#### Recap: overdispersion etc.

► saturated model:  $y_i \sim Bin(n_i, p_i)$ ,  $\tilde{p}_i = y_i/n_i$ ,

 $\ell(\tilde{p}) = \Sigma \{ y_i \log(y_i/n_i) + (n_i - y_i) \log(1 - y_i/n_i) \}$   $\overline{y_i} = \mu_i + \overline{\varepsilon}_i$ 

- ► what's the saturated model for linear regression? what is the maximized log-likelihood for this model?
- with binomial data, large-ish n<sub>i</sub>, residual deviance compares regression model to saturated model
- if it's too large, we have the wrong model
- lack of independence among individual Bernoullis; a few outliers; wrong predictors
   ELM p. 43,4
- ▶ estimate  $ilde{\phi} = X^2/(n-p)$  ELM p. 45
- inflate variance  $\hat{eta} \sim N(eta, \tilde{\phi}(X^{T}WX))$  instead of  $N(eta, X^{T}WX)$

 $\hat{\mu}_{i} = \hat{y}$ 

## ... overdispersion

```
summary (bmod
Call:
glm(formula = cbind(survive, total - survive) ~ location + period,
    family < binomial, data = troutegg)
period8
            -2 3256
                       0.2429 -9.573 < 2e-16 ***
period11 -2.4500
                       0.2341 -10.466 < 2e-16 ***
Signif. codes: 0 `***' 0.001 `**' 0.01 `*' 0.05 `.' 0.1 ` ' 1
(Dispersion parameter for binomial family taken to be
   Null deviance: 1021.469 on 19 degrees of freedom
Residual deviance: 64.495 on 12 degrees of freedom
ATC: 157.03
> summary(bmod2)
Call·
glm(formula = cbind(survive, total - survive) ~ location + period,
    family = guasibinomial, data = troutegg)
period8 -2.3256 0.5609 -4.146 0.001356 **
period11 -2.4500
                       0.5405 -4.533 0.000686 ***
Signif. codes: 0 `***' 0.001 `**' 0.01 `*' 0.05 `.' 0.1 `' 1
(Dispersion parameter for guasibinomial family taken to be 5.330358)
```

... overdispersion

 $SM ~\S10.6, \, p.512$ 

•  $Y \mid \epsilon \sim Bin(m, \epsilon p)$ 

• 
$$\mathsf{E}(\epsilon) = 1$$
,  $var(\epsilon) = \xi$ 

 $\blacktriangleright \mathsf{E}(Y) = \mathsf{E}\{\mathsf{E}(Y \mid \epsilon)\} = \mathsf{E}(mp\epsilon) = mp$ 

► 
$$\operatorname{var}(Y) = \operatorname{var}\{\operatorname{E}(Y \mid \epsilon)\} + \operatorname{E}\{\operatorname{var}(Y \mid \epsilon)\}$$
  
 $\operatorname{var}(\operatorname{hnp} \epsilon) + \operatorname{E}[\operatorname{mp} \epsilon(1 - p \epsilon)]$   
 $= \operatorname{m^2 p^2} \xi + \operatorname{mp} - \operatorname{mp^2} \epsilon^2$ 

• 
$$\operatorname{var}(Y) = m\{p(1-p) + \xi p^2(m-1)\}$$

• variance is larger than mp(1 - p)

see also ELM p.44

• can't be detected if m = 1

# ... overdispersion

•  $Y \mid \epsilon \sim Bin(m, \epsilon p)$ 

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$$\mathsf{E}(\epsilon) = 1$$
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- $\operatorname{var}(Y) = \operatorname{var}\{\mathsf{E}(Y \mid \epsilon)\} + \mathsf{E}\{\operatorname{var}(Y \mid \epsilon)\}$

$$= mp(1-p) \phi = ?$$

• 
$$\operatorname{var}(Y) = m\{p(1-p) + \xi p^2(m-1)\}$$

ntbc

- variance is larger than mp(1-p)
- can't be detected if m = 1

see also ELM p.44

*m* plays the role of  $n_i$ 

## ... matched case-control studies

- suppose we have 1 : M matching one case, M matched controls
- ► for person *i* in matched set *j*, we have  $j = l_j \dots n$  yij  $y_{ij}, x_{ij}, \quad i = 0, 1, \dots, M$ ► model:  $\log \frac{p_j(x_{ij})}{1 - p_j(x_{ij})} = \alpha_j + x_{ij}^T \beta \frac{n + p}{pars}$  Mn people
- different intercept for each matched set confounding variables
- same effect of covariates across patients and sets
- data: in matched set j, we have 1 case (person 0) and M controls (persons 1,..., M)

$$\begin{aligned} \Pr(y_{0j} = 1 \mid \Sigma_{i=1}^{M} y_{ij} = 1) &= \frac{\Pr(y_{0j} = 1, y_{1j} = 0, \dots, y_{Mj} = 0)}{\Pr(y_{1j} = 0, \dots, y_{Mj} = 0)} \\ &= \frac{\exp(x_{0j}^{T}\beta)}{\Sigma_{i=0}^{M} \exp(x_{ij}^{T}\beta)} \end{aligned}$$

## ... matched case-control studies

- suppose we have 1 : M matching one case, M matched controls
- for person i in matched set j, we have

$$y_{ij}, x_{ij}, \quad i=0,1,\ldots,M$$

model:

$$\log \frac{p_j(x_{ij})}{1 - p_i(x_{ij})} = \alpha_j + x_{ij}^{\mathrm{T}}\beta$$

- different intercept for each matched set confounding variables
- same effect of covariates across patients and sets
- data: in matched set j, we have 1 case (person 0) and M controls (persons 1,..., M)

$$\Pr(y_{0j} = 1 \mid \Sigma_{i=0}^{M} y_{ij} = 1) = \frac{\Pr(y_{0j} = 1, y_{1j} = 0, \dots, y_{Mj} = 0)}{\Pr(y_{1j} = 0, \dots, y_{Mj} = 0)}$$

$$\Pr(x_{0j} \in \mathbb{R} \setminus \mathbb{R}$$

$$\begin{array}{c|c} \text{matched case-control studies} \\ p_n \left( Y_o = 1 \middle| Y_o + Y_i = 1 \right) \\ = P \left( Y_o = 1, Y_i = 0 \right) \\ \hline \\ = \frac{P \left( Y_o = 1, Y_i = 0 \right)}{P \left( Y_o = 1, Y_i = 0 \right)} \\ = \frac{e^{2t + \beta x_o}}{P \left( Y_o = 1, Y_i = 0 \right) + P \left( Y_o = 0, Y_i = 1 \right)} \\ = \frac{e^{2t + \beta x_o}}{1 + e^{2t + \beta x_o}} \\ = \frac{e^{2t + \beta x_o}}{1 + e^{2t + \beta x_o}} \\ \hline \\ = \frac{e^{2t + \beta x_o} + e^{2t + \beta x_o}}{1 + e^{2t + \beta x_o}} \\ \hline \\ \hline \\ = \frac{e^{2t + \beta x_o} + e^{2t + \beta x_o}}{1 + e^{2t + \beta x_o}} \\ \hline \end{array}$$