Lab 2.3 - Squid

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Testisweight_{im} = $\beta_0 + \beta_1 \cdot \text{DML}_i + \beta_{2m} + \beta_{3m} \cdot \text{DML}_i + \epsilon_{im}$, $\epsilon_{im} \sim \mathcal{N}(0, \sigma_{im}^2)$ for squid i (i = 1, ..., 768) at month m (m = 1, ..., 12) with $\beta_{2m} = \beta_{3m} = 0$ if m = 1

```
library(nlme)
lm.form <- Testisweight ~ DML * fMONTH</pre>
```

Model M.lm

 $\epsilon_{im} \sim \mathcal{N}(0, \sigma^2)$ which means

$$\sigma_{i,1}^2 = \sigma_{i,2}^2 = \dots = \sigma_{i,12}^2 \quad \forall i \in \{1, \dots, 768\}$$

M.lm <- gls(lm.form, data = Squid)

Model M.gls1 - λ_i s known

 $\epsilon_{im} \sim \mathcal{N}(0, \sigma_i^2)$ where $\sigma_i^2 = \sigma^2 \cdot \text{DML}_i$, allowing for a larger residual spread if DML increases, i.e.,

$$\sigma_i^2 = \begin{cases} \sigma_1^2 \\ \dots \\ \sigma_{768}^2 \end{cases} = \begin{cases} \sigma^2 \cdot \mathrm{DML}_1 & \text{if } i = 1; \\ \dots \\ \sigma^2 \cdot \mathrm{DML}_{768} & \text{if } i = 768. \end{cases}$$

```
vf1Fixed <- varFixed(~DML)
# the variance covariate needs to be continuous!
M.gls1 <- gls(lm.form, weights = vf1Fixed, data = Squid)</pre>
```

M.lm is not nested within *varFixed*.

Model M.gls2 - $<\delta>$ -group

Month-specific variance: $\epsilon_{im} \sim \mathcal{N}(0, \sigma_m^2)$

$$\sigma_m^2 = \begin{cases} \sigma_1^2 \\ \sigma_2^2 \\ \dots \\ \sigma_{12}^2 \end{cases} = \begin{cases} \sigma^2 \cdot \delta_1^2 & \text{if } m = 1; \\ \sigma^2 \cdot 1^2 & \text{if } m = 2; \\ \dots \\ \sigma^2 \cdot \delta_{12}^2 & \text{if } m = 12. \end{cases}$$

we get: $\delta_1 = \frac{\sigma_1}{\sigma_2}$; $\delta_{12} = \frac{\sigma_{12}}{\sigma_2}$

```
vf2 <- varIdent(form= ~ 1|fMONTH)
M.gls2 <- gls(lm.form, weights = vf2, data = Squid)</pre>
```

M.lm is nested within varIdent: we can make use of likelihood ratio test.

Model M.gls3 - $<\delta>$ -group

 $\epsilon_{im} \sim \mathcal{N}(0, \sigma_{im}^2)$ with $\underline{\delta} = \delta$ (scalar) since we do not include any stratification in the model and

$$\begin{array}{lcl} \sigma_{im}^2 & = & \sigma^2 \cdot \lambda_{im}^2 \\ & = & \sigma^2 \cdot \lambda^2 (\ \delta, \ \mathrm{DML}_{im}\) & \mathrm{with}\ \lambda \ \mathrm{modeled}\ \mathrm{as}\ \mathrm{varPower}(\cdot) \\ & = & \sigma^2 \cdot |\mathrm{DML}_{im}|^{2\delta} \end{array}$$

```
vf3 <- varPower(form = DML)
M.gls3 <- gls(lm.form, weights = vf3, data = Squid)</pre>
```

M.lm is nested within varPower: if $\delta=0$ we get the "classical model". varIdent is nested within the varPower structure.

Model M.gls4 - $<\delta>$ -group

 $\epsilon_{im} \sim \mathcal{N}(0, \sigma_{im}^2)$ with $\underline{\delta} = [\delta_1, ..., \delta_{12}]'$ since we include stratification by month and

$$\begin{split} \sigma_{im}^2 &= \sigma^2 \cdot \lambda_{im}^2 \\ &= \sigma^2 \cdot \lambda^2 (\ \underline{\delta}, \ \mathrm{DML}_{im}\) \quad \mathrm{with} \ \lambda \ \mathrm{modeled} \ \mathrm{as} \ \mathrm{varPower}(\cdot) \\ &= \sigma^2 \cdot |\mathrm{DML}_{im}|^{2\delta_m} \\ &= \begin{cases} \sigma^2 \cdot |\mathrm{DML}_{i,1}|^{2\delta_1} & \text{if} \ m = 1 \\ \dots \\ \sigma^2 \cdot |\mathrm{DML}_{i,12}|^{2\delta_{12}} & \text{if} \ m = 12 \end{cases} \end{split}$$

```
# we are modeling an increase in spread for larger DML values,
# but only in certain months!
vf4 <- varPower(form=~ DML | fMONTH)
M.gls4 <- gls(lm.form, weights = vf4, data = Squid)</pre>
```

If variance covariate has values equal to 0, the variance of the residuals is 0 as well; this causes problems in the numerical estimation process and varPower() should not be used. Notice that if $\delta = 0.5$ and variance covariate assumes positive values, we get the same variance structure as $varFixed(\sim DML)$. If $\delta = 0$ we get the "classical model" **M.lm**.

Model M.gls5 - $<\delta>$ -group

 $\epsilon_{im} \sim \mathcal{N}(0, \sigma_{im}^2)$ with $\underline{\delta} = \delta$ (scalar) since we do not include any stratification in the model and

$$\sigma_{im}^{2} = \sigma^{2} \cdot \lambda_{im}^{2}$$

$$= \sigma^{2} \cdot \lambda^{2} (\delta, DML_{im}) \text{ with } \lambda \text{ modeled as } varExp(\cdot)$$

$$= \sigma^{2} \cdot e^{2\delta \cdot |DML_{im}|}$$

```
vf5 <- varExp(form = DML)
M.gls5 <- gls(lm.form, weights = vf5, data = Squid)
# as before, we could allow for different delta per month
# by specifying varExp(form = DML | fMONTH) )</pre>
```

In this case, there are no restrictions on δs or DML; moreover, this structure allows also a decrease of spread for DML values if delta is negative. If $\delta = 0$ we get the "classical model" $\mathbf{M.lm}$.

Model M.gls6 - $<\delta>$ -group

 $\epsilon_{im} \sim \mathcal{N}(0, \sigma_{im}^2)$ with $\underline{\delta_1} = \delta_1$ (scalar) and $\underline{\delta_2} = \delta_2$ (scalar), since we do not include any stratification in the model and

$$\begin{array}{lll} \sigma_{im}^2 & = & \sigma^2 \cdot \lambda_{im}^2 \\ & = & \sigma^2 \cdot \lambda^2 (\ \delta, \ \mathrm{DML}_{im}\) & \mathrm{with}\ \lambda \ \mathrm{modeled}\ \mathrm{as}\ \mathrm{varConstPower}(\cdot) \\ & = & \sigma^2 \cdot (\delta_1 + |\mathrm{DML}_{im}|^{\delta_2})^2 \end{array}$$

```
vf6 <- varConstPower(form = DML)
M.gls6 <- gls(lm.form, weights = vf6, data = Squid)</pre>
```

This variance structure should work better than the varExp() if the variance covariate has values close to zero.

Model M.gls7 - $<\delta>$ -group

 $\epsilon_{im} \sim \mathcal{N}(0, \sigma_{im}^2)$ with $\underline{\delta_1} = [\delta_{1,1}, ..., \delta_{1,12}]'$ and $\underline{\delta_2} = [\delta_{2,1}, ..., \delta_{2,12}]'$, since we include stratification by month in the model and

$$\begin{array}{lll} \sigma_{im}^2 & = & \sigma^2 \cdot \lambda_{im}^2 \\ & = & \sigma^2 \cdot \lambda^2 (\ \delta, \ \mathrm{DML}_{im}\) \quad \text{with } \lambda \ \mathrm{modeled} \ \mathrm{as} \ \mathrm{varConstPower}(\cdot) \\ & = & \sigma^2 \cdot (\delta_{1m} + |\mathrm{DML}_{im}|^{\delta_{2m}})^2 \end{array}$$

```
vf7 <- varConstPower(form = DML | fMONTH)
M.gls7 <- gls(lm.form, weights = vf7, data = Squid)</pre>
```

Model M.gls8 - $<\delta>$ -group

We can allow for both an increase in residual spread for larger DML values as well as a different spread per month:

 $\epsilon_{im} \sim \mathcal{N}(0, \sigma_{im}^2)$ with $\underline{\delta} = \delta$ (scalar), since we do not include any stratification in the model and

$$\begin{array}{lcl} \sigma_{im}^2 & = & \sigma_m^2 \cdot \lambda_{im}^2 \\ & = & \sigma_m^2 \cdot \lambda^2(\ \delta,\ \mathrm{DML}_{im}\) & \mathrm{with}\ \lambda\ \mathrm{modeled}\ \mathrm{as}\ \mathrm{varComb}(\cdot) \\ & = & \sigma_m^2 \cdot e^{2\delta \cdot |\mathrm{DML}_{im}|} \end{array}$$

```
vf8 <- varComb(varIdent(form = 1 | fMONTH), varExp(form = DML) )
M.gls8 <- gls(lm.form, weights = vf8, data = Squid)</pre>
```