

Detecting dispersion problems in GLMMs

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Ecological data analysis

Increased complexity and flexibility in ecological data modeling:

- Generalized linear modes (GLMs)
- Mixture models (e.g. zero-inflated GLMs)
- Hierarchical/Multilevel models, GLMMs

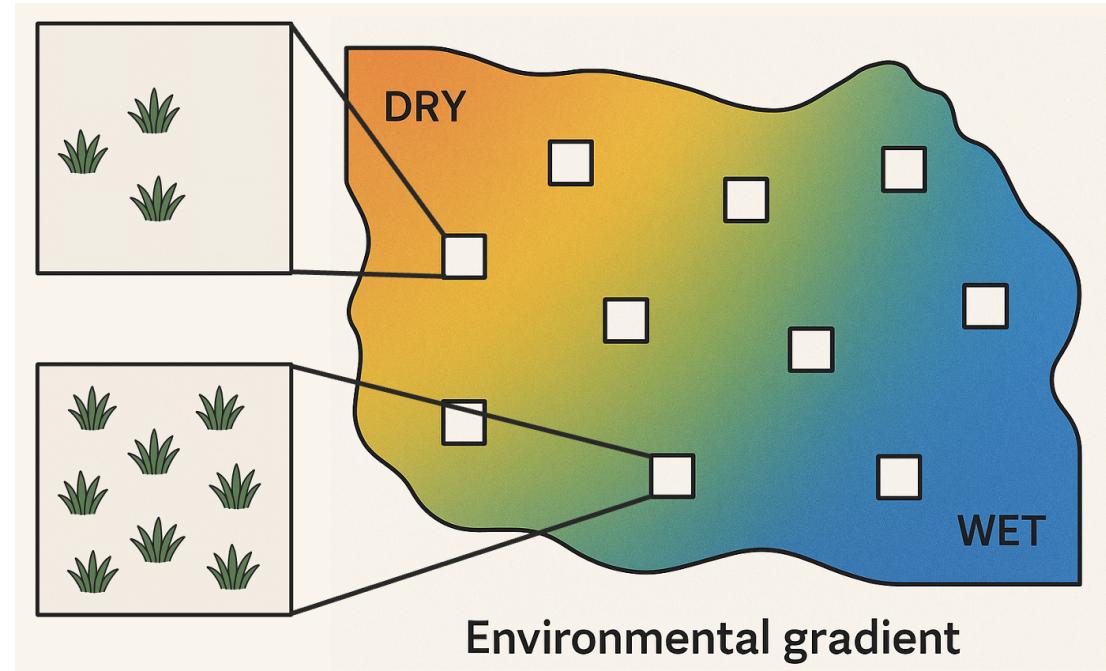


- But still few tools for model diagnostics
- Problem: **failing to check model assumptions**

Can you trust your model?

Dispersion problems in count data

- Example count data:
 - Species richness
 - Abundance of individuals
 - Number of success (K) within a number of trials (N)
- Modeling count data, GL(M)M distributions:
 - Poisson
 - Binomial (K/N) proportion



UNDER or OVERDISPERSION

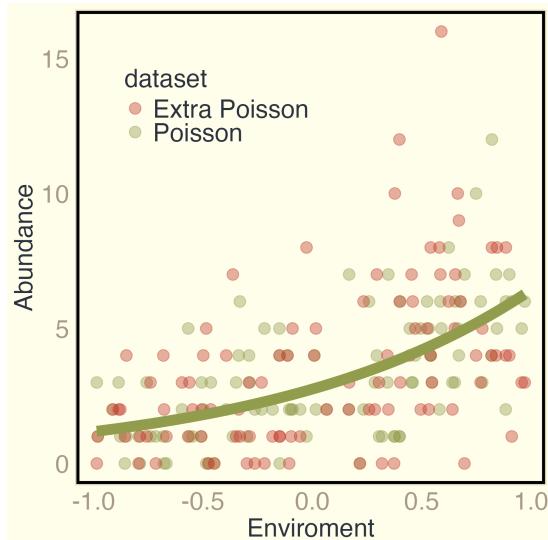
Problem when data has **more or less variability than expected by the distribution used for modeling**:

GOALS

- Aware ecologists of dispersion problems with count data
- Identify and describe the 3 main causes by using model diagnostics tools with DHARMA
- Show some modeling solutions for these causes

3 causes of dispersion problems

“Real” overdispersion:



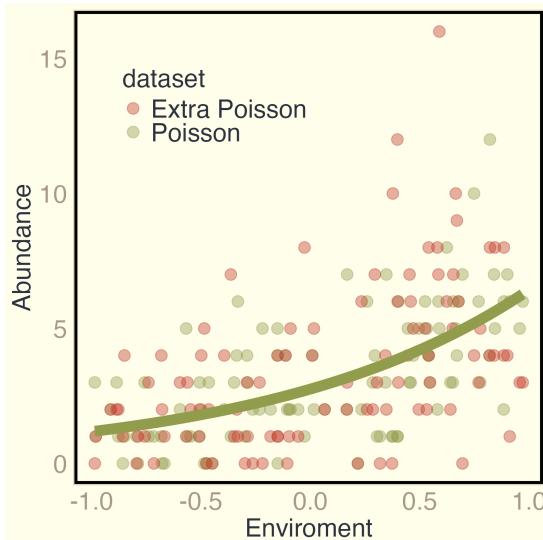
Heteroscedasticity:

Zero-inflation:

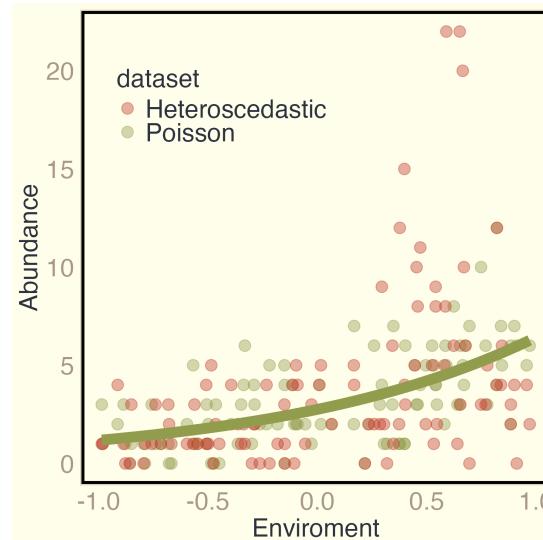
Abundances vary more than expected by the model, in general.

3 causes of dispersion problems

“Real” overdispersion:



Heteroscedasticity:



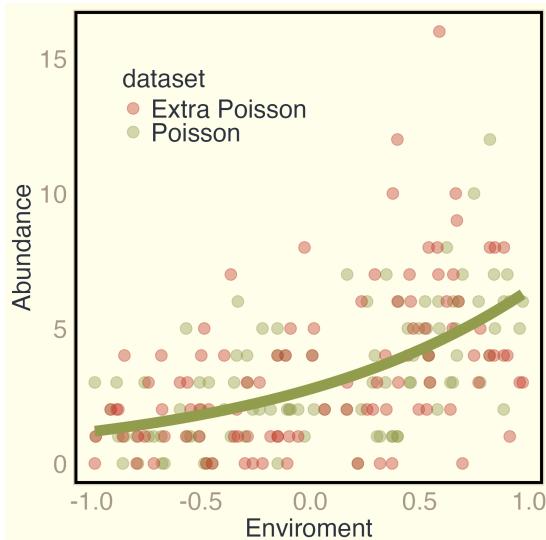
Zero-inflation:

Abundances vary more than expected by the model, in general.

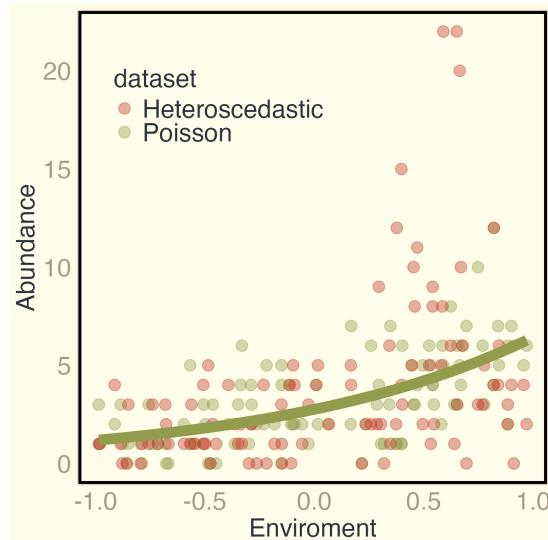
Abundances variation increases with the environmental gradient.

3 causes of dispersion problems

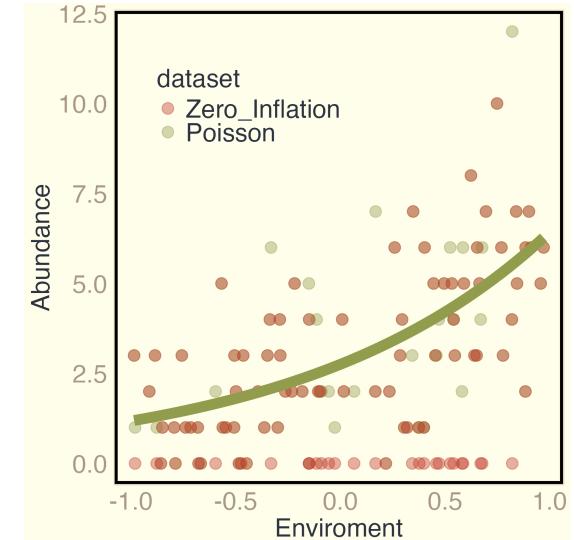
“Real” overdispersion:



Heteroscedasticity:



Zero-inflation:



Abundances vary more than expected by the model, in general.

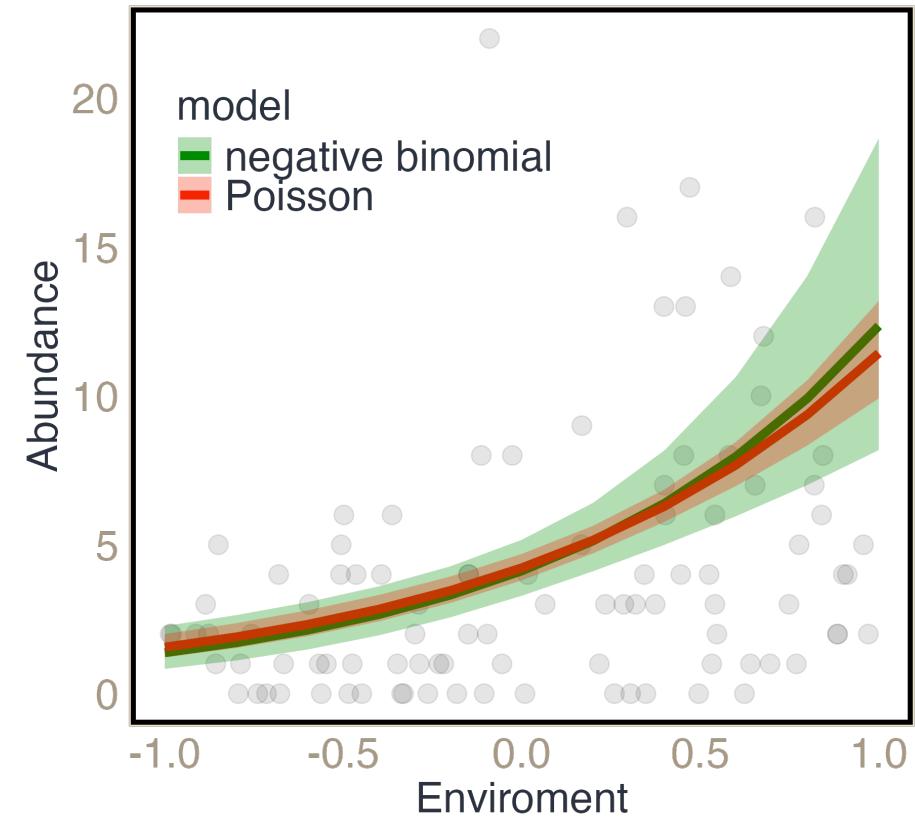
Abundances variation increases with the environmental gradient.

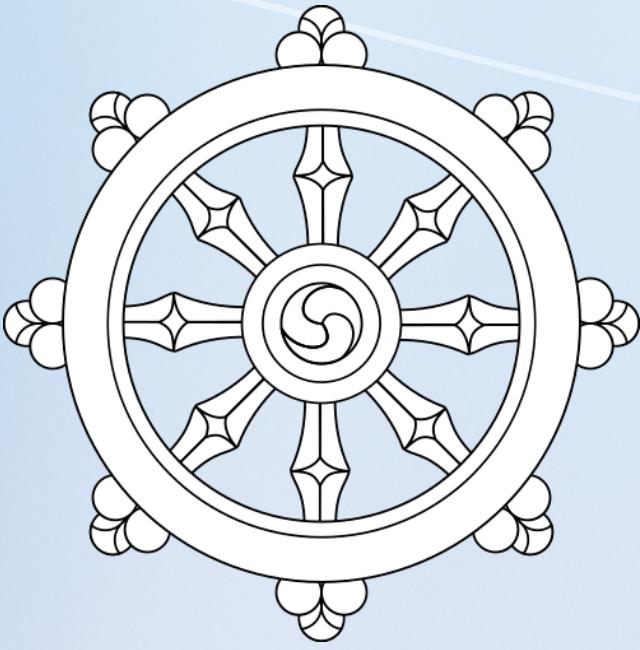
More zero abundances than expected by the model.

Consequences of dispersion problems

- Too small standard error of estimates -> narrower confidence intervals
- Larger chance of type I error: find an effect when it doesn't exist
- Wrong estimates by ignoring other processes (e.g. zero-inflation causes) in your data-generating process.
- Missing the opportunity to learn and get more info from your data. Ecological meanings for modeling/understanding unexpected variability?

Overdispersed data



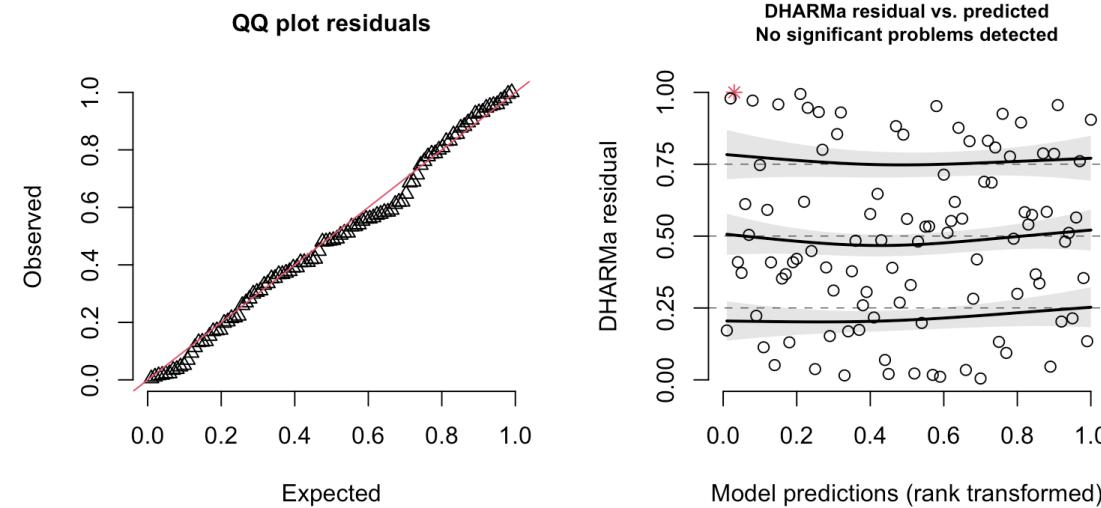


Detecting dispersion problems with DHARMa

Residual diagnostics with DHARMa

- Scaled quantile residuals -> Simulating from the model
- Residuals between 0 and 1 for ANY model complexity or distribution
- Interpreted the SAME way:

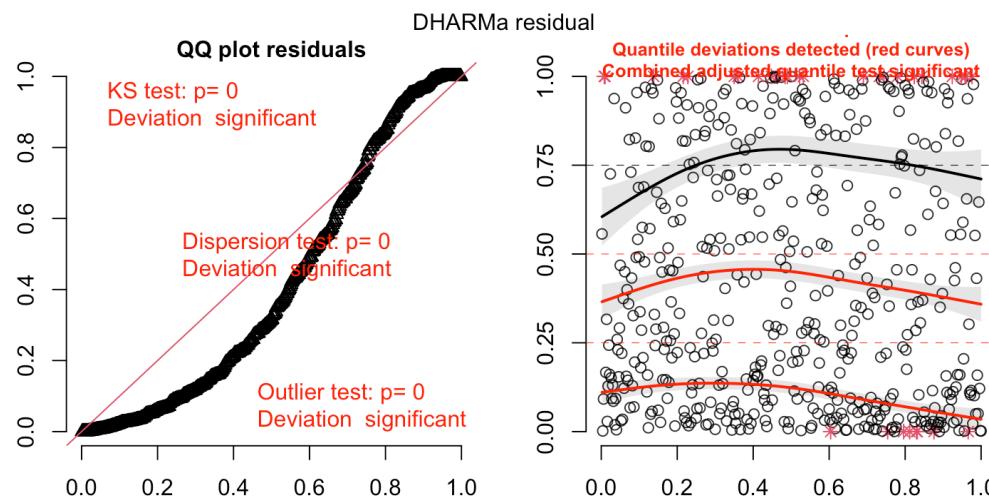
If your model is correctly specified, i.e. you have the “data-generating process”, scaled quantile residuals will present a uniform “flat” distribution between 0 and 1.



Detecting “real” overdispersion

Wrong model

```
1 m <- glmmTMB(observedResponse ~ Environment1 + (1|group),  
2     family = poisson(), data = overData)  
3 res <- simulateResiduals(m)  
4 plot(res)  
5 testDispersion(res)
```

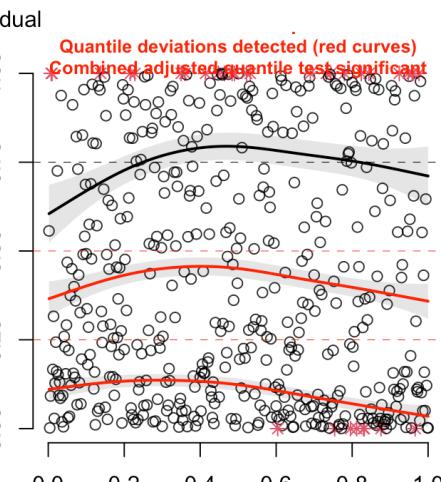
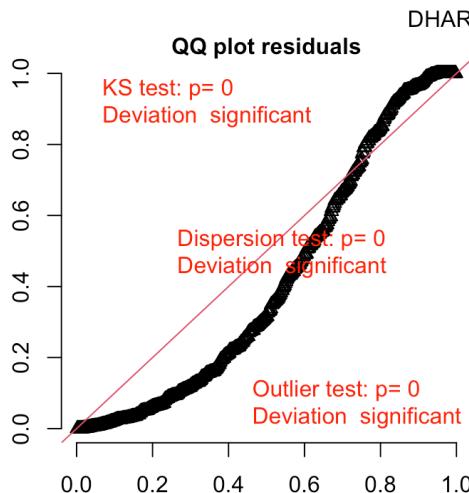


Dispersion = 5.19, p-value = 0.

Detecting “real” overdispersion

Wrong model

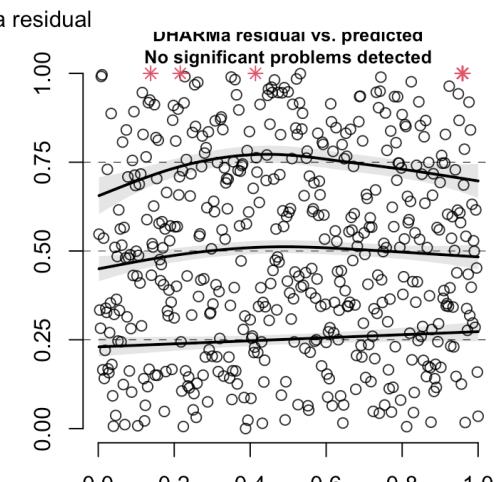
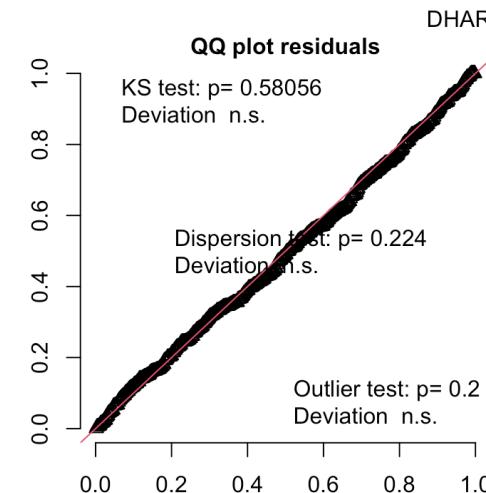
```
1 m <- glmmTMB(observedResponse ~ Environment1 + (1|group),  
2     family = poisson(), data = overData)  
3 res <- simulateResiduals(m)  
4 plot(res)  
5 testDispersion(res)
```



Dispersion = 5.19, p-value = 0.

Solution

```
1 m <- glmmTMB(observedResponse ~ Environment1 + (1|group),  
2     family = nbinom2(), data = overData)  
3 res <- simulateResiduals(m)  
4 plot(res)  
5 testDispersion(res)
```

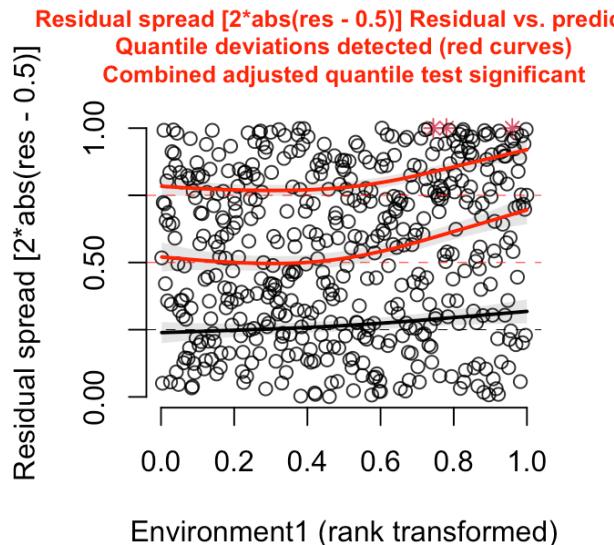


Dispersion = 1.19, p-value = 0.224.

Detecting heteroscedasticity

Wrong model

```
1 m <- glmmTMB(observedResponse ~ Environment1 + (1|group),  
2     family = poisson(), data = overData)  
3 res <- simulateResiduals(m)  
4 plotResiduals(res, form = data$Environment1,  
5                 absoluteDeviation = T)  
6 testDispersion(res)
```

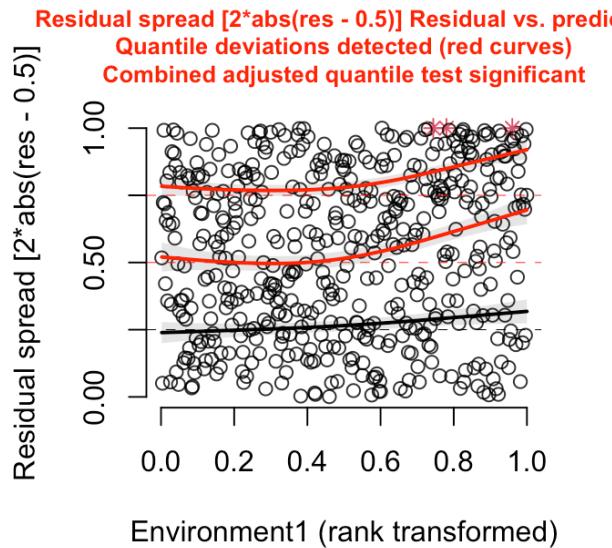


Dispersion = 1.9, p-value = 0.

Detecting heteroscedasticity

Wrong model

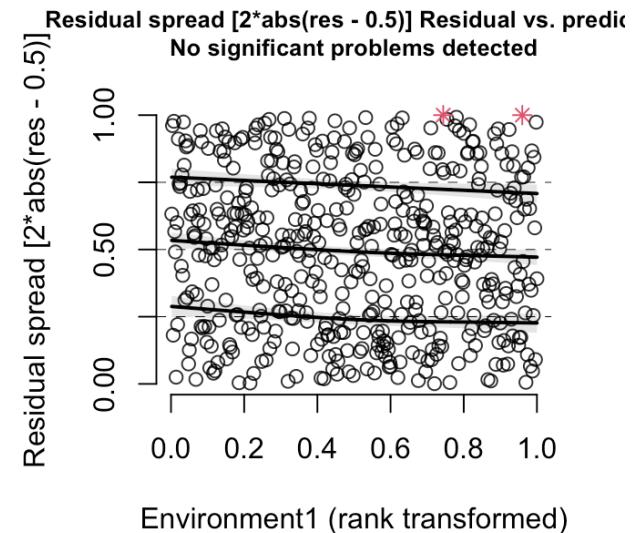
```
1 m <- glmmTMB(observedResponse ~ Environment1 + (1|group),  
2     family = poisson(), data = overData)  
3 res <- simulateResiduals(m)  
4 plotResiduals(res, form = data$Environment1,  
5                 absoluteDeviation = T)  
6 testDispersion(res)
```



Dispersion = 1.9, p-value = 0.

Solution

```
1 m <- glmmTMB(observedResponse ~ Environment1 + (1|group),  
2     dispformula = ~ Environment1, # dispersion formula  
3     family = nbinom2(), data = data) # but needs negative b  
4 res <- simulateResiduals(m)  
5 plotResiduals(res, form = data$Environment1,  
6                 absoluteDeviation = T)  
7 testDispersion(res)
```

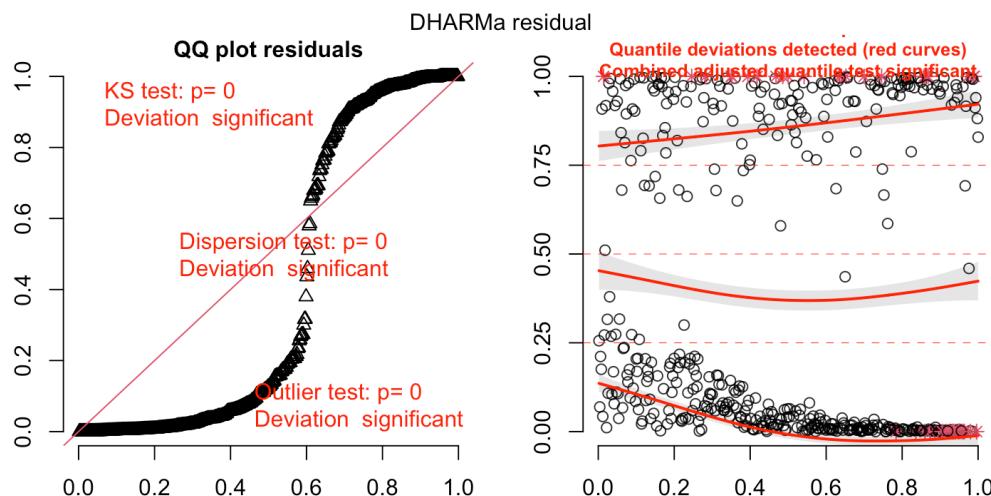


Dispersion = 1.11, p-value = 0.44.

Detecting zero-inflation

Wrong model

```
1 m <- glmmTMB(observedResponse ~ Environment1 + (1|group),  
2   family = poisson(), data = overData)  
3 res <- simulateResiduals(m)  
4 plot(res)  
5 testZeroInflation(res)
```

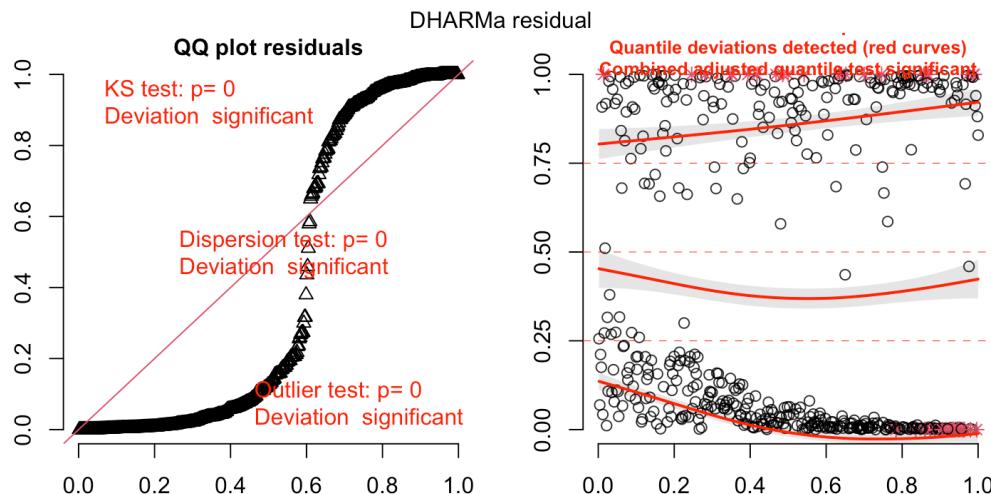


Zero-inflation = 5.16, p-value = 0.

Detecting zero-inflation

Wrong model

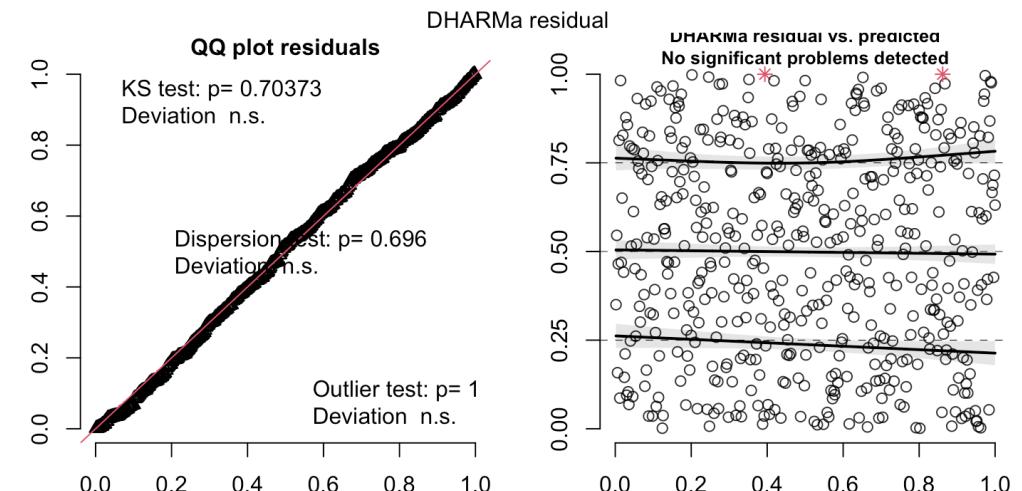
```
1 m <- glmmTMB(observedResponse ~ Environment1 + (1|group),  
2     family = poisson(), data = overData)  
3 res <- simulateResiduals(m)  
4 plot(res)  
5 testZeroInflation(res)
```



Zero-inflation = 5.16, p-value = 0.

Solution

```
1 m <- glmmTMB(observedResponse ~ Environment1 + (1|group),  
2     ziformula = ~ 1, # zero-inflation formula  
3     family = poisson(), data = data)  
4 res <- simulateResiduals(m)  
5 plotResiduals(res)  
6 testZeroInflation(res)
```



Zero-inflation = 1, p-value = 1.

Detecting dispersion problems

- Residual patterns alone will not tell you which is the cause of overdispersion. E.g.:
 - ‘Real’ overdispersion will show significant test for zero-inflation, and vice-versa.
 - ‘Real’ overdispersion and zero-inflation may have significant heteroscedasticity.
- Additional check: fit models addressing the potential problems and compare their fit (e.g. AIC, LRT) and residuals diagnostics.

Don’t always assume the most complex/complicated model is the correct one!

Conclusion

- There are many causes of dispersion problems in GLMMs
- Use [DHARMA](#) residuals tools to detect them
- Address the problem with adequate models, e.g, [glmmTMB](#)

Take-home message

- Models should ALWAYS be checked: residual diagnostics!
- Avoid an oversimplistic view of dispersion problems
- Detecting and addressing the causes of dispersion problems may also be informative for your system/data.

Comming soon:

Leite et al. *in prep.* **Dispersion tests in GLMMs: a methods comparison and practical guide.**



Thank you!

Vielen Dank!

Acknowledgements to Florian Hartig, Max Pichler, and the Theoretical Ecology Lab group