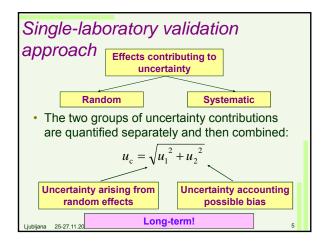
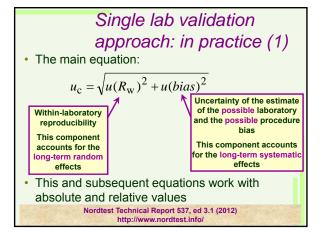


Approach based on validation and Quality Control Data

aka "the Nordtest approach"

Nordtest Technical Report 537, ed 3.1 (2012) http://www.nordtest.info/





Absolute vs relative uncertainties: Rules of Thumb

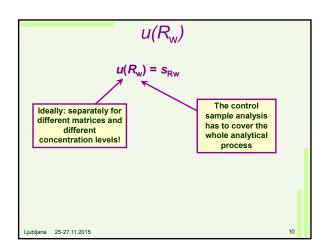
- · In general: use whichever is more constant
- Some rules of thumb:
 - At low concentrations (near detection limit, trace level) use absolute uncertainties
 - · Uncertainty is not much dependent on analyte level
 - At medium and higher concentrations use relative uncertainties
 - Uncertainty is roughly proportional to analyte level

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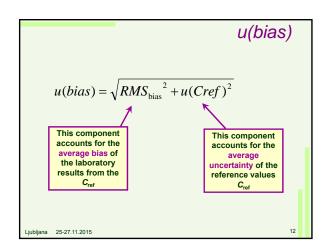
Single lab validation approach: in practice

Steps of the process:

- 1*. Specify measurand
- 2. Quantify R_{w} component $u(R_{w})$
- 3. Quantify bias component u(bias)
- 4*. Convert components to standard uncertainties u(x)
- 5*. Calculate combined standard uncertainty \boldsymbol{u}_{c}
- 6*. Calculate expanded uncertainty **U**
- * Note general step the same for modeling (i.e. ISO GUM)
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- $u(R_{\rm w})$ is the uncertainty component that $U(R_{\rm w})$ takes into account long-term variation of results within lab, that means: within-lab reproducibility (s_{Rw}) **Including sample** Ideally: preparation The same sample Sample similar to test samples – matrix, concentration, homogeneity - The same lab - The same procedure - Different days (preferably over 1 year) - Different persons - Different reagent batches Repeatability < Within-lab reproducibility < Combined uncertainty SRW



The possible bias of lab's results from the best estimate of true value is the account Including sample preparation u(bias) can be found: From analysis of the same samples with a reference procedure From analysis of certified reference materials (CRMs) - From interlaboratory comparison me Replicate - From spiking experiments measurements Ideally: several reference materials, several PTs because the bias will in most cases vary with matrix and concentration range



u(bias)

The averaging is done using the root mean square:

$$bias_{i} = Clab_{i} - Cref_{i} \qquad RMS_{bias} = \sqrt{\frac{\sum (bias_{i})^{2}}{n}}$$
$$u(Cref_{i}) = \sqrt{\frac{\sum u(Cref_{i})^{2}}{n}} \qquad u(Cref_{i}) = \frac{s_{i}}{\sqrt{n_{i}}}$$

- Each bias, is obtained as an average of replicate measurements
- Only this way is it possible to reduce the random effects
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u(bias): only one CRM

· If only one single CRM is used:

$$u(bias) = \sqrt{RMS_{\text{bias}}^2 + s_{\text{bias}}^2 / n + u(Cref)^2}$$

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Uncertainty due to possible bias

Evaluation of uncertainty due to bias, ideally:

- Separately for different sample matrices
- Separately for different concentration levels

This approach is rather demanding in terms of availability of sample data

Single-lab validation approach in practice:

Determination of acrylamide in snacks by LC-MS

- Concentration level 998 μg/kg
- Laboratory has analysed two certified reference materials (CRMs) with similar matrixes
 - Potato chips and crisp bread
 - The crisp bread CRM is also used as a control sample

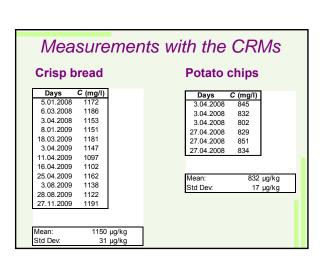
Certified reference material (CRM)

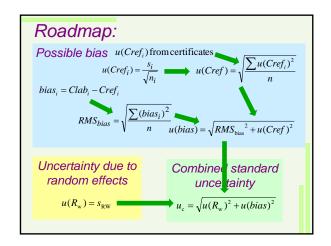
 The crisp bread CRM has the following acrylamide content:

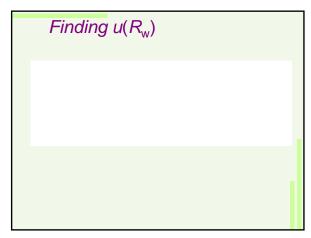
 $C_{\text{acrylamide}} = (1179 \pm 68) \, \mu \text{g/kg} \, (k = 2, \text{norm.})$

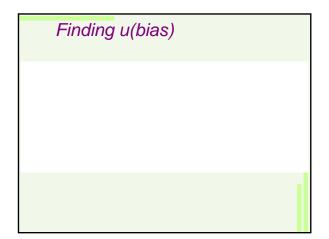
 The potato chips CRM has the following acrylamide content:

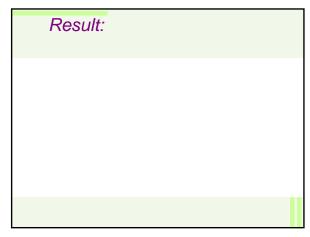
 $C_{\text{acrylamide}} = (860 \pm 42) \,\mu\text{g/kg}$ (k = 2, norm.)













Thank you for your participation! The materials are available from: http://tera.chem.ut.ee/~ivo/Temp/QA Hg Ljubljana 2015/ More explanations and examples: http://sisu.ut.ee/measurement/ You are always welcome to contact me: ivo.leito@ut.ee

