURNED TO OWN	ER:200				3rd Qu.: 9.00		
					Max. :78.00		

hist(shelter\_02\$time\_at\_shelter)

# Histogram of shelter\_02\$time\_at\_shelter



## Call:

```
glm(formula = time_at_shelter ~ year + month + animal_type +
   intake_type + outcome_type + chip_status, family = poisson(),
   data = shelter_02)
```

#### Coefficients:

	Estimate	Std. Error	z value	Pr(> z )
(Intercept)	209.563215	219.298918	0.956	0.33927
year	-0.108790	0.043770	-2.486	0.01294 *
month	-0.016839	0.005842	-2.882	0.00395 **
${\tt animal\_typeCAT}$	13.253664	200.734972	0.066	0.94736
${\tt animal\_typeDOG}$	13.354757	200.734971	0.067	0.94696
${\tt animal\_typeLIVESTOCK}$	-0.191216	348.317912	-0.001	0.99956
${\tt animal\_typeWILDLIFE}$	12.834001	200.735017	0.064	0.94902
intake_typeOWNER SURRENDER	-1.367180	0.049511	-27.614	< 2e-16 ***
intake_typeSTRAY	-0.856870	0.044964	-19.057	< 2e-16 ***

```
-0.469573
                                    0.113310 -4.144 3.41e-05 ***
outcome_typeDIED
                          outcome_typeEUTHANIZED
outcome_typeFOSTER
                          outcome_typeRETURNED TO OWNER -1.621092
                                     0.050170 -32.312 < 2e-16 ***
                                     0.031581 -8.190 2.62e-16 ***
chip_statusSCAN NO CHIP
                          -0.258643
chip_statusUNABLE TO SCAN
                          Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for poisson family taken to be 1)
   Null deviance: 8495.8 on 1134 degrees of freedom
Residual deviance: 6544.7 on 1120
                               degrees of freedom
AIC: 9670.3
Number of Fisher Scoring iterations: 10
  model_poi <- step(glm_model_poi)</pre>
Start: AIC=9670.31
time_at_shelter ~ year + month + animal_type + intake_type +
   outcome_type + chip_status
             Df Deviance
                          AIC
                 6544.7 9670.3
<none>
             1 6550.9 9674.5
- year
            1 6553.0 9676.6
- month
- animal_type 4 6587.0 9704.6
- chip_status 2 6651.2 9772.8
- intake_type 2 7270.5 10392.1
- outcome_type 4
                 8056.1 11173.7
  library(MASS)
Attaching package: 'MASS'
The following object is masked from 'package:dplyr':
   select
```

```
glm_model_nb <- glm.nb(time_at_shelter ~ animal_type + month + year + intake_type + outcom
summary(glm_model_nb)
```

#### Call:

```
glm.nb(formula = time_at_shelter ~ animal_type + month + year +
   intake_type + outcome_type + chip_status, data = shelter_02,
   init.theta = 0.9633756977, link = log)
```

#### Coefficients:

	Estimate	Std. Error	z value	Pr(> z )
(Intercept)	3.078e+02	1.333e+05	0.002	0.9982
animal_typeCAT	2.619e+01	1.333e+05	0.000	0.9998
animal_typeDOG	2.631e+01	1.333e+05	0.000	0.9998
animal_typeLIVESTOCK	-3.120e-01	2.315e+05	0.000	1.0000
${\tt animal\_typeWILDLIFE}$	2.574e+01	1.333e+05	0.000	0.9998
month	-2.029e-02	1.613e-02	-1.258	0.2084
year	-1.638e-01	1.217e-01	-1.345	0.1785
intake_typeOWNER SURRENDER	-1.703e+00	1.600e-01	-10.640	< 2e-16 ***
intake_typeSTRAY	-1.295e+00	1.506e-01	-8.602	< 2e-16 ***
outcome_typeDIED	-4.871e-01	3.005e-01	-1.621	0.1050
${\tt outcome\_typeEUTHANIZED}$	-6.033e-01	7.598e-02	-7.940	2.02e-15 ***
outcome_typeFOSTER	-4.783e-01	2.175e-01	-2.199	0.0279 *
outcome_typeRETURNED TO OWNER	-1.843e+00	1.108e-01	-16.638	< 2e-16 ***
chip_statusSCAN NO CHIP	-1.717e-01	9.032e-02	-1.901	0.0573 .
chip_statusUNABLE TO SCAN	-7.708e-01	1.816e-01	-4.244	2.20e-05 ***
01 10 1 0 1 1 0 004	1 1 . 0 . 0.4			

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for Negative Binomial(0.9634) family taken to be 1)

Null deviance: 1640.1 on 1134 degrees of freedom Residual deviance: 1312.5 on 1120 degrees of freedom

AIC: 6252.2

Number of Fisher Scoring iterations: 1

Theta: 0.9634 Std. Err.: 0.0542

2 x log-likelihood: -6220.2480

### Start: AIC=6250.25 time\_at\_shelter ~ animal\_type + month + year + intake\_type + outcome\_type + chip\_status Df Deviance AIC 1 1314.0 6249.7 - month 1 1314.3 6250.1 - year <none> 1312.5 6250.2 - animal\_type 4 1325.0 6254.8 - chip\_status 2 1330.3 6264.1 2 1439.1 6372.9 intake\_type - outcome\_type 4 1573.6 6503.4 Step: AIC=6249.74 time\_at\_shelter ~ animal\_type + year + intake\_type + outcome\_type + chip\_status AIC Df Deviance 1 1313.0 6248.2 - year 1312.6 6249.7 <none> - animal\_type 4 1325.5 6254.6 - chip\_status 2 1329.9 6263.0 - intake\_type 2 1439.1 6372.2 - outcome\_type 4 1574.3 6503.4 Step: AIC=6248.17 time\_at\_shelter ~ animal\_type + intake\_type + outcome\_type + chip\_status Df Deviance AIC 1312.6 6248.2 <none> - animal\_type 4 1325.4 6253.0 - chip\_status 2 1330.1 6261.7 intake\_type 2 1439.7 6371.3 - outcome\_type 4 1577.9 6505.5

model\_nb <- step(glm\_model\_nb)</pre>