Response modeled with explanatory variables.

- Tarus length continuous
 - Hatch island factor, 5 levels
 - Sex factor, 2 levels
 - NAO covariate
- Oispersal, categorical, binary
 - Hatch island factor, 5 levels
 - Sex factor, 2 levels
 - NAO covariate
 - Wing length covariate
- Number of off-spring categorical, counts
 - Dispersal factor, 2 levels
 - Hatch island factor, 5 levels
 - Body mass covariate

- We recommend binary data with at least one continuous and one nominal/ordinal explanatory variable.
- But any data / model that fit the course is OK.
- Discuss with Ingelin and Alessandro!

Find a friend and start today!

Example: Chronically medical conditions

- Women in rural area see GP less then women in urban area.
- Why? Less sick or less accessible?

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- Why? Less sick or less accessible?

Data

- Group 1: No. of chronically conditions for 26 town women with \leq 3 GP visits.
- Group 2: No. of chronically conditions for 23 town women with \leq 3 GP visits.

Do women in the two groups with the same number of visits have the same need?

 $\begin{array}{l} y_{jk} \colon \text{Woman } j \text{ from group } k. \\ H_0 \text{ Same need: } Y_{jk} \sim Po(\theta) \\ H_1 \text{ Different needs: } Y_{jk} \sim Po(\theta_k) \end{array}$

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Standardized residuals

Normal model: $E(Y_i) = \mu_i$, $Y_i = N(\mu_i, \sigma^2)$ Poisson model: $E(Y_i) = \theta_i$, $Y_i \sim Po(\theta_i)$

•
$$r_i = \frac{y_i - \hat{\mu}_i}{\hat{\sigma}}$$

• $r_i = \frac{y_i - \hat{\theta}}{\sqrt{\hat{\theta}}}$

If model is correct: Approximately: $r_i \sim N(0, 1)$ Plots for r_i

- qq-plot
- against each explanatory variable
- other potential explanatory variables
- plot r_i vs ŷ_i (check assumption of constant variance / homoscedasity)
- plot r_i in order y_i was measured.

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