

Negative Binomial Regression with R

Data Set

- ▶ The table below shows the average numbers of days absent by program type and seems to suggest that program type is a good candidate for predicting the number of days absent, our outcome variable, because the mean value of the outcome appears to vary by **prog**.

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```
with(dat, tapply(daysabs, prog, function(x) {  
  sprintf("M (SD) = %1.2f (%1.2f)", mean(x), sd(x))  
}))  
##                General  
## "M (SD) = 10.65 (8.20)"  
##                Academic  
## "M (SD) = 6.93 (7.45)"  
##                Vocational  
## "M (SD) = 2.67 (3.73)"
```

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- ▶ The variances within each level of **prog** are higher than the means within some of the levels.
- ▶ These are the conditional means and variances. These differences suggest that over-dispersion is present and that a Negative Binomial model would be appropriate.

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Negative binomial regression analysis

We will use the `glm.nb` function from the MASS package to estimate a negative binomial regression.

```
summary(m1 <- glm.nb(daysabs ~ math + prog,  
  data = negbinom))  
##  
## Call:  
## glm.nb(formula = daysabs ~ math + prog,  
          data = dat, init.theta = 1.032713156,  
##        link = log)  
##
```

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- ▶ R first displays the call and the deviance residuals.
- ▶ Next, we see the regression coefficients for each of the variables, along with standard errors, z-scores, and p-values.

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Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	2.61527	0.19746	13.24	< 2e-16 ***
math	-0.00599	0.00251	-2.39	0.017 *
progAcademic	-0.44076	0.18261	-2.41	0.016 *
progVocational	-1.27865	0.20072	-6.37	1.9e-10 ***

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

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- ▶ The variable **math** has a coefficient of -0.006, which is statistically significant.
- ▶ This means that for each one-unit increase in **math**, the expected log count of the number of days absent decreases by 0.006.
- ▶ The indicator variable shown as **progAcademic** is the expected difference in log count between group 2 and the reference group (prog=1).

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- ▶ The expected log count for level 2 of prog is 0.44 lower than the expected log count for level 1.
- ▶ The indicator variable for **progVocational** is the expected difference in log count between group 3 and the reference group.

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- ▶ The expected log count for level 3 of prog is 1.28 lower than the expected log count for level 1.
- ▶ To determine if prog itself, overall, is statistically significant, we can compare a model with and without **prog**.
- ▶ The reason it is important to fit separate models, is that unless we do, the overdispersion parameter is held constant.

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```
m2 <- update(m1, . ~ . - prog)
anova(m1, m2)
## Likelihood ratio tests of Negative Binomial Models
##
## Response: daysabs
##
```

	Model	theta	Resid. df	2 x log-lik.	Test
## 1	math	0.8559	312	-1776	
## 2	math + prog	1.0327	310	-1731	1 vs 2

```
## Pr(Chi)
## 1
## 2 1.652e-10
```

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- ▶ The two degree-of-freedom chi-square test indicates that prog is a statistically significant predictor of daysabs.
- ▶ The null deviance is calculated from an intercept-only model with 313 degrees of freedom.
- ▶ Then we see the residual deviance, the deviance from the full model.
- ▶ We are also shown the AIC and $2 \cdot \log$ likelihood.

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- ▶ The theta parameter shown is the dispersion parameter.

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```
## (Dispersion parameter for Negative Binomial(1.033) fami
##
##      Null deviance: 427.54  on 313  degrees of freedom
## Residual deviance: 358.52  on 310  degrees of freedom
## AIC: 1741
##
## Number of Fisher Scoring iterations: 1
##
##
##              Theta:   1.033
##              Std. Err.:  0.106
##
## 2 x log-likelihood:  -1731.258
```

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Checking model assumption

- ▶ As we mentioned earlier, negative binomial models assume the conditional means are not equal to the conditional variances.
- ▶ This inequality is captured by estimating a dispersion parameter (not shown in the output) that is held constant in a Poisson model.
- ▶ Thus, the Poisson model is actually nested in the negative binomial model.
- ▶ We can then use a likelihood ratio test to compare these two and test this model assumption.
- ▶ To do this, we can run our model as a Poisson,

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Confidence Intervals

We can get the confidence intervals for the coefficients by profiling the likelihood function.

```
(est <- cbind(Estimate = coef(m1), confint(m1)))  
## Waiting for profiling to be done...  
##           Estimate    2.5 %    97.5 %  
## (Intercept)    2.615265  2.2421  3.012936  
## math          -0.005993 -0.0109 -0.001067  
## progAcademic  -0.440760 -0.8101 -0.092643  
## progVocational -1.278651 -1.6835 -0.890078
```

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Incidence Rate Ratios

- ▶ We might be interested in looking at incident rate ratios rather than coefficients.
- ▶ To do this, we can exponentiate our model coefficients.
- ▶ The same applies to the confidence intervals.

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```
exp(est)
##              Estimate  2.5 %  97.5 %
## (Intercept)    13.6708  9.4127 20.3470
## math           0.9940  0.9892  0.9989
## progAcademic    0.6435  0.4448  0.9115
## progVocational  0.2784  0.1857  0.4106
```

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- ▶ The output above indicates that the incident rate for $\text{prog} = 2$ is 0.64 times the incident rate for the reference group ($\text{prog} = 1$).
- ▶ Likewise, the incident rate for $\text{prog} = 3$ is 0.28 times the incident rate for the reference group holding the other variables constant.
- ▶ The percent change in the incident rate of **daysabs** is a 1% decrease for every unit increase in **math**.

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- ▶ The form of the model equation for negative binomial regression is the same as that for Poisson regression.
- ▶ The log of the outcome is predicted with a linear combination of the predictors:

$$\ln(\widehat{daysabs}_i) = \text{Intercept} + b_1(\text{prog}_i = 2) + b_2(\text{prog}_i = 3) + b_3\text{math}_i$$

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$$\begin{aligned}\widehat{daysabs}_i &= e^{Intercept + b_1(prog_i=2) + b_2(prog_i=3) + b_3math_i} \\ &= e^{Intercept} e^{b_1(prog_i=2)} e^{b_2(prog_i=3)} e^{b_3math_i}\end{aligned}$$

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- ▶ The coefficients have an additive effect in the $\ln(y)$ scale and the IRR have a multiplicative effect in the y scale.
- ▶ The dispersion parameter in negative binomial regression does not effect the expected counts, but it does effect the estimated variance of the expected counts.

