

ZERO-INFLATED STATISTICAL MODELS FOR ANIMAL BEHAVIOR STUDIES

Jay Harrison

Too many zeroes?

Hypothetical data: Which is the better treatment?

Treatment 1

0
0
0
200

Average: 50

Average without zeroes: 200

Sample size without zeroes: 1

Treatment 2

100
120
180
200

Average: 150

Average without zeroes: 150

Sample size without zeroes: 4

AFB International manufactures palatants for pet food, treats, and toys.

- <http://afbinternational.com/research-and-development>
- Palatability Assessment Resource Center
- Standard test: two bowls, two days

Which science is hardest?



Edward Tufte @EdwardTufte · 18 Sep 2016

Rocket science made difficult.

[#thinking](#) [#teaching](#) [#dataviz](#) [#analytics](#)



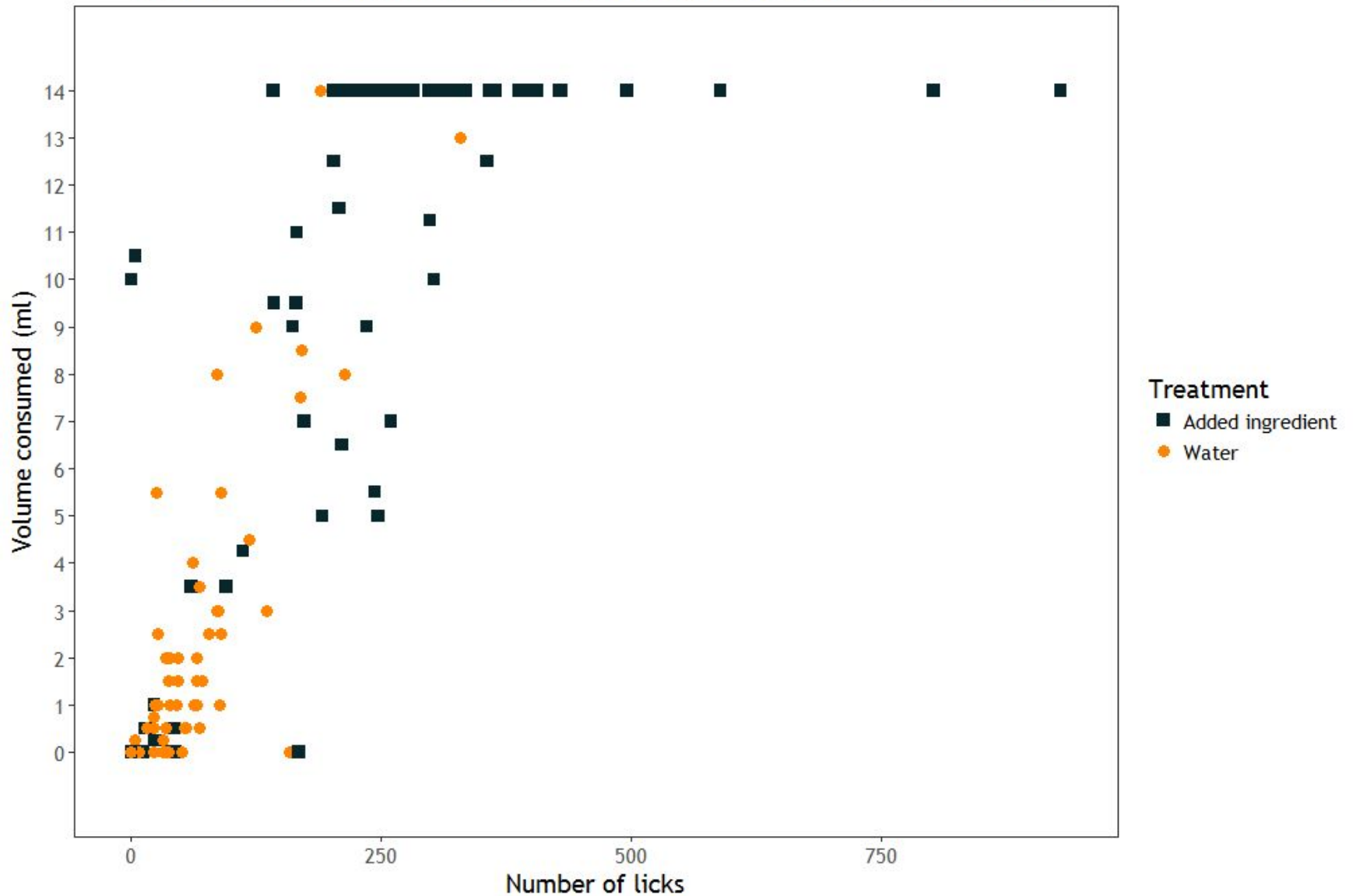
Analysis of human behavior
isn't rocket science.

It's harder than rocket science.



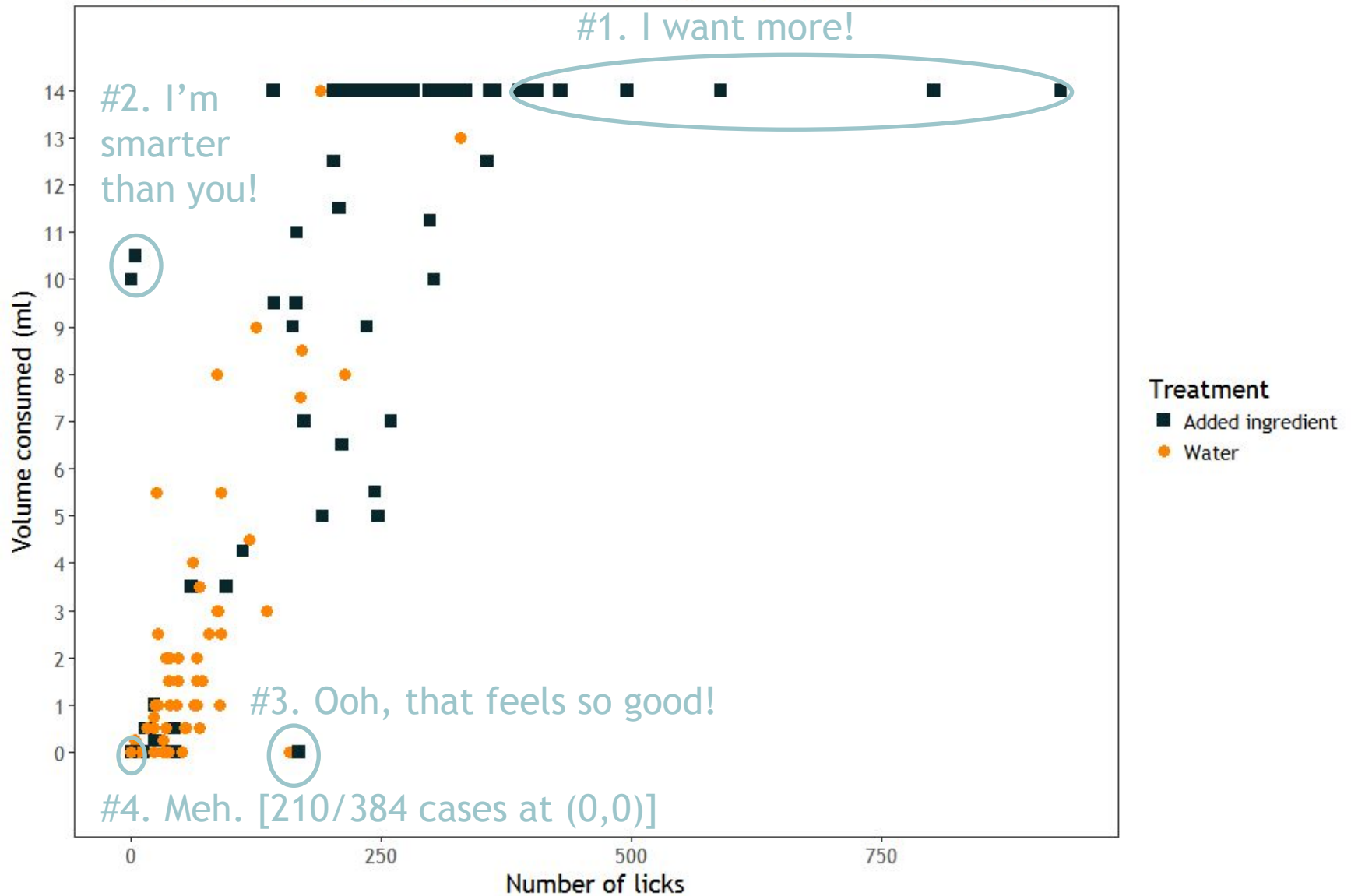
Example - Lickometer study

Licks and consumption for tests comparing added ingredient to water



Special cases

Licks and consumption for tests comparing added ingredient to water



Displaying extra zeroes

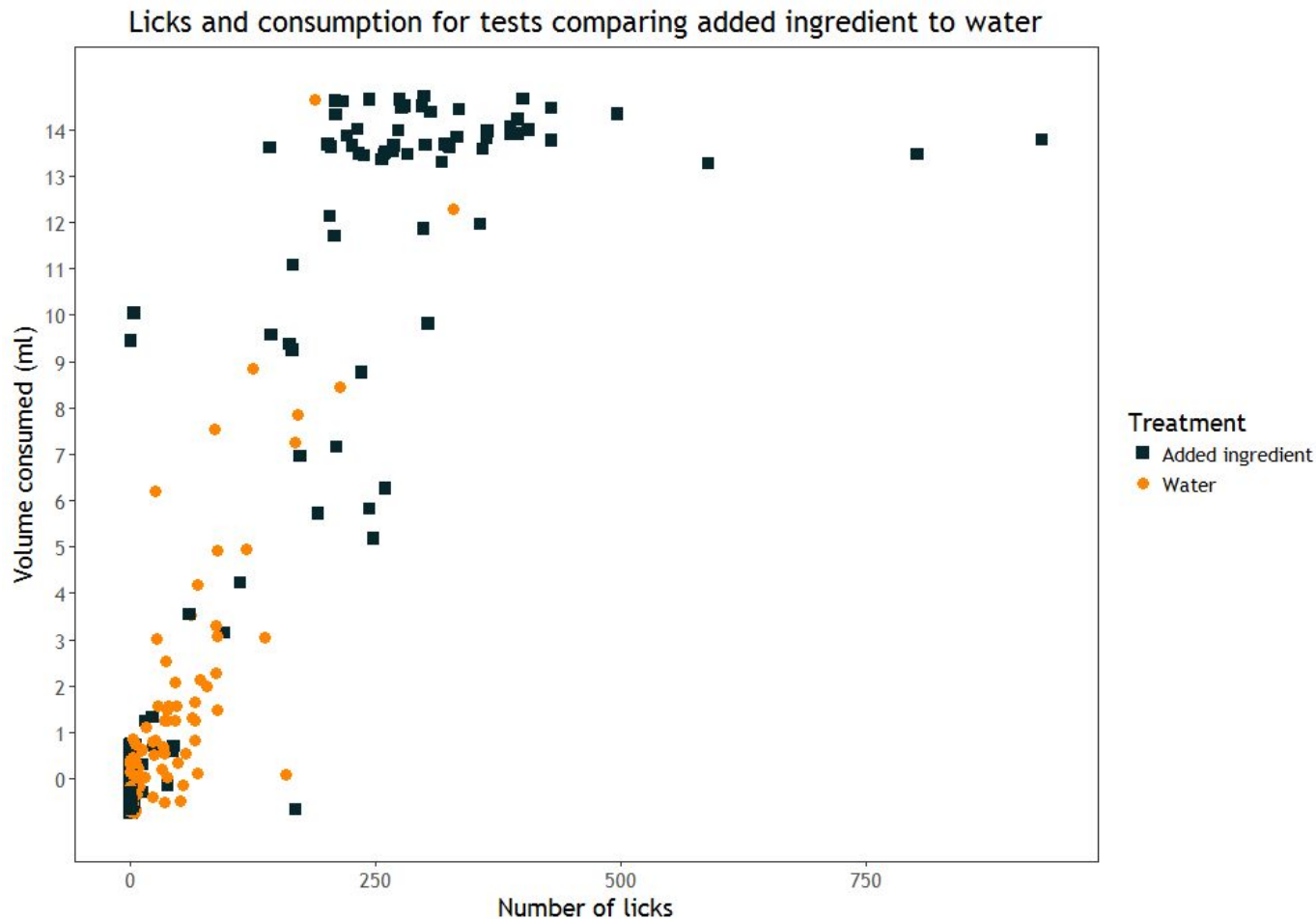
1. Use jittering to indicate clusters of points.

Data frame licks:

Test <chr>	Animal <chr>	Day <dbl>	Ration <chr>	Licks <dbl>	Consumed <dbl>
Test1	Cat1	1	A	236	9
Test1	Cat1	1	B	0	0
Test1	Cat1	2	A	211	6.5
Test1	Cat1	2	B	0	0
Test1	Cat2	1	A	0	0
Test1	Cat2	1	B	0	0
Test1	Cat2	2	A	256	14
Test1	Cat2	2	B	68	0.5
etc.					

Displaying extra zeroes

1. Use jittering to indicate clusters of points.



```
set.seed(79311)  
library(ggplot2)
```

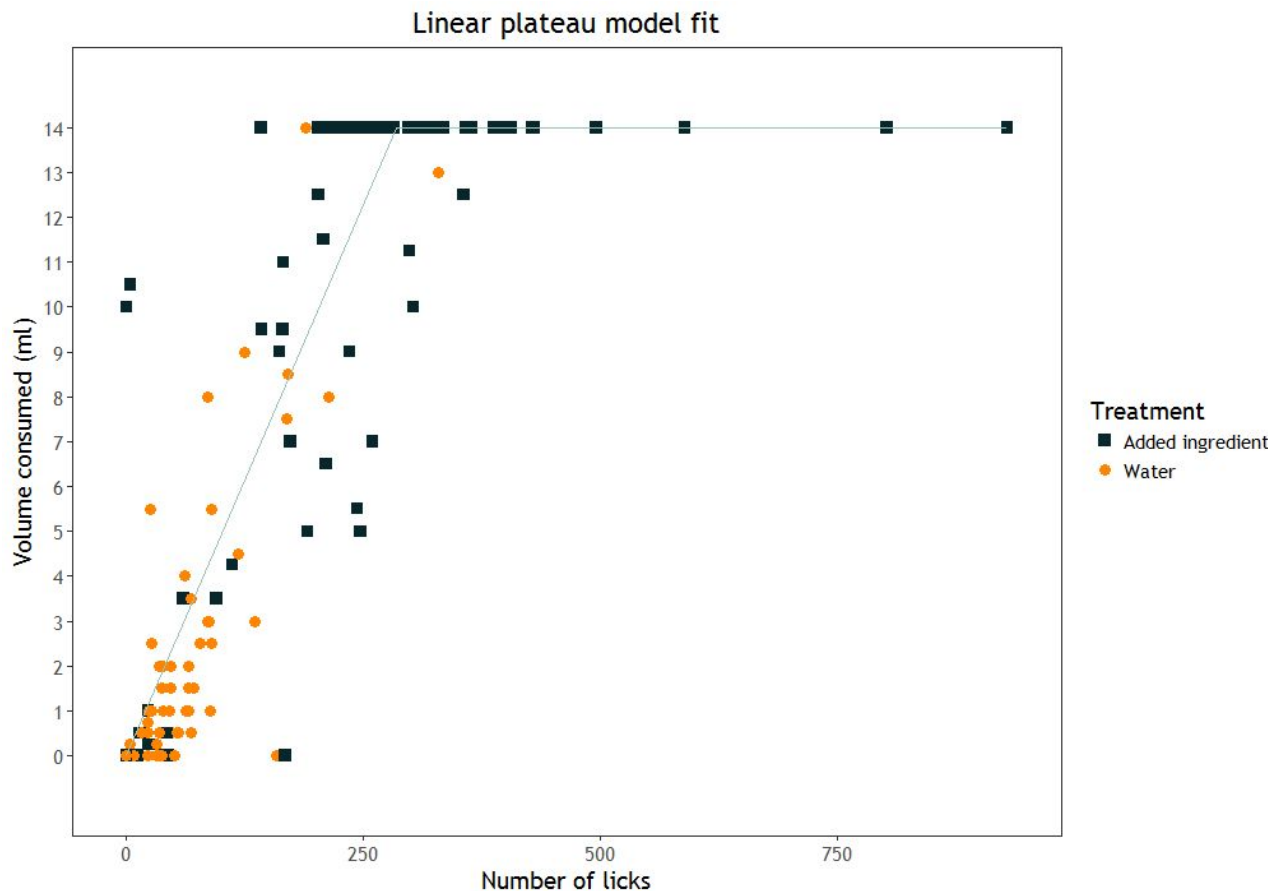
```
plot1 <- ggplot(licks,  
  aes(x=Licks,  
    y=Consumed)) +  
  geom_jitter(size=3,  
    height=.75,width=.75,  
    aes(shape=Ration,  
      colour=Ration))
```

plot1

This illustrates
density at
the expense of
accuracy.

Displaying extra zeroes

2. Use a regression line that can be influenced by the zeroes to indicate the trend.



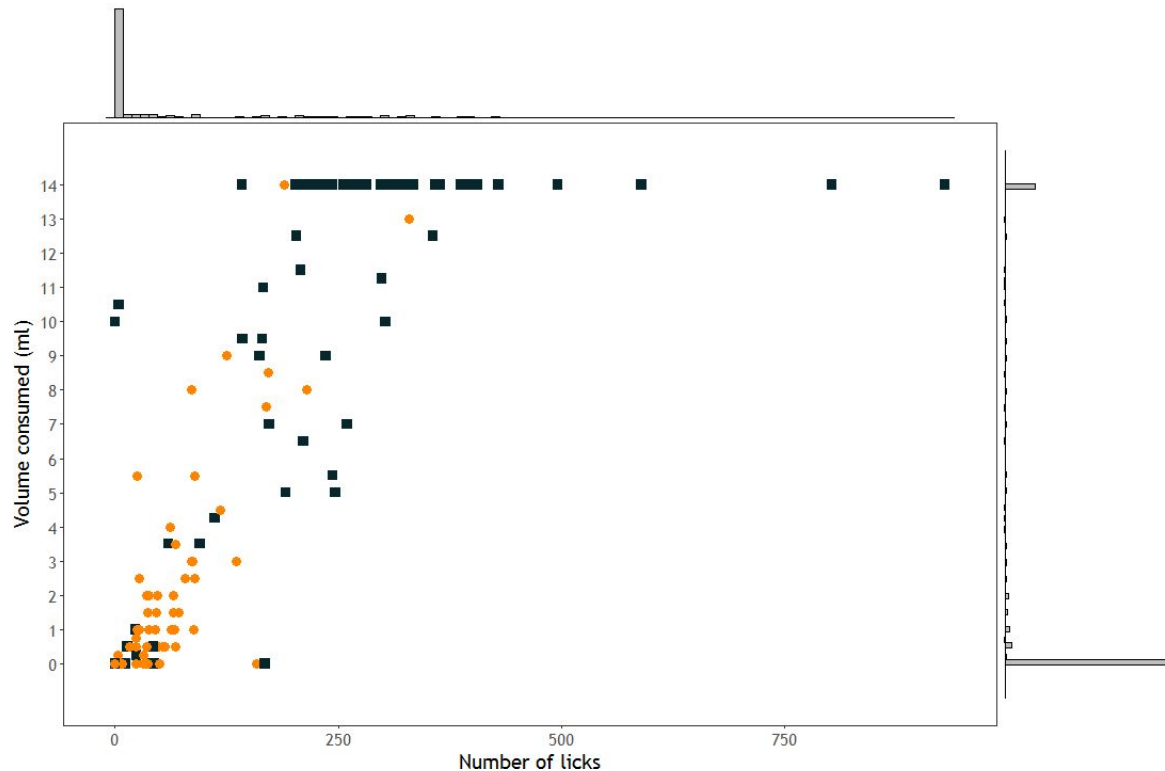
```
library(stats)
```

```
fornls <- licks %>%  
select(Licks,  
Consumed)
```

```
plateau <- nls(Consumed ~  
(b*Licks)*(b*Licks<=14) +  
14*(b*Licks>14),  
data=fornls,  
start=list(b=1))  
splateau <-  
summary(plateau)  
joinpoint <- 14/  
splateau$parameters[1,1]
```

Displaying extra zeroes

3. Use marginal histograms in the ggExtra package to indicate large frequencies at (0,0) and at 14 ml.



```
library(ggExtra)
```

```
plot1b <- ggplot(licks,  
  aes(x=Licks,  
    y=Consumed)) + ...
```

```
ggMarginal(plot1b,  
  type="histogram",  
  margins="both",  
  bins=100)
```

Can the numbers of zeroes be reduced?

Example: Use two-day feeding totals instead of single days

One or two zeroes do not cause excessive violations of ordinary statistical assumptions.

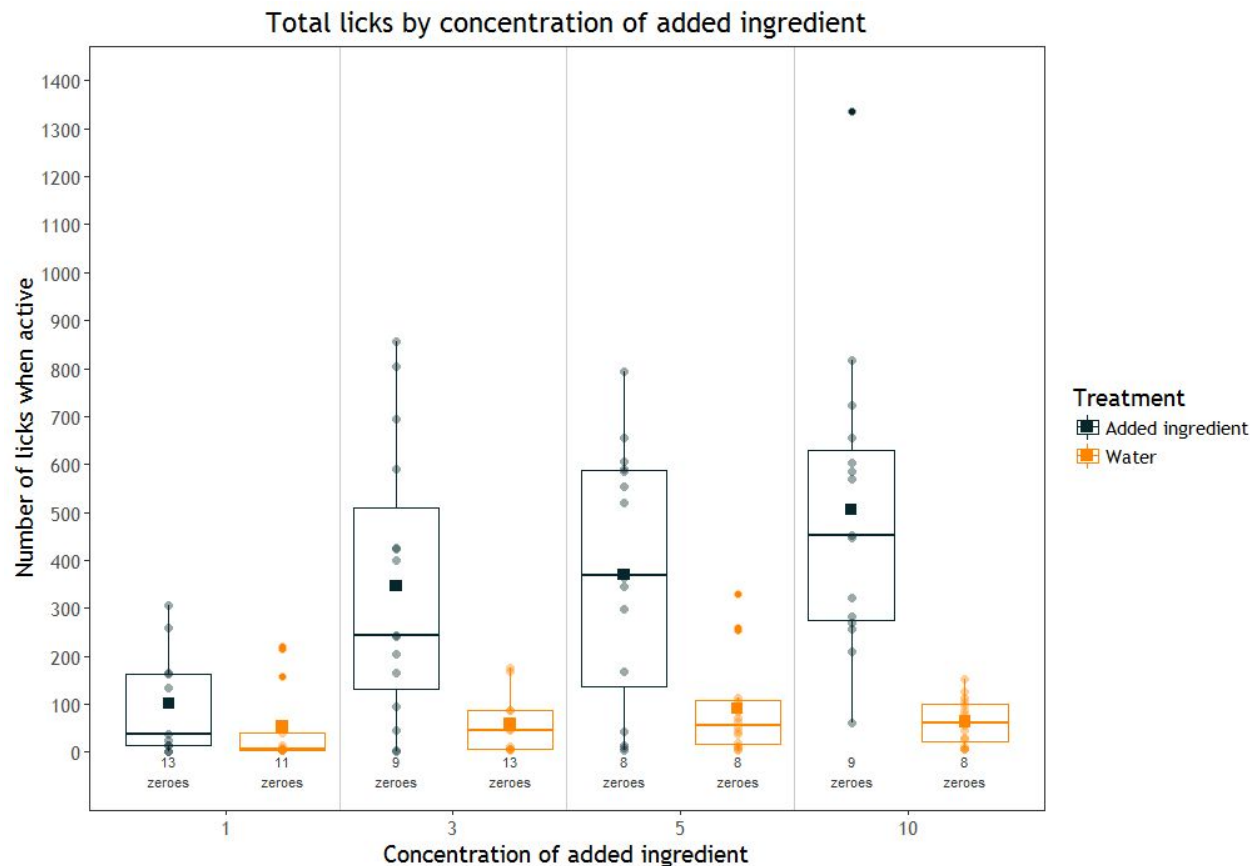
ZINB - Zero-inflated negative binomial model

Data frame licks2 counts the total number of licks over 2 days (Test.Conc identifies the pairing and the concentration)

Animal. <fctr>	Test.Conc <fctr>	Ration <chr>	Licktotal <dbl>	xconc <dbl>
Cat1	Test1.1	A	0	1
Cat1	Test1.0	B	4	0
Cat2	Test1.1	A	0	1
Cat2	Test1.0	B	0	0
Cat3	Test1.1	A	25	1
Cat3	Test1.0	B	0	0
etc.				

Displaying extra zeroes

4. Consider the total number of licks over 2 exposures to be the primary response of interest and display univariate data with annotated boxplots.



```
plot1 <- ggplot(licks2 %>%  
  filter(Licktotal>0),  
  aes(x=xconc,y=Licktotal,  
  colour=Ration)) +
```

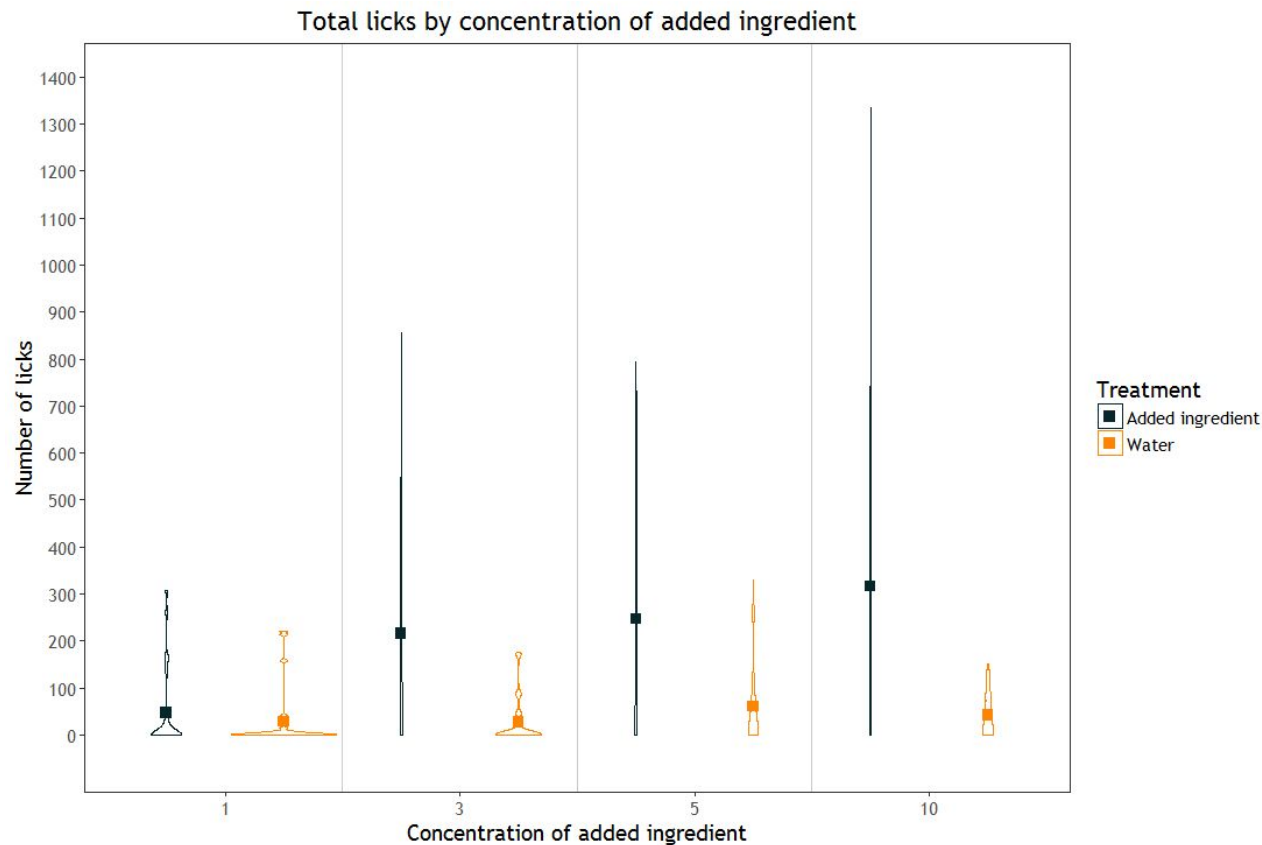
```
  geom_boxplot(position=  
    position_dodge(1)) +
```

```
  geom_point(position=  
    position_dodge(1)) +
```

```
  stat_summary(fun.y=mean,  
    geom="point") +
```

```
  geom_text(data=lickdata,  
    aes(x=seq(.75,4.25,by=.5),  
    y=rep(-40,8)),  
    label=paste(lickdata$num0s,  
    "zeroes",sep="\n"))
```

5. Violin plots

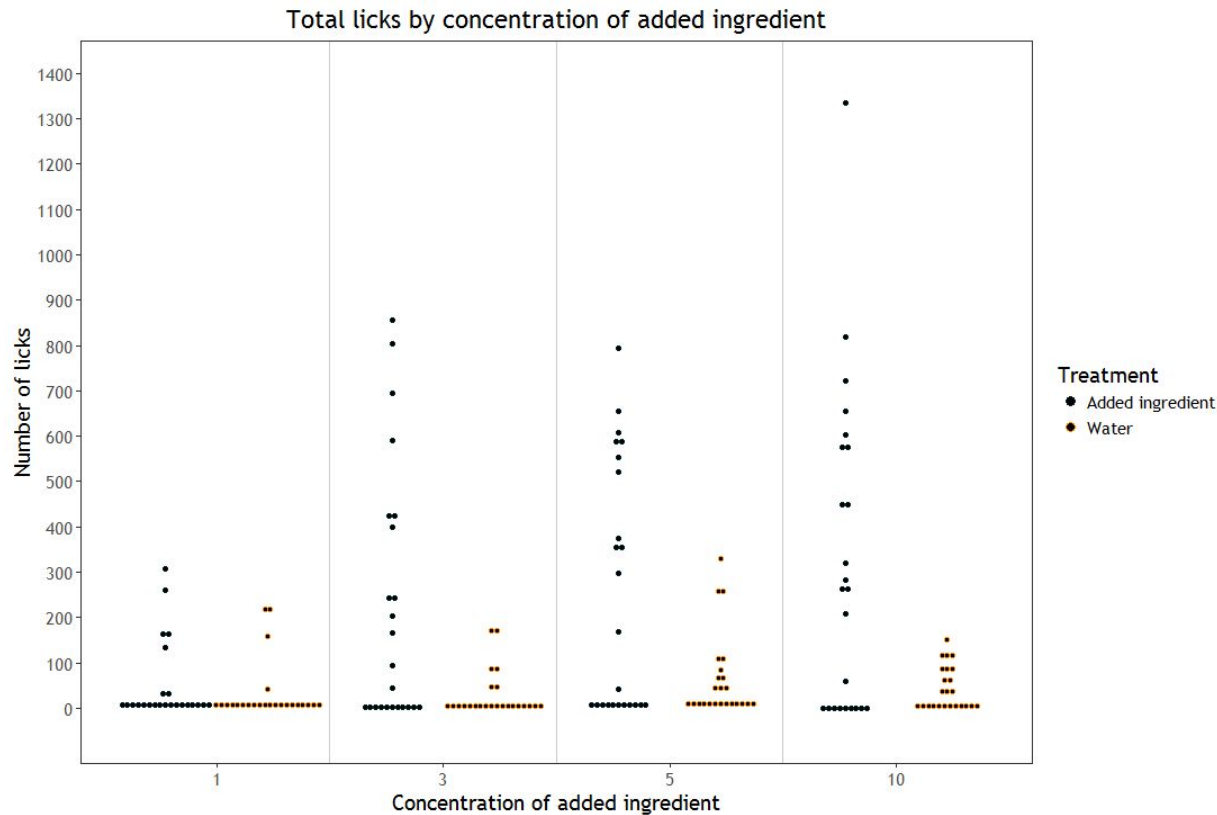


```
plot2 <- ggplot(licks2,  
aes(x=xconc,y=Licktotal,  
colour=Ration)) +
```

```
geom_violin(position=  
position_dodge(1)) +
```

```
stat_summary(fun.y=mean,  
geom="point", shape=15,  
size=3,position=  
position_dodge(width=1))
```

6. Alternative: Stacked dot plots



```
plot3 <- ggplot(licks2,  
  aes(x=xconc, y=Licktotal,  
  colour=Ration)) +
```

```
  geom_dotplot(binaxis='y',  
  stackdir='center',  
  dotsize=.6,  
  binwidth=20,  
  position="dodge")
```

Tabular summaries with excessive zeroes

A table of descriptive statistics was used to explicitly draw attention to the high numbers of zeroes.

Test	Ration	Concentration of added ingredient	No. of cats	No. of cats with 0 licks	% of cats with 0 licks	No. of cats with >0 licks	Avg. licks (when positive)	Min. no. of licks (when positive)	Median no. of licks (when positive)	Max. no. of licks (when positive)	Total no. of licks for all cats
1	A	1x	24	13	54	11	102	1	38	307	1121
1	B	0	24	11	46	13	53	3	6	221	686
2	A	3x	24	9	38	15	346	1	244	855	5191
2	B	0	24	13	54	11	58	3	45	175	641
3	A	5x	24	8	33	16	370	4	369	793	5923
3	B	0	24	8	33	16	91	3	56	329	1459
4	A	10x	24	9	38	15	506	60	452	1336	7583
4	B	0	24	8	33	16	63	6	62	152	1014

Note that $\text{Max} > 2(\text{Avg})$

Tabular summaries with excessive zeroes

The table was generated using dplyr:

```
lickdata <- licks2 %>%  
  group_by(Test, Ration, xconc) %>%  
  summarize(num.cats=n(), num.zeroes=sum(Licktotal==0),  
    pct.zero=round(100*num.zeroes/num.cats),  
    num.positive=sum(Licktotal>0),  
    avg.positive=mean(Licktotal[Licktotal>0]),  
    min.licks=min(Licktotal[Licktotal>0]),  
    median.licks=median(Licktotal[Licktotal>0]),  
    max.licks=max(Licktotal[Licktotal>0]),  
    total.licks=sum(Licktotal))
```

- Hurdle models combine a left-truncated count component with a right-censored hurdle component
- Zeroes can only occur in the count component.

Example: A laboratory instrument reports 0 if the amount of a substance falls below a detection limit.

Modeling data with excessive zeroes

- Zero-inflated models combine a count component and a point mass at zero.
- Zeroes can arise in both situations.

Example: A participant may not be able to complete a task; or, if able, may choose not to do it.

Both hurdle and zero-inflated models are provided in the pscl package in R (Zeileis, Kleiber, and Jackman, “Regression Models for Count Data in R,” available at cran.r-project.org/web/packages/pscl/vignettes/countreg.pdf).

Other applications of these models include:

- econometrics
- political science (PSCL @ Stanford)
- agriculture

Models for count portion of a zero-inflated model

1. Poisson

- a) Traditional application: Number of occurrences of an event per unit of time or space
- b) $\text{Variance} = \text{mean}$

Models for count portion of a zero-inflated model

2. Negative binomial

- a) Traditional application: Number of binomial trials needed to achieve a specified number of successes
- b) Variance $>$ mean
- c) With two parameters for mean and variance, and multiple equivalent parameterizations for a negative binomial distribution, results are harder to interpret

ZINB - Zero-inflated negative binomial model

Data frame licks2 (Test.Conc identifies the pairing and the concentration)

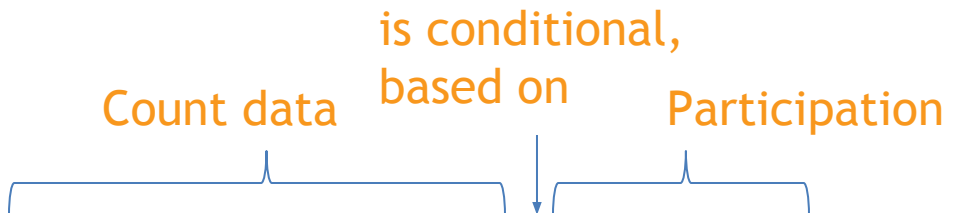
Animal.	Test.Conc	Ration	Licktotal
<fctr>	<fctr>	<chr>	<dbl>
Cat1	Test1.1	A	0
Cat1	Test1.0	B	4
Cat2	Test1.1	A	0
Cat2	Test1.0	B	0
Cat3	Test1.1	A	25
Cat3	Test1.0	B	0
etc.			

ZINB - Zero-inflated negative binomial model

Model in R (zeroinfl is in the pscl library)

Count data is conditional, based on Participation

```
zinb0 <- zeroinfl(Licktotal~Test.Conc + Animal. | Test.Conc,  
data=licks2, dist="negbin",link="logit")
```



The chance that a cat will “decide to participate” is assumed to be related to the ingredients in the bottles, using (by default) a logit link.

Given that a cat chooses to participate, the mean number of licks is related to an ingredient effect and a cat effect via a log link.

ZINB - Zero-inflated negative binomial model

```
> summary(zinb0)
```

```
Pearson residuals:
```

	Min	1Q	Median	3Q	Max
	-1.003604	-0.654252	-0.001159	0.209992	4.304403

```
Count model coefficients (negbin with log link):
```

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-0.5302	0.6813	-0.778	0.436481
Test.concTest1.0	-0.1333	0.4449	-0.300	0.764509
Test.concTest0.0	0.1343	0.3999	0.336	0.737021
...				
Animal.Cat1	4.4654	0.7964	5.607	2.06e-08 ***
Animal.Cat2	4.2511	0.7805	5.447	5.13e-08 ***
...				
Log(theta)	0.1848	0.1526	1.211	0.226046

```
Zero-inflation model coefficients (binomial with logit link):
```

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-0.8903	0.5348	-1.665	0.0959 .
Test.concTest1.0	0.3002	0.7527	0.399	0.6900
Test.concTest1.1	-17.0435	3440.1875	-0.005	0.9960
...				

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

```
Theta = 1.2029
```

```
Number of iterations in BFGS optimization: 57
```

```
Log-likelihood: -710.2 on 40 Df
```

ZINB - Zero-inflated negative binomial model

```
> names(zinb0)
```

"coefficients"	"residuals"	"fitted.values"	"optim"	"method"
"control"	"start"	"weights"	"offset"	"n"
"df.null"	"df.residual"	"terms"	"theta"	"SE.logtheta"
"loglik"	"vcov"	"dist"	"link"	"linkinv"
"converged"	"call"	"formula"	"levels"	"contrasts"
"model"	"y"			

ZINB - Zero-inflated negative binomial model

```
zinb0 <- zeroinfl(Licktotal~Test.Conc + Animal. | Test.Conc,  
data=licks2, dist="negbin", link="logit")
```

```
zinb1<- zeroinfl(Licktotal~Test.Conc + Animal. | 1,  
data=licks2, dist="negbin", link="logit")
```

```
vuong(zinb0,zinb1) #From package pscl  
lrtest(zinb0,zinb1) #From package lmttest
```

If there was not a significant feature (such as color or scent) that attracted the cats to one bottle instead of the other, then the Test.Conc term on the right can be a simple intercept (1).

ZINB - Zero-inflated negative binomial model

```
zinb0 <- zeroinfl(Licktotal~  
Test.Conc + Animal. |  
Test.Conc,  
data=licks2, dist="negbin", link="logit")
```

```
zinb2<- zeroinfl(Licktotal~  
Test.Conc + Animal. |  
Test.Conc + Animal.,  
data=licks2, dist="negbin", link="logit")
```

The second model did not converge, because some cats never registered a lick.

ZINB - Zero-inflated negative binomial model

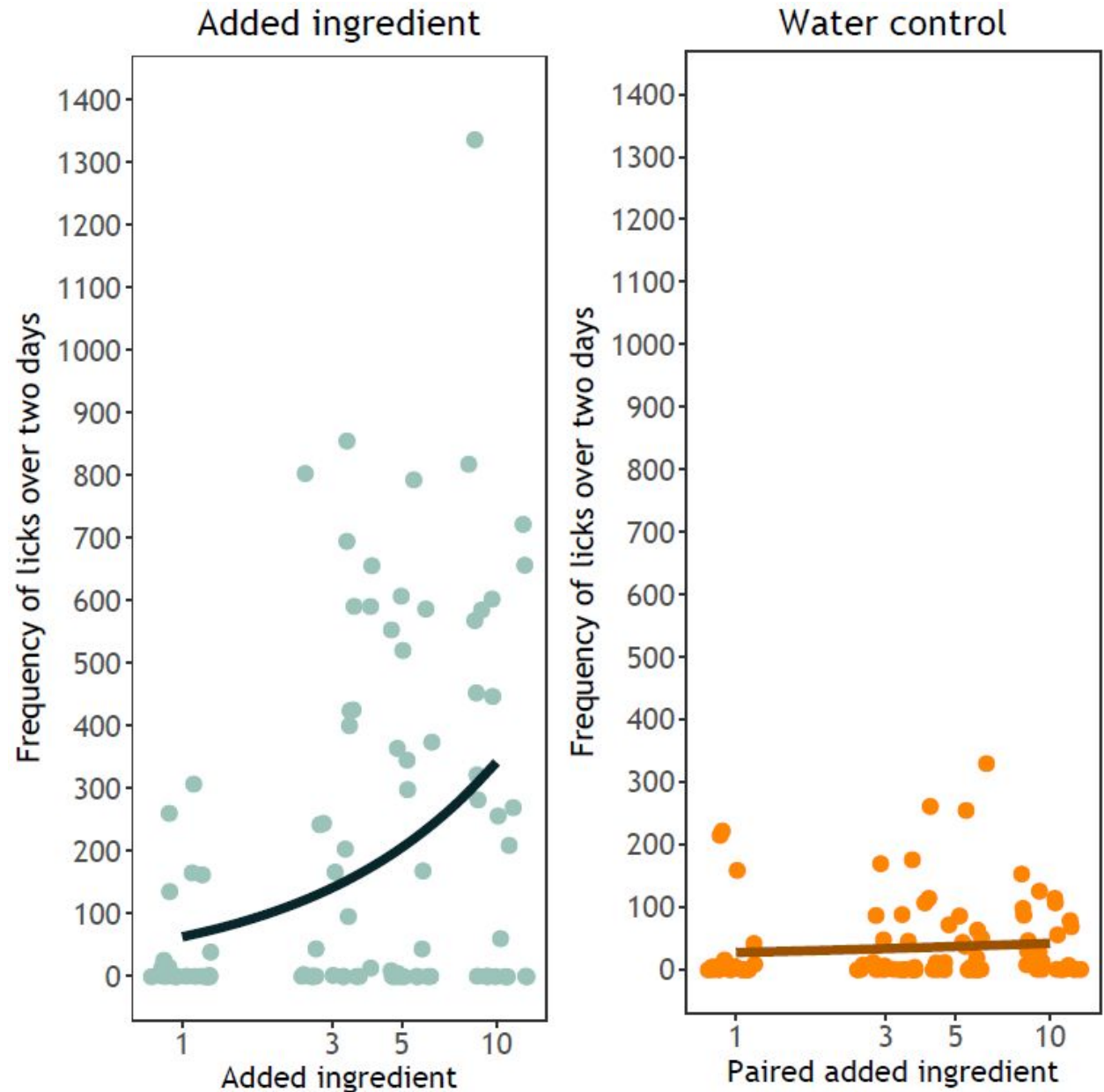
```
zinb0 <- zeroinfl(Licktotal~Test.Conc +  
Animal.|Test.Conc, data=licks2, dist="negbin",  
link="logit")
```

```
zinb3 <- zeroinfl(Licktotal~1 + controlintercept +  
logConc + controlslope + Animal.|1,  
data=licks2, dist="negbin", link="logit")
```

Idea: Replace the fixed effect of the different concentrations (1x, 3x, 5x, and 10x) with a slope with covariate values of $\log(1)$, $\log(3)$, $\log(5)$, and $\log(10)$ to obtain a dose-response relationship.

ZINB - Zero-inflated negative binomial model

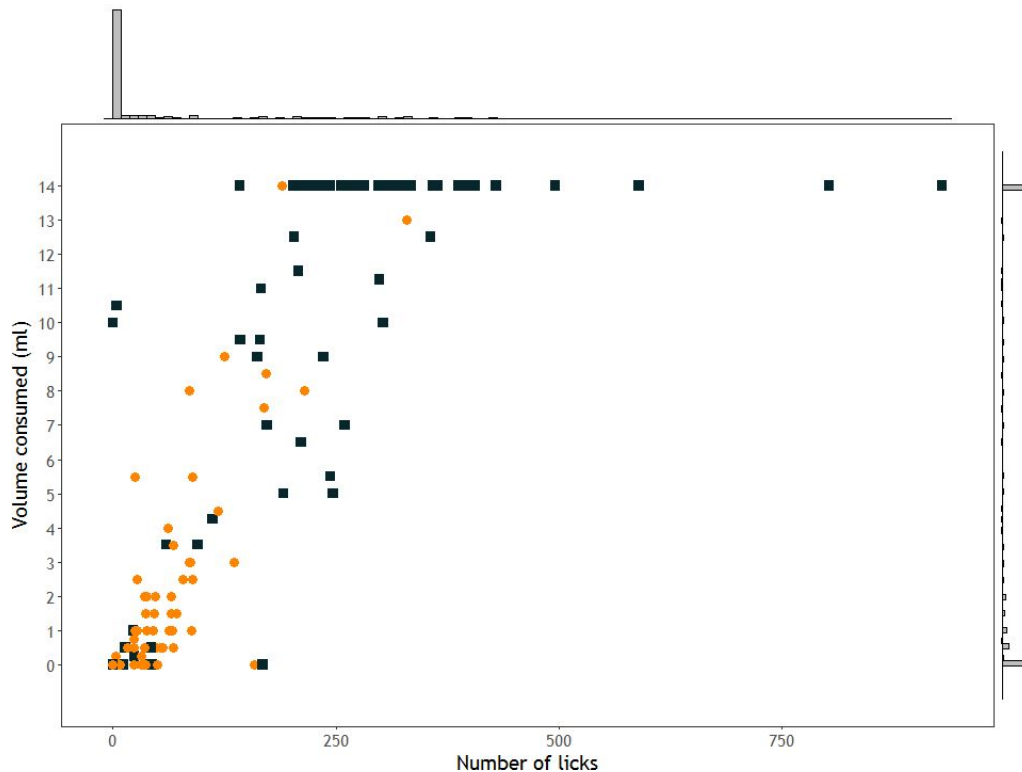
Overlays with regression lines worked with `geom_jitter` but not with `geom_dotplot`.



ANOVA-like comparisons of treatments can be performed using the delta method with the model estimates and their variance-covariance matrix.

R packages `msm` and `car` have delta method approximations that require entry of a function to estimate, but no derivatives are needed.

Two inflated points



What if we want to model the volume consumed instead of the number of licks?

There are many zeroes and many ones.

First, convert the volume consumed (0 ml to 14 ml) to a fraction consumed (0 to 1).

The R package zoib fits models using a beta distribution with inflation at 0, at 1, or both.

ZOIB - Zero-one inflated beta distribution

- Instead of a “count” side and a “participation” side, the ZOIB model can include up to 5 different sections for the mean and shape of the beta distribution, zero and one inflation, and random effects
- Uses Bayesian estimation with Gibbs sampling via JAGS

Convergence may be difficult.

Kumaraswamy? $f(x|a,b) = abx^{a-1}(1 - x^a)^{b-1}$

Aroma tests

Cats are given two minutes to explore two perforated cans that contain different aromatic agents.

Responses of interest: Lengths of time that the cat sniffs Aromas A and B.

For some length of time, the cat will not be sniffing either container. This has ranged from 44 seconds (37%) to 120 seconds (100%).

See references from John Cornell

In a 3-component mixture experiment:

- Three components A, B, and C are blended in specified proportions
- These amounts are linearly dependent:
$$A + B + C = 1$$
- Additionally, there may be other constraints, e.g.
$$A < .1 \text{ and}$$
$$A + B < .3$$
- Goal: Optimize some feature of interest (taste, viscosity)

Ternary plot

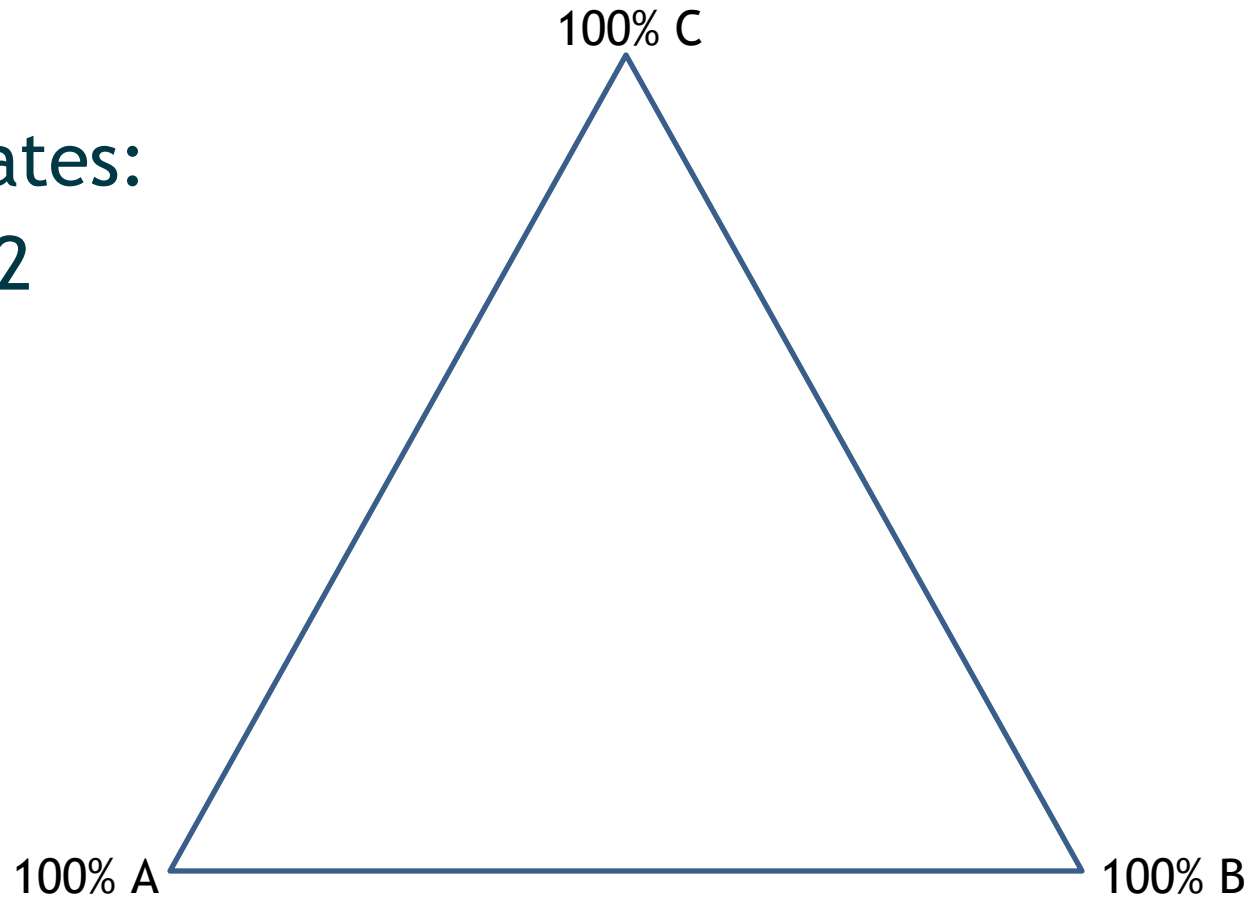
A 3-component mixture can be represented with an equilateral triangle in 2 dimensions, since

$$A + B + C = 1$$

New coordinates:

$$X = (2B + C)/2$$

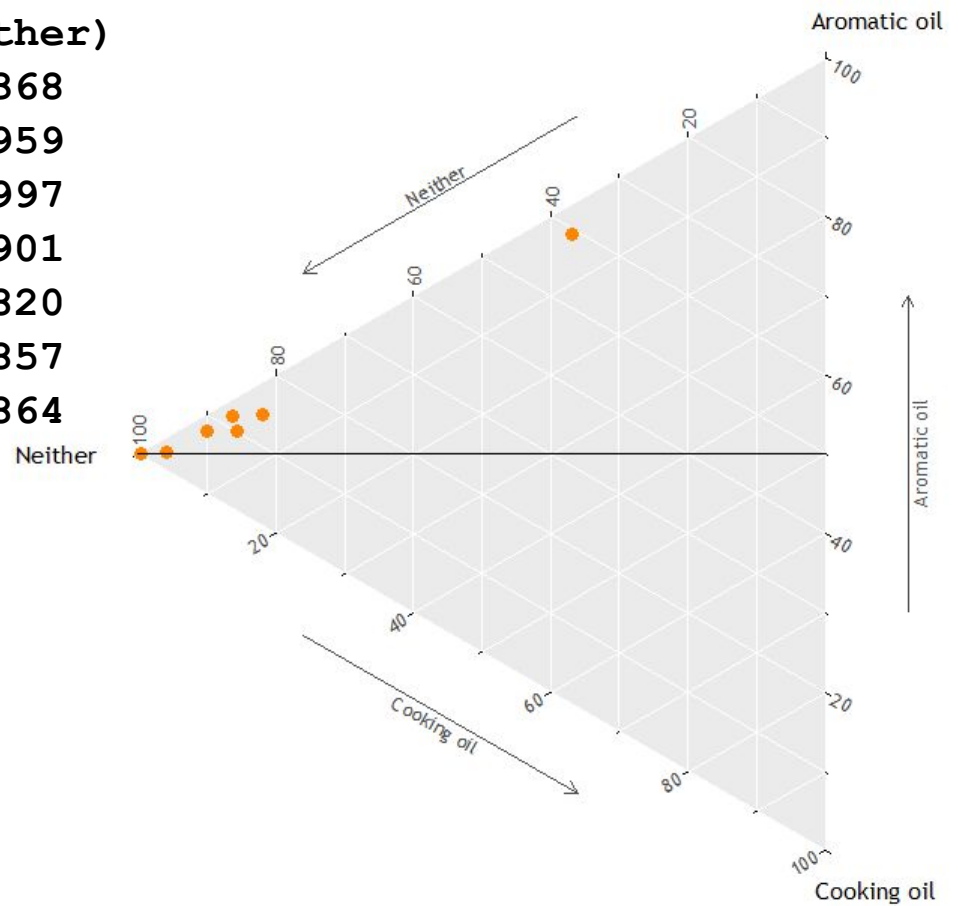
$$Y = (C\sqrt{3})/2$$



Ternary plot example

Data frame catdata:

Cat	Cooking (oil)	Aromatic (oil)	NS (Neither)
1	0.038	0.594	0.368
2	0.018	0.023	0.959
3	0.000	0.003	0.997
4	0.019	0.080	0.901
5	0.039	0.141	0.820
6	0.042	0.100	0.857
7	0.020	0.116	0.864

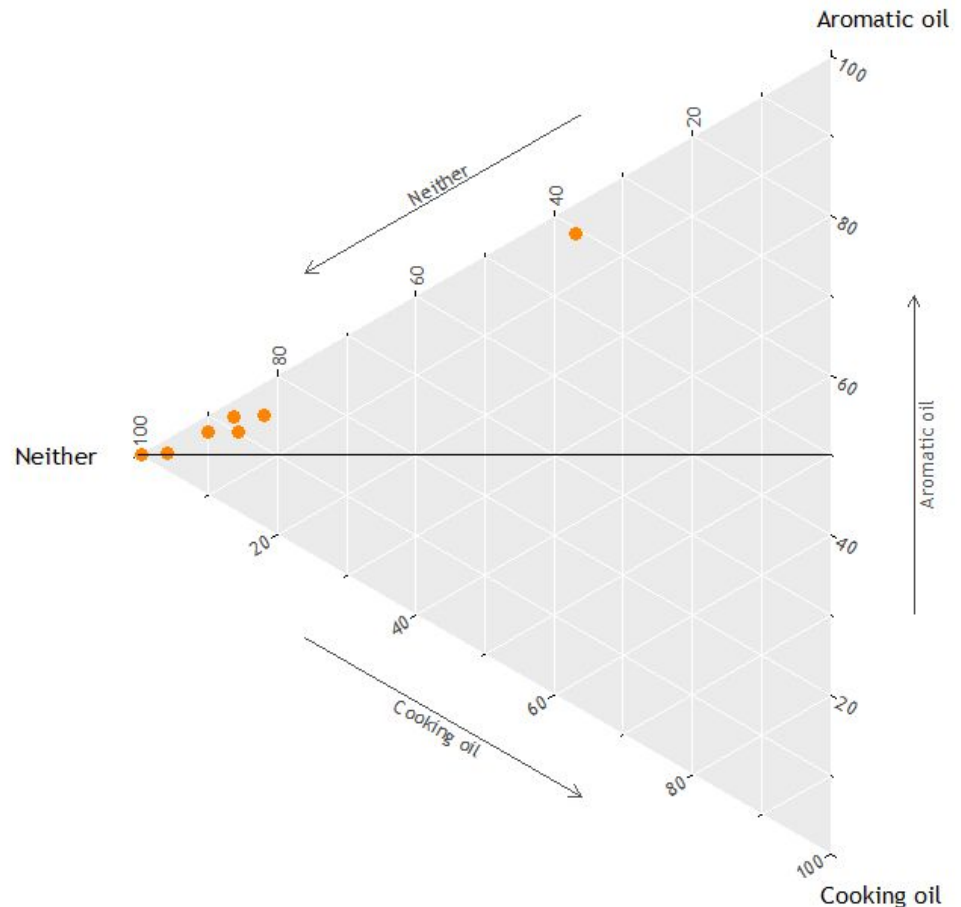
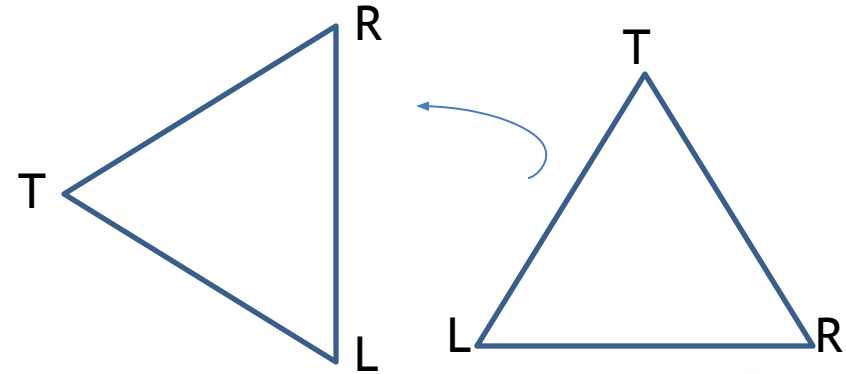


Ternary plot example

The R package ggtern makes ternary plots using ggplot2-like syntax.

```
triplot <- ggtern(data=catdata,  
aes(x=Cooking, y=NS, z=Aromatic) +  
geom_Tisoprop(value=0.5) +
```

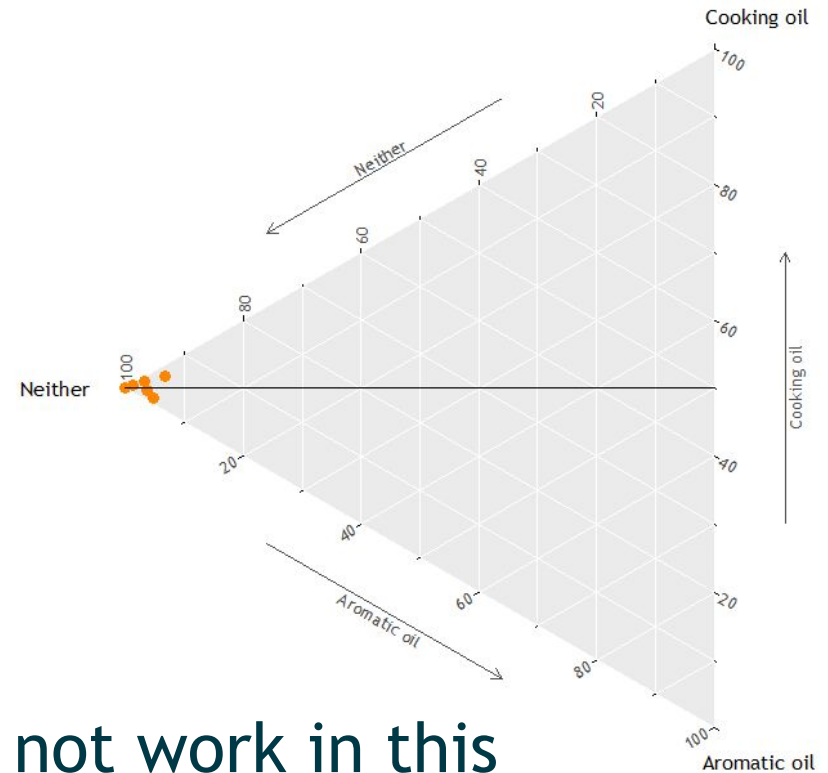
```
  Tlab("Neither",  
  labelarrow="Neither") +  
  Llab("Cooking oil",  
  labelarrow="Cooking oil") +  
  Rlab("Aromatic oil",  
  labelarrow="Aromatic oil") +  
  theme_gray() +  
  theme_showarrows() +  
  theme_nomask() +  
  theme_rotate(degrees=90)
```



Multiple zeroes in a ternary plot

Data from another trial:

Cat	Cooking oil	Aromatic oil	Neither
1	0.010	0.002	0.987
2	0.027	0.006	0.967
3	0.000	0.000	1.000
4	0.009	0.038	0.952
5	0.000	0.000	1.000
6	0.051	0.015	0.934
7	0.015	0.021	0.965



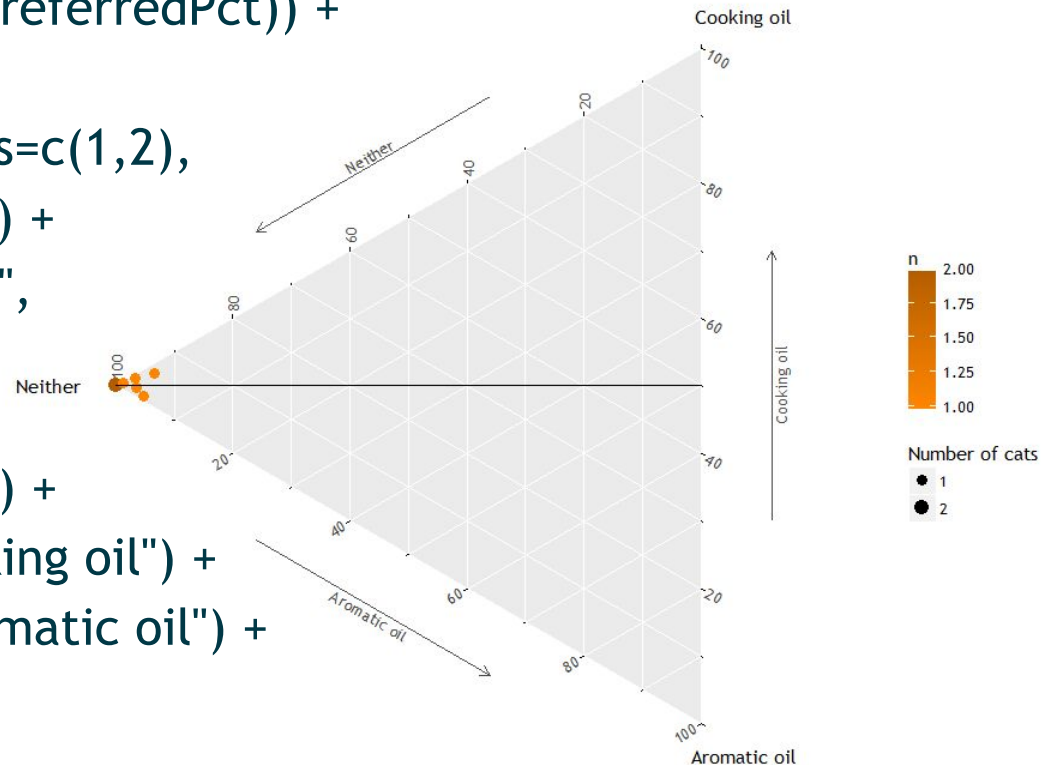
Jittering, violin plots, etc. do not work in this coordinate system

Zooming in the left vertex changes the “room map”

Contour and density plots try to interpolate outside the region where cats actually had data

Multiple zeroes in a ternary plot

```
triplot2 <- ggtern(data=catdata,  
  aes(x=NotPreferredPct,y=NSPct,z=PreferredPct)) +  
→ geom_count(aes(colour=..n..)) +  
→ scale_size_area(limits=c(1,2),breaks=c(1,2),  
  name="Number of cats",max_size=4) +  
→ scale_colour_gradient(low="#ff8500",  
  high="#b35d00") +  
geom_Tisoprop(value=0.5) +  
Tlab("Neither",labelarrow="Neither") +  
Rlab("Cooking oil",labelarrow="Cooking oil") +  
Llab("Aromatic oil",labelarrow="Aromatic oil") +  
theme_gray(base_size = 12) +  
theme_showarrows() +  
theme_nomask() +  
theme_rotate(degrees=90)
```



Thank you!

Jay Harrison
jharrison@afbinternational.com

