

# Quasi-Poisson models

## Recap: Quasi-Poisson regression

A model for overdispersed Poisson-like counts, using an estimated dispersion parameter  $\hat{\phi}$ , is called a *quasi-Poisson* model.

```
m1 <- glm(nv ~ region, offset = log(enroll1000),
          data = crimes, family = quasipoisson)
summary(m1)
```

```
...
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -1.30445    0.34161  -3.818 0.000274 ***
## regionMW     0.09754    0.48893   0.199 0.842417
## regionNE     0.76268    0.42117   1.811 0.074167 .
## regionSE     0.87237    0.42175   2.068 0.042044 *
## regionSW     0.50708    0.50973   0.995 0.323027
...

```

## Recap: Poisson vs. quasi-Poisson

### Poisson:

```
...  
##           Estimate Std. Error z value Pr(>|z|)  
## (Intercept) -1.30445    0.12403  -10.517  < 2e-16 ***  
## regionMW    0.09754    0.17752   0.549  0.58270  
## regionNE    0.76268    0.15292   4.987  6.12e-07 ***  
## regionSE    0.87237    0.15313   5.697  1.22e-08 ***  
...
```

### Quasi-Poisson:

```
...  
##           Estimate Std. Error t value Pr(>|t|)  
## (Intercept) -1.30445    0.34161  -3.818  0.000274 ***  
## regionMW    0.09754    0.48893   0.199  0.842417  
## regionNE    0.76268    0.42117   1.811  0.074167 .  
...
```

# Quasi-likelihood models

# Pros and cons of quasi-Poisson

## Pros:

- + Estimated coefficients are the same as the Poisson model
- + Just need to get  $\mu$  and  $V(\mu)$  correct
- + Easy to use and interpret estimated dispersion  $\hat{\phi}$

**Cons:** Uses a quasi-likelihood, not a full likelihood. So we don't get

- + AIC or BIC (these require log-likelihood)
- + Quantile residuals (these require a defined CDF)

# Inference with quasi-Poisson models

```
m1 <- glm(nv ~ region, offset = log(enroll1000),
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summary(m1)
```

```
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## regionSW     0.50708    0.50973   0.995 0.323027
## regionW      0.20934    0.51242   0.409 0.684055
...
```

How can we test whether there is a difference between crime rates for Western and Central schools?

# $t$ -tests for single coefficients

# Inference with quasi-Poisson models

```
m1 <- glm(nv ~ region, offset = log(enroll1000),
          data = crimes, family = quasipoisson)
summary(m1)
```

```
...
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -1.30445    0.34161  -3.818 0.000274 ***
## regionMW     0.09754    0.48893   0.199 0.842417
## regionNE     0.76268    0.42117   1.811 0.074167 .
## regionSE     0.87237    0.42175   2.068 0.042044 *
## regionSW     0.50708    0.50973   0.995 0.323027
## regionW      0.20934    0.51242   0.409 0.684055
...

```

How can we test whether there is any relationship between Region and crime rates?



# $F$ -tests for multiple coefficients

## $F$ -test example

## *F*-test example

```
m1 <- glm(nv ~ region, offset = log(enroll1000),
          data = crimes, family = quasipoisson)
m0 <- glm(nv ~ 1, offset = log(enroll1000),
          data = crimes, family = quasipoisson)
```

```
deviance_change <- m0$deviance - m1$deviance
df_numerator <- m0$df.residual - m1$df.residual
numerator <- deviance_change/df_numerator
denominator <- m1$deviance/m1$df.residual

numerator/denominator
```

```
## [1] 2.003533
```

```
pf(numerator/denominator, df_numerator,
   m1$df.residual, lower.tail=F)
```

```
## [1] 0.0878041
```

# An alternative to quasi-Poisson

## Poisson:

- + Mean =  $\lambda_i$
- + Variance =  $\lambda_i$

## quasi-Poisson:

- + Mean =  $\lambda_i$
- + Variance =  $\phi\lambda_i$
- + Variance is a linear function of the mean

What if we want variance to depend on the mean in a different way?

# The negative binomial distribution

If  $Y_i \sim NB(\theta, p)$ , then  $Y_i$  takes values  $y = 0, 1, 2, 3, \dots$  with probabilities

$$P(Y_i = y) = \frac{\Gamma(y + \theta)}{\Gamma(y + 1)\Gamma(\theta)} (1 - p)^\theta p^y$$

- +  $\theta > 0, \quad p \in [0, 1]$
- +  $\mathbb{E}[Y_i] = \frac{p\theta}{1 - p} = \mu$
- +  $Var(Y_i) = \frac{p\theta}{(1 - p)^2} = \mu + \frac{\mu^2}{\theta}$
- + Variance is a *quadratic* function of the mean

# Mean and variance for a negative binomial variable

If  $Y_i \sim NB(\theta, p)$ , then

$$+ \mathbb{E}[Y_i] = \frac{p\theta}{1-p} = \mu$$

$$+ \text{Var}(Y_i) = \frac{p\theta}{(1-p)^2} = \mu + \frac{\mu^2}{\theta}$$

How is  $\theta$  related to overdispersion?

# Negative binomial regression

$$Y_i \sim NB(\theta, p_i)$$

$$\log(\mu_i) = \beta^T X_i$$

- +  $\mu_i = \frac{p_i \theta}{1 - p_i}$
- + Note that  $\theta$  is the same for all  $i$
- + Note that just like in Poisson regression, we model the average count
  - + Interpretation of  $\beta$ s is the same as in Poisson regression

## In R

```
library(MASS)
m3 <- glm.nb(nv ~ region + offset(log(enroll1000)),
             data = crimes)
```

...

```
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -1.33404      0.28137  -4.741 2.12e-06 ***
## regionMW      0.14230      0.44824   0.317  0.75089
## regionNE      0.94567      0.36641   2.581  0.00985 **
## regionSE      1.18534      0.39736   2.983  0.00285 **
## regionSW      0.33449      0.45666   0.732  0.46387
## regionW       0.06466      0.47628   0.136  0.89201
##
## (Dispersion parameter for Negative Binomial(1.0662) fami
```

...

$$\hat{\theta} = 1.066$$