

# Statistics and experimental design in silage research: Some comments on design and analysis of comparative silage experiments

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# **1. Introduction**





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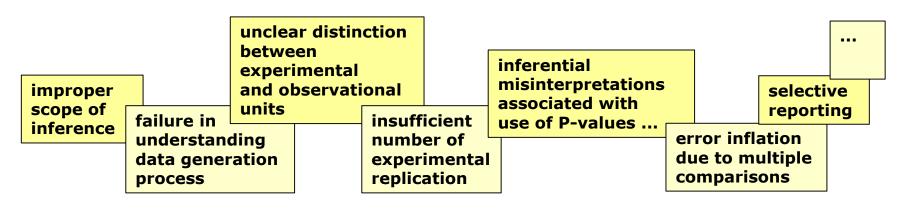
#### *Invited review:* Reproducible research from noisy data: Revisiting key statistical principles for the animal sciences

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"Reproducible results define the very core of scientific integrity in modern research"

.. but across all scientific disciplines only too a little number of trials generate reproducible results.



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# Aim: Critical findings on design, analysis, and interpretation of results will be addressed based on comparative silage experiments.

#### Scientific papers on silage trials during the last 8 years published in

- Journal of Dairy Science
- Grass and Forage Science
- Agricultural and Food Science
- Experiments with **1 up to 3 factors** (sometimes even more),
- 3 to 6 replicates per treatment,
- Statistical analyses:

frequently performed by **parametric analysis of variance**, followed by pairwise comparisons (**LSD, Tukey**, Bonferroni, Sidak, Duncan, ...),

sometimes by non-parametric procedures (Wilcoxon and others).



Aim: Critical findings on design, analysis, and interpretation of results will be addressed based on comparative silage experiments.

### Lab-scale ensiling trial on biostatistical questions in 2017

(1) What can the scope of inference for one ensiling experiment with mini-silos be?

- material taken from different field locations versus a composite sample,
- impact of location and fermentation process on silage traits.

(2) Is the frequently used (low) number of replications sufficient regarding significance and relevance of results?

Do the traits meet the assumptions of normal distribution and variance homogeneity?

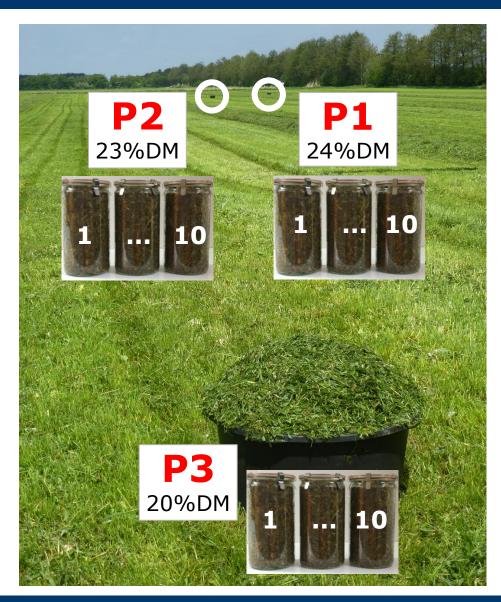
- three treatments: untreated control, biological, and chemical silage additive,
- ten mini-silos as replicates (composite sample = restricting experimental input),
- samples of smaller size (n=3, n=6) extracted:
  to contrast the results of statistical analyses for different sample sizes.

# 2. Description of the grassland trial



# Lab-scale ensiling trial

- (1) with respect to field sampling locations
  - material taken from three randomly selected sampling points (P1, P2, P3) of natural grassland
  - per sampling point: ten 1.5-L jars (=mini-silos) filled with grass material
  - without silage additives (= CON)





# Lab-scale ensiling trial

### (2) with grass mixture

- composite grassland sample, mixing identical quantities from three sampling points
- three treatments (silage additive)
- ten 1.5-L jars per treatment



without silage additive



P1+P2+P3 = MIX



chemical additive



#### biological additive

Lactobacillus buchneri CNCM-I 4323 and Pediococcus acidilactici DSM 11673

sodium nitrite, hexamethylene tetramine and potassium sorbate

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# 2. Description of the grassland trial



# Lab-scale ensiling trial

#### Traits $\rightarrow$ fresh forage



per sampling point (P1, P2, P3) n=5

DM	dry matter
WSC	water-soluble carbohydates
NO <sub>3</sub> -	nitrate
BC	buffering capacity
Yeasts	yeast count
Lactobac	lactic acid bacteria

known to have an influence on the fermentation process Traits  $\rightarrow$  silage (after 121 days of storage at 22°C)

- per sampling point (P1, P2, P3)
- mixture (MIX): per treatment (CON, CHEM, LAB)
- n=10 -> total sample size N=60

рН	
LA	lactic acid
AA	acetic acid
1,2-PD	1,2-propanediol
WSC	water-soluble carbohydates
ЕТОН	ethanol
PROP	n-propanol
ASTA	aerobic stability
DML	anaerobic DM losses

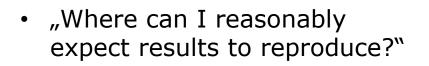
butyric acid, counts of yeasts and moulds: small values



# Scope of inference



- Population, to which the results from a research study are applicable.
- Ideally, this population is sampled at random.



= Degree of generalization.





# Scope of inference

#### (1) One field sampling point

Forage material represents exactly this field location.

 $\overline{\mathbf{D}}$ 

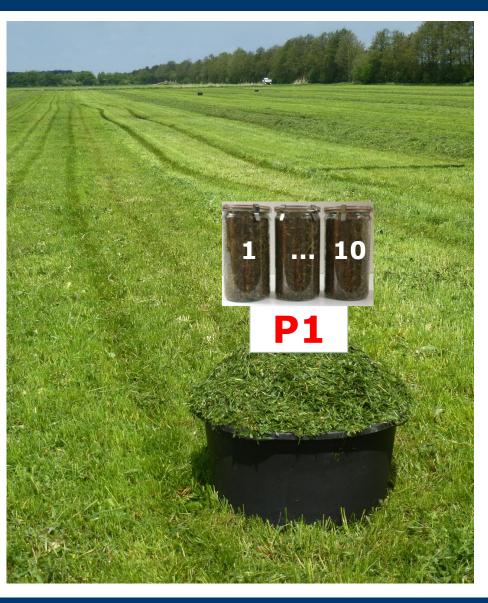
Variability of measurements reflects only the different fermentation processes in the replicated mini-silos per treatment for one material.

#### different location









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Ρ1

10

P1+P2+P3 = MIX

# Scope of inference

#### (2) Mixture of field sampling points

Composite sample of forage material represents an average field situation.

 $\overline{\mathbf{D}}$ 

Variability of measurements reflects

only the different fermentation processes

in the replicated mini-silos per treatment for one composed material.

#### different location



### farm silo



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РЗ

Ρ2

П





# Scope of inference

#### (3) Several field sampling points, n=1

Forage material represents the field.

Variability of measurements reflects arbitrary field locations and the different fermentation processes in the replicated mini-silos per treatment.

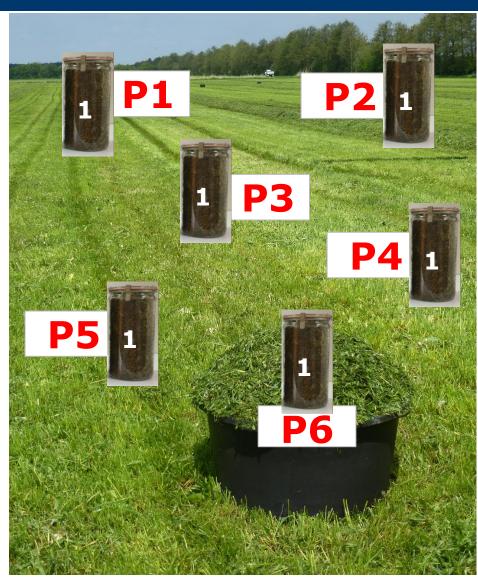
Both effects are confounded (no separation possible).

#### arbitrary location



#### farm silo





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# Scope of inference

#### (4) Several field sampling points, n>1

Forage material represents the field.

Variability of measurements reflects arbitrary field locations and the different fermentation processes in the replicated mini-silos per treatment.

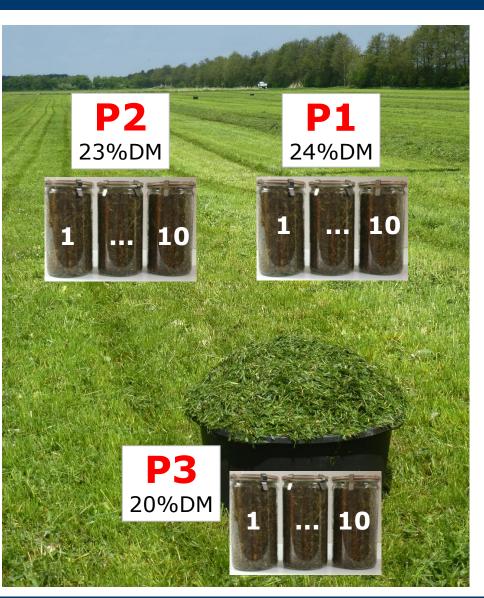
Both effects are not confounded (separation is possible).

#### arbitrary location



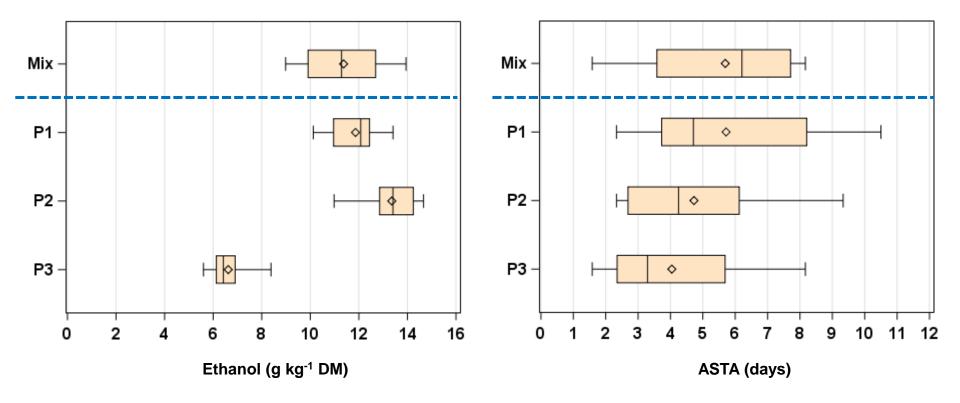
#### farm silo





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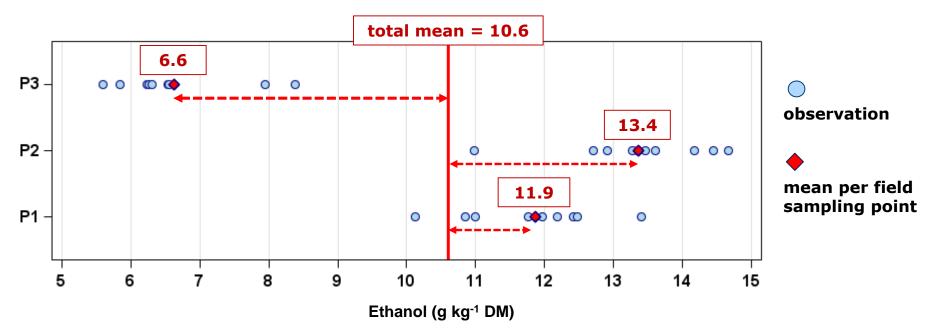
#### Box-Whisker-Plot for three sampling points and grass mixture (MIX), n=10



- except ASTA: observations more or less different between sampling points (e.g. ethanol)
- all traits: values from grassland mixture reflect average situation



### Variation between field sampling points and within sampling points for Ethanol



#### **Decomposition of total variability** (random effects model)

Source of variation	Variance component	percentage %
Field sampling point	12.42	93
Residual (Fermentation process)	0.93	7
Total	13.35	100





#### Decomposition of total variability of observed values Variance Component (%) 100 All traits (except ASTA): 90 largest fraction of variation was 80 caused by field sampling point, 70 ٠ 60 50 process of the ten replicates 40 per sampling point. 30 ASTA: 20 10 by the fermentation process. 0 1,2-PD WSC ETOH PROP ASTA pН LA AA DML **Field** sampling point Residual (=Fermentation process)

- remaining residual variation was related to effects of fermentation
- was affected almost completely

- Note: How far the efficacy of silage additives will be affected by sampling points cannot be shown in our study.
- Final evaluation of silage additive effects should request more than one trial (EFSA). But:

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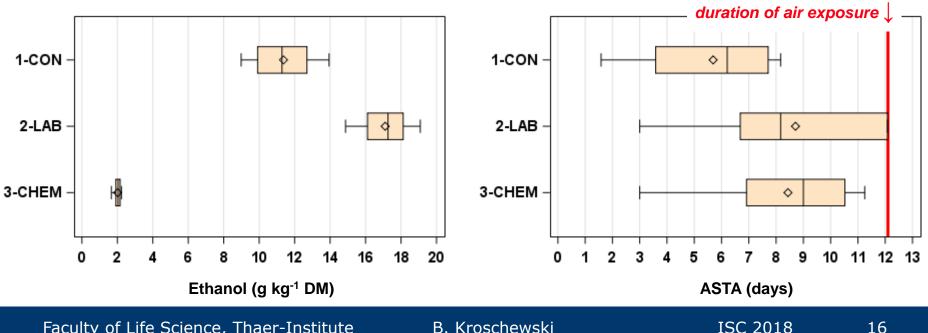


### Comparison of treatments

- fixed effects model, •
- Anova + Tukey's test procedure, ٠
- assuming normally distributed data, ٠ variance homogeneity,
- scope of inference: ٠ one composed material, mini-silos.

Ethanol	ASTA			
11.4 <mark>b</mark>	5.7 <mark>a</mark>			
17.1 <mark>c</mark>	8.7 <mark>a</mark>			
2.0 <mark>a</mark>	8.4 <mark>a</mark>			
1.4	3.1			
12.2	37.1			
	Ethanol 11.4 b 17.1 c 2.0 a 1.4			

Table of I SMeans n-10

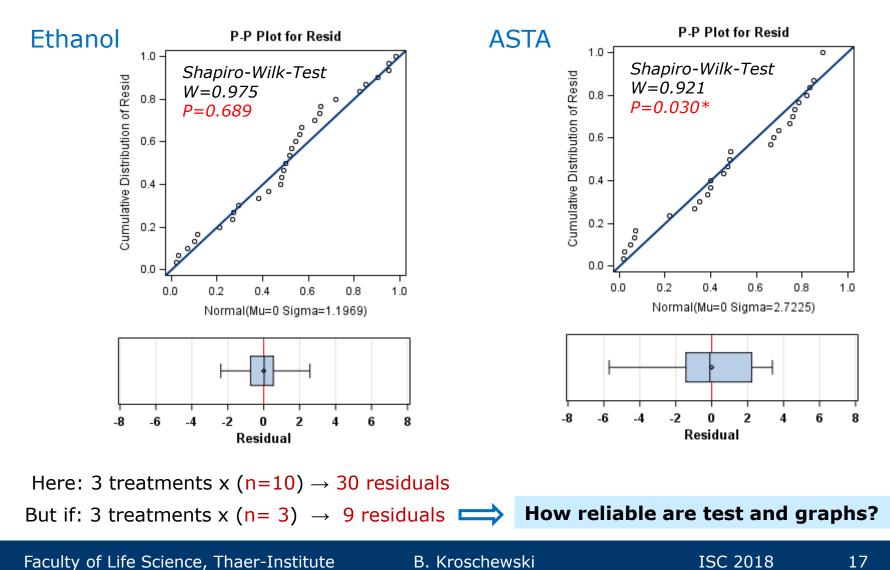


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### Observations come from populations with normally distributed data?





### Observations come from populations with homogeneous variances?

Trait		Residual	variance	AIC (fit criteria) "smaller is better"		
	CON	LAB	CHEM	Average	Var.hom.	Var.het.
Acetic acid	4.0	31.1	5.1	13.4	155.6	147.6
Ethanol	2.61	1.98	0.03	1.54	97.2	73.1
ASTA	4.9	11.3	7.7	8.0	141.5	144.0
DM losses	0.002	0.090	0.007	0.033	-6.3	-30.3
	(				1	Λ

Anova approach, assuming

variance homogeneity,

average residual variance used

Here: treatment variances estimated from n=10(as basis for inferences).

With n=3/6 also reliable estimations?

ANOVA approach, assuming

variance heterogeneity,

individual residual variances used



### Extraction of subsets from the whole sample (n=10) - e.g. Ethanol

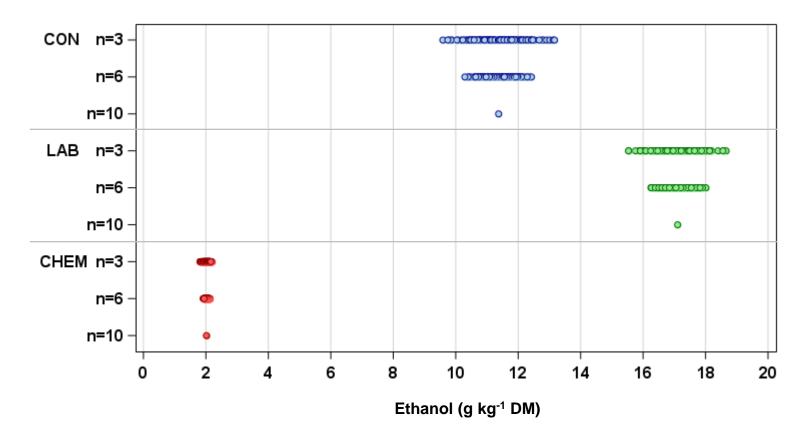
	Treatment	Replication									
n=3		1	2	3	4	5	6	7	8	9	10
	CON	9.9	10.7	10.4	13.9	9.0	11.8	12.5	9.8	12.6	12.9
	LAB	16.4	17.6	19.1	15.4	14.9	17.8	17.0	16.4	19.1	17.5
	CHEM	2.0	2.0	1.7	1.8	2.2	2.0	2.1	2.1	2.1	2.2
		subset 1						-	sı	ıbset	120

Treatment		Replication								
	1	2	3	4	5	6	7	8	9	10
CON	9.9	10.7	10.4	13.9	9.0	11.8	12.5	9.8	12.6	12.9
LAB	16.4	17.6	19.1	15.4	14.9	17.8	17.0	16.4	19.1	17.5
CHEM	2.0	2.0	1.7	1.8	2.2	2.0	2.1	2.1	2.1	2.2
	subset 1					•	sub	set 2	10	
	CON LAB	1        CON      9.9        LAB      16.4	12CON9.910.7LAB16.417.6CHEM2.02.0	123CON9.910.710.4LAB16.417.619.1CHEM2.02.01.7	1234CON9.910.710.413.9LAB16.417.619.115.4CHEM2.02.01.71.8	12345CON9.910.710.413.99.0LAB16.417.619.115.414.9CHEM2.02.01.71.82.2	123456CON9.910.710.413.99.011.8LAB16.417.619.115.414.917.8CHEM2.02.01.71.82.22.0	1234567CON9.910.710.413.99.011.812.5LAB16.417.619.115.414.917.817.0CHEM2.02.01.71.82.22.02.1	12345678CON9.910.710.413.99.011.812.59.8LAB16.417.619.115.414.917.817.016.4CHEM2.02.01.71.82.22.02.12.1	123456789CON9.910.710.413.99.011.812.59.812.6LAB16.417.619.115.414.917.817.016.419.1CHEM2.02.01.71.82.22.02.12.12.1

separate data analyses for all subsamples



#### LSMeans for total sample and subsets – Ethanol

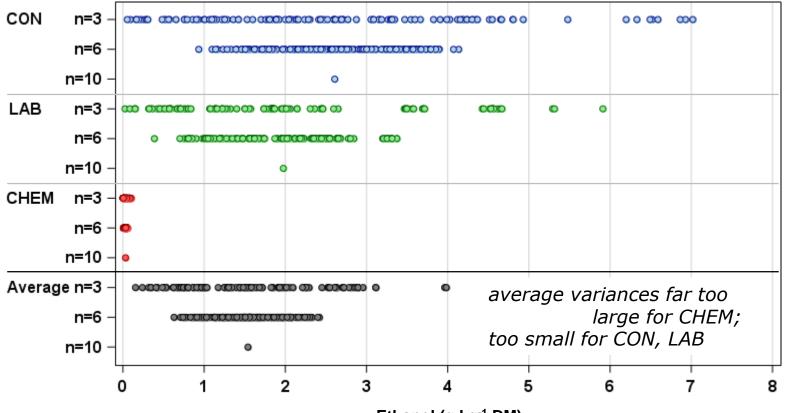


Compared to CON, LAB increases ethanol content, whereas CHEM decreases ...

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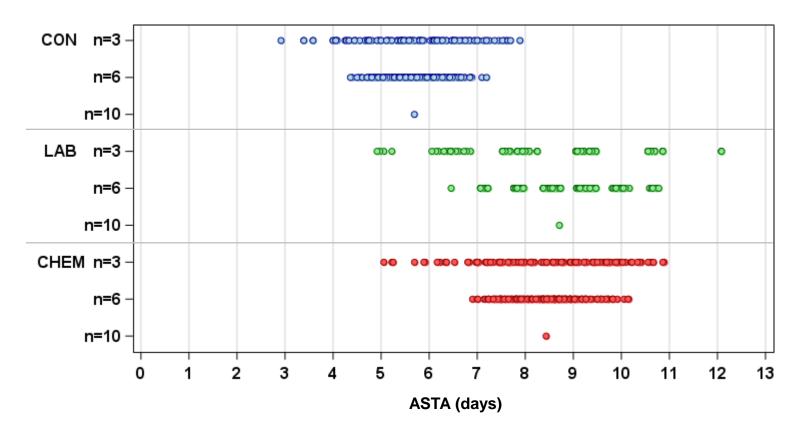
#### **Residual variances** for total sample and subsets – **Ethanol**



Ethanol (g kg<sup>-1</sup> DM)

Compared to CON, LAB increases ethanol content, whereas CHEM decreases ... CHEM reduces the variability dramatically!

#### **LSMeans** for total sample and subsets – **ASTA**

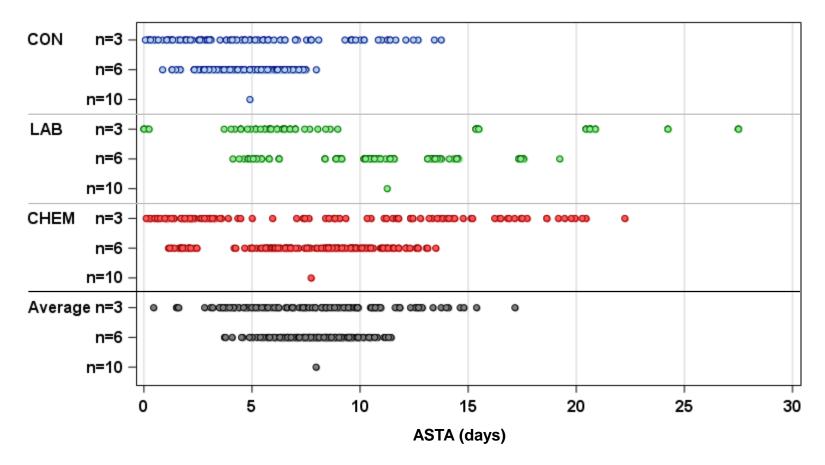


Compared to CON, LAB and CHEM show slightly higher aerobic stability ...

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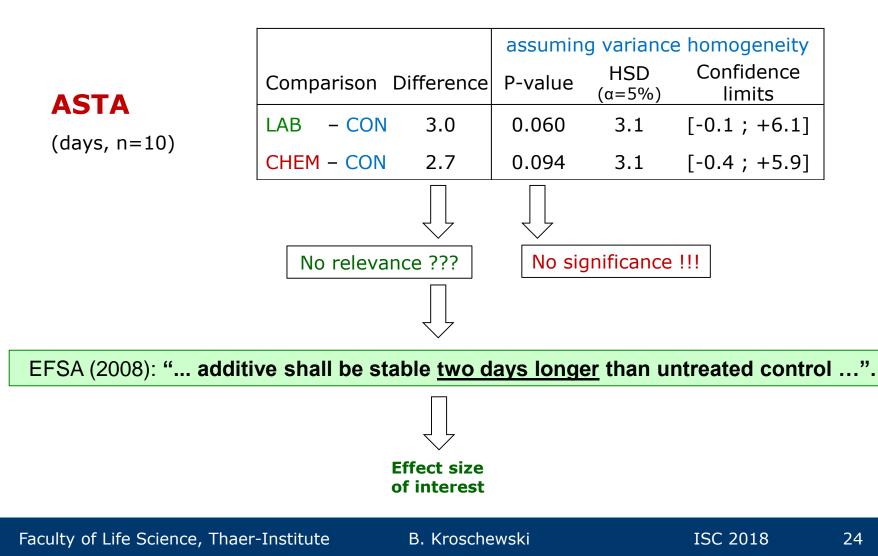
#### **Residual variances** for total sample and subsets – **ASTA**



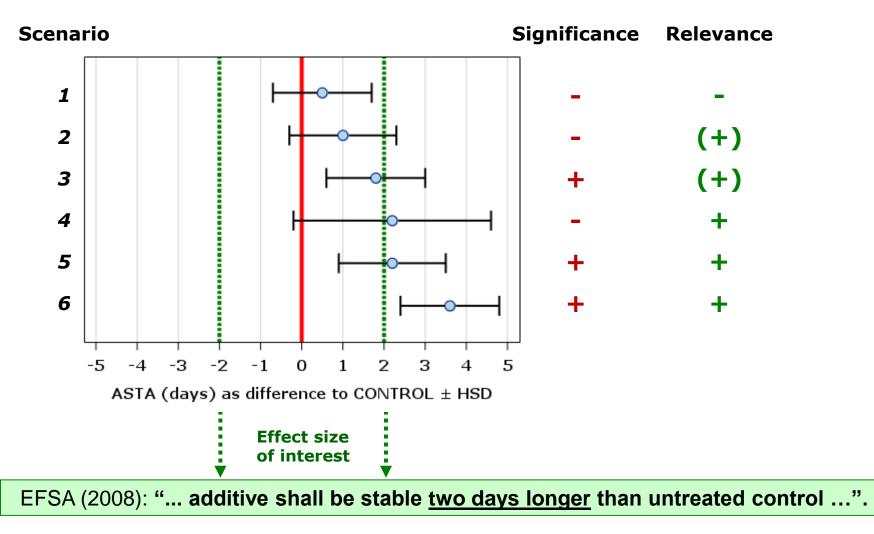
Compared to CON, LAB and CHEM show slightly higher stability ....

variability similar.









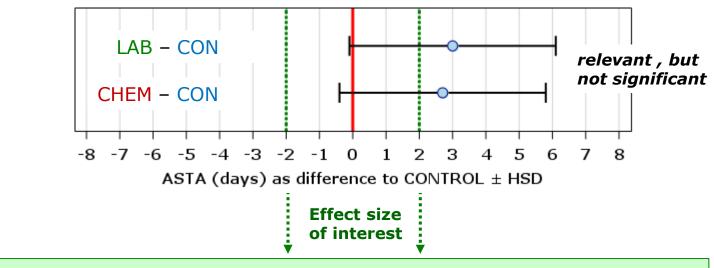
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ASTA

(days, n=10)

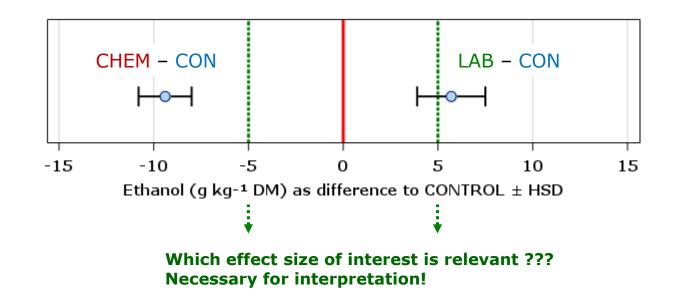
		assumin	g variance	e homogeneity
Comparison	Difference	P-value	HSD (α=5%)	Confidence limits
LAB – CON	3.0	0.060	3.1	[-0.1;+6.1]
CHEM – CON	2.7	0.094	3.1	[-0.4 ; +5.9]



EFSA (2008): "... additive shall be stable two days longer than untreated control ...".



Ethanol			assuming	g variance	heterogeneity
(g kg <sup>-1</sup> DM, n=10)	Comparison	Difference	P-value	HSD (α=5%)	Confidence limits
	LAB – CON	5.7	<0.001	1.8	[3.9 ; 7.5]
	CHEM – CON	-9.4	<0.001	1.4	[-10.7 ; -8.0]



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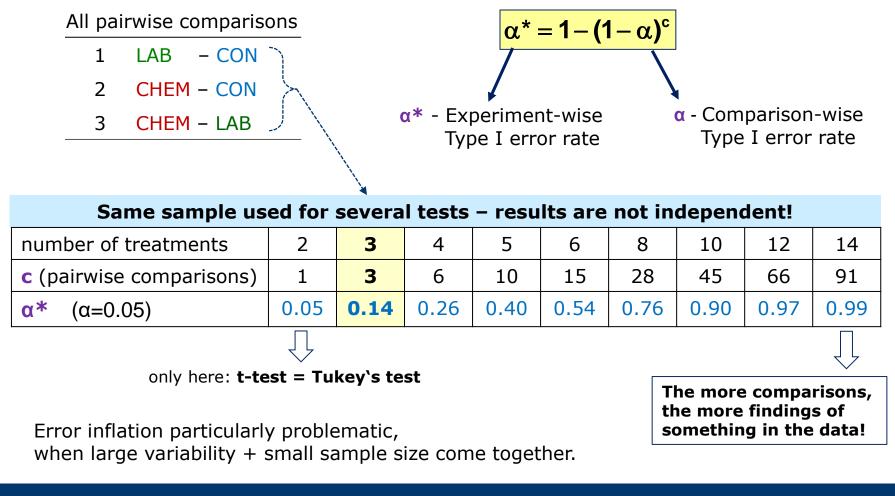
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# 3.2 Results – grass mixture (CON, LAB, CHEM)



### Comparison of treatments – problem of multiplicity

(caused by number of comparisons + numer of response variables)

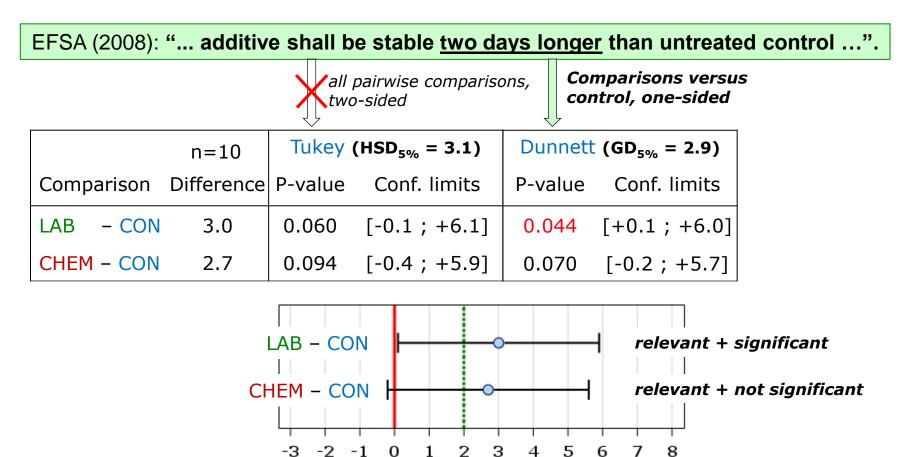


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# ASTA as one of the most important responses

(1) assuming normally distributed values  $\rightarrow$  parametric data analysis



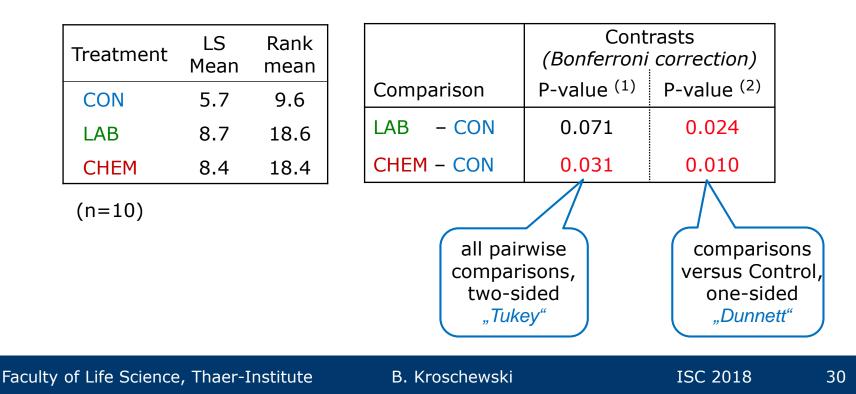
ASTA (days) as difference to CONTROL ± GD



# ASTA as one of the most important responses

(2) assuming non-normally distributed values  $\rightarrow$  nonparametric data analysis

- $\rightarrow$  rank procedure with ANOVA-Typ-Statistics (SAS, Proc Mixed)
  - for  $\geq$  1 treatment factor, variance heterogeneity of ranks considered, identical observations no problem, ...
  - but: minimal sample size for reliable results about  $\underline{n=10}$





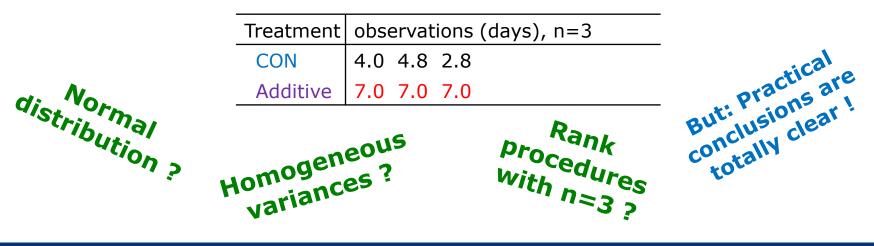
# ASTA as one of the most important responses

(3) How to consider identical observations for treatments?

• grass mixture: duration of air exposure 12.1 days

Treatment	observations (days), n=10				
CON	1.6 8.2				
LAB	3.0 8.4 12.1 12.1 12.1 12.1				
CHEM	3.0 11.3				

• often situation more extreme (Weiss et. al 2016): duration of air exposure 7 days



# 4. Summary

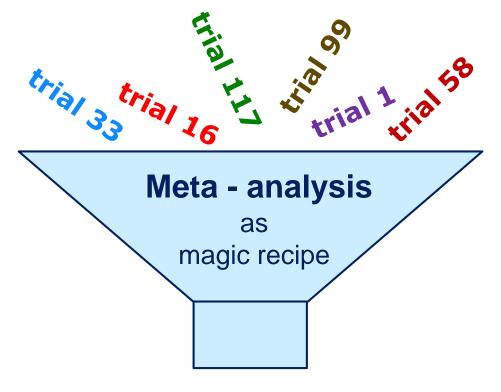


- Comparative silage experiments are most frequently performed with few replications.
- The traits of interest for the evaluation of treatments do **not meet** in each case the **assumptions** for the chosen statistical analysis procedures.
- Moreover, checking the assumptions by statistical preliminary tests and performing the analysis on the same data are problematic.
- Often, the experimenter has **information beforehand** about treatments which have an impact not only on the magnitude but also on variation.
- Both, rank procedures in case of non-normality as well as reliable estimations of individual treatment variances in case of variance heterogeneity demand higher sample sizes.
- Together with <u>well justified</u> effect sizes of interest, the trials should be designed in such a way that **significance** and **relevance** of results **come together**.

# 4. Summary



Problematic single trials? Don't worry! Put together!



Are aggregated results correct ?????

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