




COMPARISON OF SPMDs AND BIOTIC SAMPLERS USING GHOSTIC ANALYSIS

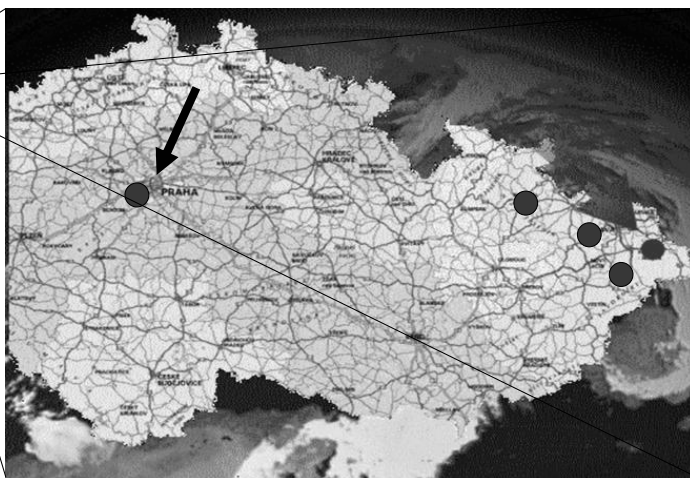
Institute of Public Health, Ostrava, Czech Republic
National reference laboratory for POPs

Tomas Ocelka, Pavel Kovanic

Tomas.Ocelka@zuova.cz

	TOPICS
	<ul style="list-style-type: none"> ▪ Sampling methods to be compared ▪ Objects of measuring ▪ Problems of analysis ▪ Gnostic analysis ▪ Methods' features to be compared ▪ Results of comparison

Geographic location



Centre laboratories, accreditation



EA

- Personnel: over 140, 5+2 workplaces
- According to ČSN EN ISO/IEC 17 025
 - Over 200 parameters, PCDD/Fs, PCBs, OCPs, PBDE, ...
 - Recognized by ILAC, EA, IAF
- Sampling and Testing
 - Integral - water
 - SPMDs
 - DGTs
 - POCIS
 - Biotic organisms
- Intercalibration
 - Czech + International
- Data analysis (univariate/multivariate)
 - Statistical
 - Gnostic



**Instrumentation
(worth over 6 mil. USD)**



GC-MS/MS (ion-trap)

- GCQ, Polaris
- Since 1996 (starting to POPs issue)

GC-HRMS (POPs)

- MAT 95XP
- since 2003



**LC-MS/MS (pharmacy,
pesticides)**


- ThermoFinnigan
- since 2006





Data source for comparison of methods


- All rivers within Czech Republic scale (15)
- 21 sampling profiles
- Complementary to biotic sampling system (since 1999) with abiotic (SPMDs, DGTs, POCIS) – since 2003
- Aims
 - Pilot application 2 years before routine application
 - Parallel exposure of *Dreissena Polymorpha*, Benthos, Plants
 - POPs (basic: OCPs, PCBs)
 - POPs (other: PCBs – cong., PCDD/Fs, PAHs, PBDEs)

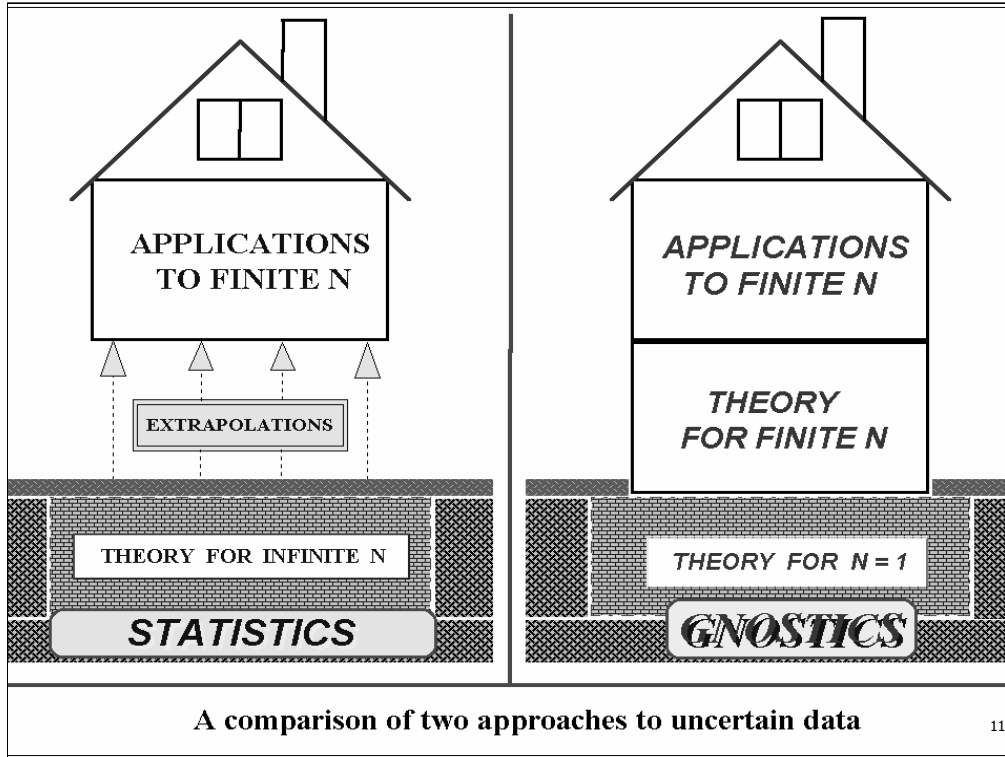


	<h2 style="text-align: center;">SAMPLING METHODS TO BE COMPARED</h2>
	<p>Three biotic methods:</p> <ul style="list-style-type: none"> ❑ Benthos ❑ Dreissena ❑ Plants <p>One abiotic method: SPMD (Semipermeable Membrane Measuring Device)</p>

	<p>The selection</p>
	<p>Concentrations of selected permanent organic pollutants (POPs) in several locations of Elbe river in Czech Republic:</p> <p>p.p.DDE, PCB138, PCB180, PCB101, PCB28.31, p.p.DDT, p.p.DDD, PCB52, PCB118</p>

	<h2>PROBLEMS OF ANALYSIS</h2>
	<ul style="list-style-type: none"> ❑ Small data samples ❑ Different mean concentrations ❑ Strong variability ❑ Different length of data vectors ❑ Data censoring (eg data below the LOD) ❑ Non-homogeneous and outlying data

	<p style="text-align: center;">SPECIFICS of MATHEMATICAL GNOSTICS</p>
	<ul style="list-style-type: none"> ❑ Theory of individual data and small data samples ❑ Realistic assumptions ❑ Uncertainty: a lack of knowledge ❑ “Let data speak for themselves” ❑ Results maximizing information ❑ Natural robustness <p style="text-align: right;">10</p>



GNOSTIC DISTRIBUTION FUNCTIONS

- No a priori model (everything from data)
- Maximum information
- Robustness in estimation of probability, quantiles, scale and location parameters, bounds of data support, and membership interval
- Robust correlations




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GNOTIC DISTRIBUTION FUNCTIONS II

- Data homogeneity tests
- Marginal cluster analysis
- Cross-section filtering
- Applicability to censored data
- Applicability to heteroscedastic data



	<h2 style="text-align: center;">QUALITY OF METHODS TO BE COMPARED</h2>
	<ul style="list-style-type: none"> <input type="checkbox"/> Relative sensitivity (threshold, range) <input type="checkbox"/> Homogeneity of results <input type="checkbox"/> Consistency of results <ul style="list-style-type: none"> ▪ Internal (of method's own results) ▪ External (mutual consistency of methods) <input type="checkbox"/> Informativeness of results <input type="checkbox"/> Precision

RELATIVE SENSITIVITY

Method's relative sensitivity depends:


- On the pollutant's concentration
- On the method's measuring domain

$$RS = (1 - NC/N) \times 100 (\%)$$

NC ... number of data in the interval
[sensitivity threshold, max(range)]

N ... all data of the sample



	<h1 style="text-align: center;">HOMOGENIZATION</h1> <p style="text-align: center;"><i>TO BE OR NOT TO BE?</i></p>
	<p>Homogeneous data:</p> <ul style="list-style-type: none"> the same origin of true values the same nature of the uncertainty <p>To homogenize?</p> <ul style="list-style-type: none"> ❑ Pros: <ul style="list-style-type: none"> More certain main cluster ❑ Cons: <ul style="list-style-type: none"> Possible loss of information <p style="text-align: center;"><i>Rule: homogenize and verify</i></p>

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MEASURABILITY

Homogenization ... elimination of outliers

$$Meas = (1 - (NL + NU)/N) \times 100 (\%)$$

NL ... number of lower outliers

NU ... number of upper outliers

N ... number of the sample's data

$N - NL - NU$... data of the main cluster



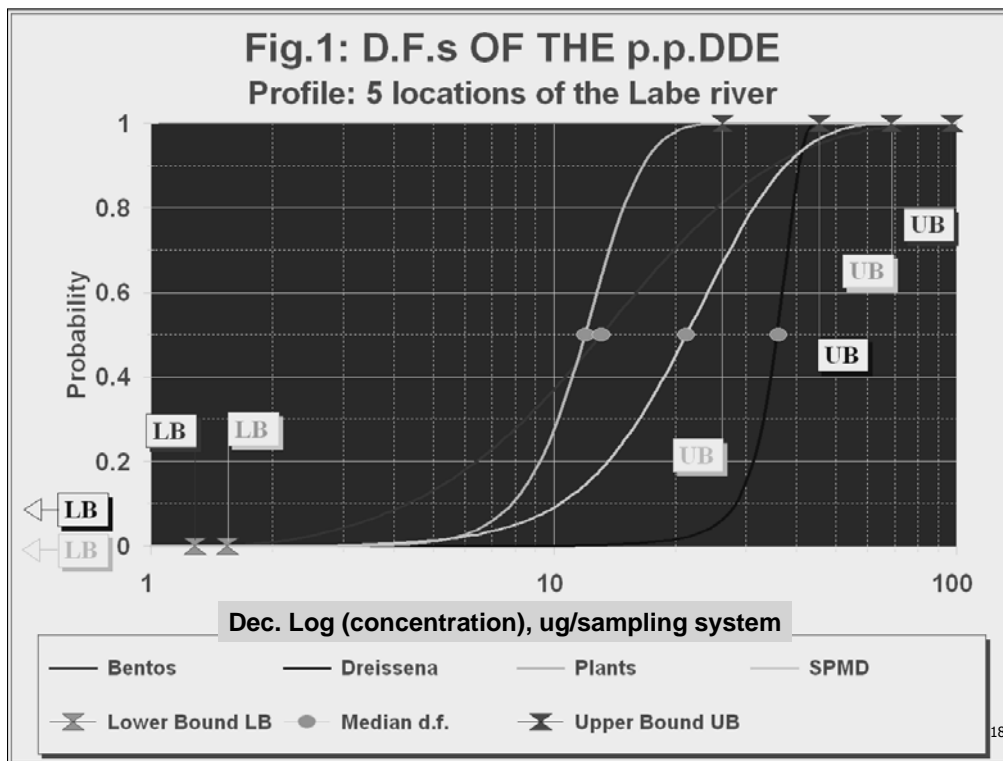


Fig.2: D.F.s OF THE p.p.DDE

Profile: 5 locations of the Labe river

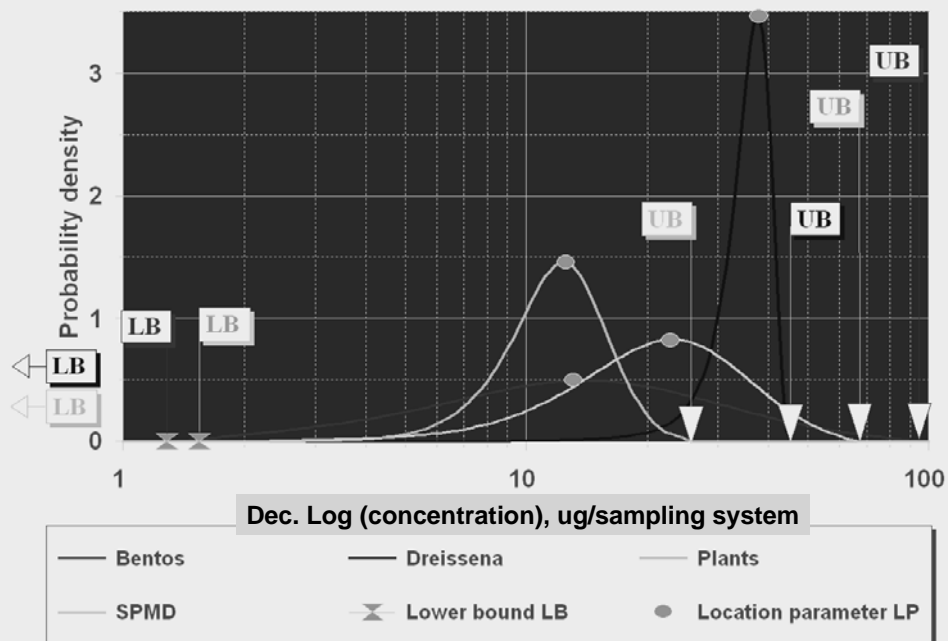
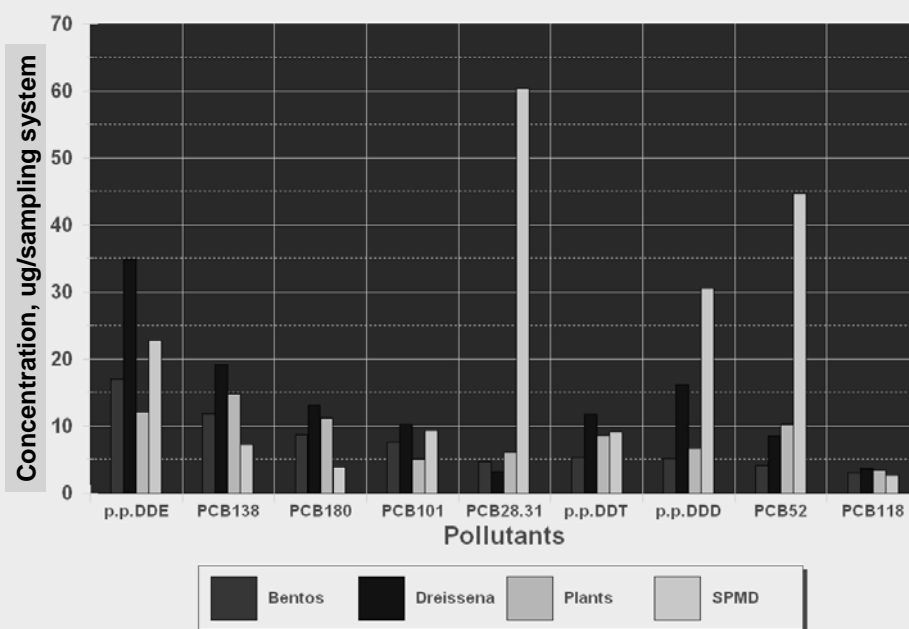



Fig.3: ROBUST MEANS OF CONCENTRATIONS
Profile: 5 locations of the river Elbe




DIFFERENCES IN METHODS

- ❑ Different accumulation of pollutants:
 - different mean concentrations
 - different variabilities
- ❑ Different relations between means
- ❑ Rare exception: agreement in PCB118
- ❑ Impact of outliers to SPMD? ***NO!***



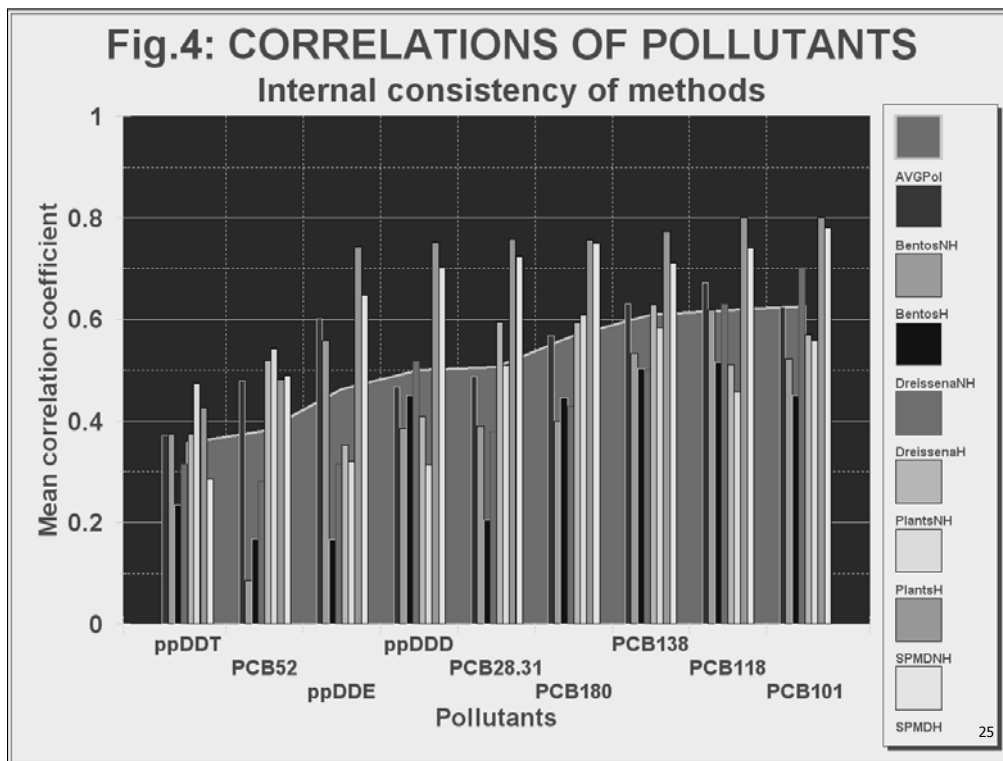
	<h2>METHOD'S CONSISTENCY</h2>
	<p>Methods are <i>consistent</i> when they give similar results</p> <p>Measuring of similarity: Correlations, or (more generally) mean angles between vectors of results</p> $SIM_{cc} = 100 \times \text{correl.coefficient} (\%)$ $SIM_{qa} = 100 \times (1 - Ang /180) (\%)$

	<h2 style="text-align: center;">GNOSTIC CORRELATIONS</h2>
	<p>Data error in gnostic: <i>irrelevance</i></p> $ir = (2p - 1)/2$ <p>p ... probability of the data item.</p> <p>Correlation coefficient of two samples:</p> $Gcc(M,N) = cc\{ir(m),ir(n)\}$ <p>$(m \text{ in } M, n \text{ in } N), cc\{.. \}$ statist. cor.coef.</p> <p>Robustness:</p> $- 1 \leq ir \leq + 1$

SIGNIFICANCE OF CORRELATIONS

- ❑ Problems: false statistical model (normality?!, finite data support), small data samples, unrobustness
- ❑ Gnostic estimating of significance:
 - fast, auxiliary: using Spearman's robust estimate of significance
 - carefully: distribution function of correlation coefficients





QUANTILE VECTORS

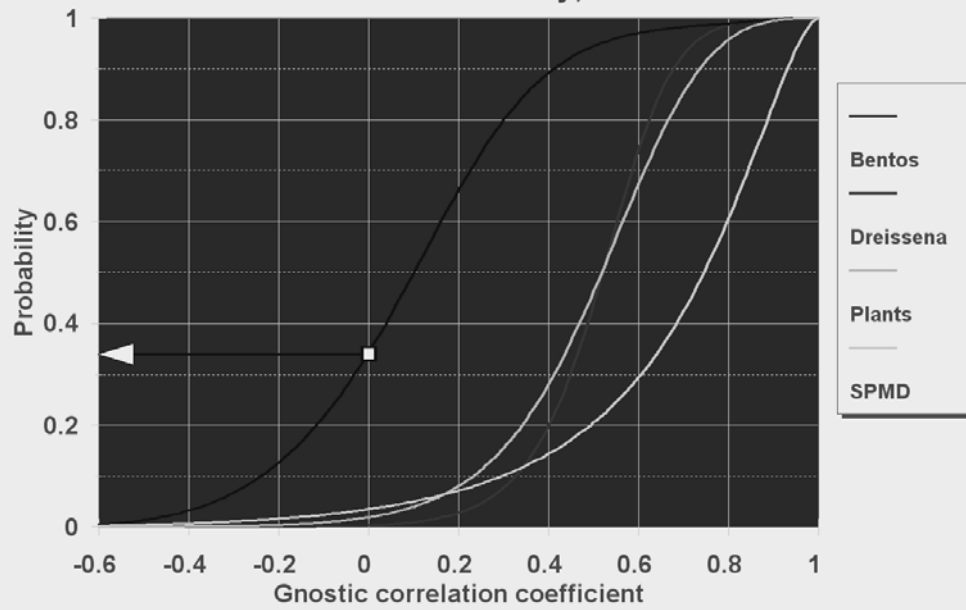
- Make sample's distribution function
- Set a series of probabilities p_1, \dots, p_N
- Find quantiles q_1, \dots, q_N so that $P\{q_k\} = p_k$
- Take q_1, \dots, q_N as a quantile vector

Advantages:

Robustness, making use of censored data, independence of data amount and of mean data value, filtering effect.



Fig.5: DISTRIBUTIONS OF CORREL. COEFS
Internal consistency, NH data



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Fig.6: DIVERGENCE OF DISTRIBUTIONS

Design of the quantile vectors

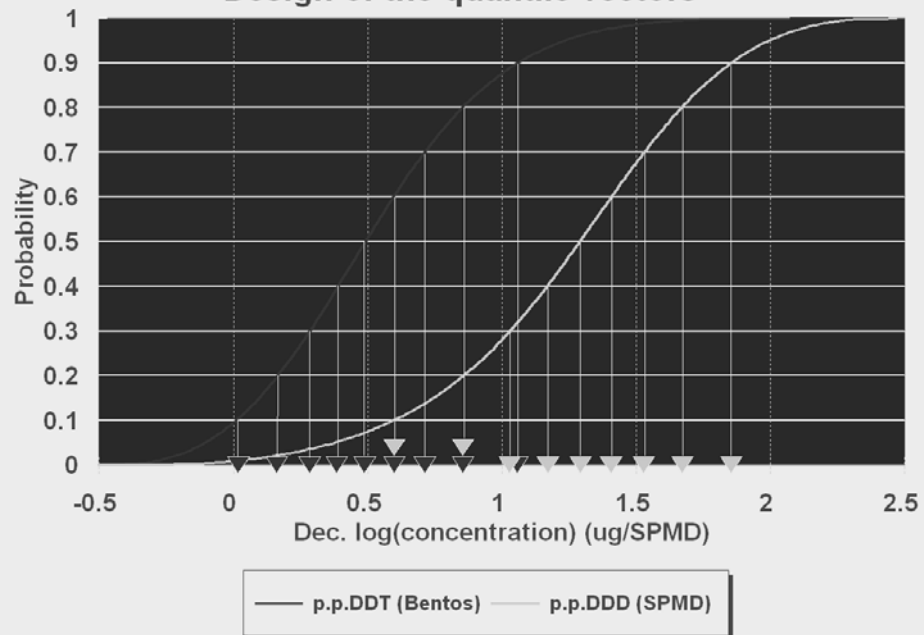
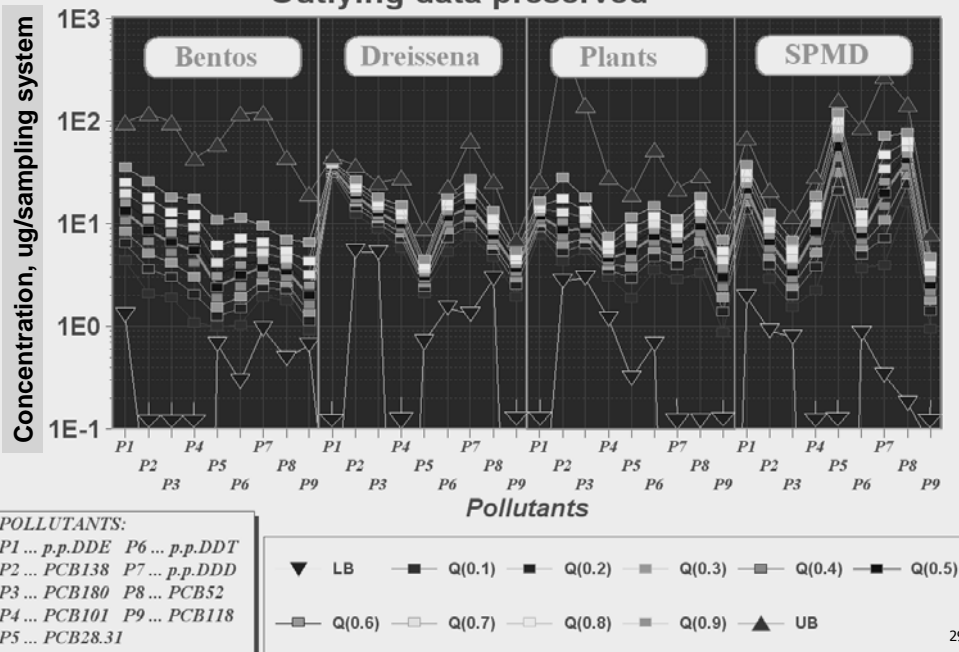


Fig.7: DIVERGENCE OF QUANTILES
Outlying data preserved



EXTERNAL CONSISTENCY

Approaches:

- ❑ Correlations
- ❑ Angles between MD-vectors of means
- ❑ Angles between quantile vectors
- ❑ Conjunction of typical data intervals
- ❑ Conjunction of data supports



INTERVAL ANALYSIS

1) Distribution functions


2) Interval analysis:

- a) Data support (LB, UB)
- b) Membership interval (LSB, USB)
- c) Interval of typical data (ZL, UL)
- d) Tolerance interval ($Z0L, Z0U$)

3) Overlapping:

$100 \times \text{conjunction}(I1, I2) / \text{union}(I1, I2) (\%)$



	<h2 style="text-align: center;">INFORMATIVENESS</h2>
	<ol style="list-style-type: none"> 1) Data sample 2) Distribution function 3) Probability p of an individual data item 4) Information of the data item: $Info = (p \log(p) + (1-p) \log(1-p)) / \log(1/2)$ 5) Informativeness of a data sample: $100 \times Mean(Info) (\%)$

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EVALUATION OF PRECISION

- Weak variability:

$$Prec = 100 \times (1 - STD/AVG) (\%)$$

(*STD* ... standard deviation, *AVG* ... mean)

- Strong uncertainty:

$$Prec = 100 \times (1 - Mean(GW)) (\%)$$

(*GW* ... gnostic weight of data; entropy change caused by the uncertainty)

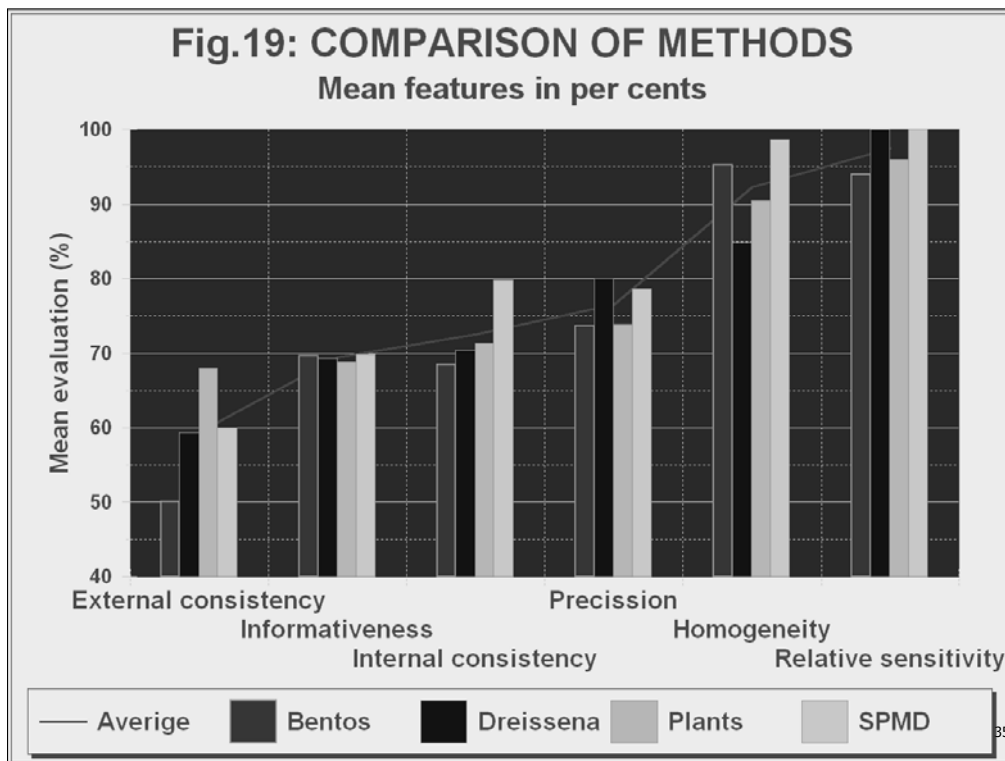
$$0 \leq GW \leq 1$$



SUMMARY COMPARISON


	Average of 14 evaluations	
Method	Non-hom.data	Homog. data
Bentos	60.9 %	62.7 %
Dreissena	64.5 %	67.5 %
Plants	64.2 %	68.9 %
SPMD	67.5 %	69.5 %






RATING OF METHODS


Feature	Bentos	Dreiss.	Plants	SPMD
Ext.consistency	4	3	1	2
Int.consistency	4	3	2	1
Informativeness	1	3	4	2
Precision	3	1	4	2
Homogeneity	2	4	3	1
Rel.sensitivity	3	1	2	1
Mean rating	2.8	2.5	2.7	1.5


 ZDRAVOTNÍ ÚSTAV
 JIHOMORAVSKÉHO ÚZEMÍ

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	<h2>Conclusions</h2>
	<ul style="list-style-type: none"> ■ Passive sampling, like SPMDs shown the best results; if there are no legal requirements for biota, biotic organisms can be replaced ■ Do not forget to analyze data precisely, independently, before your interpretation <ul style="list-style-type: none"> – Do not rely ONLY on functionality of any processing package – Statistical approach has some limitations on small data sets (majority of monitoring studies) ■ Any headache from analytical tools can be eliminated by experience <ul style="list-style-type: none"> – Try it!

	<h2>Further intentions</h2>
	<ul style="list-style-type: none"> ■ Finalization of Gnostic analytical tool, with GUI (S-Plus) ■ Extension to other platforms by interface ■ Linking to databases (LIMS, GIS, ...) ■ Training and dissemination ■ Projects solutions and participations <ul style="list-style-type: none"> – Join us: 2-FUN project, www.2-fun.org

... **thank you for your attention!**

