# Friday, Mar 24

This in-class demonstration will feature the estimation of Poisson and logistic regression models, and the interpretation of rate and odds ratios.

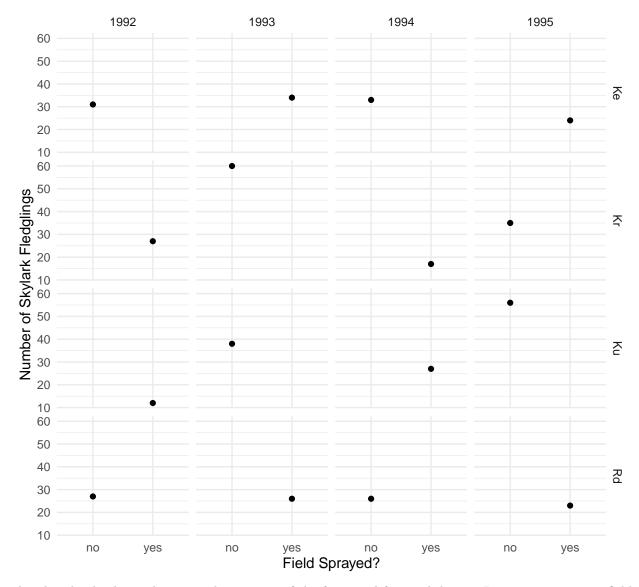
## Impact of Pesticides on Skylark Reproductivity

During the four summers from 1992 to 1995 researchers from the National Environmental Research Institute in the Ministry of Environment and Energy in Denmark conducted a study to examine how pesticide use impacts skylark reproduction in barley fields. The study used a fractional factorial design in which each year two of four fields were sprayed with pesticides while the other two fields were not. Which fields were sprayed was alternated so that a field was sprayed every other year. The number of fledgling skylarks produced in each field each year was recorded. The data are in the skylark data frame from the trtools package. The data are plotted below.

```
library(trtools)
library(ggplot2)
p <- ggplot(skylark, aes(x = spray, y = count)) +
    geom_point() + facet_grid(field ~ year) +
    labs(shape = "Field", x = "Field Sprayed?",
        y = "Number of Skylark Fledglings") + theme_minimal()
plot(p)</pre>
```

<sup>&</sup>lt;sup>1</sup>Odderskær, P., Prang, A., Eknegaard, N., & Andersen, P. N. (1997). Skylark reproduction in pesticide treated fields (Comparative studies of Alauda arvensis breeding performance in sprayed and unsprayed barley fields). Bekæmpelsesmiddelforskning fra Miljøstyrelsennr, 32, National Environmental Research Institute, Ministry of the Environment and Energy, Denmark: Danish Environmental Protection Agency.

<sup>&</sup>lt;sup>2</sup>A fractional factorial design is a design in which observations are made at only a subset of the possible combinations of levels of two or more factors. Such designs are quite economical but can preclude the estimation of interactions. This does not mean that such interactions are not present, but rather that if they are they are confounded with the main effects. For this particular design it is only possible to fully estimate a model with "main effects" for each of the three factors. Ideally factional factorial designs are used when interactions are negligible.



The plot clearly shows the incomplete nature of the fractional factorial design. In any given year, a field either was or was not sprayed. The objective is to investigate the effect of spraying on the number of skylarks while controlling for the effects of year and field.

1. Estimate a Poisson regression model for the number of skylark fledglings as your response variable that will reproduce the following results.

## cbind(summary(m)\$coefficients, confint(m))

	Estimate	Std. Error	z value	Pr(> z )	2.5 %	97.5 %
(Intercept)	3.430943	0.13262	25.86999	1.450e-147	3.16352	3.68367
sprayyes	-0.456126	0.09385	-4.86011	1.173e-06	-0.64141	-0.27324
fieldKr	0.049089	0.12672	0.38738	6.985e-01	-0.19929	0.29806
fieldKu	0.004964	0.12800	0.03879	9.691e-01	-0.24611	0.25625
fieldRd	-0.179048	0.13417	-1.33452	1.820e-01	-0.44342	0.08326
year1993	0.462623	0.13064	3.54108	3.985e-04	0.20868	0.72149
year1994	0.060018	0.14149	0.42420	6.714e-01	-0.21735	0.33816
year1995	0.327281	0.13411	2.44041	1.467e-02	0.06596	0.59240

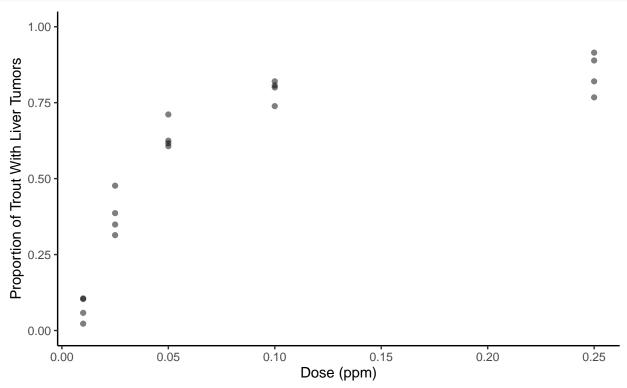
Note that here m is a model object created using the glm function.

- 2. What is the estimated rate ratio for the effect of spraying? How can this be interpreted?
- 3. What is the estimated expected number of fledgelings for each condition?

#### Aflatoxicol and Liver Tumors in Trout

The data in the data frame ex2116 in the Sleuth3 package are from an experiment that investigated the relationship between aflatoxicol and liver tumors in trout. The figure below shows the proportion of trout in each tank that developed liver tumors as well as the dose of aflatoxicol to which the trout were exposed. Aflatoxicol is a metabolite of Aflatoxin B1, a toxic by-product produced by a mold that infects some nuts and grains. Twenty tanks of rainbow trout embryos were exposed to one of five doses of aflatoxicol for one hour. The number of fish in each tank that developed liver tumors one year later was then observed. The plot below shows the data.

```
library(Sleuth3)
library(ggplot2)
p <- ggplot(ex2116, aes(x = Dose, y = Tumor/Total)) +
    geom_point(alpha = 0.5) + theme_classic() + ylim(0, 1) +
    labs(x = "Dose (ppm)", y = "Proportion of Trout With Liver Tumors")
plot(p)</pre>
```



The goal here is to estimate the effect of aflatoxicol on the risk of liver tumors in trout. Here we will consider three different logistic regression models.

1. Consider the logistic regression model

$$\log\left[\frac{E(Y_i)}{1 - E(Y_i)}\right] = \beta_0 + \beta_1 d_i,$$

where  $Y_i$  is the *i*-th observation of the proportion of trout from a tank that developed liver tumors and  $d_i$  is the corresponding dose of aflatoxicol to which those trout were exposed. This model can also be written as

$$E(Y_i) = \frac{e^{\eta_i}}{1 + e^{\eta_i}},$$

where  $\eta_i = \beta_0 + \beta_1 d_i$ . Note that by definition  $E(Y_i)$  is also the *probability* that a trout from a given tank will develop liver tumors, and  $E(Y_i)/[1 - E(Y_i)]$  is the *odds* that a trout from a given tank will develop liver tumors. Estimate this model using glm. You should be able to replicate the following results.

### cbind(summary(m)\$coefficients, confint(m))

```
Estimate Std. Error z value Pr(>|z|) 2.5 % 97.5 % (Intercept) -0.867 0.07673 -11.3 1.321e-29 -1.019 -0.7179 Dose 14.334 0.93695 15.3 7.838e-53 12.558 16.2346
```

Next estimate the odds ratio for the effect of increasing dose by 0.05 ppm using the contrast function.<sup>3</sup> How is this odds ratio interpreted?

2. Now consider a model where we use the base-2 logarithm of dose as the explanatory variable so that

$$\eta_i = \beta_0 + \beta_1 \log_2(d_i).$$

Recall that the function  $\log_2$  is known to R as  $\log 2$ . Estimate this model using glm. You should be able to replicate the following results.

#### cbind(summary(m)\$coefficients, confint(m))

```
Estimate Std. Error z value Pr(>|z|) 2.5 % 97.5 % (Intercept) 4.1634 0.2085 19.97 9.564e-89 3.7631 4.581 log2(Dose) 0.8997 0.0446 20.17 1.628e-90 0.8141 0.989
```

Here increasing the base-2 logarithm of dose by one unit is the same thing as *doubling* dose, and so the effect on the odds ratio of doubling dose will be the same regardless of what you double (e.g., 0.05 to 1 ppm, 0.1 to 0.2 ppm, etc.).<sup>4</sup> How is this odds ratio interpreted?

3. Rather than trying to decide between using dose or some transformation of dose in the model, we can instead define dose as a 5-level factor. There are two ways we could specify dose as a factor. One would be to create a new variable.

```
ex2116$Dosef <- factor(ex2116$Dose)
```

The levels of Dosef will be the original values of Dose but converted to strings, which we can see if we use the levels function.

#### levels(ex2116\$Dosef)

```
[1] "0.01" "0.025" "0.05" "0.1" "0.25"
```

Another approach is to replace Dose in the model formula with factor(Dose). Use the contrast function to estimate the odds ratio for the odds of tumor development at 0.025 ppm versus 0.01 ppm, 0.05 ppm versus 0.01 ppm, 0.1 ppm versus 0.01 ppm, and 0.25 ppm versus 0.01 ppm.<sup>5</sup> How are these odds ratios interpreted?

 $<sup>^3</sup>$ Here  $e^{\beta_1}$  would be the odds ratio for the effect of increasing dose by 1 ppm. However that is probably not a realistic effect as it would be a relatively large increase in dose. The study only considered up to 0.25 ppm. Using contrast is convenient here to estimate the odds ratio for the effect of an arbitrary change in dose.

<sup>&</sup>lt;sup>4</sup>We do not have to use the odds ratio for the effect of doubling dose just because we are using the base-2 logarithm of dose as our explanatory variable. We could also estimate the odds ratio for the effect of increasing dose from, say, 0.05 ppm to 0.1 ppm. But we would need to remember that because we are using the base-2 logarithm of dose as an explanatory variable that this would *not* be the same odds ratio as increasing dose the same amount from, say, 0.1 ppm to 0.15 ppm. Similarly for the previous model where we did not use the base-2 logarithm of dose, we could still estimate the odds ratio for the effect of doubling dose. But here we would need to remember that the odds ratio of doubling from, say, 0.05 ppm to 0.1 ppm would *not* be the same as the odds ratio for doubling from 0.1 ppm to 0.2 ppm.

<sup>&</sup>lt;sup>5</sup>Note that how you specify the levels of dose will depend on whether you created a new variable like Dosef or converted it to a factor within the model formula with factor(Dose). For the latter you will need to specify dose as a *number* but if you created it to a new variable you will need to specify it as a *string* by enclosing it in quotes.

4. Estimate the odds and probability of tumor development at each value of dose used in the study for

any of the three models.