y: Nopation pilo): sample proportions

Z(h) = n(h) + (y-fu)  $\frac{1}{w!} = \left(\frac{dM_{i}^{2}}{d\mu_{i}}\right) \cdot \left(\frac{\hat{\mu}(\hat{\mu})}{\hat{\mu}(\hat{\mu})}\right)$ wi = nimi(1-Mi) var(B)=(XWX) ver (B) = (xTWX) Compare Standard CM: var B = (XTWX/ 52)
(with weights) Standard errors of LM Slandard errors y GLM: beignes

Deviances D = deviance  $\frac{V}{\phi}$ : scaled deviance Recall:  $l_i(y_i|\Phi, M_i) = y_i l_i - b(0_i) + \dots$ Fitted model: defines  $\hat{\mu}_i, \hat{\partial}_i$ Saturated model: typically  $\mu_i = \hat{\mu}_i = y_i$   $\hat{\partial}_i = log \underbrace{\hat{\mu}_i}_{l-\mu_i}$ Person  $\hat{D} = log \underbrace{\hat{\mu}_i}_{l-\mu_i}$ Define RESELTUID, MI - CHALLIS  $D = 2 \sum_{\phi} \{l_i | y_i | \phi, \widetilde{\mu}_i \} - l_i | y_i | \phi, \widehat{\mu}_i \}$  $= \frac{2}{9} \sum_{i} w_{i} \left\{ y_{i} \widetilde{\theta}_{i} - b(\widetilde{\theta}_{i}) - y_{i} \widehat{\theta}_{i} + b(\widetilde{\theta}_{i}) \right\}$ Bigramial  $0 = \log 4$   $b(0) = \log (1 + e^{0}) = -\log(1 - \mu)$   $\Phi = 1$   $\left[ p + 4 \text{ of GM handowl} \right]$ D= 2 \( \text{Nif y: log \( \frac{\f 

Residuals Standard LM: Vi = Yi - Mi Response Residu "Pearson Residual" yi-Mi

Vi (Mi) Awn "Deviance Residuel" D = Zd? di = Tai · sign (yi- Mi) "Working Residual": The residual that comes from the LM fit inside the iteration