DAS2021-Group-19

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1 Introduction

1.1 Background Information

Dataset 19 comes from the Dallas animal shelter. You will have access to the following variables, recorded by animal admission:

- Animal type The type of animal admitted to the shelter
- Month Month the animal was admitted, recorded numerically with January=1
- Year Year the animal was admitted to the shelter.
- Intake_type Reason for the animal being admitted to the shelter
- Outcome_type Final outcome for the admitted animal
- Chip_Status Did the animal have a microchip with owner information?
- Time_at_Shelter Days spent at the shelter between being admitted and the final outcome.

1.2 Question of Interest

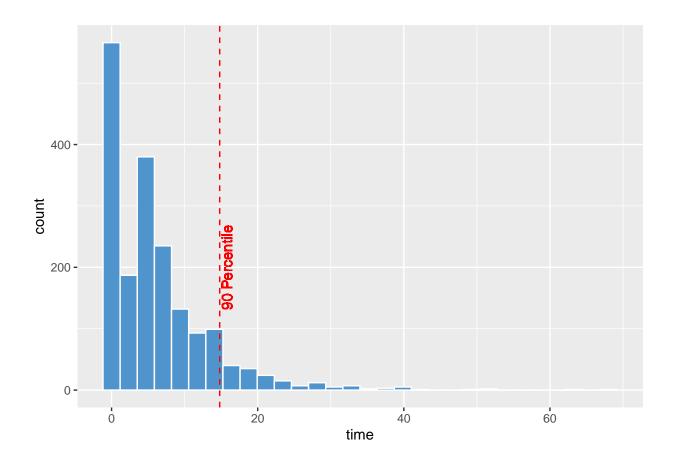
Imagine you have been asked by the shelter management to investigate the following questions of interest:

 Which factors influence the number of days an animal spends in the shelter before their final outcome is decided?

You should conduct an analysis to answer your question using a Generalised Linear Model (GLM). Following your analyses, you should then summarise your results in the form of a presentation.

2 Exploratory Data Analysis

2.1 Tidying Data



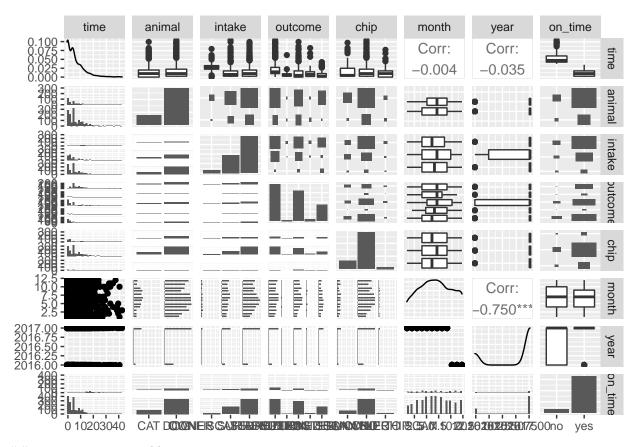
mean s.d min Q1 med Q3 max 1 6.132758 7.130583 0 1 4 9 68

The time spent in the shelter is skewed - most animals (90%) stay no longer than two weeks and very few after three. The majority seem leave the shelter within one week. We could speculate that there are factors (possibly categorical variables) that contribute to the time at the shelter being shorter, and therefore if these factors do not occur then the animal stays longer.

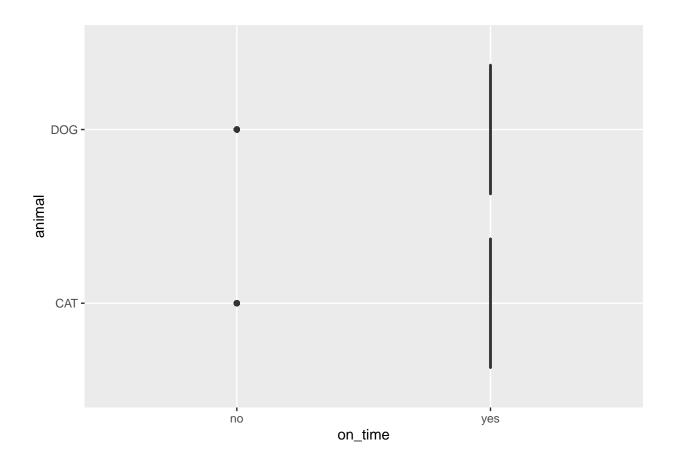
We see that 90% of animals' outcome is decided by 14.8 days at the shelter - this could be our research: what makes animals stay beyond 14 days?

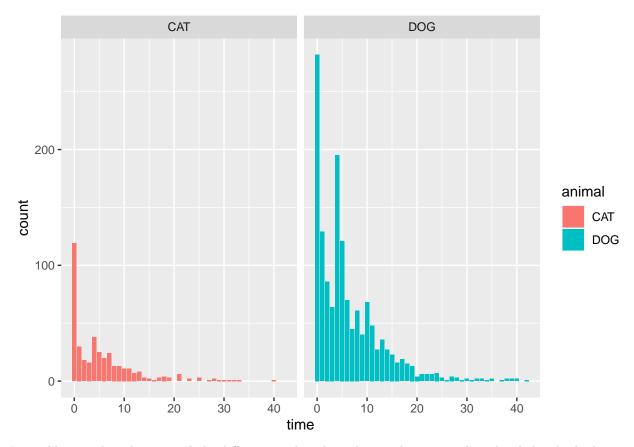
If we consider staying longer than 14 days as a failure, then we can reduce our problem to a binary GLM.

There are only two birds in the data and 13 wildlife. Therefore the impact these animal types will have on the other animal types will be minimal as they do not even contribute to 0.01% of the data, so they can be removed.

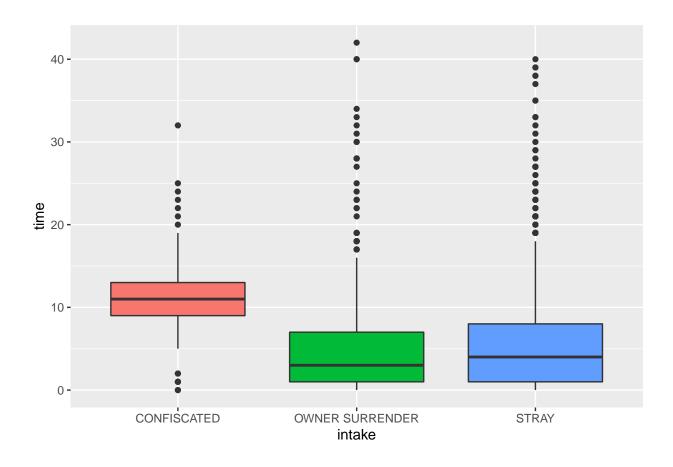


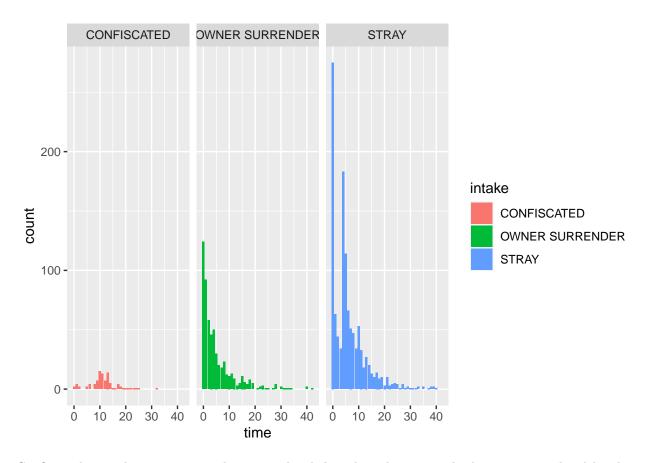
Time response variable



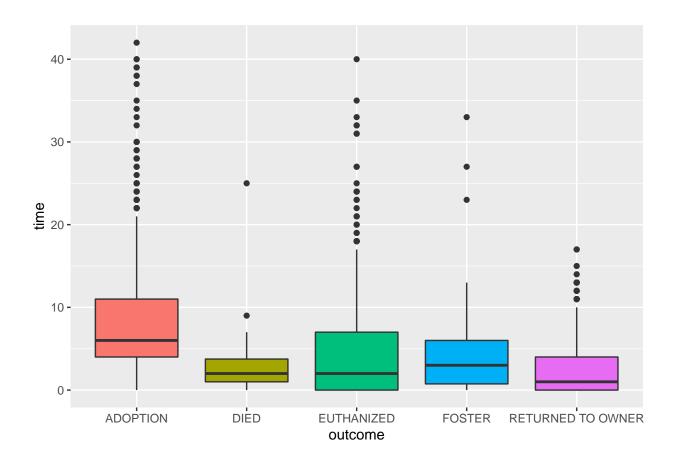


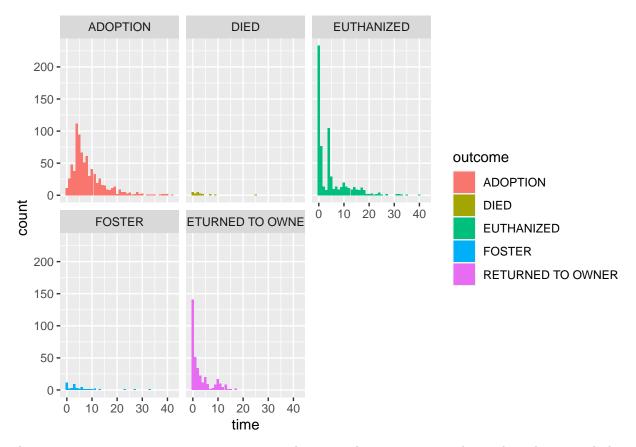
It would seem that there is a slight difference in how long dogs and cats spend at the shelter by looking at the histograms, but from the boxplots we can see that both animals have very similar distributions: the medians are in similar places, their ranges are the similar, both have long tails (skewed right) and lots of outliers. It could be possible that this variable will be omitted from the final model.



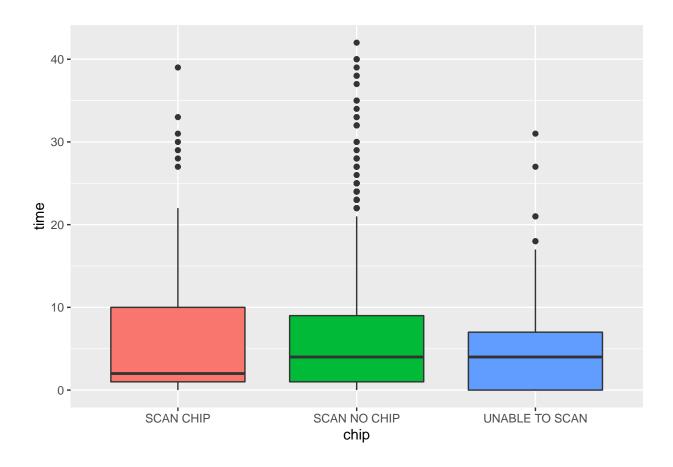


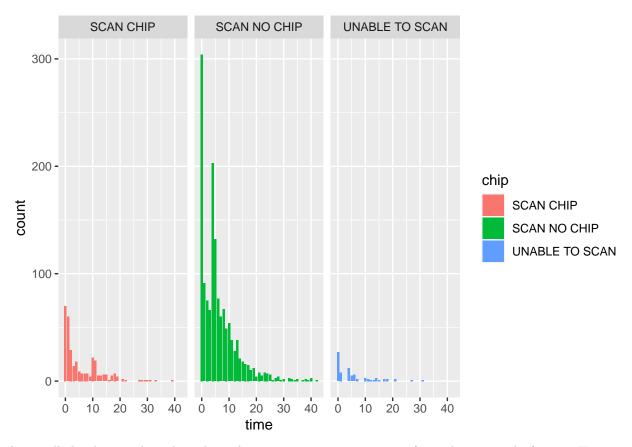
Confiscated animals seems to stay longer at the shelter than those animals that are surrendered by their owner or found as strays. There is a clear separation of one group from the other two. Strays seem to stay slighlty longer than animals that are surrendered.



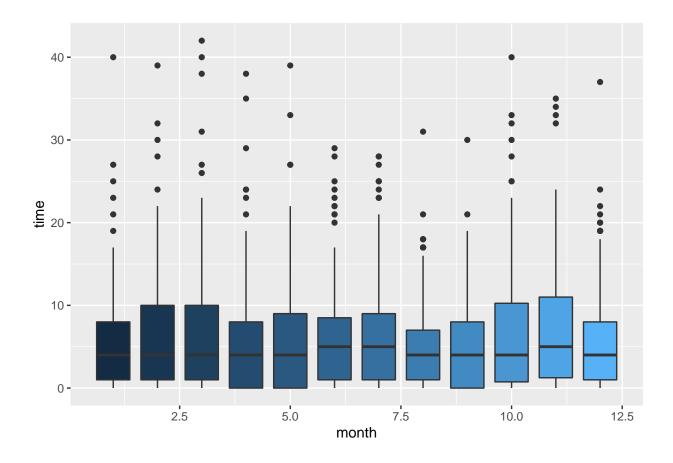


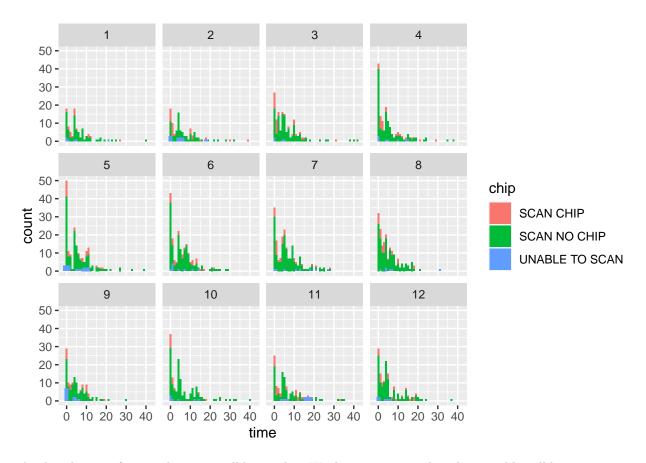
There is some variation across outome - animals returned to owner or euthanized tend to spend short times at the shelter. Those that are adopted tend to be at the shelter for a slightly longer time and the distribution of this subgroup is more normally distributed. There is less data for died and fostered to make an assumption.



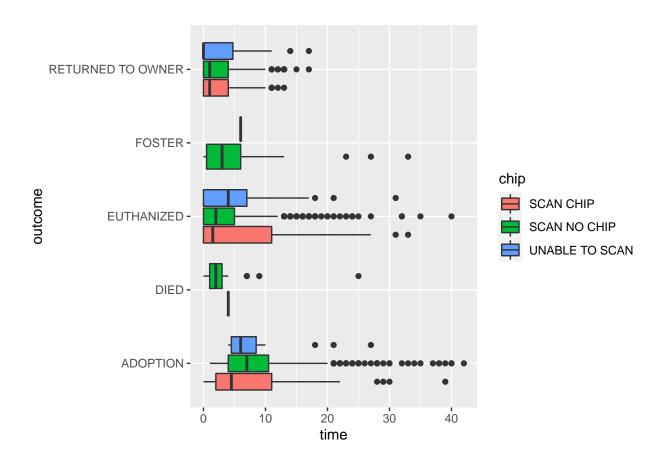


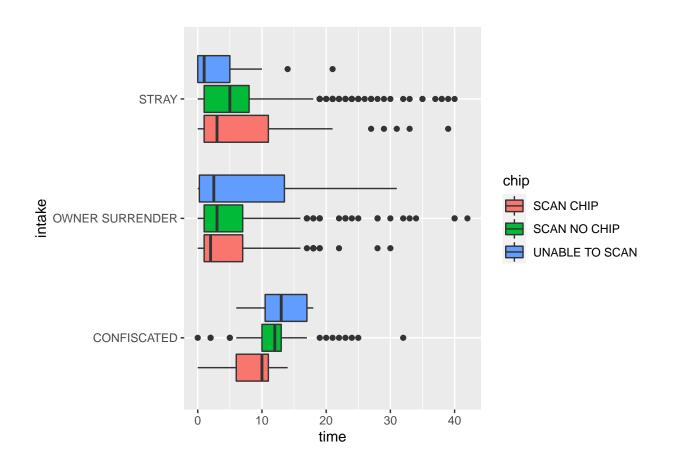
Again all the data is skewed to the right, meaning most types stay for a short period of time. However, it would seem that animals with a chip is a lot more skewed that those that are not. We could speculate that these animals are rescued early and if not then held onto for longer, thus this variable could be a good predictor if combined with outcome perhaps.

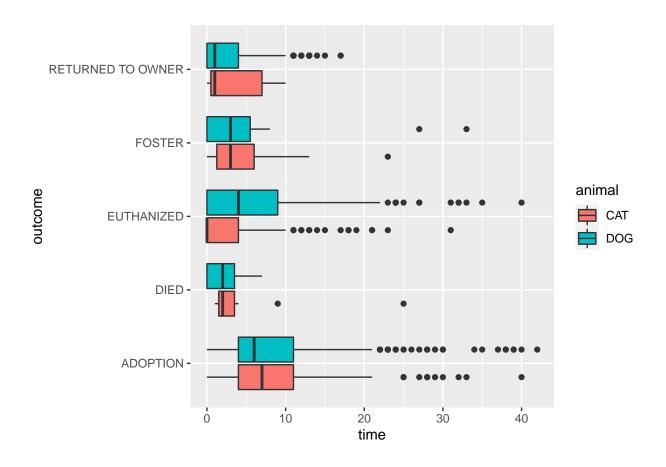


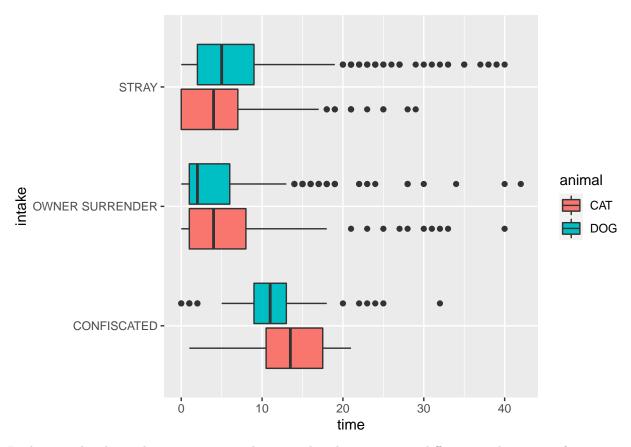


The distributions for month seem to all be similar. We do not envisage that this variable will have an impact on the time variable. Similarly the variable year, which only has 2 levels, may not be a good predictor of time. that means we could potentially have 4 categorical variables explaining one continuous variable **possible loglinear model**.

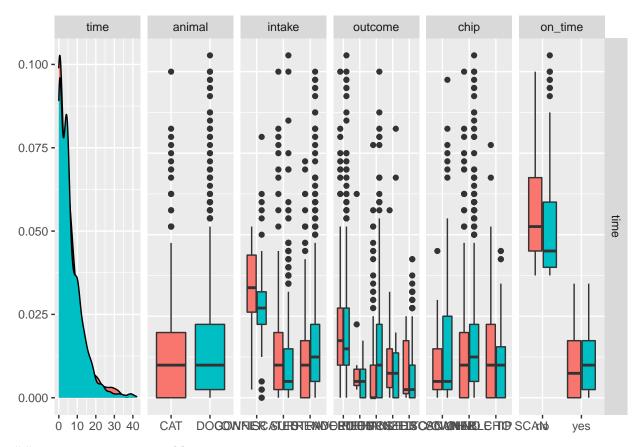




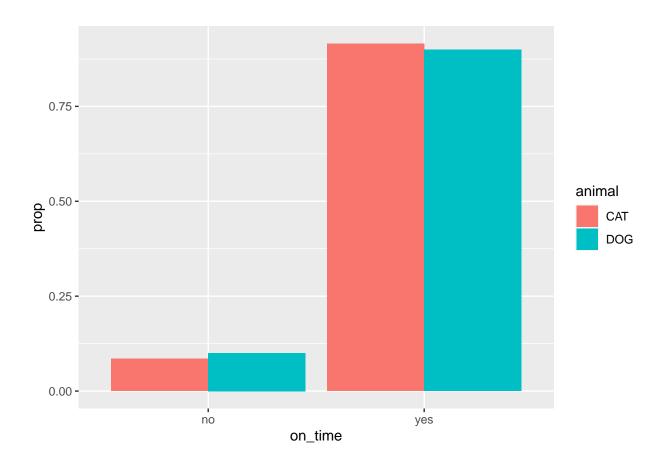


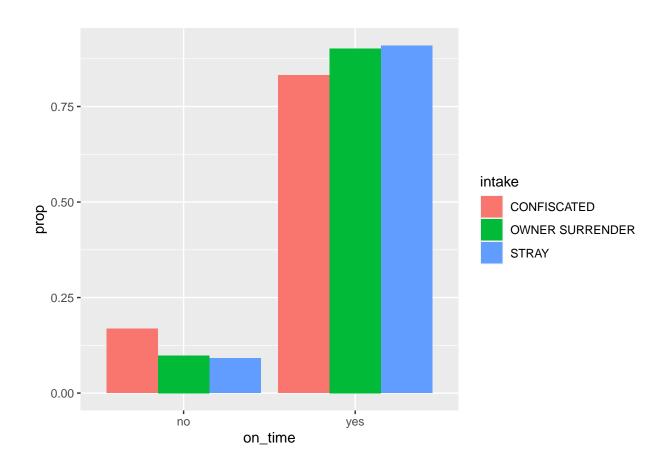


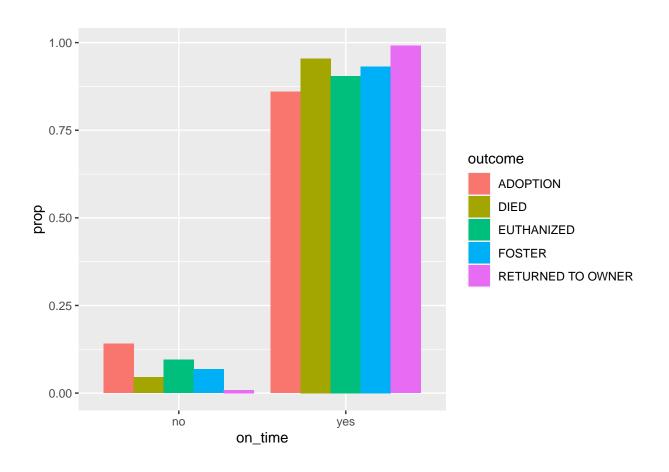
Looking at the above plots, we see some change in distributions across different combinations of categories.

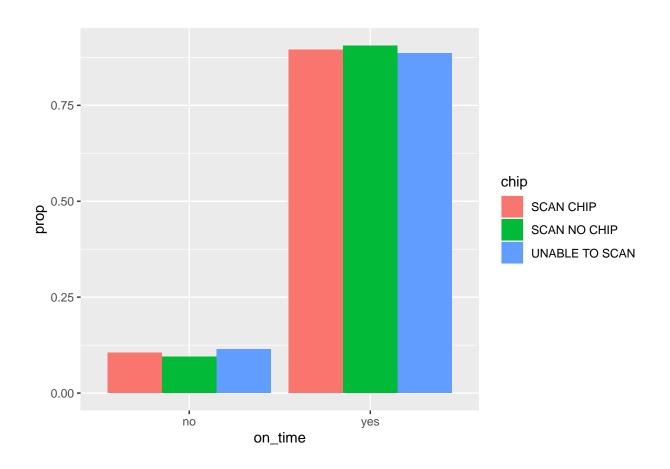


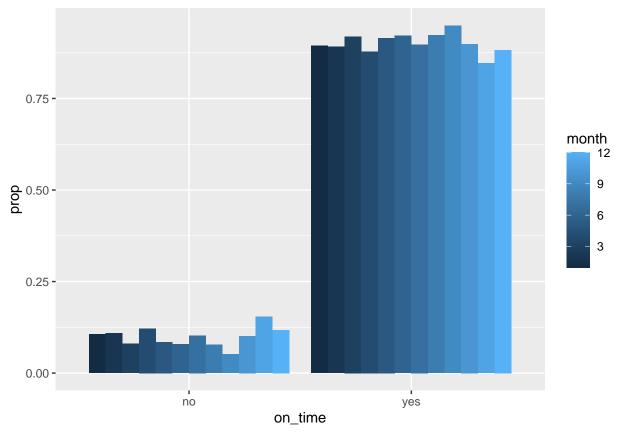
on_time response variable





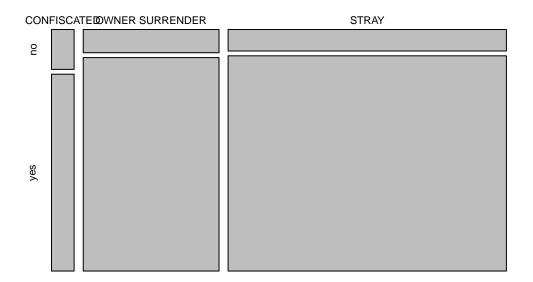




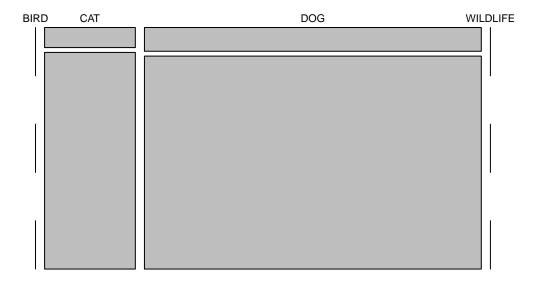


There does not seem to be much difference in proportions between the two types of animals. There is some difference in proportion between confiscated and the other levels of intake. There is some differences from one group to another in the variable outcome. There does not seem to be much differences in proportions for chip. For month, the proportions seem fairly even - cannot see any trend that could relate to seasonal activity.

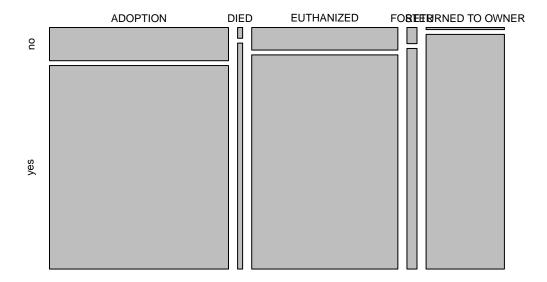
ct_int



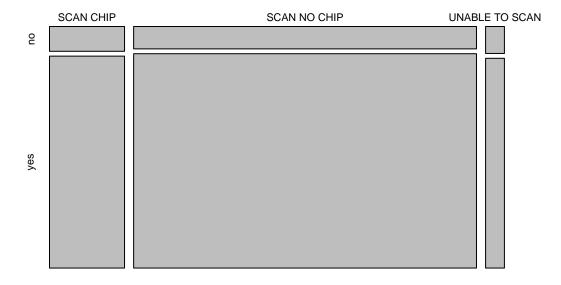
ct_anml



ct_outcome



ct_chip



Very little difference in proportions for yes/no across different levels of each categorical explanatory - maybe do not go for log-odds model.

3 Formal Data Analysis

3.1 Binomial models for on_time

Observations	1833
Dependent variable	on_time
Type	Generalized linear model
Family	binomial
Link	logit

$\chi^{2}(9)$	92.20
Pseudo-R ² (Cragg-Uhler)	0.10
Pseudo-R ² (McFadden)	0.08
AIC	1096.09
BIC	1151.23

BINOMIAL MODEL: From the exploratory analysis, it seemed fitting to try models based on intake and/or outcome, and then build up from there. Having fitted some models, model1 seems to be the best, which is simply intake as the only explanatory variable for the log-odds.

	Est.	S.E.	z val.	p
(Intercept)	0.24	0.45	0.54	0.59
animalDOG	-0.31	0.21	-1.44	0.15
intakeOWNER SURRENDER	1.69	0.35	4.80	0.00
intake STRAY	1.39	0.33	4.18	0.00
outcomeDIED	1.13	1.03	1.09	0.27
outcomeEUTHANIZED	0.53	0.18	2.99	0.00
outcomeFOSTER	0.59	0.62	0.96	0.34
outcomeRETURNED TO OWNER	3.51	0.61	5.77	0.00
chipSCAN NO CHIP	0.44	0.23	1.93	0.05
chipUNABLE TO SCAN	-0.02	0.44	-0.05	0.96

Standard errors: MLE

Observations	1833
Dependent variable	on_time
Type	Generalized linear model
Family	binomial
Link	logit

$\chi^2(4)$	65.23
Pseudo-R ² (Cragg-Uhler)	0.07
Pseudo-R ² (McFadden)	0.06
AIC	1113.07
BIC	1140.63

	Est.	S.E.	z val.	р
(Intercept)	1.81	0.10	17.63	0.00
outcomeDIED	1.23	1.03	1.20	0.23
${\it outcome} {\it EUTHANIZED}$	0.44	0.17	2.58	0.01
outcomeFOSTER	0.80	0.61	1.32	0.19
outcomeRETURNED TO OWNER	2.92	0.59	4.95	0.00

Standard errors: MLE

Observations	1833
Dependent variable	on_time
Type	Generalized linear model
Family	binomial
Link	logit

$\chi^{2}(2)$	5.18
Pseudo-R ² (Cragg-Uhler)	0.01
Pseudo-R ² (McFadden)	0.00
AIC	1169.11
BIC	1185.65

3.2 Poisson models for time

Call:

	Est.	S.E.	z val.	p
(Intercept)	1.60	0.27	5.82	0.00
$intake OWNER\ SURRENDER$	0.62	0.31	2.01	0.04
intake STRAY	0.71	0.29	2.42	0.02

Standard errors: MLE

Observations	1833
Dependent variable	on_time
Type	Generalized linear model
Family	binomial
Link	logit

$\chi^{2}(6)$	84.96
Pseudo-R ² (Cragg-Uhler)	0.10
Pseudo-R ² (McFadden)	0.07
AIC	1097.33
BIC	1135.93

	Est.	S.E.	z val.	p
(Intercept)	0.32	0.33	0.97	0.33
outcomeDIED	1.25	1.03	1.21	0.23
${\bf outcome EUTHANIZED}$	0.53	0.17	3.07	0.00
outcomeFOSTER	0.79	0.61	1.31	0.19
outcomeRETURNED TO OWNER	3.34	0.60	5.52	0.00
$intake OWNER\ SURRENDER$	1.65	0.35	4.78	0.00
intakeSTRAY	1.45	0.33	4.41	0.00

Standard errors: MLE

Observations	1833
Dependent variable	on_time
Type	Generalized linear model
Family	binomial
Link	logit

$\chi^2(4)$	5.42
Pseudo-R ² (Cragg-Uhler)	0.01
Pseudo-R ² (McFadden)	0.00
AIC	1172.88
BIC	1200.45

	Est.	S.E.	z val.	p
(Intercept)	1.55	0.32	4.86	0.00
chipSCAN NO CHIP	0.08	0.21	0.37	0.71
chipUNABLE TO SCAN	-0.05	0.40	-0.13	0.89
$intake OWNER\ SURRENDER$	0.61	0.31	1.98	0.05
intake STRAY	0.69	0.29	2.34	0.02

Standard errors: MLE

Observations	1833
Dependent variable	on_time
Type	Generalized linear model
Family	binomial
Link	$\log it$

$\chi^{2}(3)$	5.77
Pseudo-R ² (Cragg-Uhler)	0.01
Pseudo-R ² (McFadden)	0.00
AIC	1170.52
BIC	1192.58

	Est.	S.E.	z val.	p
(Intercept)	1.74	0.33	5.24	0.00
animalDOG	-0.15	0.20	-0.76	0.45
intakeOWNER SURRENDER	0.60	0.31	1.94	0.05
intake STRAY	0.69	0.29	2.35	0.02

Standard errors: MLE

glm(formula = time ~ intake + animal + chip + outcome, family = poisson(),
 data = shelter)

Deviance Residuals:

Min 1Q Median 3Q Max -5.0878 -2.0942 -0.8433 0.5840 9.8204

Coefficients:

Estimate Std. Error z value Pr(>|z|)0.04922 66.151 < 2e-16 *** (Intercept) 3.25588 intakeOWNER SURRENDER -1.40129 0.03806 -36.820 < 2e-16 *** intakeSTRAY -1.01833 0.03393 -30.013 < 2e-16 *** animalDOG 0.15174 0.02523 6.015 1.80e-09 *** chipSCAN NO CHIP 0.02783 -5.056 4.28e-07 *** -0.14071chipUNABLE TO SCAN -0.10393 0.05517 -1.884 0.0596 . outcomeDIED -0.87212 0.11426 -7.633 2.29e-14 *** outcomeEUTHANIZED -0.64037 0.02225 -28.775 < 2e-16 *** -0.39698 0.06786 -5.850 4.92e-09 *** outcomeFOSTER outcomeRETURNED TO OWNER -1.46341 0.03663 -39.953 < 2e-16 ***

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

(Dispersion parameter for poisson family taken to be 1)

Null deviance: 12201.6 on 1832 degrees of freedom Residual deviance: 9378.3 on 1823 degrees of freedom

AIC: 14517

Number of Fisher Scoring iterations: 6

2.5 % 97.5 % (Intercept) 3.1590085 3.35194721

```
intakeOWNER SURRENDER
                         -1.4755537 -1.32635806
intakeSTRAY
                         -1.0843667 -0.95135276
animalDOG
                          0.1025139 0.20141301
chipSCAN NO CHIP
                         -0.1950108 -0.08591295
chipUNABLE TO SCAN
                         -0.2131352 0.00317712
outcomeDIED
                         -1.1044422 -0.65595751
outcomeEUTHANIZED
                         -0.6840986 -0.59686078
outcomeFOSTER
                         -0.5326087 -0.26647242
outcomeRETURNED TO OWNER -1.5357062 -1.39211393
```

Call:

glm(formula = time ~ intake + outcome, family = poisson(), data = shelter)

Deviance Residuals:

Min 1Q Median 3Q Max -5.0660 -2.0874 -0.9047 0.5466 9.9237

Coefficients:

Estimate Std. Error z value Pr(>|z|)(Intercept) 3.26169 0.03392 96.147 < 2e-16 *** intakeOWNER SURRENDER 0.03731 -37.042 < 2e-16 *** -1.38194 intakeSTRAY -1.030470.03372 -30.561 < 2e-16 *** 0.11401 -8.107 5.18e-16 *** outcomeDIED -0.92431outcomeEUTHANIZED -0.65207 0.02207 -29.551 < 2e-16 *** $\verb"outcomeFOSTER"$ 0.06674 -7.303 2.81e-13 *** -0.48740 outcomeRETURNED TO OWNER -1.40548 0.03578 -39.284 < 2e-16 ***

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

(Dispersion parameter for poisson family taken to be 1)

Null deviance: 12201.6 on 1832 degrees of freedom Residual deviance: 9448.5 on 1826 degrees of freedom

AIC: 14581

Number of Fisher Scoring iterations: 6

2.5 % 97.5 % (Intercept) 3.1946881 3.3276782 intakeOWNER SURRENDER -1.4547129 -1.3084578 intakeSTRAY -1.0960888 -0.9639035 outcomeDIED -1.1561811 -0.7086556 outcomeEUTHANIZED -0.6954363 -0.6089368 outcomeFOSTER -0.6209137 -0.3591875 outcomeRETURNED TO OWNER -1.4761247 -1.3358652

[1] FALSE

[1] FALSE

POISSON MODEL: looking at model6, using it has a very high pearson residual which is higher than the chi-squared statistic meaning it is probably suffering from overdispersion. To deal with this could be to introduce a dispersion parameter and have a quasi-poisson model OR we go for a negative binomial model.

```
Call:
glm(formula = time ~ intake + outcome, family = poisson(), data = shelter)
Deviance Residuals:
                Median
   Min
            1Q
                             3Q
                                    Max
-5.0660 -2.0874 -0.9047 0.5466
                                  9.9237
Coefficients:
                       Estimate Std. Error z value Pr(>|z|)
(Intercept)
                       3.26169 0.08524 38.265 < 2e-16 ***
intakeOWNER SURRENDER
                                 0.09374 -14.742 < 2e-16 ***
                       -1.38194
intakeSTRAY
                      -1.03047 0.08472 -12.163 < 2e-16 ***
                       outcomeDIED
outcomeEUTHANIZED
                       outcomeFOSTER
                       -0.48740
                                  0.16769 -2.907 0.00365 **
outcomeRETURNED TO OWNER -1.40548 0.08990 -15.634 < 2e-16 ***
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
(Dispersion parameter for poisson family taken to be 6.313406)
   Null deviance: 12201.6 on 1832 degrees of freedom
Residual deviance: 9448.5 on 1826 degrees of freedom
AIC: 14581
Number of Fisher Scoring iterations: 6
Single term deletions
Model:
time ~ intake + outcome
       Df Deviance
                   AIC F value
                                 Pr(>F)
           9448.5 14581
intake 2 10587.1 15715 110.03 < 2.2e-16 ***
outcome 4 11750.8 16875 111.24 < 2.2e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
Coefficients still significant (by F-tests) but deviance still above chi-squared.
Call:
glm.nb(formula = time ~ intake + outcome + chip + animal, data = shelter,
   init.theta = 1.114925073, link = log)
Deviance Residuals:
   Min
           1Q
                Median
                             ЗQ
                                    Max
-2.1487 -1.0149 -0.3325 0.2207
                                  3.6358
Coefficients:
                       Estimate Std. Error z value Pr(>|z|)
```

3.48306

(Intercept)

0.14330 24.306 < 2e-16 ***

```
-1.62161
intakeOWNER SURRENDER
                             0.11844 -13.692 < 2e-16 ***
intakeSTRAY
                    -1.25740 0.11014 -11.416 < 2e-16 ***
outcomeDIED
                   -0.71229 0.05612 -12.691 < 2e-16 ***
outcomeEUTHANIZED
outcomeFOSTER
                    -0.43965
                             0.16496 -2.665 0.00769 **
chipSCAN NO CHIP -0.12566 0.07004 -1.794 0.07279 .
                   -0.07656 0.13781 -0.556 0.57852
chipUNABLE TO SCAN
animalDOG
                    0.17273
                            0.06356 2.718 0.00658 **
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
(Dispersion parameter for Negative Binomial(1.1149) family taken to be 1)
   Null deviance: 2660.0 on 1832 degrees of freedom
Residual deviance: 2156.3 on 1823 degrees of freedom
AIC: 10079
Number of Fisher Scoring iterations: 1
           Theta: 1.1149
        Std. Err.: 0.0507
2 x log-likelihood: -10057.4320
Call:
glm.nb(formula = time ~ intake + outcome, data = shelter, init.theta = 1.103703043,
   link = log)
Deviance Residuals:
   Min
       1Q
              Median
                         3Q
                                Max
-2.1038 -0.9345 -0.4035 0.2407
                             3.3850
Coefficients:
                    Estimate Std. Error z value Pr(>|z|)
(Intercept)
                    3.51491 0.11394 30.849 < 2e-16 ***
intakeOWNER SURRENDER
                    -1.62341 0.11796 -13.762 < 2e-16 ***
intakeSTRAY
                    0.23616 -4.030 5.58e-05 ***
outcomeDIED
                    -0.95171
                    outcomeEUTHANIZED
outcomeFOSTER
                    -0.52936
                             0.16266 -3.254 0.00114 **
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
(Dispersion parameter for Negative Binomial(1.1037) family taken to be 1)
   Null deviance: 2642.4 on 1832 degrees of freedom
Residual deviance: 2154.8 on 1826 degrees of freedom
AIC: 10085
```

Number of Fisher Scoring iterations: 1

Theta: 1.1037 Std. Err.: 0.0500

2 x log-likelihood: -10069.4000

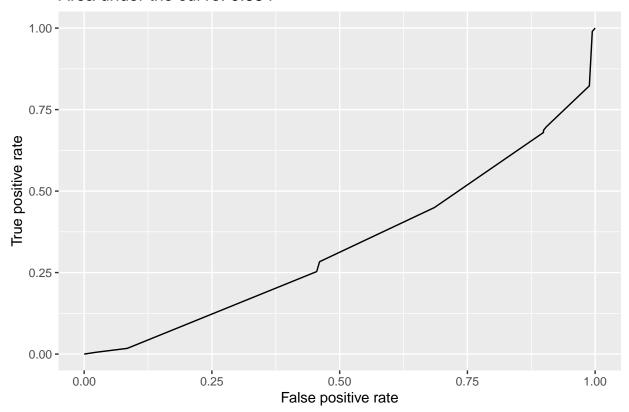
intakeSTRAY	intakeOWNER SURRENDER	(Intercept)
0.2831305	0.1972248	33.6128993
outcomeFOSTER	${\tt outcomeEUTHANIZED}$	${\tt outcomeDIED}$
0.5889826	0.4933791	0.3860798

outcomeRETURNED TO OWNER

0.2110144

	2.5 %	97.5 %
(Intercept)	26.8856526	42.0234173
intakeOWNER SURRENDER	0.1565133	0.2485260
intakeSTRAY	0.2281128	0.3514178
outcomeDIED	0.2430290	0.6133324
outcomeEUTHANIZED	0.4422130	0.5504654
outcomeFOSTER	0.4282019	0.8101330
outcomeRETURNED TO OWNER	0.1815460	0.2452663

Area under the curve: 0.334



$$\widehat{on\ time}\ =\ 33.613 + 0.197 \cdot I_{Surrender}\ (i) + 0.283 \cdot I_{Stray}\ (i) + 0.386 \cdot I_{Died}\ (j) + 0.493 \cdot I_{Euthanized}\ (j) + 0.0.589 \cdot I_{Foster}\ (j)$$

$$I_{\hbox{Surrender}}(i) = \left\{ \begin{array}{ll} 1 & \hbox{if Position of ith observation is Surrender}, \\ 0 & \hbox{Otherwise}. \end{array} \right.$$

```
I_{\mathrm{Stray}}(i) = \left\{ \begin{array}{l} 1 & \mathrm{if\ Position\ of}\ ith\ \mathrm{observation\ is\ Stray}, \\ 0 & \mathrm{Otherwise}. \end{array} \right. I_{\mathrm{Died}}(j) = \left\{ \begin{array}{l} 1 & \mathrm{if\ Position\ of}\ jth\ \mathrm{observation\ is\ Died}, \\ 0 & \mathrm{Otherwise}. \end{array} \right. I_{\mathrm{Euthanized}}(j) = \left\{ \begin{array}{l} 1 & \mathrm{if\ Position\ of}\ jth\ \mathrm{observation\ is\ Euthanized}, \\ 0 & \mathrm{Otherwise}. \end{array} \right. I_{\mathrm{Foster}}(j) = \left\{ \begin{array}{l} 1 & \mathrm{if\ Position\ of}\ jth\ \mathrm{observation\ is\ Foster}, \\ 0 & \mathrm{Otherwise}. \end{array} \right. I_{\mathrm{Returned}}(j) = \left\{ \begin{array}{l} 1 & \mathrm{if\ Position\ of}\ jth\ \mathrm{observation\ is\ Returned}, \\ 0 & \mathrm{Otherwise}. \end{array} \right. I_{\mathrm{Returned}}(j) = \left\{ \begin{array}{l} 1 & \mathrm{if\ Position\ of}\ jth\ \mathrm{observation\ is\ Returned}, \\ 0 & \mathrm{Otherwise}. \end{array} \right.
```

Conclusions $\{\#sec:Conc\}$