

University of Stuttgart Germanv



Analytische Qualitätssicherung Baden-Württemberg

Proficiency Test 5/16 TW S2 – pharmaceuticals in drinking water

sulfadiazine, sulfadimidine, sulfamethoxazole, sulfaethoxypyridazine, sulfamerazine, sulfathiazole sulfadoxine, sulfamethoxypyridazine, sulfachloropyridazine, sulfadimethoxine, trimethoprime

provided by AQS Baden-Württemberg at Institute for Sanitary Engineering, Water Quality and Solid Waste Management, University of Stuttgart Bandtäle 2, 70569 Stuttgart-Büsnau, Germany

> and **IWW Water Center** Moritzstr. 26, 45476 Mülheim an der Ruhr, Germany





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General

This PT was provided by AQS Baden-Württemberg in cooperation with IWW Water Center in Mülheim an der Ruhr and with the network "NORMAN" (Network of reference laboratories for monitoring of emerging environmental pollutants). In this round sulfadiazine, sulfadimidine, sulfamethoxazole, sulfaethoxypyridazine, sulfamerazine, sulfathiazole, sulfadoxine, sulfamethoxypyridazine, sulfachloropyridazine, sulfadimethoxine and trimethoprime were to be determined.

The PT was executed and evaluated according to the requirements of DIN 38402-A45 and ISO/TS 20612.

PT design

Each participant received the following samples:

• 3 samples for the determination of above mentioned parameters in 1000-mlground bottles. The samples were preserved by adding 40 mg/l sodium azide. The samples also contained acetonitrile as solubility promoter.

3 different concentration levels/batches were produced. All participants received the same samples.

Sample preparation

The samples for the determination of the pharmaceuticals were based on a real ground water matrix from the northern part of the region Ruhr in North Rhine-Westphalia. The ground water was used without treatment for the sample preparation.

The ground water was spiked with stock solutions and the concentrations covered drinking and ground water relevant ranges.

Sample distribution

The samples were dispatched on 11 October 2016 by express service.

Analytical methods

The participants were free to choose a suitable method, but following limits of quantification were required:

Parameter	limit of quantification
 sulfadiazine 	0.05 µg/l
sulfadimidine	0.05 μg/l
 sulfamethoxazole 	0.05 µg/l
 sulfaethoxypyridazine 	0.05 µg/l
sulfamerazine	0.05 µg/l
sulfathiazole	0.05 µg/l
sulfadoxine	0.05 μg/l
 sulfamethoxypyridazine 	0.05 µg/l
sulfachloropyridazine	0.05 µg/l
 sulfadimethoxine 	0.05 μg/l
trimethoprime	0.05 μg/l

The samples had to be analysed in duplicate over the complete method (sample preparation and measurement). The participants were asked to report the results as average values in μ g/l with three significant digits.

Submission of results

The deadline for the submission of results was on 07 November 2016.

Evaluation procedure

The statistical evaluation was executed according to DIN 38402-A45 and ISO TS 20612 "Interlaboratory comparisons for proficiency testing of analytical chemistry laboratories". From the participants' results a relative standard deviation was calculated for each concentration level and parameter using the Q-method. The reference values (see chapter "Traceable reference values") were used as assigned values x_{pt} . The standard deviation resulting from the Q-method was used as σ_{pt} .

 σ_{pt} was limited as follows:

- lower limit: 5%
- upper limit: 25%

A z-score was calculated for each measurement result derived from the assigned value x_{pt} and the standard deviation for proficiency assessment σ_{pt} :

$$z - Score = \frac{\left(x - x_{pt}\right)}{\rho_{pt}}$$

The z-score was modified to a z_U -score with a correction factor for proficiency assessment (as described in the standards mentioned above). The tolerance limit was defined as $Iz_UI=2.0$.

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Assessment

There was no overall assessment of the proficiency test round, but the single parameters were assessed.

A parameter was assessed as successful, if more than half of the values were correctly determined (2 out of 3 values are within the tolerance limits).

According to ISO 13528 (2015) the single results were assessed as follows:

$ z_u \leq 2.0$	successful
$2.0 < z_u < 3.0$	questionable
$ z_u \ge 3.0$	unsatisfactory

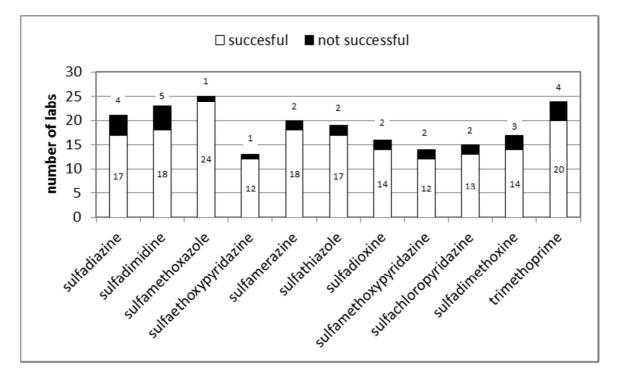
Not successful were:

- 1) Values which were not determined (if the other samples of this parameters were analysed),
- 2) Values, which were indicated with "lower than limit of quantification",
- 3) Values, which were subcontracted,
- 4) Values, which were submitted after the deadline of submission of results.

Evaluation

Number of participants:	26	
1 laboratory did not report any result.		
Number of reported values:	621	
Number of accepted values:	544	(87,6%)

In the following figure the successful and not successful laboratories for each parameter are illustrated.



Explanation of Appendix A

Appendix A contains for each parameter

- parameter tables
- a figure of participants' means versus the spiked amounts for the determination of the recovery rate
- a figure of the relative standard deviations versus the concentrations
- a figure of the tolerance limits in the PT versus the concentrations
- the frequency of application of analytical methods
- the method specific evaluation
- a comparison of mean and reference values for each concentration level
- a comparison of the relative standard deviations of the different methods
- the statistical characteristics of the method specific evaluation
- a tabular comparison of the means with the reference values and their uncertainties

Parameter tables

In these tables the following values for each concentration level are listed:

- assigned value
- expanded uncertainty of the assigned value in %, calculated from an uncertainty budget (see chapter "Traceable reference values")
- standard deviation, calculated using robust statistical method
- standard deviation for proficiency assessment for the calculation of z_U-scores
- rel. standard deviation for proficiency assessment
- tolerance limits above and below
- permitted deviations above and below in %
- number of values in this level
- number of not satisfactory values below and above the assigned value and the percentage of these values in total.

Determination of recovery rate

In the diagrams of the assigned values versus the spiked amount of analyte a linear regression line was calculated using a generalized least square regression which takes into account the uncertainties of the values in both directions. From these values the recovery rate for each parameter was determined (see diagrams). The slope of the line indicates the average recovery rate. The diagrams also contain the expanded uncertainty (k=2) of the concentrations from the spike and the assigned values.

Relative standard deviations and tolerance limits

The diagrams for the relative standard deviation vs. the assigned value show the concentration dependency of the standard deviation and the tolerance limits in percent. The relative standard deviations calculated from participants' data are the stars connected by an interrupted line, the rel. standard deviation taken from the variance function (and sometimes limited by the upper or lower limit) are given by squares, connected by a continuous line.

Method specific evaluation

For each parameter the methods used by the participants are shown in a diagram. In a second diagram for each method with a frequency of more than 5 %, values are sorted in 5 categories:

too low	results with z_U -score < -2
low	results with $-2 \le z_U$ -score < -1
correct	results with $-1 \le z_U$ -score $\le +1$
high	results with +1 < z_U -score \leq +2
too high	results with z_U -score > +2

Comparison of means and reference values for each concentration level

Finally the mean value calculated from all results, the reference value (see chapter Traceable reference values) are compared with mean values calculated for all methods separately. All mean values were calculated using the Hampel estimator described in ISO/TS 20612. Mean values were calculated only, if more than 8 results were within a z-score-range of ± 2 . The means are reported with their expanded uncertainty, calculated according to ISO 13528.

All mean values and their expanded uncertainties are additionally compared with the reference values and their expanded uncertainties.

Explanation of Appendix B

Participants were asked to report expanded uncertainties of their results on a voluntary basis. In this diagram for each parameter the reported uncertainties for all concentration levels with the reproducibility standard deviation (horizontal line) are displayed. Values which deviate from the reproducibility standard deviation with a factor more than 2 are usually not realistic.

Explanation of Appendix C

In the last part of the report, for all concentration levels the results of all participants are illustrated. Confidentiality of participants is ensured by using lab codes. The lab codes were sent to participants with the certificates.

In detail Appendix C contains:

- a table with all data
- figures with
 - o all reported results
 - \circ all z_U-scores
 - o all reported expanded uncertainties
 - \circ all ζ -scores

Table with all data

The assigned value with the expanded uncertainty and the tolerance limits for the concentration level is illustrated in the table. For each participant the following data are given:

- lab code
- reported result
- measurement uncertainty of the value (if reported)
- ζ -score for this value, calculated with the following formula

$$\zeta = \frac{X - X_{pt}}{\sqrt{u_{lab}^2 + u_{x_{pt}}^2}}, \text{ with}$$

 $X - X_{pt}$

•

- $\begin{array}{l} & \begin{array}{l} & \begin{array}{l} & \end{array} \\ & \end{array} \end{array} = difference from the measured value and the assigned value \\ & \begin{array}{l} & \\ & \end{array} \\ & \begin{array}{l} & \\ & \\ & \end{array} \\ & \begin{array}{l} & \\ & \\ & \end{array} \\ & \begin{array}{l} & \\ & \end{array} \end{array} = standard uncertainty of the value, reported by the participant \end{array}$
- $U_{x_{pt}}$ = standard uncertainty of the assigned value

- z_U-score for proficiency assessment
- assessment of the value according to its z-score

Meaning of ζ-scores:

The assessment of ζ -scores is similar to that of z_U -scores. If the data are normally distributed and the uncertainties are well estimated, ζ -scores will lie between -2 and +2 with a probability of around 95 %.

 ζ -scores are mainly influenced by the measurement uncertainties reported by the laboratory. Therefore ζ -scores are usually not appropriate for the assessment of the reported results, unless the reported measurement uncertainty is checked for fitness-for-purpose.

Therefore we do not use the ζ -scores for the assessment of the laboratories. Nevertheless ζ -scores are appropriate to check the plausibility of the reported measurement uncertainty:

If the z_U -score of a result is within the tolerance limit and the ζ -score is outside, then the measurement uncertainty is underestimated.

If the z_U -score is outside the tolerance limits and the absolute value of the ζ -score is lower than two, then the requirements of the proficiency test were stronger compared with the reported measurement uncertainty.

Diagrams of uncertainty data

In the first figure for all lab codes the measurement uncertainty (together with the reproducibility standard deviation) is illustrated. The second figure shows the associated ζ -scores.

Measurement uncertainty

10 (40%) out of 25 laboratories with valid values reported measurement uncertainties. In total 231 (37.2%) out of 621 valid values were given with the measurement uncertainty. The following table displays the number of values with measurement uncertainty against the accreditation status.

Accreditation status of	Number of	Number of values with meas-
the values	values	urement uncertainty
accredited	339	192 (56.6%)
not accredited	66	33 (50%)
not specified	216	6 (2.8%)

We would like to put emphasis on the fact that reporting of measurement uncertainties in our PT scheme is absolutely voluntary. The only objective is to help all participants to reasonably handle measurement uncertainties and their estimation.

The diagrams show that the spread of reported uncertainties in some cases is vast, from unrealistic low values up to very high. A plausibility check using reproducibility standard deviations of the PT round could be helpful here.

If measurement uncertainties are underestimated values assessed as "satisfactory" in the PT ($|z_U| \le 2$), will have a large ζ -score. $|\zeta| > 2$ means that the "own" requirements (defined in terms of estimated uncertainty) are not fulfilled.

27 (13.9%) of the 194 values reported with uncertainties and having a z_U -score $|z_U| \le 2.0$ had a ζ -score > 2.0. This means that the requirements of the PT scheme have been fulfilled , but not the "own" requirements, the uncertainty is underestimated.

Traceable reference values

Traceability of analytical results to national and international references is of increasing importance in all laboratories. This is not easy to realise for chemical analyses and often can only be done by analysing certified reference materials. But availability of these reference materials in the water sector is very limited. Therefore we try to provide reference values for the proficiency test samples, traceable to national and international references.

Since our PT samples without exception are spiked, real water samples, reference values can be calculated from the sum of matrix content and spiked amount of analyte. For both summands traceable values and their uncertainty have to be determined. Thereby we assume that no unrecognised bias resulting from sample preparation and transport is present and that we recognise all uncertainty components.

All spiking of samples was controlled gravimetrically and volumetrically. This procedure allows the preparation of a complete uncertainty budget.

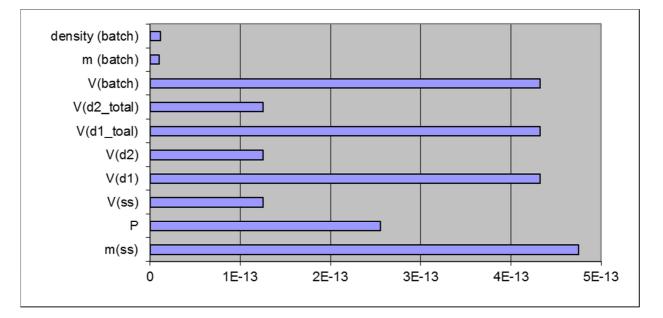
The first step is the specification of the measurand with a formula. This shows the links between the result and all influence quantities for the parameter trimethoprime.

$$c_{lot} = \frac{m_{ss} \cdot P \cdot V_{d1} \cdot V_{d2} \cdot V_{lot} \cdot \rho_{lot}}{V_{ss} \cdot V_{d1_total} \cdot V_{d2_total} \cdot m_{lot_total}}$$

with:

Clot	concentration of the analyte in the lot resulting from the spike
m _{ss}	mass of substance added for preparation of the stock solution
V _{d1}	volume of stock solution added into the dilution A
V _{d2}	volume of dilution A added into the dilution B
V _{lot}	volume of dilution B added into the lot
V _{ss}	volume of stock solution
V _{d1 total}	total volume of dilution A
V _{d2 total}	total volume of dilution B
m _{lot_total}	total mass of the lot
ρ _{lot}	density of the lot in g/l
P	purity of the used substance

Based on this formula the uncertainty budget can be prepared and all components can be quantified. The following figure shows a typical distribution of the contributions for trimethoprime. The main contributions result from the expanded uncertainty (0,5%), of the purity of the chemical (99,1%) and the pipette steps.

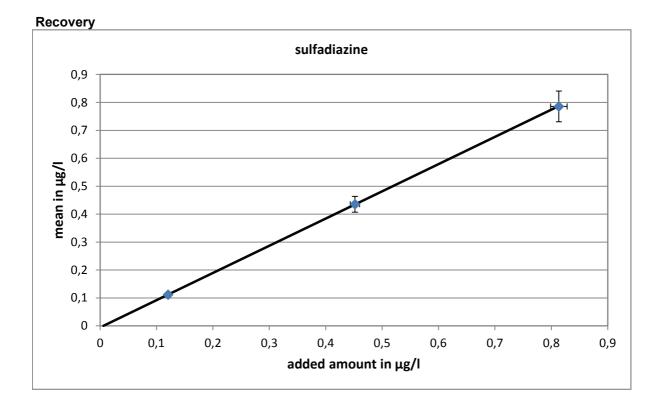


Attention was paid to use a ground water which did not contain any of the analytes. Therefore no matrix content had to be considered and the reference values could be calculated directly from the spikes.

Internet

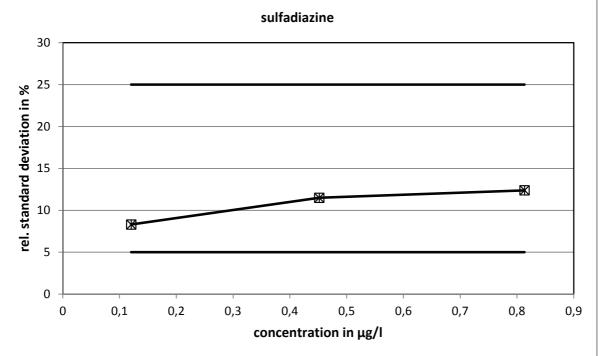
The report is available on the following webpage: http://www.aqsbw/pdf/report516.pdf

	sulfadiazine												
level	assigned value [µg/l]	expanded uncertainty of the assigned value [%]	standard deviation, calculated using robust statistics [µg/l]	standard deviation for proficiency assessment [µg/l]	standard deviation for proficiency assessment [%]	upper tolerance limit [µg/l]	lower tolerance limit [µg/l]	upper tolerance limit [%]	lower tolerance limit [%]	number of results	out below	out above	out [%]
1	0,1205	1,80	0,0100	0,0100	8,31	0,1415	0,1013	17,43		21	1	3	19,0
2	,	1,80	0,0519	0,0519	11,49	0,5626	0,3534	24,53		21	0	4	19,0
3	0,8133	1,80	0,1008	0,1008	12,39	1,029	0,6231	26,58	-23,39	21	1	1	9,5
									sum	63	2	8	15,9

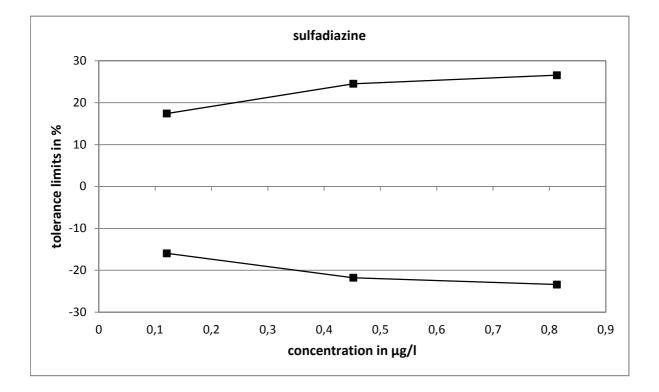


slope of the regression:0,975average recovery:97,5%

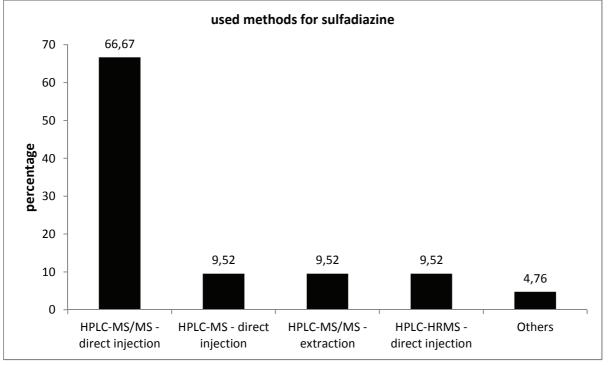


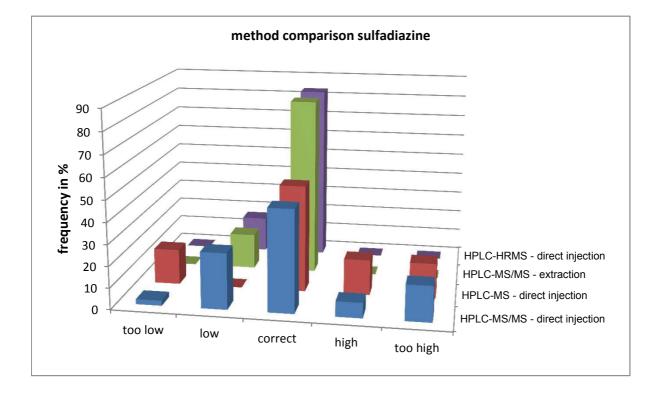


The relative standard deviations, calculated with the Q-method, did not reach the limits.

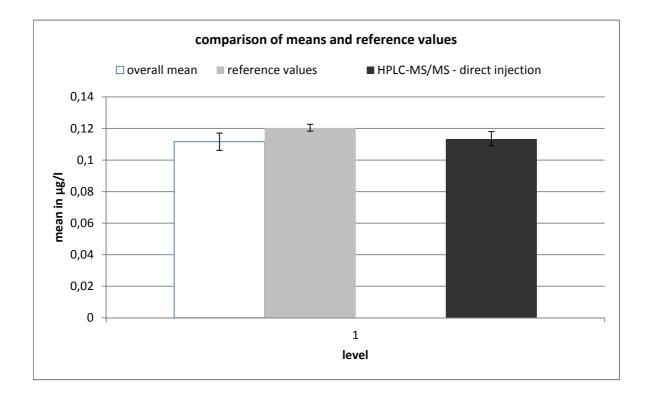


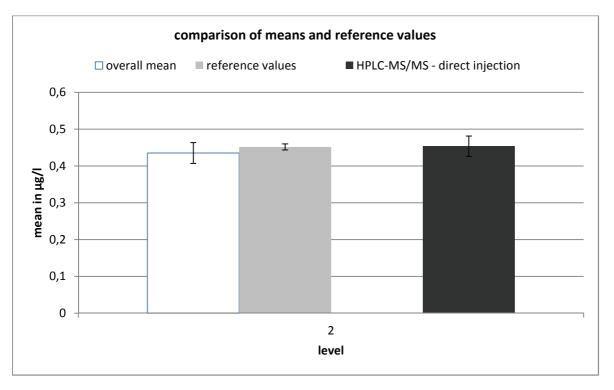
Method specific evaluation

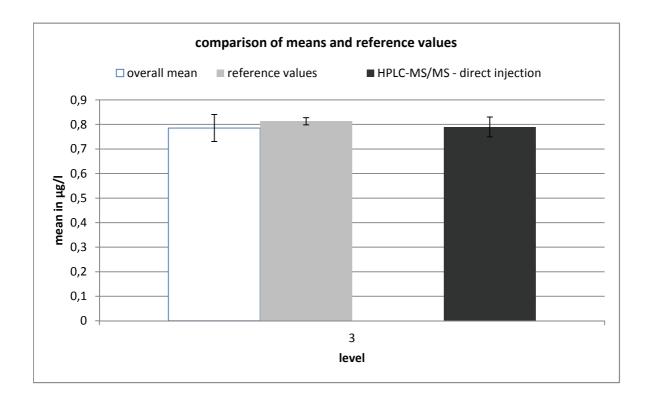


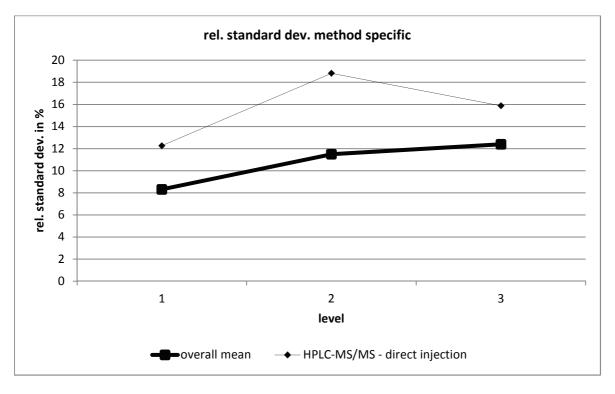


level	mean [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]	reference value [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]
1	0,1116	0,0055	4,9	0,1205	0,0022	1,8
2	0,4353	0,0283	6,5	0,4518	0,0082	1,8
3	0,7855	0,0550	7,0	0,8133	0,0147	1,8



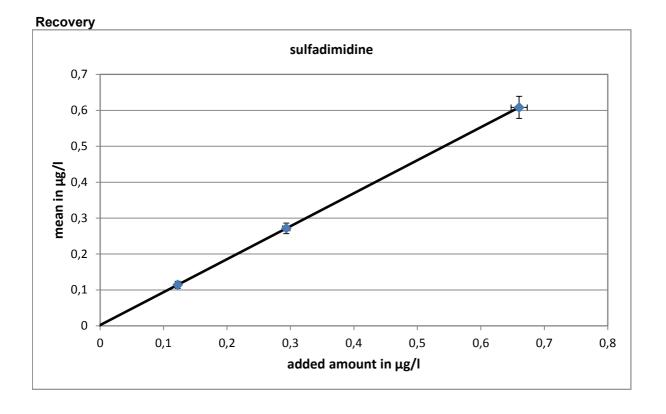






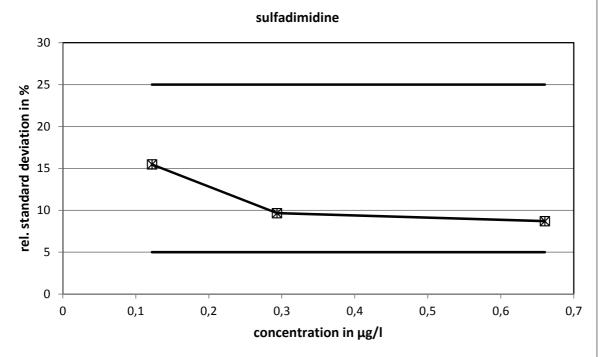
HPI	LC-MS/N	NS - dire	ect injec	tion					
level	robust mean [µg/l]	exp. unc. of the mean [µg/l]	exp. unc. of the mean [%]	robust standard deviation [µg/l	robust standard deviation [%]	number of results	out below	out above	out [%]
1	0,113	0,004	3,958	0,014	12,26	15	1	3	26,67
2	0,454	0,028	6,075	0,085	18,82	15	1	1	13,33
3	0,79	0,04	5,124	0,125	15,88	15	1	1	13,33

	sulfadimidine												
level	assigned value [µg/l]	expanded uncertainty of the assigned value [%]	standard deviation, calculated using robust statistics [µg/l]	standard deviation for proficiency assessment [µg/l]	standard deviation for proficiency assessment [%]	upper tolerance limit [µg/l]	lower tolerance limit [µg/l]	upper tolerance limit [%]	lower tolerance limit [%]	number of results	out below	out above	out [%]
1	0,1223	1,88	0,0189	0,0189	15,48	0,1638	0,0871	33,97	-28,80	24	0	3	13,0
2	0,2935	1,90	0,0284	0,0284	9,66	0,3535	0,2394	20,44	-18,42	23	4	2	26,1
3	0,6604	1,91	0,0575	0,0575	8,71	0,7813	0,5502	18,32	-16,67	22	3	1	17,4
									sum	69	7	6	18,8

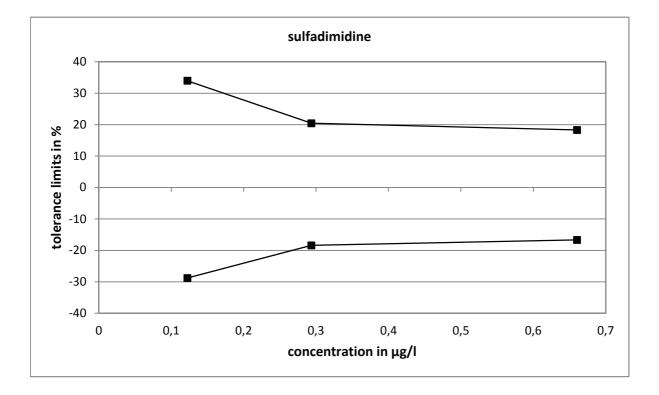


slope of the regression:0,919average recovery:91,9%



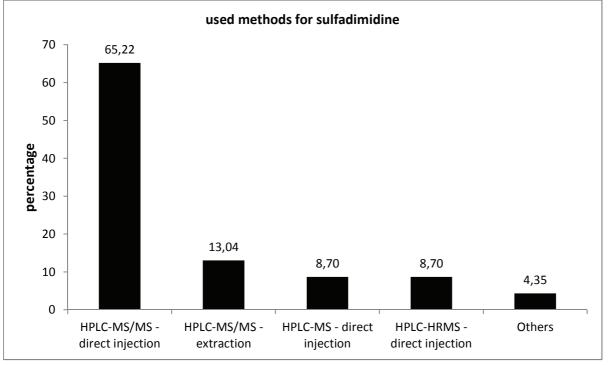


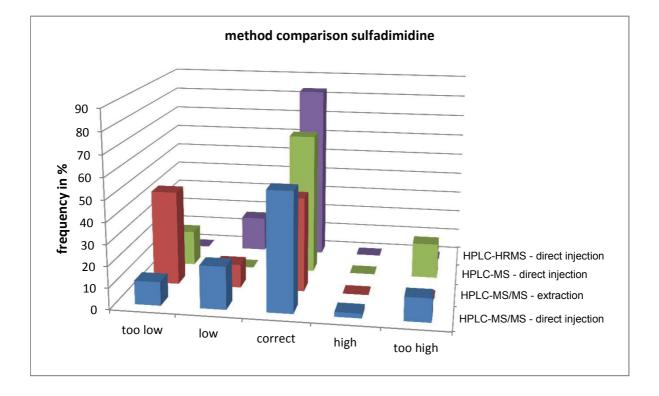
The relative standard deviations, calculated with the Q-method, did not reach the limits.



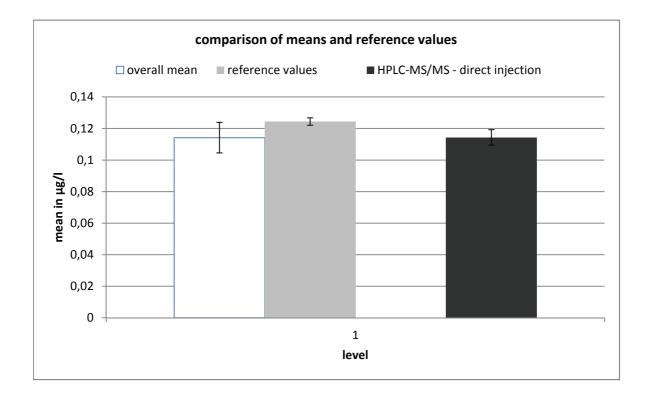
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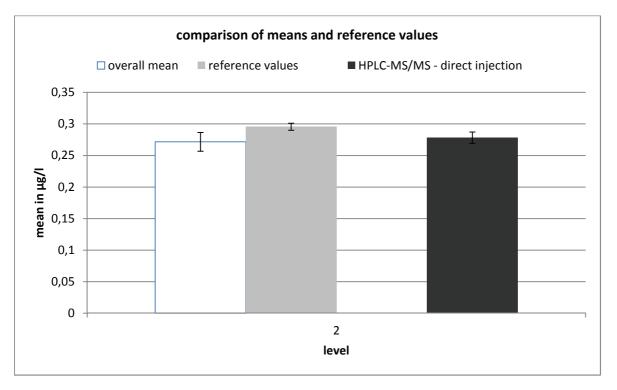
Method specific evaluation

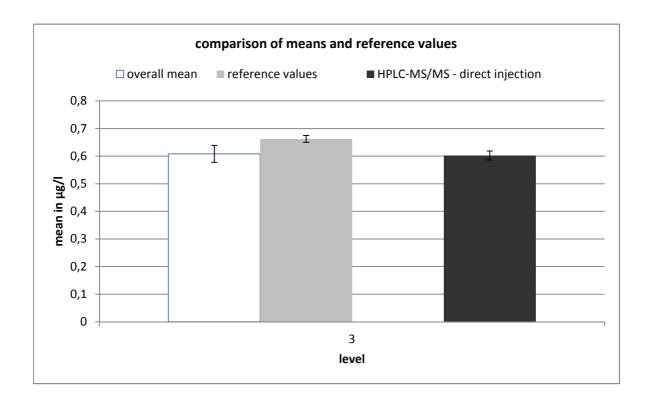


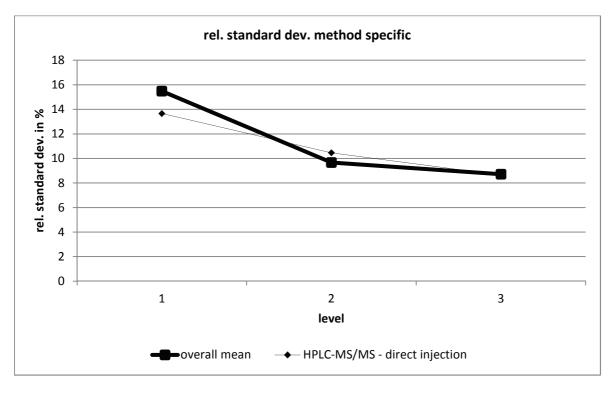


level	mean [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]	reference value [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]
1	0,1142	0,0097	8,5	0,1244	0,0023	1,9
2	0,2716	0,0148	5,4	0,2956	0,0056	1,9
3	0,6083	0,0307	5,0	0,6624	0,0126	1,9



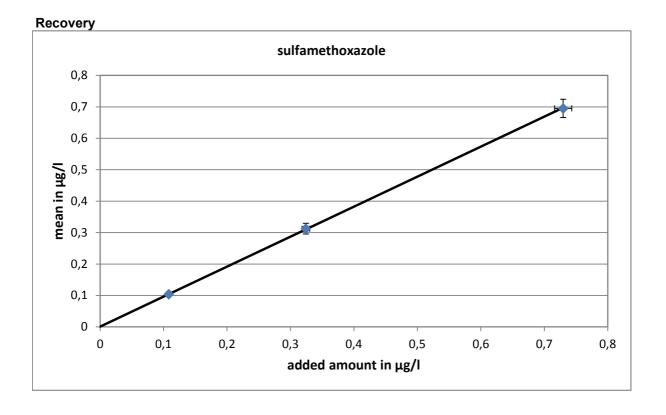






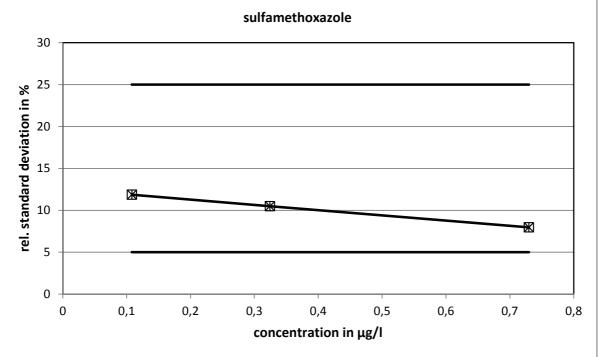
HPI	LC-MS/N	NS - dire	ect injec	tion	-		_		
level	robust mean [µg/l]	exp. unc. of the mean [µg/l]	exp. unc. of the mean [%]	robust standard deviation [µg/l	robust standard deviation [%]	number of results	out below	out above	out [%]
1	0,114	0,005	4,269	0,016	13,66	16	1	2	18,75
2	0,278	0,009	3,269	0,029	10,46	16	1	2	18,75
3	0,602	0,016	2,7	0,052	8,639	16	1	2	18,75

	sulfamethoxazole												
level	assigned value [µg/l]	expanded uncertainty of the assigned value [%]	standard deviation, calculated using robust statistics [µg/l]	standard deviation for proficiency assessment [µg/l]	standard deviation for proficiency assessment [%]	upper tolerance limit [µg/l]	lower tolerance limit [µg/l]	upper tolerance limit [%]	lower tolerance limit [%]	number of results	out below	out above	out [%]
1	0,1081	1,83	0,0128	0,0128	11,87	0,1356	0,0838	25,40	-22,45	25	0	1	4,0
2	0,3243	1,84	0,0340	0,0340	10,50	0,3965	0,2596	22,26	-19,97	25	0	0	0,0
3	0,7298	1,84	0,0582	0,0582	7,98	0,8514	0,6178	16,68	-15,34	25	2	1	12,0
									sum	75	2	2	5,3

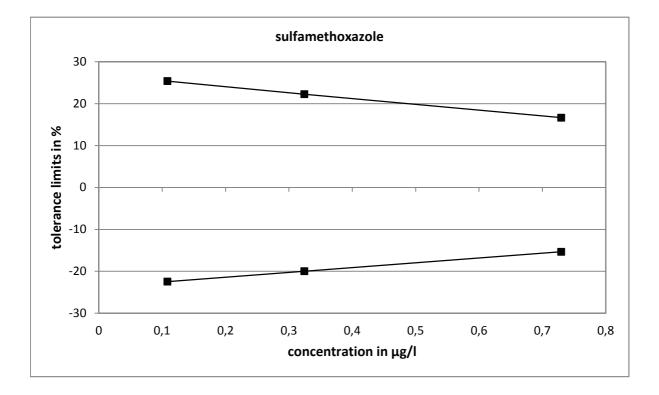


slope of the regression:0,955average recovery:95,5%



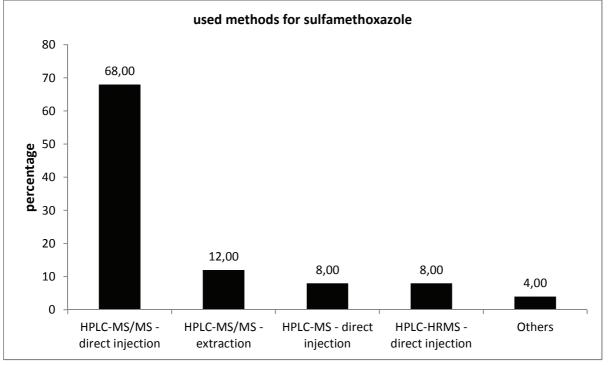


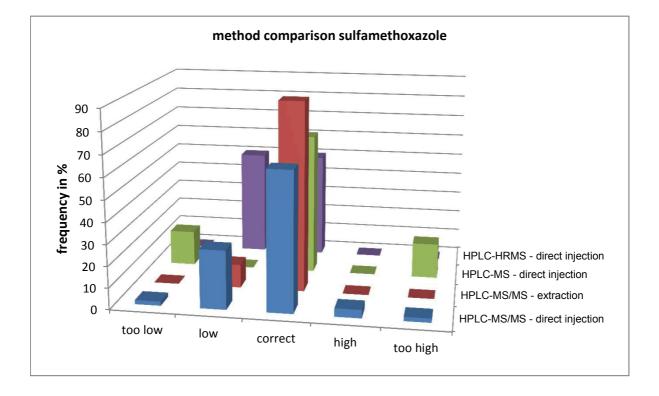
The relative standard deviations, calculated with the Q-method, did not reach the limits.



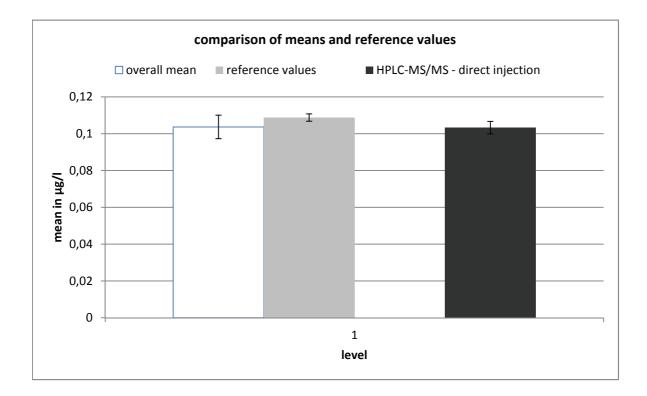
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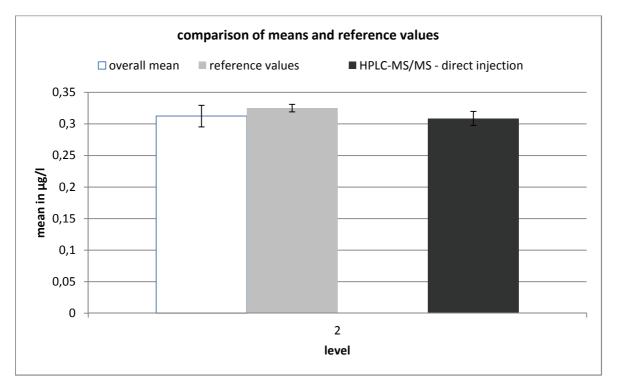
Method specific evaluation

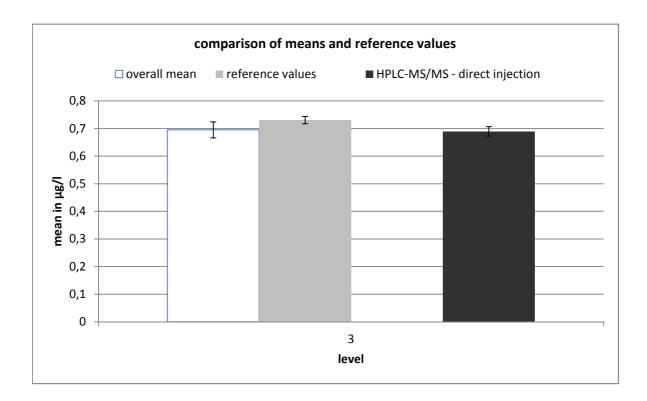


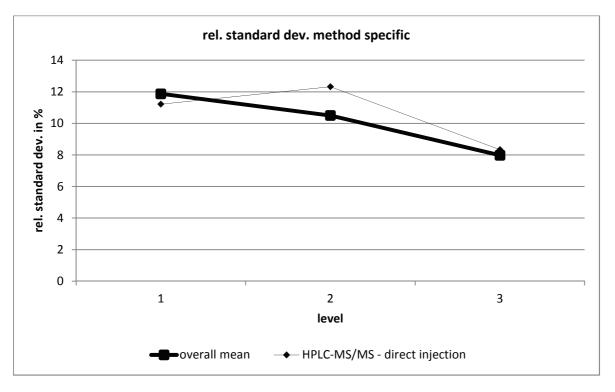


level	mean [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]	reference value [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]
1	0,1037	0,0064	6,2	0,1088	0,0020	1,8
2	0,3124	0,0170	5,4	0,3250	0,0060	1,8
3	0,6952	0,0291	4,2	0,7304	0,0135	1,8



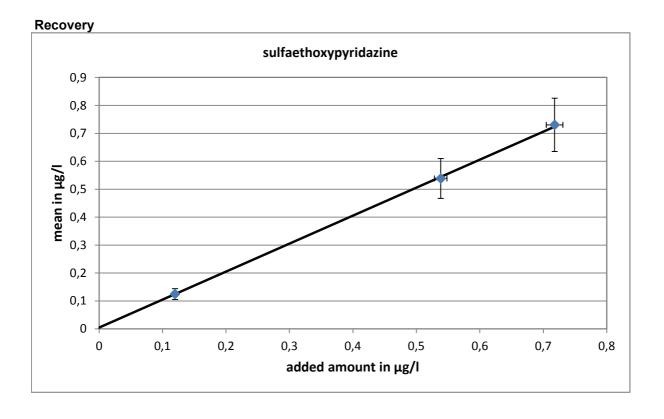






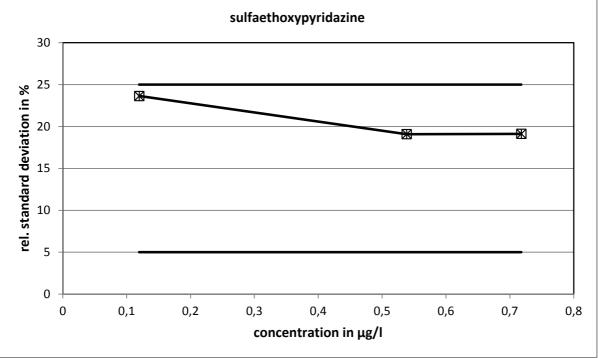
HPI	LC-MS/N	HPLC-MS/MS - direct injection											
level	robust mean [µg/l]	exp. unc. of the mean [µg/l]	exp. unc. of the mean [%]	robust standard deviation [µg/]	robust standard deviation [%]	number of results	out below	out above	out [%]				
1	0,103	0,003	3,306	0,012	11,22	18	1	0	5,556				
2	0,309	0,011	3,63	0,038	12,32	18	1	0	5,556				
3	0,69	0,017	2,456	0,057	8,337	18	1	1	11,11				

	sulfaethoxypyridazine												
level	assigned value [µg/l]	expanded uncertainty of the assigned value [%]	standard deviation, calculated using robust statistics [µg/l]	standard deviation for proficiency assessment [µg/l]	standard deviation for proficiency assessment [%]	upper tolerance limit [µg/l]	lower tolerance limit [µg/l]	upper tolerance limit [%]	lower tolerance limit [%]	number of results	out below	out above	out [%]
	0,1196	1,76	0,0283	0,0283	23,65	0,1843	0,0681	54,01	-43,06	13	0	1	7,7
2	0,5384	1,81	0,1028	0,1028	19,09	0,7676	0,3489	42,57	-35,20	13	1	0	7,7
3	0,7179	1,82	0,1373	0,1373	19,12	1,023	0,4645	42,57	-35,29	13	0	0	0,0
									sum	39	1	1	5,1

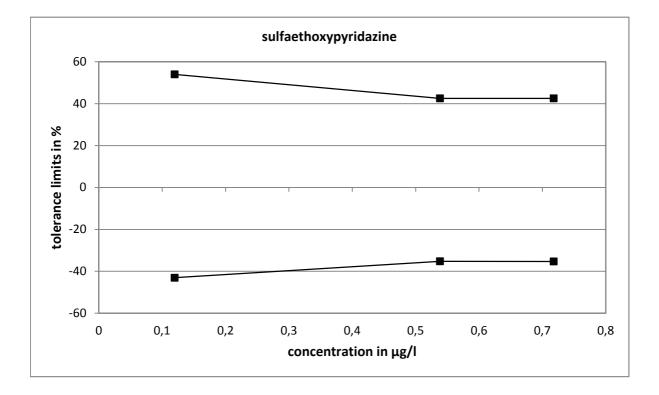


slope of the regression:0,1002average recovery:100,2%

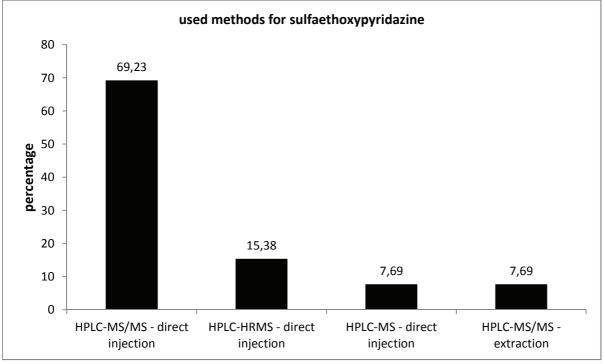


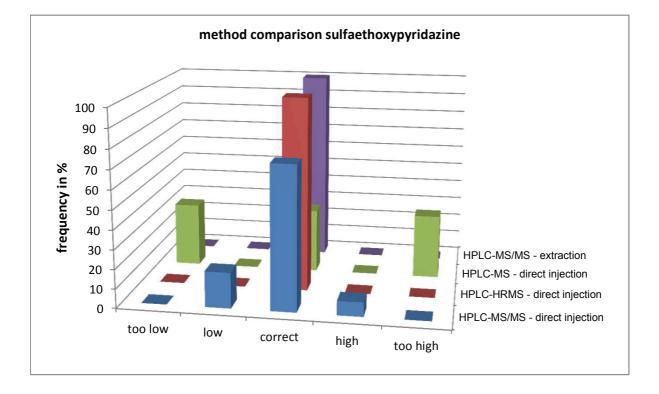


The relative standard deviations, calculated with the Q-method, did not reach the limits.

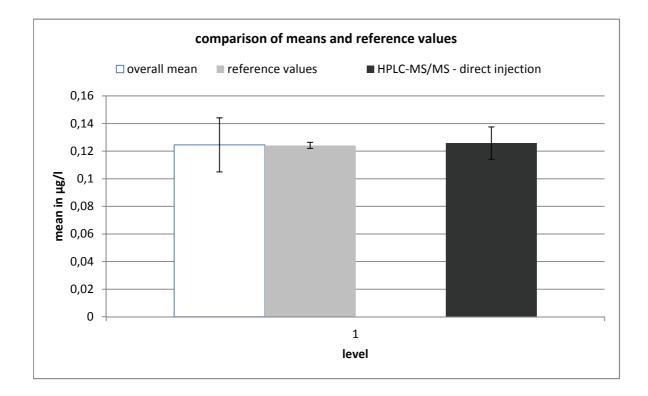


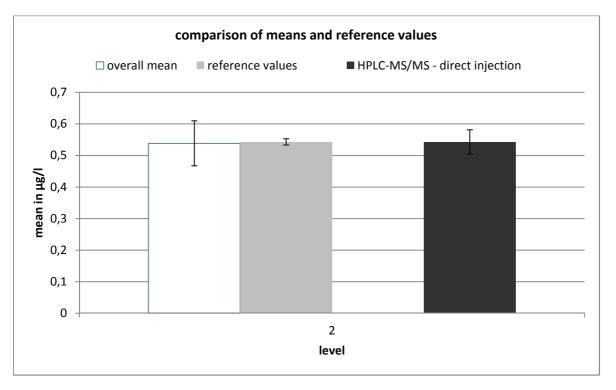
Method specific evaluation

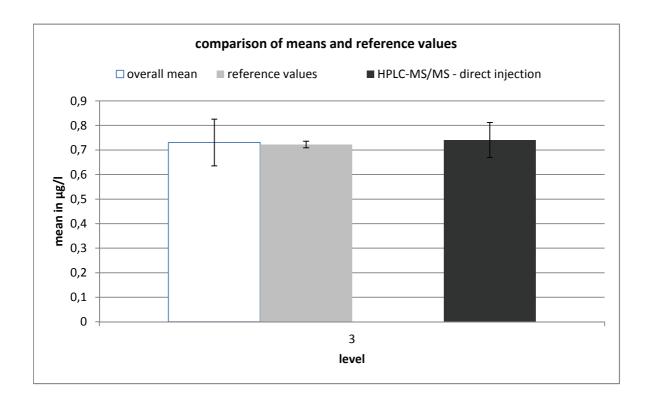


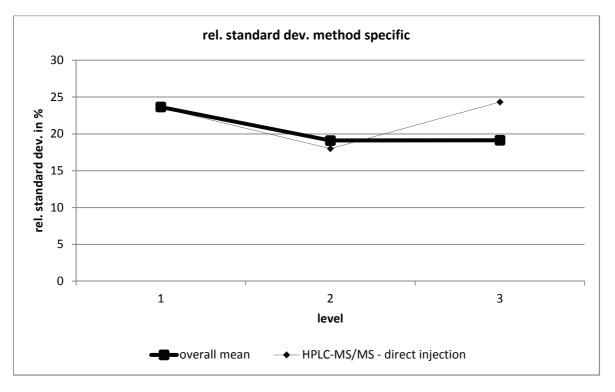


level	mean [µg/l]	exp. uncertainty [µg/l]	ୟ exp. uncertainty [%] ଝ	reference value [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]
1	0,1245	0,0196	15,8	0,1242	0,0022	1,8
2	0,5385	0,0713	13,2	0,5429	0,0099	1,8
3	0,7304	0,0952	13,0	0,7224	0,0131	1,8



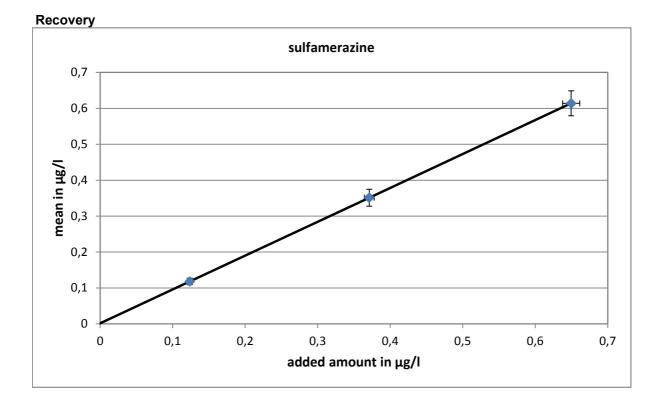






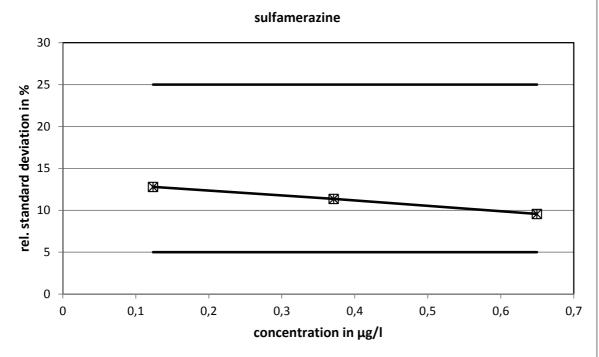
HPLC-MS/MS - direct injection									
level	robust mean [µg/l]	exp. unc. of the mean [µg/l]	exp. unc. of the mean [%]	robust standard deviation [µg/]	robust standard deviation [%]	number of results	out below	out above	out [%]
1	0,126	0,012	9,299	0,03	23,52	10	0	1	10
2	0,543	0,039	7,114	0,098	18	10	0	0	0
3	0,741	0,071	9,619	0,18	24,33	10	0	0	0

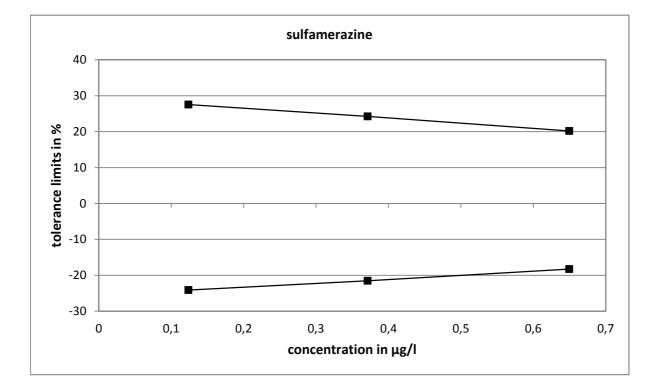
sulfamerazine													
level	assigned value [µg/l]	expanded uncertainty of the assigned value [%]	standard deviation, calculated using robust statistics [µg/l]	standard deviation for proficiency assessment [µg/l]	standard deviation for proficiency assessment [%]	upper tolerance limit [µg/l]	lower tolerance limit [µg/l]	upper tolerance limit [%]	lower tolerance limit [%]	number of results	out below	out above	out [%]
1	0,1237	1,80	0,0158	0,0158	12,80	0,1578	0,0939	27,55	-24,10	20	0	0	0,0
2	0,3712	1,81	0,0422	0,0422	11,36	0,4612	0,2913	24,25		20	2	1	15,0
3	0,6496	1,82	0,0621	0,0621	9,57	0,7808	0,5309	20,20	-18,26	20	2	1	15,0
									sum	60	4	2	10,0

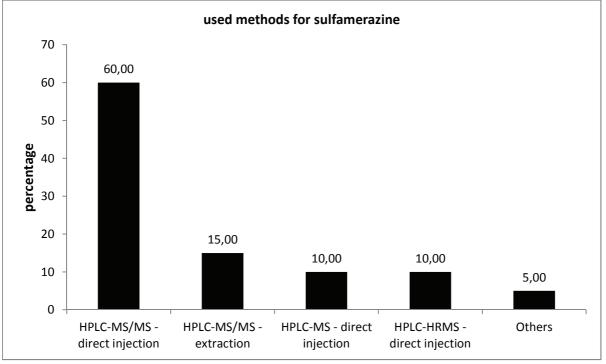


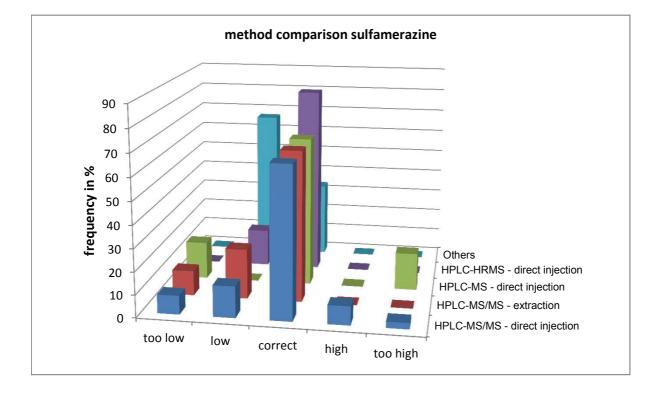
slope of the regression:0,943average recovery:94,3%



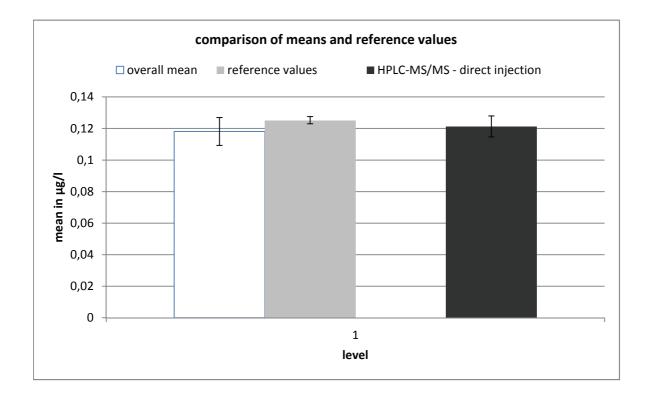


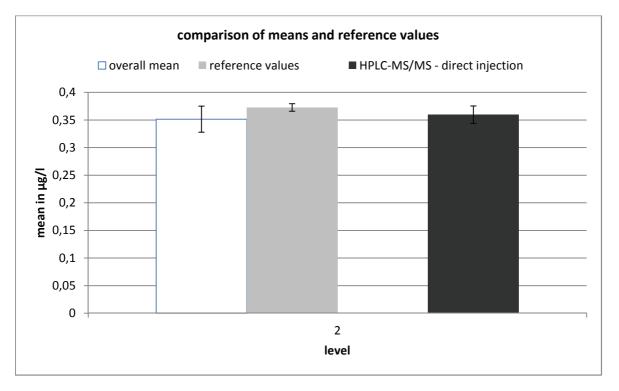


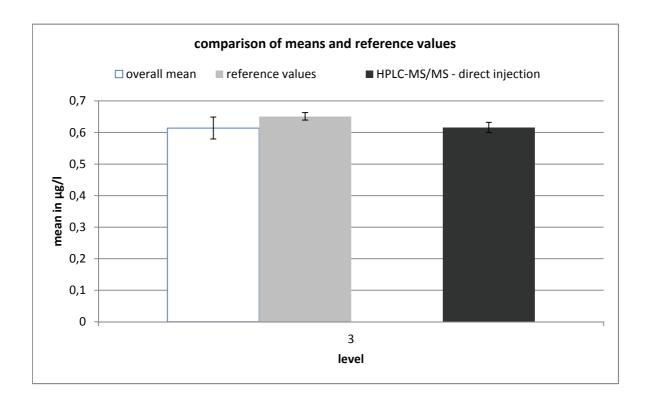


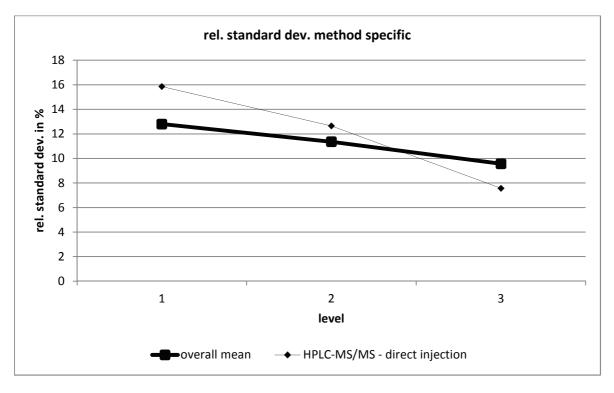


level	mean [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]	reference value [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]
1	0,1181	0,0089	7,5	0,1252	0,0023	1,8
2	0,3514	0,0236	6,7	0,3727	0,0068	1,8
3	0,6140	0,0347	5,7	0,6511	0,0118	1,8



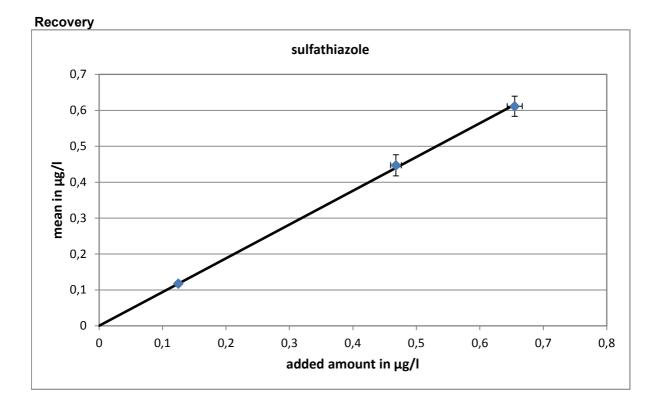






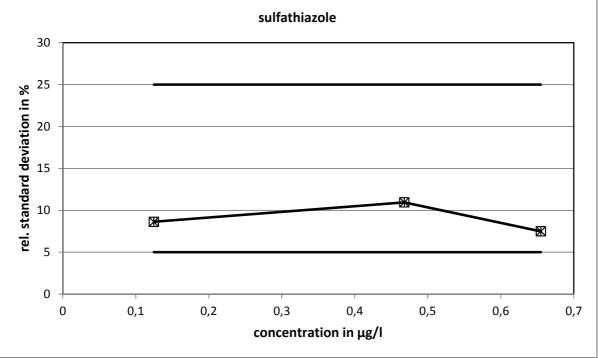
HPI	LC-MS/	NS - dire	ect injec	tion					
level	robust mean [µg/l]	exp. unc. of the mean [µg/l]	exp. unc. of the mean [%]	robust standard deviation [µg/l	robust standard deviation [%]	number of results	out below	out above	out [%]
1	0,121	0,007	5,497	0,019	15,86	13	0	1	7,692
2	0,36	0,016	4,386	0,045	12,65	13	0	1	7,692
3	0,616	0,016	2,624	0,047	7,569	13	1	1	15,38

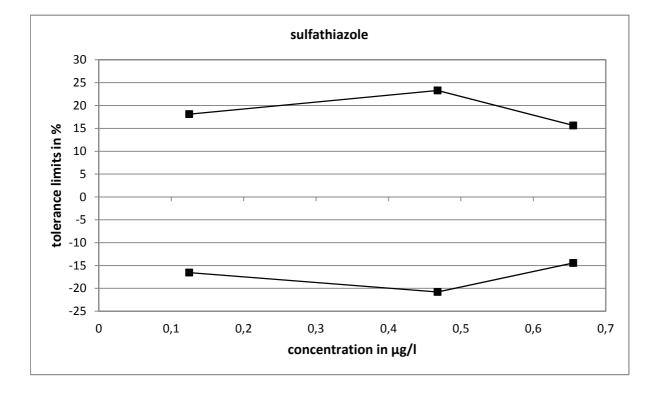
	sulfathiazole												
level	assigned value [µg/l]	expanded uncertainty of the assigned value [%]	standard deviation, calculated using robust statistics [µg/l]	standard deviation for proficiency assessment [µg/l]	standard deviation for proficiency assessment [%]	upper tolerance limit [µg/l]	lower tolerance limit [µg/l]	upper tolerance limit [%]	lower tolerance limit [%]	number of results	out below	out above	out [%]
1	0,1248	1,79	0,0108	0,0108	8,63	0,1474	0,1041	18,12	-16,54	19	0	2	10,5
2	0,4678	1,79	0,0512	0,0512	10,95	0,5768	0,3706	23,30	-20,78	19	0	2	10,5
3	0,6550	1,79	0,0491	0,0491	7,50	0,7574	0,5604	15,64	-14,44	19	1	1	10,5
									sum	57	1	5	10,5

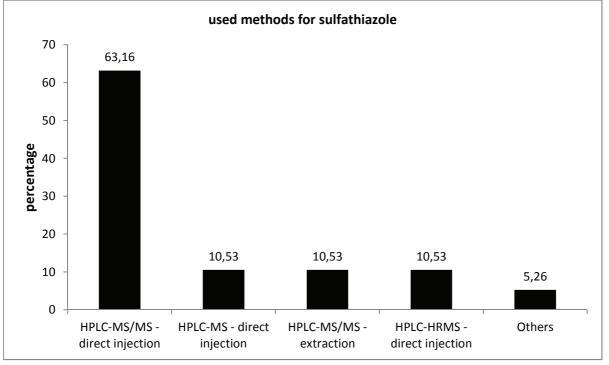


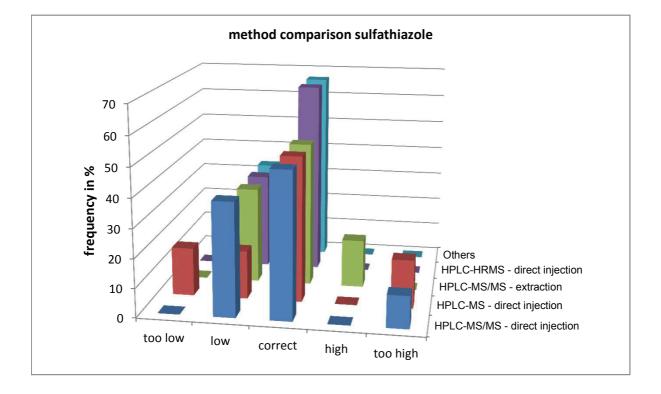
slope of the regression:0,94average recovery:94,0%



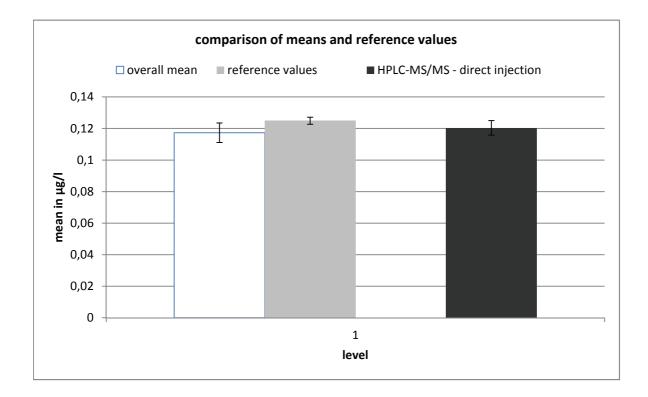


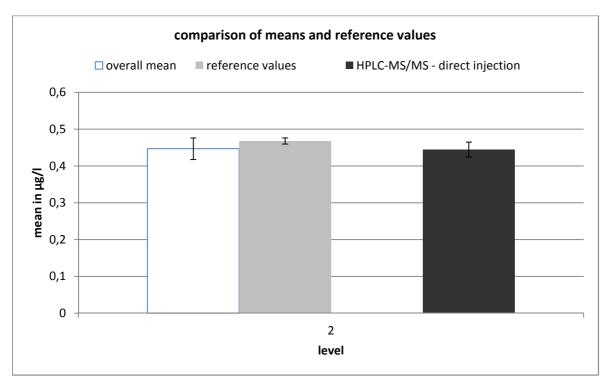


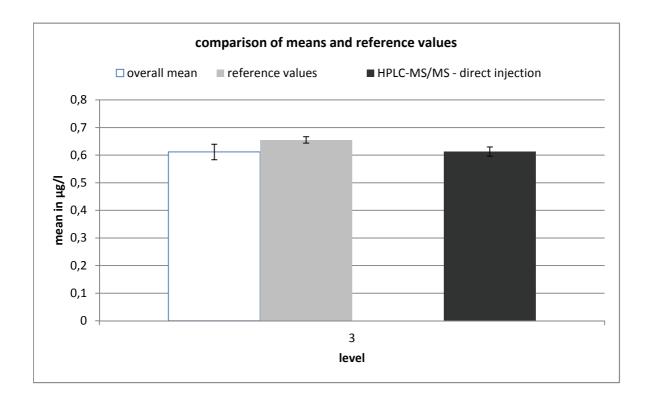


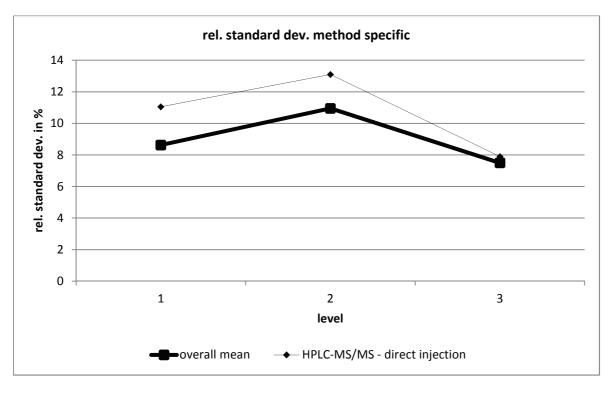


level	mean [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]	reference value [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]
1	0,1173	0,0062	5,3	0,1249	0,0022	1,8
2	0,4470	0,0294	6,6	0,4680	0,0084	1,8
3	0,6114	0,0282	4,6	0,6551	0,0117	1,8



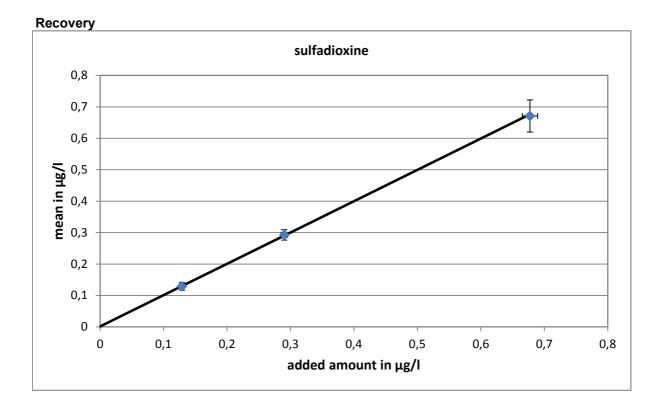






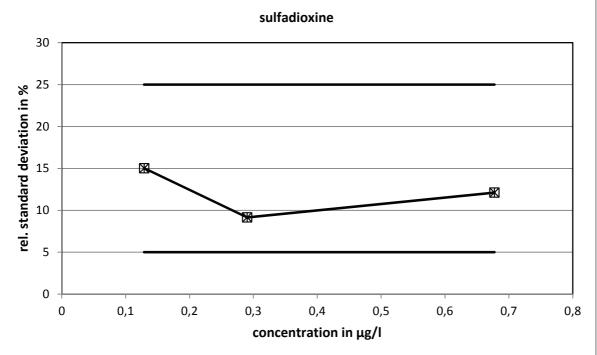
HPI	LC-MS/	NS - dire	ect injec	tion					
level	robust mean [µg/l]	exp. unc. of the mean [µg/l]	exp. unc. of the mean [%]	robust standard deviation [µg/l	robust standard deviation [%]	number of results	out below	out above	out [%]
	0,12	0,005	3,831	0,013	11,05	13	0	3	23,08
2	0,445	0,02	4,542	0,058	13,1	13	0	2	15,38
3	0,613	0,017	2,737	0,048	7,895	13	0	1	7,692

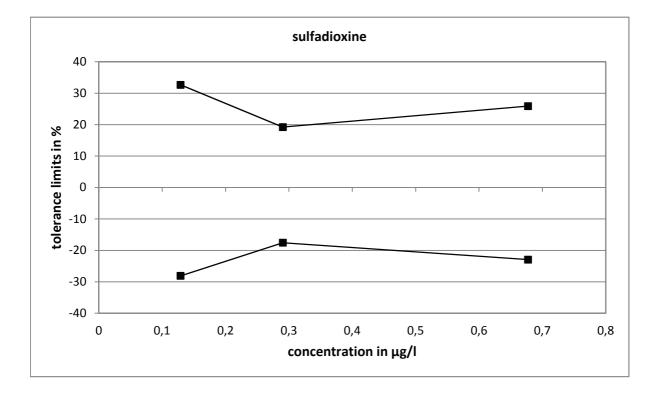
	sulfadioxine												
level	assigned value [µg/l]	expanded uncertainty of the assigned value [%]	standard deviation, calculated using robust statistics [µg/l]	standard deviation for proficiency assessment [µg/l]	standard deviation for proficiency assessment [%]	upper tolerance limit [µg/l]	lower tolerance limit [µg/l]	upper tolerance limit [%]	lower tolerance limit [%]	number of results	out below	out above	out [%]
1	0,1290	1,77	0,0194	0,0194	15,03	0,1712	0,0927	32,66	-28,12	16	0	1	6,3
2	0,2903	1,78	0,0266	0,0266	9,16	0,3461	0,2393	19,22	-17,56	16	0	2	12,5
3	0,6774	1,78	0,0821	0,0821	12,12	0,8529	0,5220	25,91	-22,94	16	1	1	12,5
									sum	48	1	4	10,4

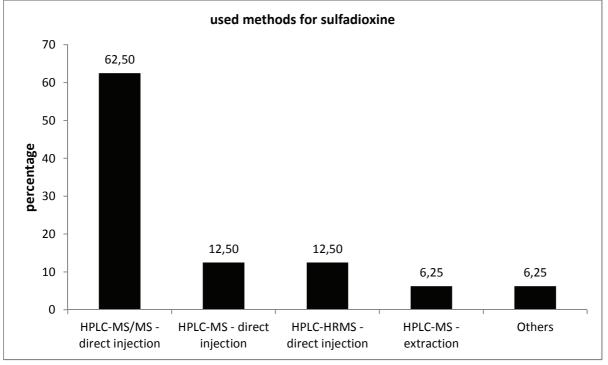


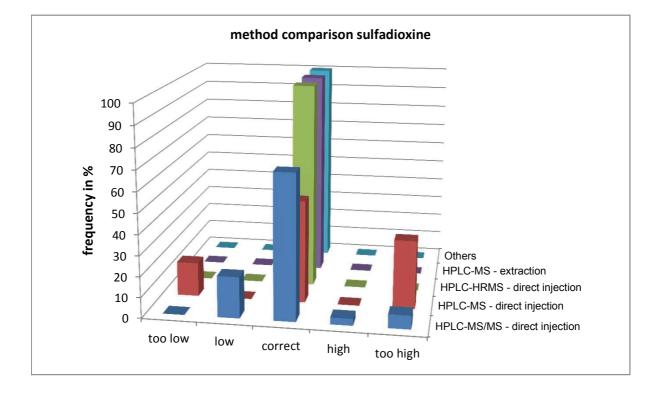
slope of the regression:0,997average recovery:99,7%



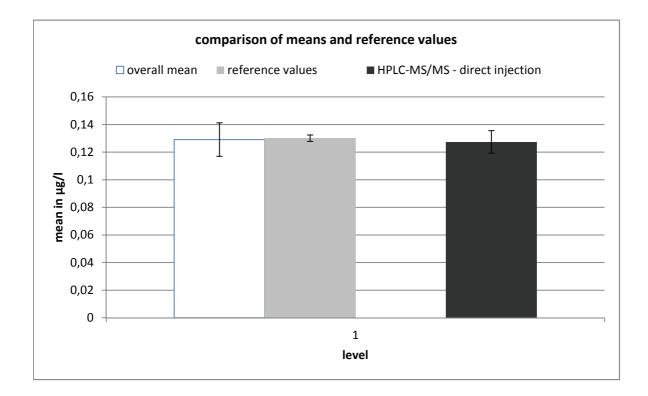


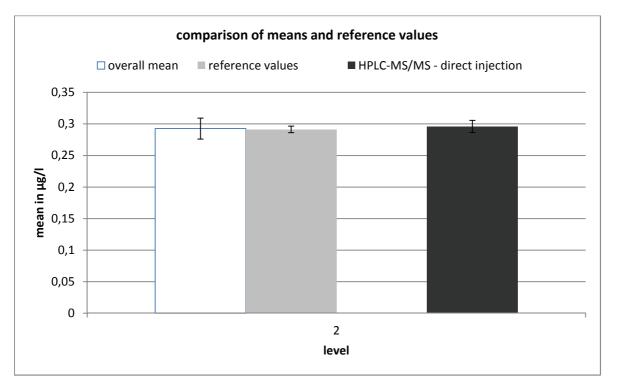


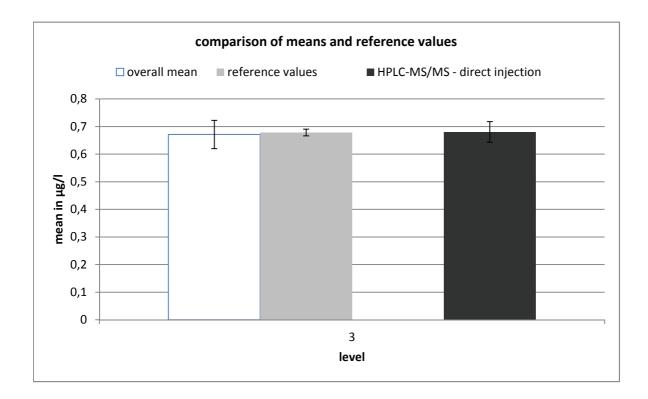


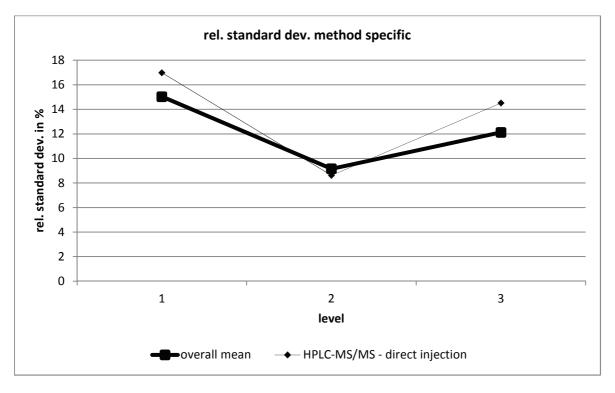


level	mean [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]	reference value [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]
1	0,1291	0,0121	9,4	0,1301	0,0023	1,8
2	0,2926	0,0166	5,7	0,2914	0,0052	1,8
3	0,6712	0,0513	7,6	0,6785	0,0121	1,8



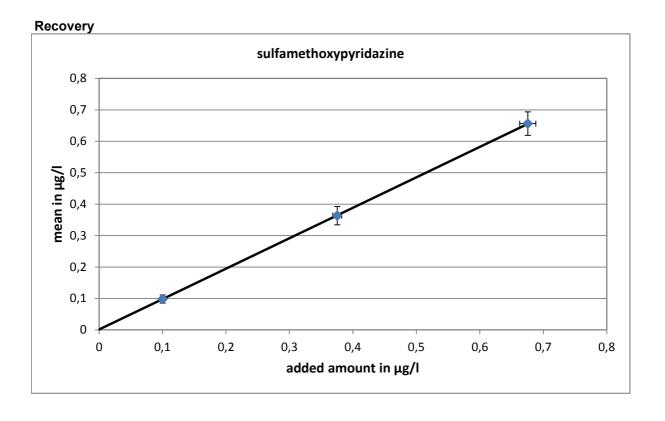






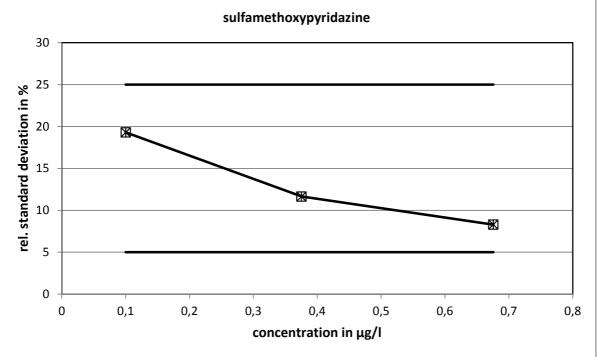
HPI	LC-MS/N	NS - dire	ect injec	tion					
level	robust mean [µg/l]	exp. unc. of the mean [µg/l]	exp. unc. of the mean [%]	robust standard deviation [µg/]	robust standard deviation [%]	number of results	out below	out above	out [%]
1	0,127	0,008	6,401	0,022	16,98	11	0	1	9,091
2	0,296	0,01	3,248	0,026	8,619	11	0	2	18,18
3	0,68	0,037	5,47	0,099	14,51	11	0	1	9,091

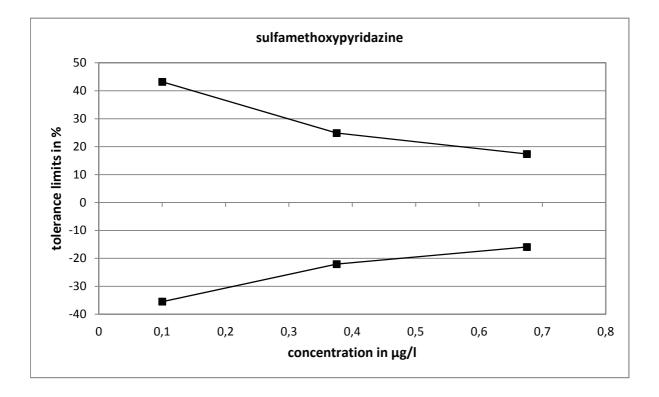
	sulfamethoxypyridazine												
level	assigned value [µg/l]	expanded uncertainty of the assigned value [%]	standard deviation, calculated using robust statistics [µg/l]	standard deviation for proficiency assessment [µg/l]	standard deviation for proficiency assessment [%]	upper tolerance limit [µg/l]	lower tolerance limit [µg/l]	upper tolerance limit [%]	lower tolerance limit [%]	number of results	out below	out above	14 56 out [%]
1	0,1001 0,3753	1,86 1,87	0,0193 0,0437	0,0193 0,0437	19,30 11,65	0,1433 0,4687	0,0646 0,2925	43,19 24,87	-35,50 -22,06	14 14	0	2	14,3 14,3
3	0,6756	1,88	0,0437	0,0437	8,29	0,7928		17,36	-15,94	14	0	1	7,1
	0,0700	1,00	0,0000	0,0000	0,20	0,1020	0,0070	11,00	sum	42	1	4	11,9

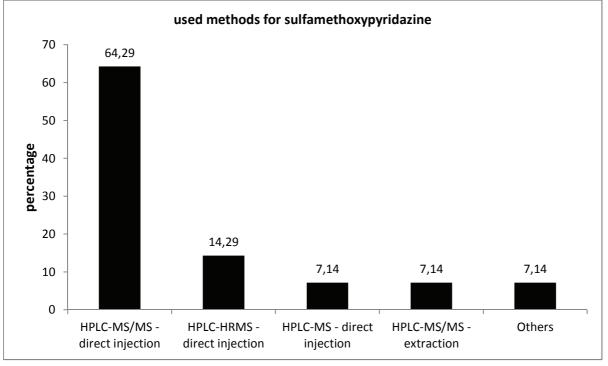


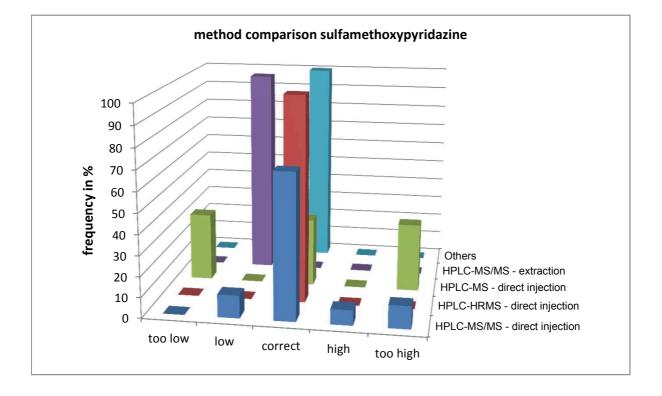
slope of the regression:0,969average recovery:96,9%



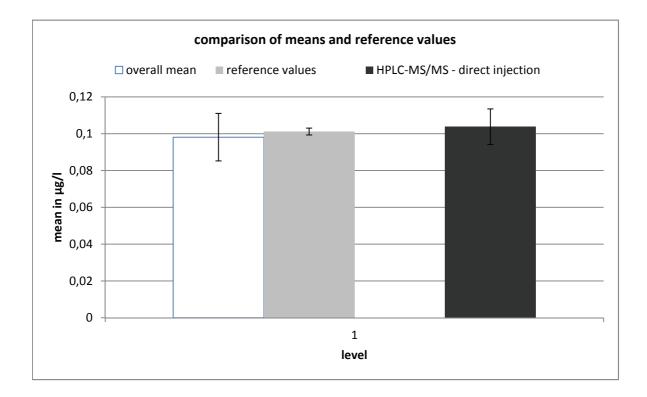


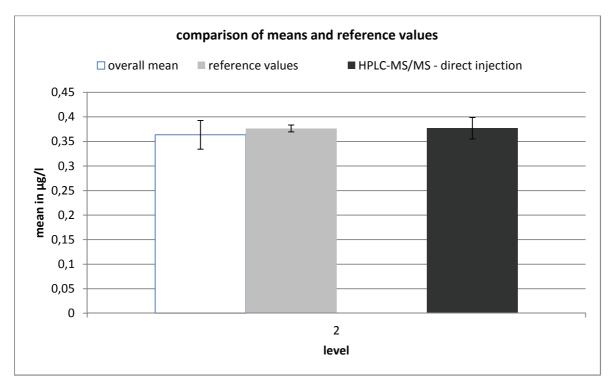


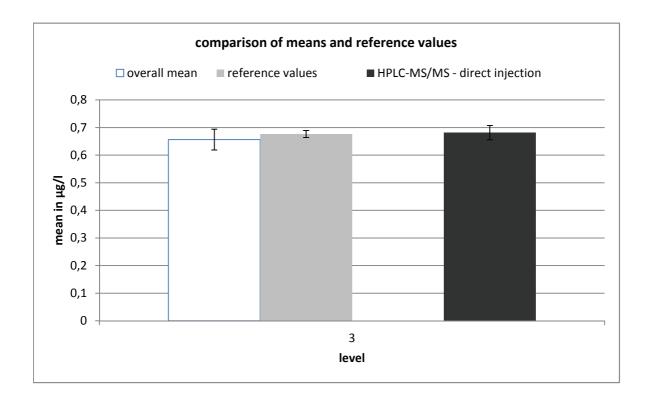


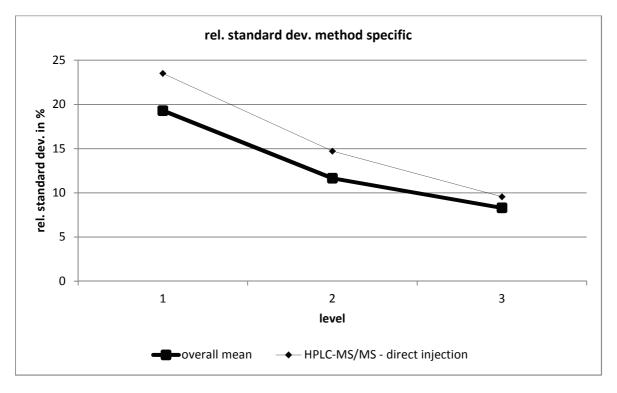


ے اevel	g mean [µg/l]	c exp. uncertainty [µg/l]	င္တဲ့ exp. uncertainty [%] လ	5 reference value [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]
1	0,0981	0,0129	13,2	0,1012	0,0019	1,9
2	0,3636	0,0292	8,0	0,3764	0,0071	1,9
3	0,6564	0,0374	5,7	0,6767	0,0127	1,9



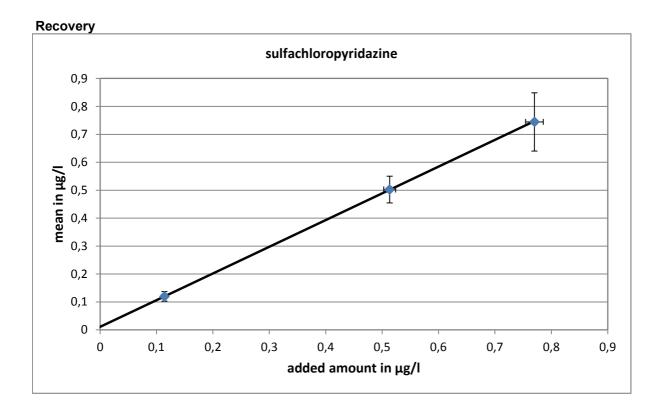






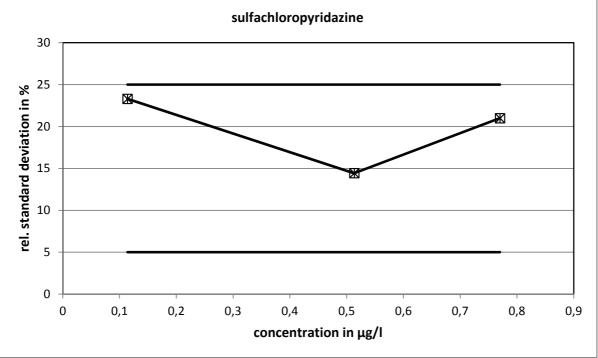
HPI	HPLC-MS/MS - direct injection											
level	robust mean [µg/l]	exp. unc. of the mean [µg/l]	exp. unc. of the mean [%]	robust standard deviation [µg/]	robust standard deviation [%]	number of results	out below	out above	out [%]			
1	0,104	0,01	9,295	0,024	23,51	10	0	1	10			
2	0,377	0,022	5,817	0,055	14,71	10	0	1	10			
3	0,681	0,026	3,776	0,065	9,552	10	0	1	10			

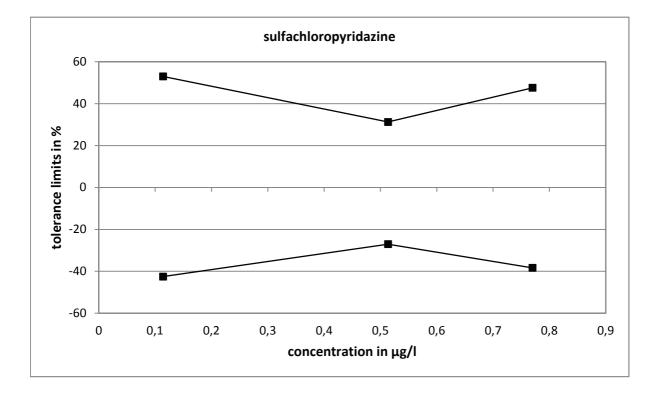
	sulfachloropyridazine												
level	assigned value [µg/l]	expanded uncertainty of the assigned value [%]	standard deviation, calculated using robust statistics [µg/l]	standard deviation for proficiency assessment [µg/l]	standard deviation for proficiency assessment [%]	upper tolerance limit [µg/l]	lower tolerance limit [µg/l]	upper tolerance limit [%]	lower tolerance limit [%]	number of results	out below	out above	۲۵ 1%] 1%]
1	0,1141	1,84	0,0266	0,0266	23,30	0,1746	0,0656	53,02	-42,50	15	0	2	13,3
2	0,5134	1,98	0,0741	0,0741	14,43	0,6741	0,3746	31,31	-27,03	15	0	1	6,7
3	0,7701	1,99	0,1616	0,1616	20,99	1,137	0,4748	47,60	-38,34	15	1	1	13,3
									sum	45	1	4	11,1

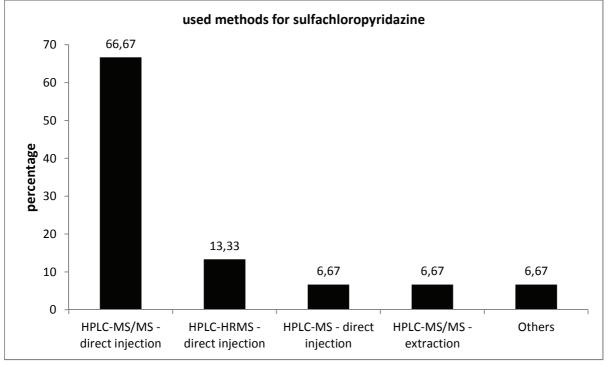


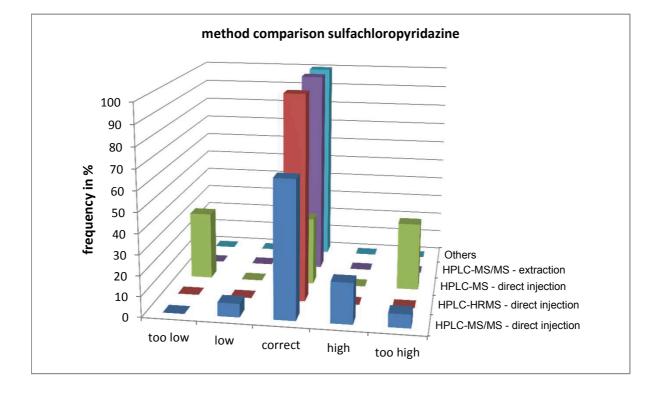
slope of the regression:0,956average recovery:95,6%



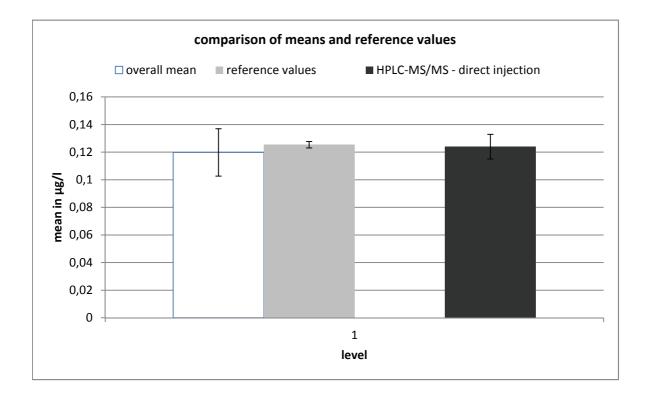


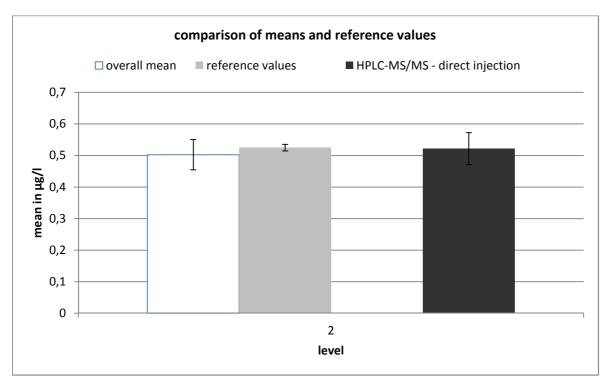


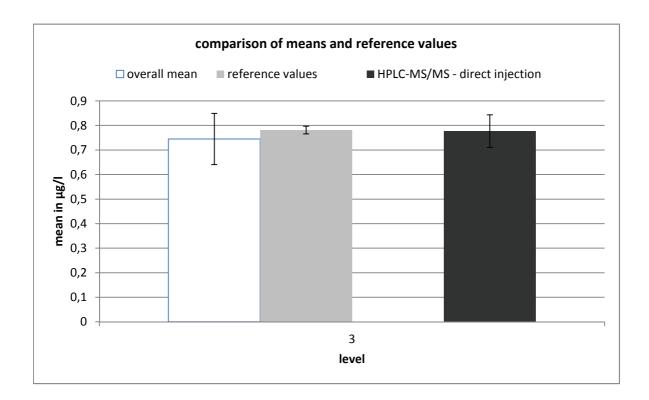


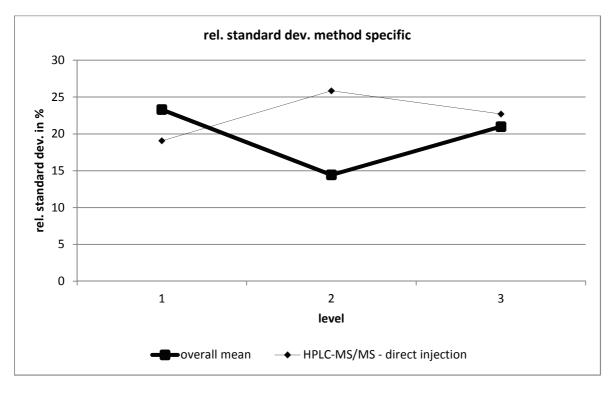


Comparison of means and reference values



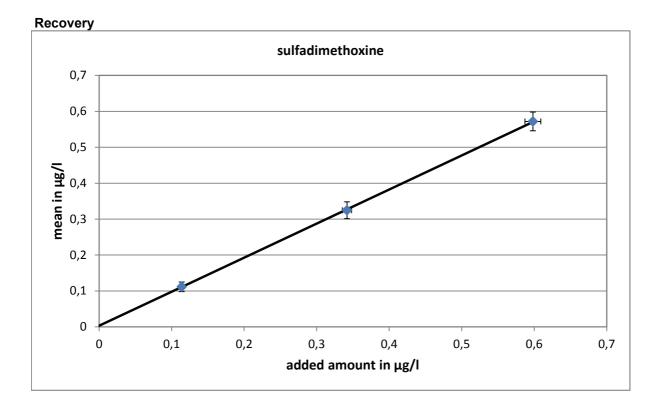






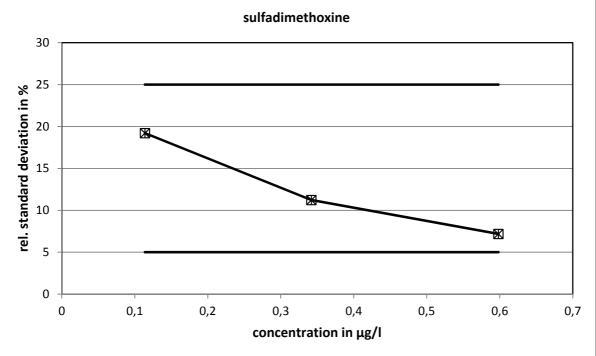
HPI	LC-MS/N	NS - dire	ect injec	tion					
level	robust mean [µg/l]	exp. unc. of the mean [µg/l]	exp. unc. of the mean [%]	robust standard deviation [µg/l	robust standard deviation [%]	number of results	out below	out above	out [%]
	0,124	0,009	7,184	0,024	19,06	11	0	2	18,18
2	0,521	0,051	9,744	0,135	25,85	11	0	0	0
3	0,777	0,066	8,558	0,176	22,71	11	0	0	0

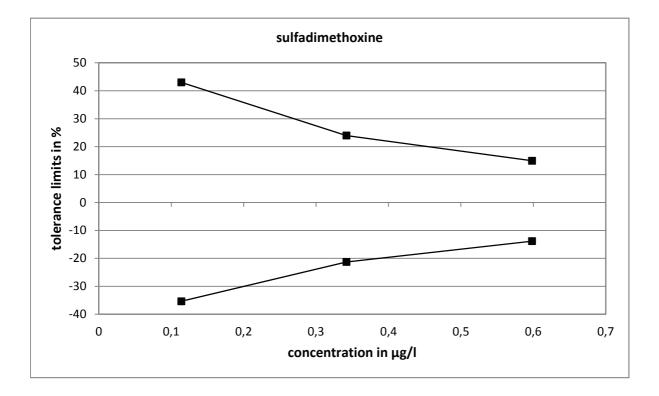
	sulfadimethoxine												
level	assigned value [µg/l]	expanded uncertainty of the assigned value [%]	standard deviation, calculated using robust statistics [µg/l]	standard deviation for proficiency assessment [µg/l]	standard deviation for proficiency assessment [%]	upper tolerance limit [µg/l]	lower tolerance limit [µg/l]	upper tolerance limit [%]	lower tolerance limit [%]	number of results	out below	out above	9) out [%]
1	0,1139	1,78	0,0219	0,0219	19,22	0,1629	0,0737	43,00	-35,36	17	1	2	17,6
2	0,3418	1,81	0,0384	0,0384	11,23	0,4237	0,2690	23,96	-21,29	17	3	0	17,6
3	0,5982	1,82	0,0429	0,0429	7,18	0,6875	0,5153	14,94	-13,86	17	3	1	23,5
									sum	51	7	3	19,6

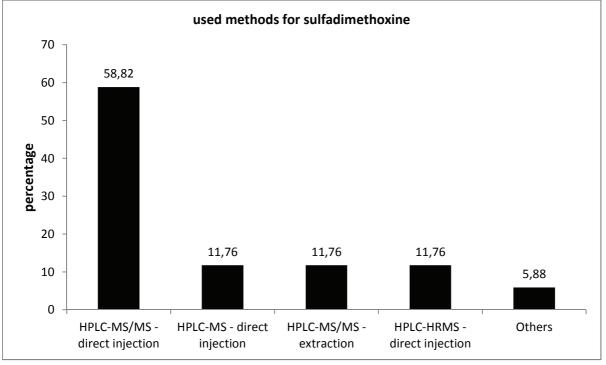


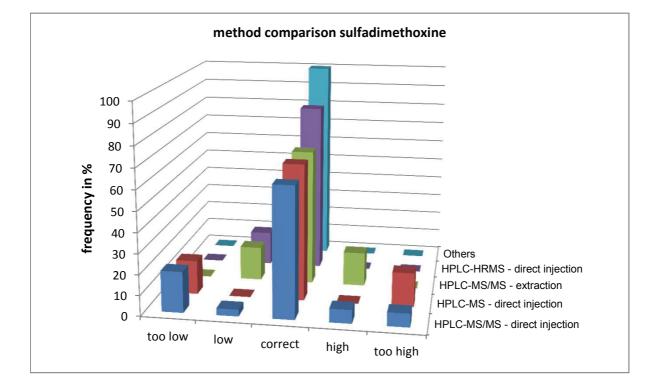
slope of the regression:0,949average recovery:94,9%



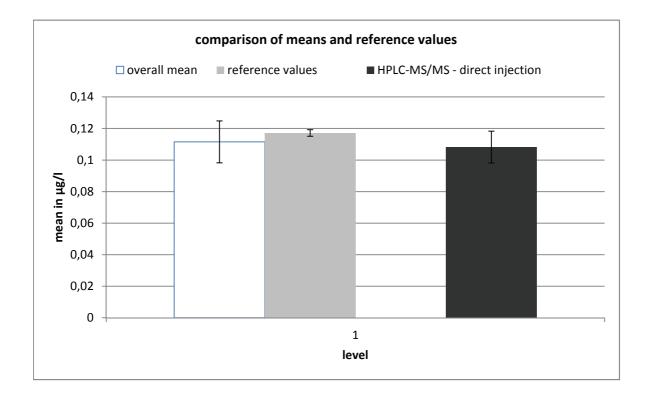


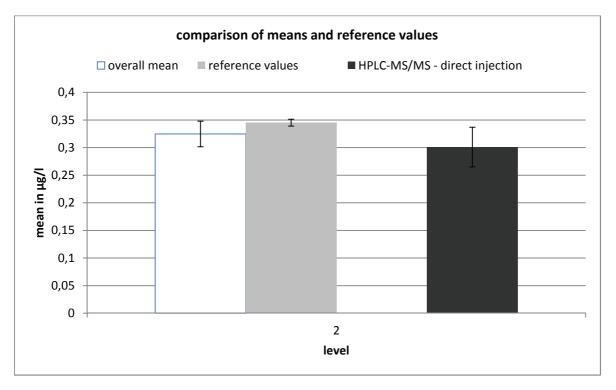


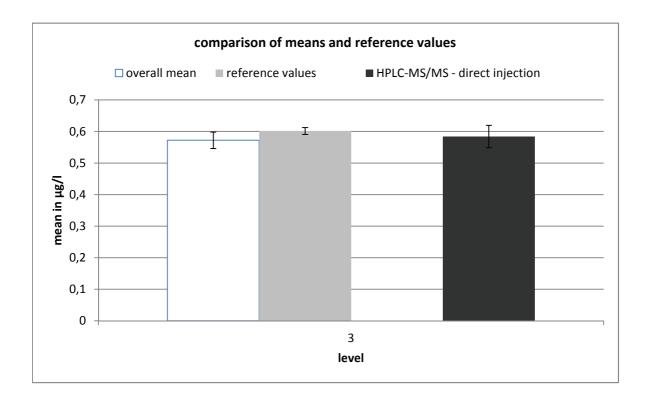


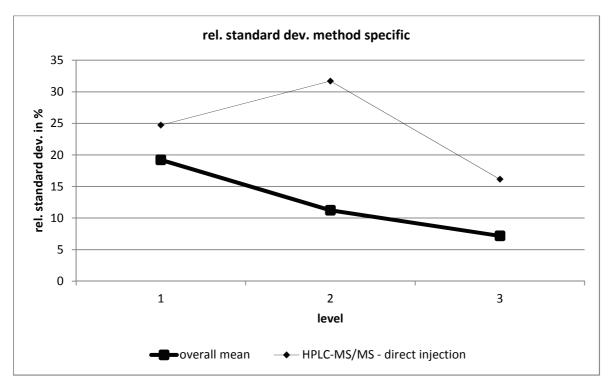


level	mean [µg/l]	exp. uncertainty [µg/l]	는 exp. uncertainty [%] 	reference value [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]
1	0,1115	0,0133	11,9	0,1171	0,0021	1,8
2	0,3247	0,0233	7,2	0,3450	0,0062	1,8
3	0,5720	0,0260	4,6	0,6014	0,0109	1,8









HPI	HPLC-MS/MS - direct injection											
level	robust mean [µg/l]	exp. unc. of the mean [µg/l]	exp. unc. of the mean [%]	robust standard deviation [µg/l	robust standard deviation [%]	number of results	out below	out above	out [%]			
1	0,108	0,01	9,315	0,027	24,72	11	0	2	18,18			
2	0,301	0,036	11,95	0,095	31,7	11	1	1	18,18			
3	0,584	0,036	6,088	0,094	16,15	11	1	0	9,091			

evel

1 2

3

0,3793

0,6827

1,81

1,86

1,87

0,0385

0,0788 0,0788

0,0385

trimethoprime astandard deviation for proficiency assessment [%] assessment [%] assessment [%] by the second sec © © standard deviation for proficiency © assessment [µg/l] ອ B standard deviation, calculated ຝ using robust statistics [µg/l] expanded uncertainty of the upper tolerance limit [%] lower tolerance limit [%] 0.0 1101 10,10 11 assigned value [µg/l] assigned value [%] assessment [µg/l] Number of results A out above out below out [%]

10,14 0,4605

11,54 0,8509 0,5335

-18,70

-19,34

-21,86

sum

24

24

72

0

2 2 16,7

8 2

8,3

16,7

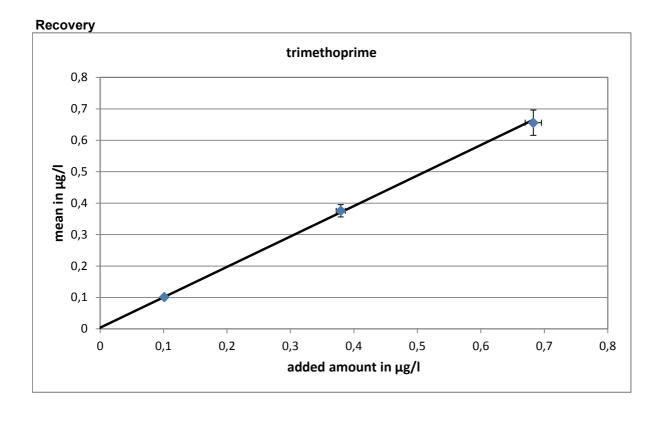
13,9

20,61

21,42

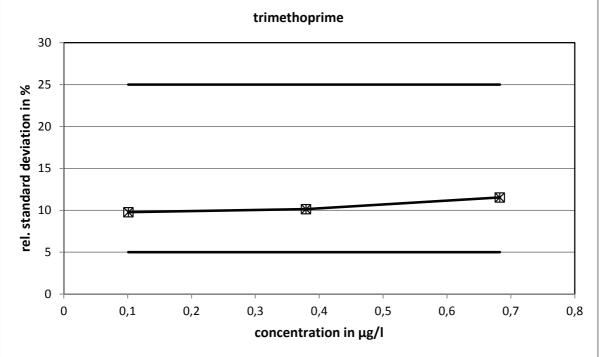
24,64

0,3059

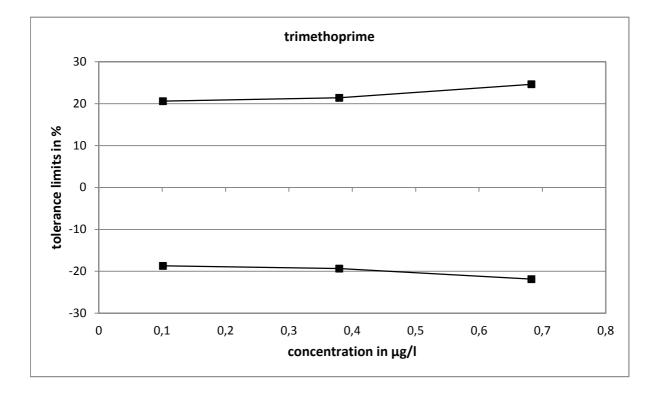


slope of the regression: 0,969 average recovery: 96,9%

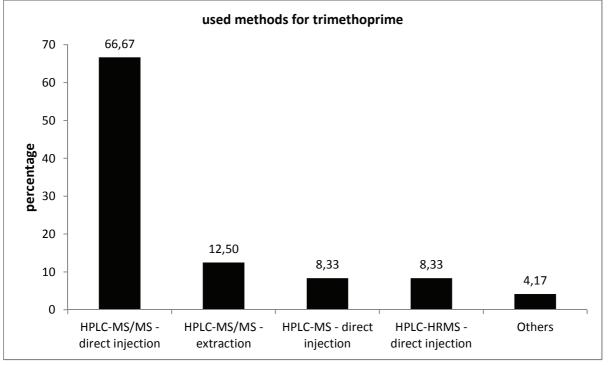


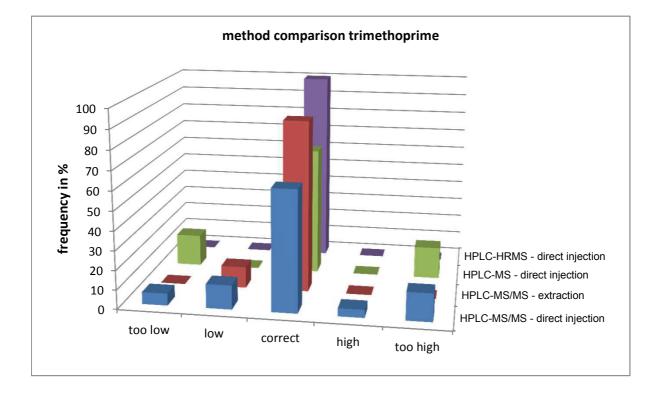


The relative standard deviations, calculated with the Q-method, did not reach the limits.

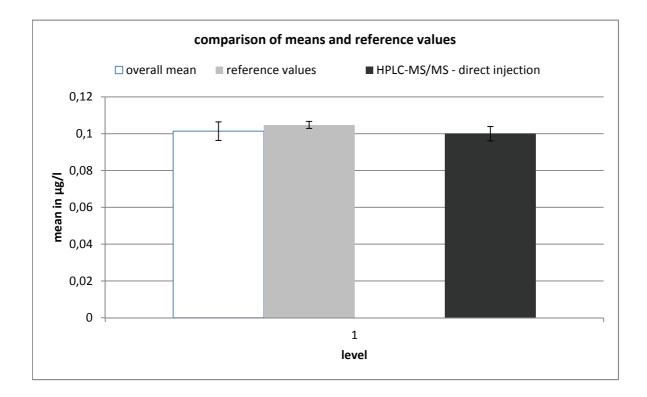


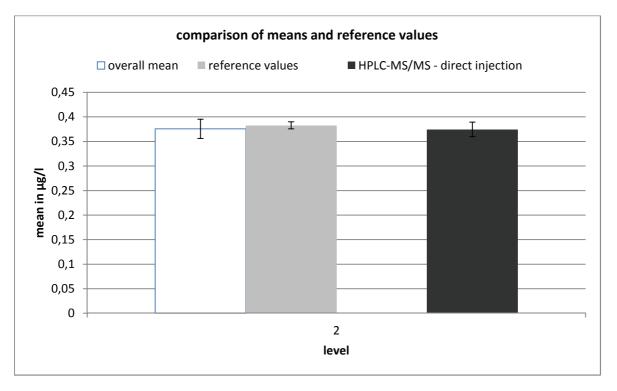
Method specific evaluation

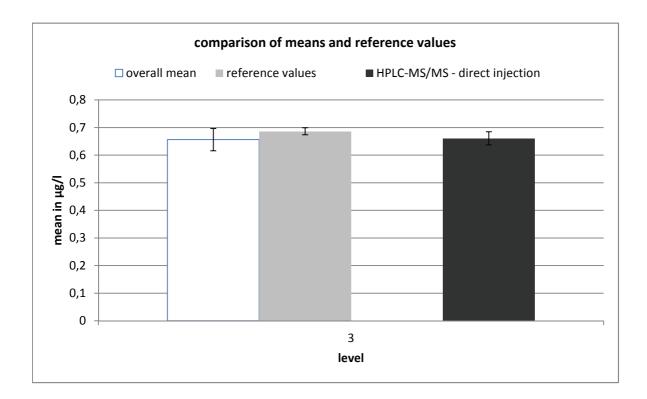


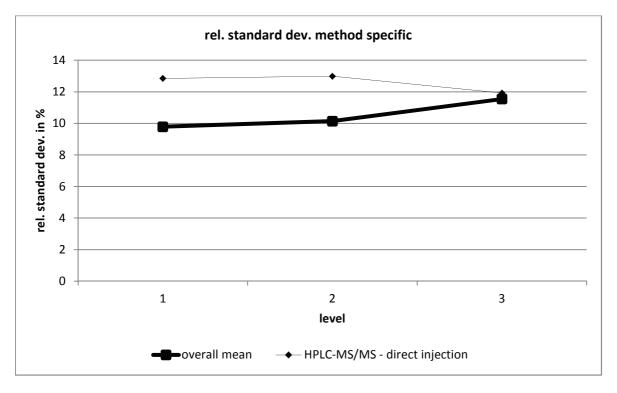


level	mean [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]	reference value [µg/l]	exp. uncertainty [µg/l]	exp. uncertainty [%]
1	0,1014	0,0050	5,0	0,1048	0,0019	1,8
2	0,3757	0,0196	5,2	0,3829	0,0071	1,9
3	0,6561	0,0402	6,1	0,6863	0,0128	1,9

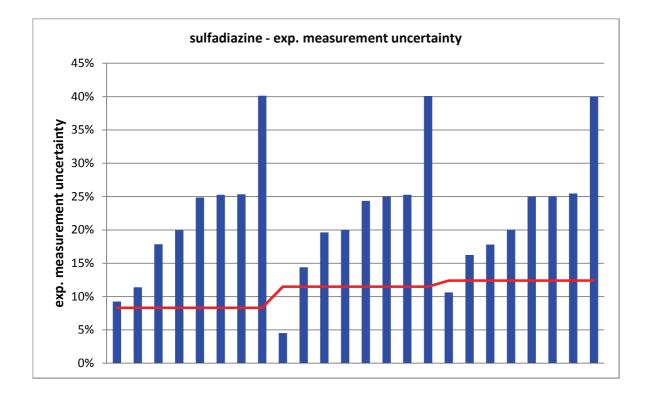


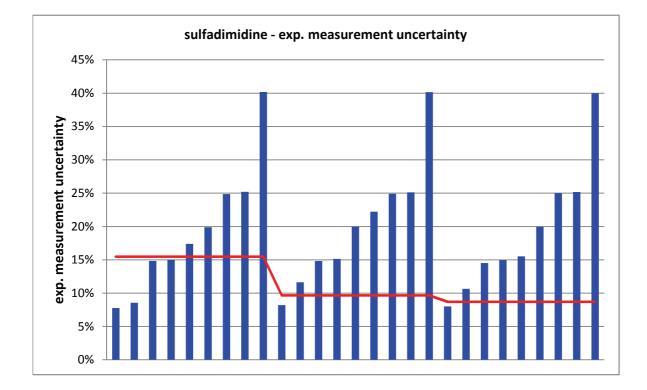


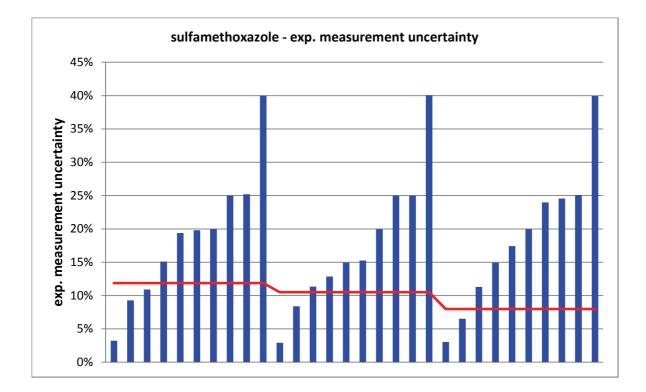


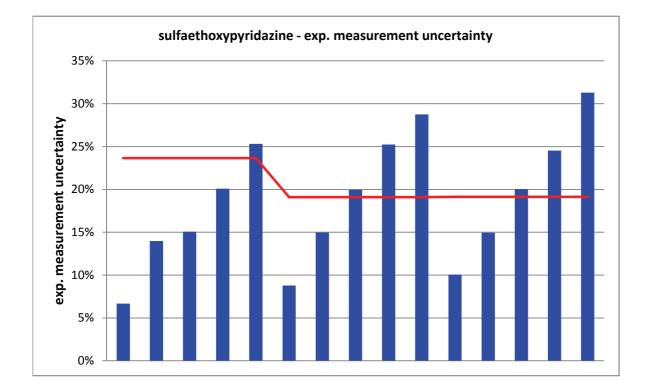


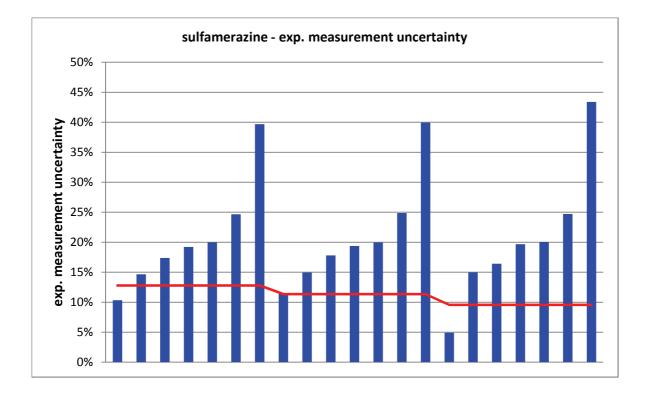
HPI	HPLC-MS/MS - direct injection								
level	robust mean [µg/l]	exp. unc. of the mean [µg/l]	exp. unc. of the mean [%]	robust standard deviation [µg/l	robust standard deviation [%]	number of results	out below	out above	out [%]
1	0,1	0,004	3,897	0,013	12,85	17	0	3	17,65
2	0,375	0,015	3,937	0,049	12,99	17	0	3	17,65
3	0,661	0,024	3,614	0,079	11,92	17	1	2	17,65

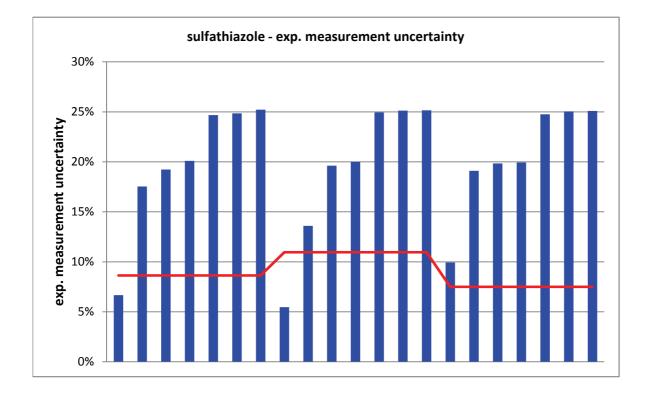


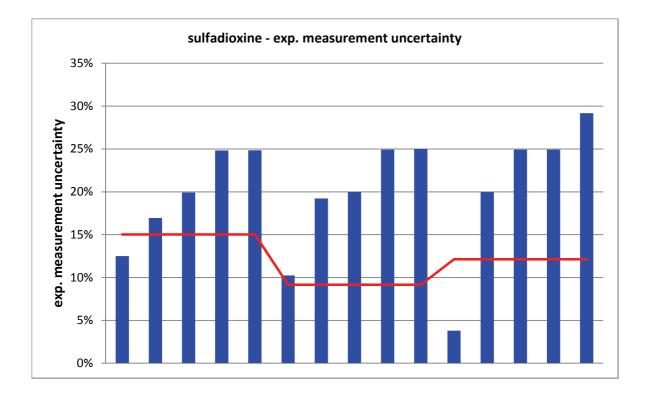


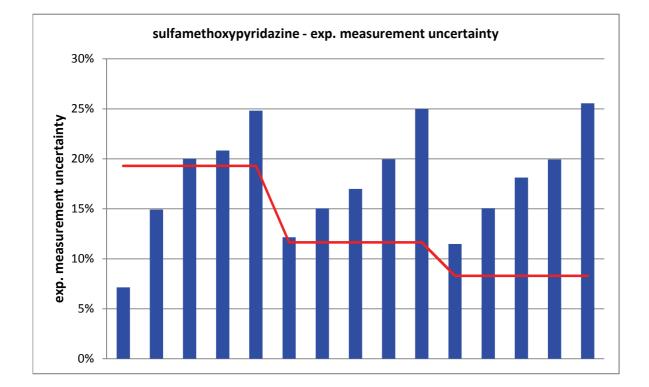


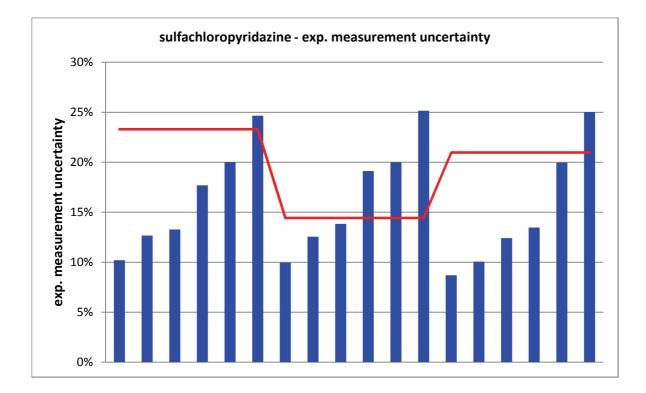


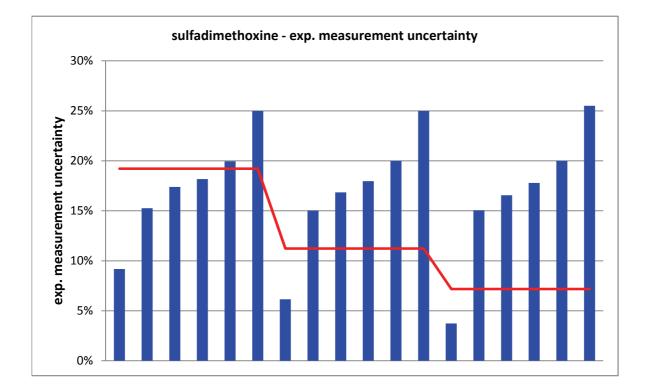


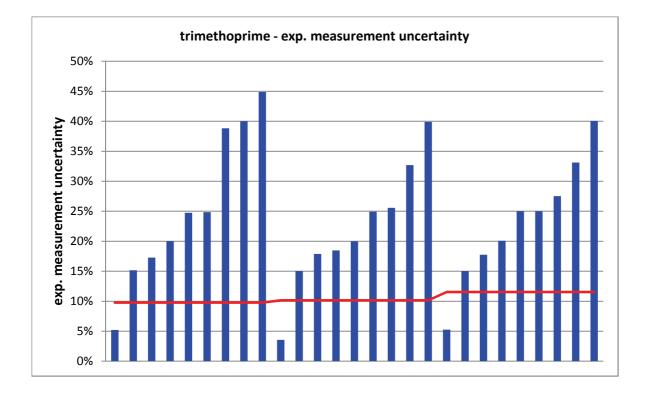




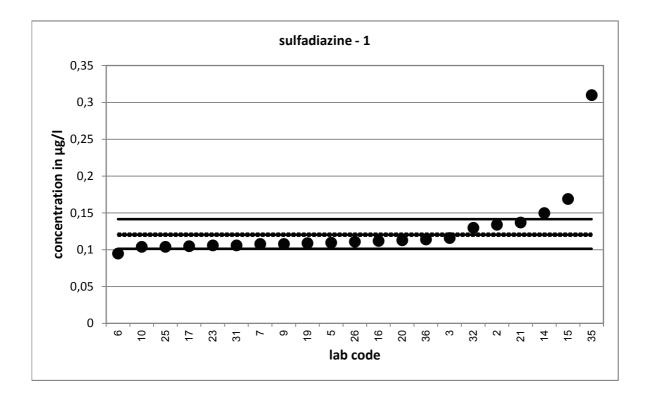


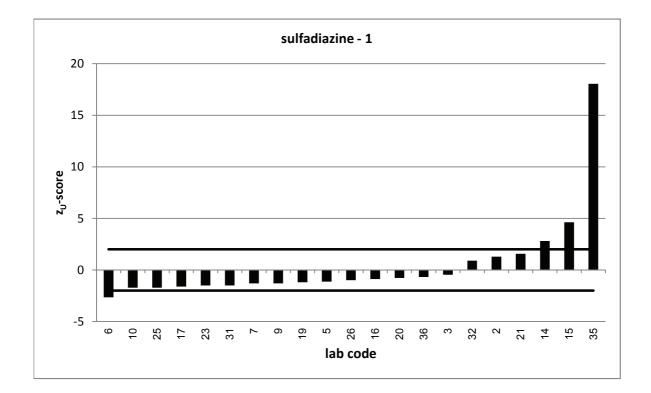


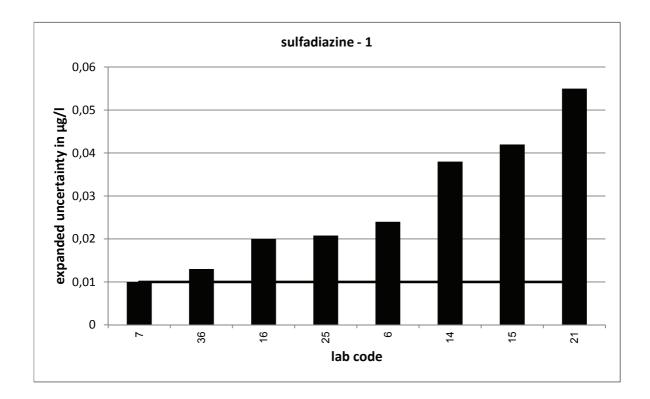


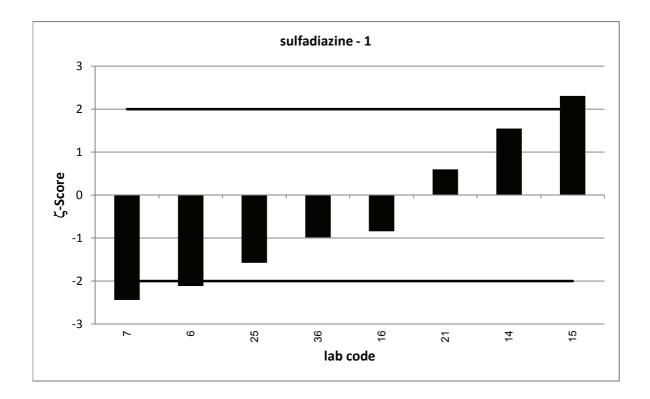


PT 5/16 - T	sulfadiazine - 1				
assigned va				± 0,0022	
	ance limit [µg/l]		0,1415		
	nce limit [µg/l]		0,1013		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
2	0,134			1,3	s
3	0,116			-0,5	S
5	0,1096			-1,1	S
6	0,095	0,024	-2,1	-2,7	q
7	0,108	0,01	-2,4	-1,3	S
9	0,108			-1,3	S
10	0,104			-1,7	S
14	0,15	0,038	1,6	2,8	q
15	0,169	0,042	2,3	4,6	u
16	0,112	0,02	-0,8	-0,9	S
17	0,105			-1,6	S
19	0,109			-1,2	S
20	0,113			-0,8	S
21	0,137	0,055	0,6	1,6	S
23	0,106			-1,5	S
25	0,104	0,021	-1,6	-1,7	S
26	0,111			-1,0	S
31	0,106			-1,5	S
32	0,13			0,9	S
35	0,31			18,1	u
36	0,114	0,013	-1,0	-0,7	S

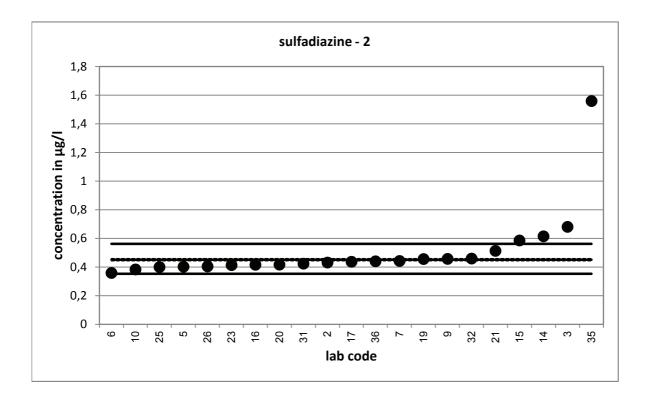


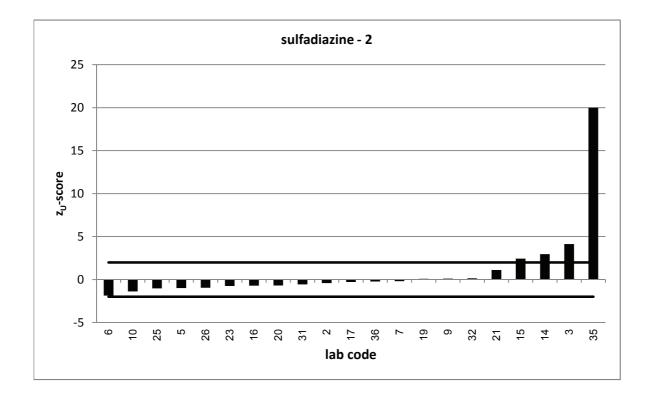


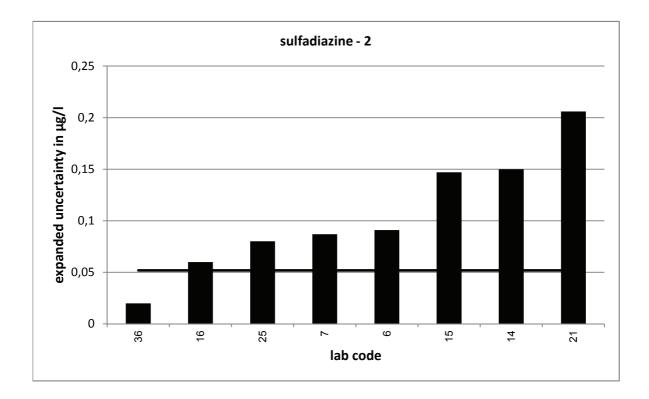


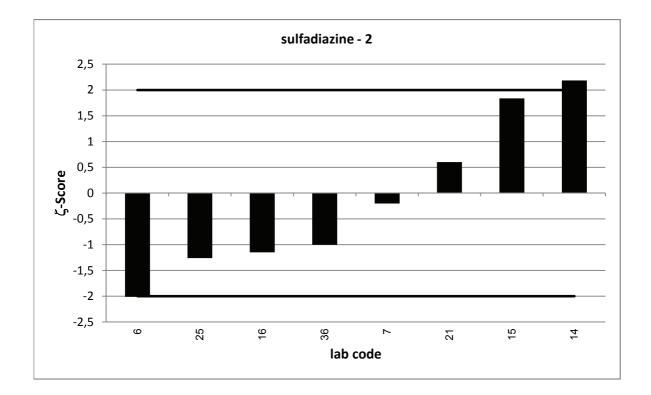


PT 5/16 - T	sulfadi	azine - 2			
assigned va	alue [µg/l]*	0,4518 ± 0,0082			
upper tolera	ance limit [µg/l]		0,5626		
lower tolera	nce limit [µg/l]		0,3534		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
2	0,432			-0,4	s
3	0,681			4,1	u
5	0,4032			-1,0	S
6	0,36	0,091	-2,0	-1,9	S
7	0,443	0,087	-0,2	-0,2	S
9	0,458			0,1	S
10	0,384			-1,4	S
14	0,616	0,15	2,2	3,0	u
15	0,587	0,147	1,8	2,4	q
16	0,417	0,06	-1,1	-0,7	S
17	0,438			-0,3	S
19	0,457			0,1	S
20	0,418			-0,7	S
21	0,514	0,206	0,6	1,1	S
23	0,415			-0,7	S
25	0,401	0,08	-1,3	-1,0	S
26	0,405			-1,0	S
31	0,424			-0,6	S
32	0,46			0,1	S
35	1,56			20,0	u
36	0,441	0,02	-1,0	-0,2	S

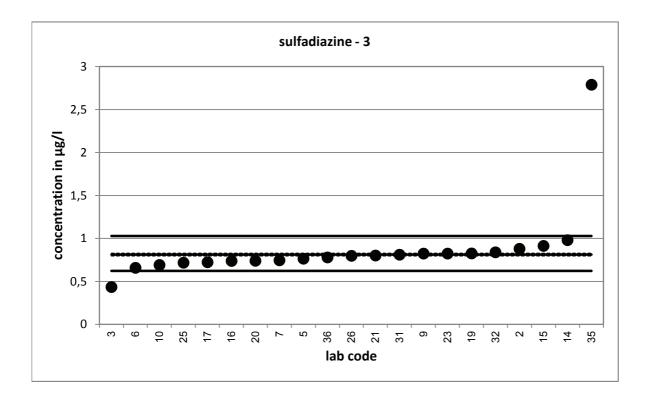


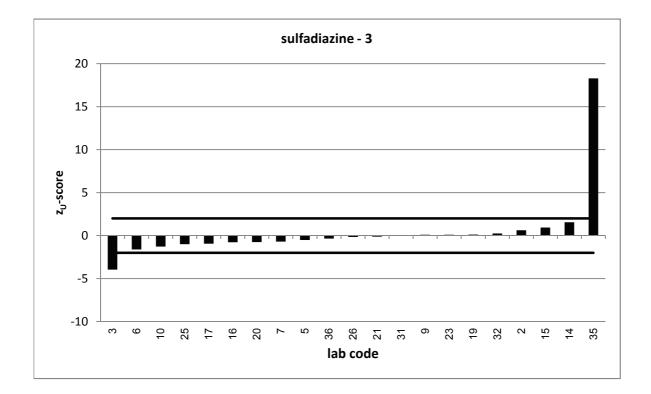


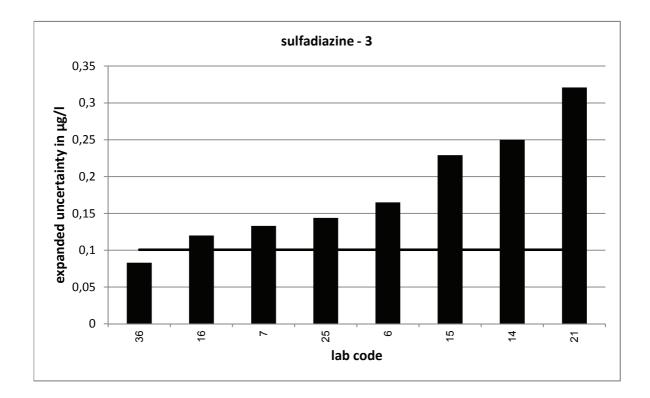


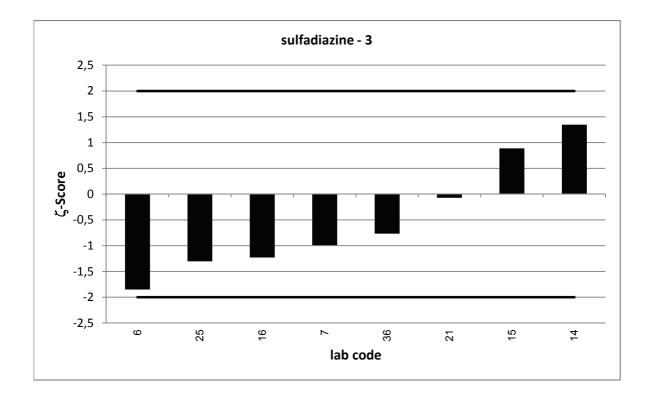


PT 5/16 - T	sulfadi	azine - 3			
assigned va	alue [µg/l]*	0,8133 ± 0,0147			
upper tolera	ance limit [µg/l]		1,029		
lower tolera	nce limit [µg/l]		0,6231		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
2	0,881			0,6	S
3	0,436			-4,0	u
5	0,7651			-0,5	S
6	0,66	0,165	-1,9	-1,6	S
7	0,747	0,133	-1,0	-0,7	S
9	0,825			0,1	S
10	0,692			-1,3	S
14	0,982	0,25	1,3	1,6	S
15	0,915	0,229	0,9	0,9	S
16	0,739	0,12	-1,2	-0,8	S
17	0,724			-0,9	S
19	0,827			0,1	S
20	0,741			-0,8	S
21	0,802	0,321	-0,1	-0,1	S
23	0,825			0,1	S
25	0,719	0,144	-1,3	-1,0	S
26	0,798			-0,2	S
31	0,811			0,0	S
32	0,84			0,2	S
35	2,79			18,3	u
36	0,781	0,083	-0,8	-0,3	S

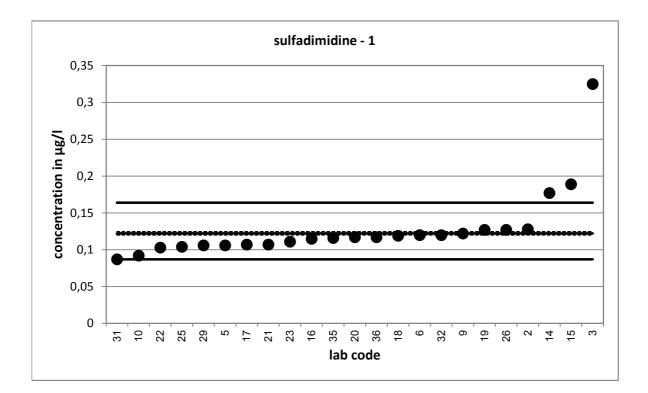


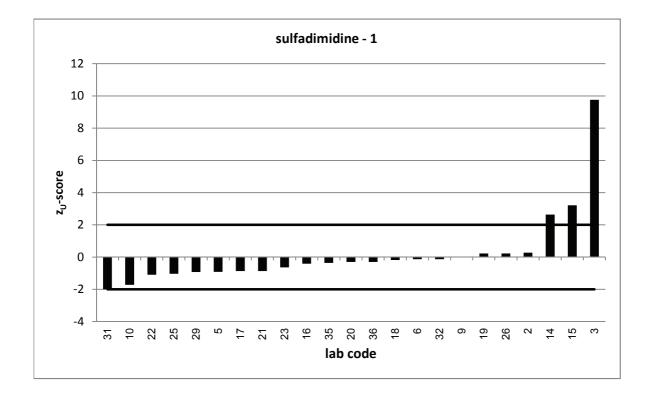


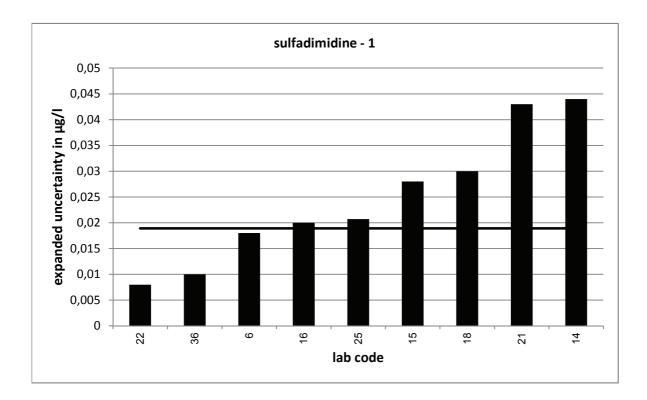


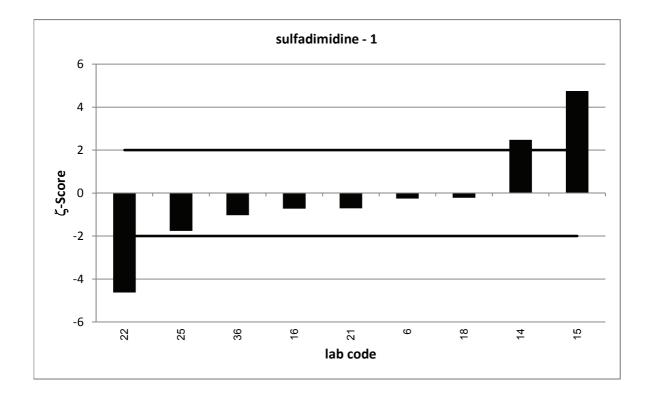


PT 5/16 - T	sulfadimidine - 1					
assigned va	alue [µg/l]*	0,1223 ± 0,0023				
upper tolera	ance limit [µg/l]		0,1638			
lower tolera	nce limit [µg/l]		0,08707			
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**	
2	0,128			0,3	S	
3	0,325			9,8	u	
5	0,1061			-0,9	S	
6	0,12	0,018	-0,3	-0,1	S	
9	0,122			0,0	S	
10	0,092			-1,7	S	
14	0,177	0,044	2,5	2,6	q	
15	0,189	0,028	4,7	3,2	u	
16	0,115	0,02	-0,7	-0,4	S	
17	0,107			-0,9	S	
18	0,119	0,03	-0,2	-0,2	S	
19	0,127			0,2	S	
20	0,117			-0,3	S	
21	0,107	0,043	-0,7	-0,9	S	
22	0,103	0,008	-4,6	-1,1	S	
23	0,111			-0,6	S	
25	0,104	0,021	-1,8	-1,0	S	
26	0,127			0,2	S	
29	0,106			-0,9	S	
31	0,087			-2,0	S	
32	0,12			-0,1	S	
35	0,116			-0,4	S	
36	0,117	0,01	-1,0	-0,3	S	

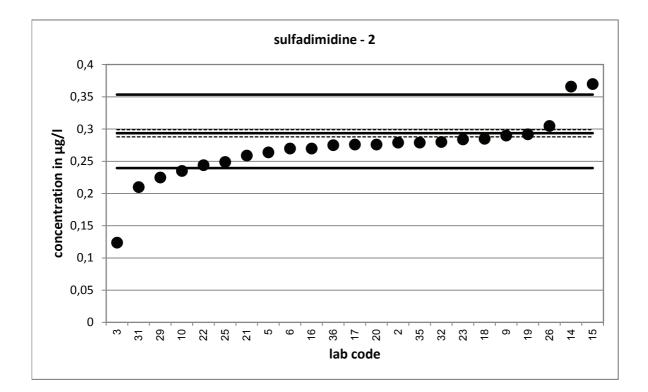


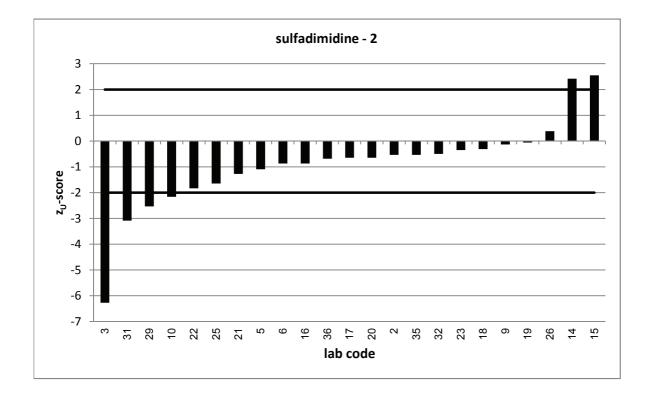


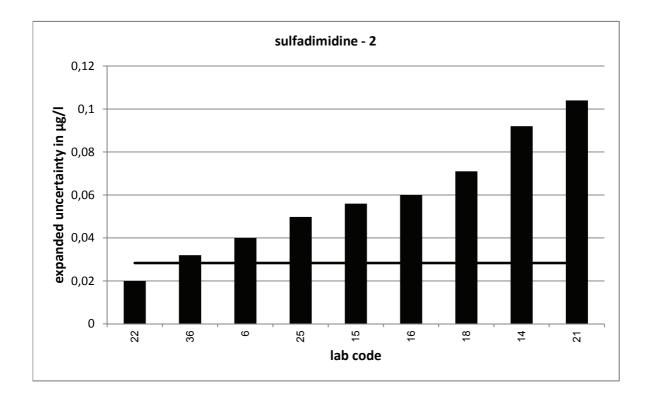


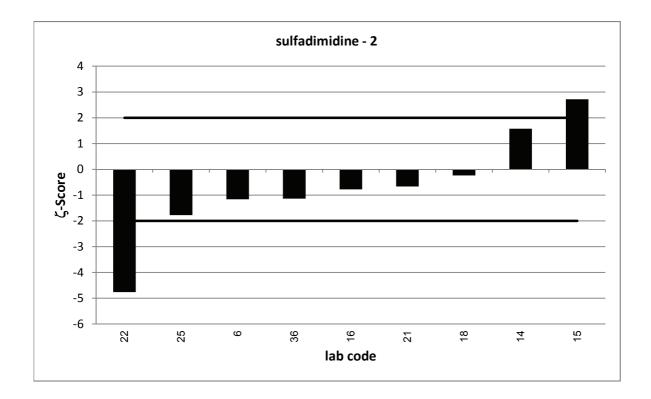


PT 5/16 - T	sulfadimidine - 2				
assigned va	0,2935 ± 0,0056				
upper tolera	ance limit [µg/l]		0,3535		
lower tolera	nce limit [µg/l]		0,2394		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
2	0,279			-0,5	S
3	0,124			-6,3	u
5	0,2639			-1,1	S
6	0,27	0,04	-1,2	-0,9	S
9	0,29			-0,1	S
10	0,235			-2,2	q
14	0,366	0,092	1,6	2,4	q
15	0,37	0,056	2,7	2,6	q
16	0,27	0,06	-0,8	-0,9	S
17	0,276			-0,6	S
18	0,285	0,071	-0,2	-0,3	S
19	0,292			-0,1	S
20	0,276			-0,6	S
21	0,259	0,104	-0,7	-1,3	S
22	0,244	0,02	-4,8	-1,8	S
23	0,284			-0,4	S
25	0,249	0,05	-1,8	-1,6	S
26	0,305			0,4	S
29	0,225			-2,5	q
31	0,21			-3,1	u
32	0,28			-0,5	S
35	0,279			-0,5	S
36	0,275	0,032	-1,1	-0,7	S

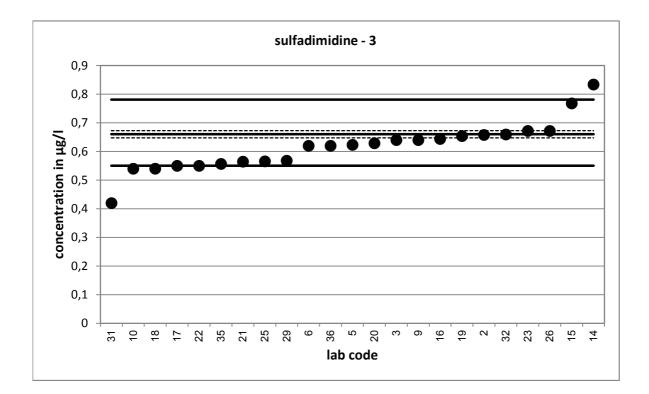


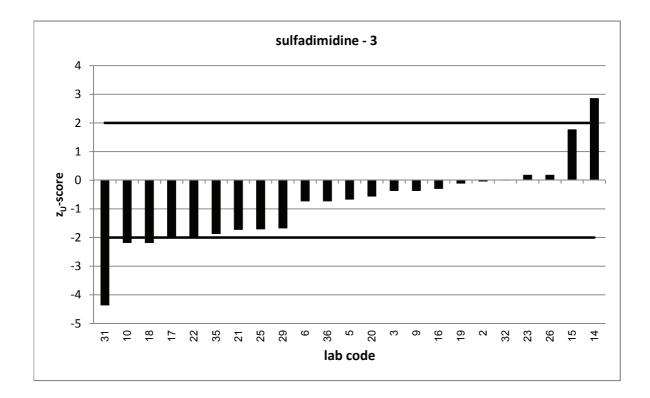


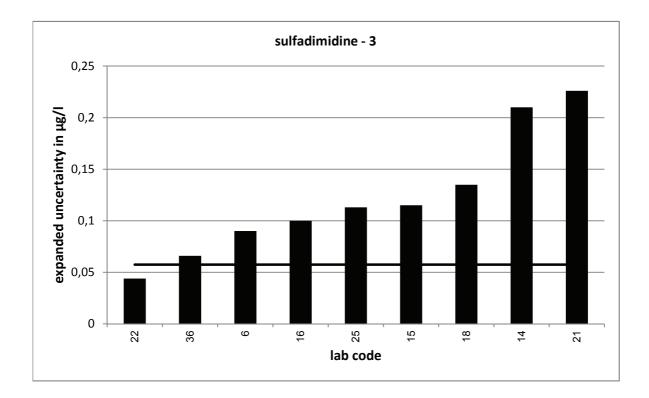


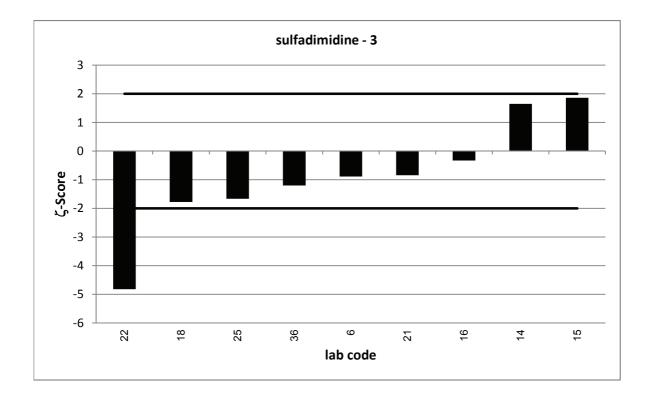


PT 5/16 - T	sulfadimidine - 3					
assigned va		0,6604 ± 0,0126				
	ance limit [µg/l]		0,7813			
lower tolera	nce limit [µg/l]		0,5502			
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**	
2	0,658			0,0	s	
3	0,64			-0,4	S	
5	0,6233			-0,7	S	
6	0,62	0,09	-0,9	-0,7	S	
9	0,64			-0,4	S	
10	0,54			-2,2	q	
14	0,834	0,21	1,7	2,9	q	
15	0,768	0,115	1,9	1,8	S	
16	0,644	0,1	-0,3	-0,3	S	
17	0,55			-2,0	S	
18	0,54	0,135	-1,8	-2,2	q	
19	0,654			-0,1	S	
20	0,629			-0,6	S	
21	0,565	0,226	-0,8	-1,7	S	
22	0,55	0,044	-4,8	-2,0	S	
23	0,672			0,2	S	
25	0,566	0,113	-1,7	-1,7	S	
26	0,672			0,2	S	
29	0,568			-1,7	S	
31	0,42			-4,4	u	
32	0,66			0,0	S	
35	0,557			-1,9	S	
36	0,62	0,066	-1,2	-0,7	S	

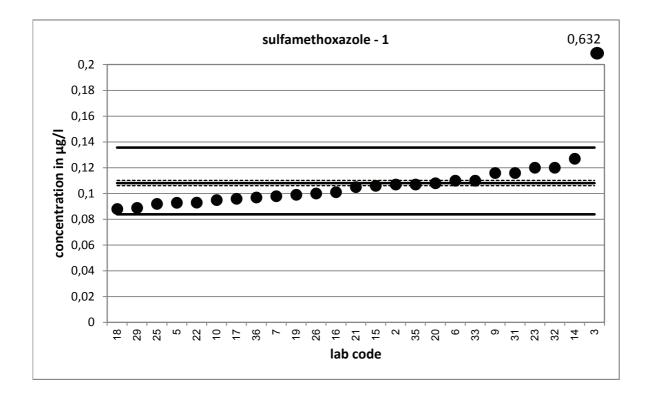


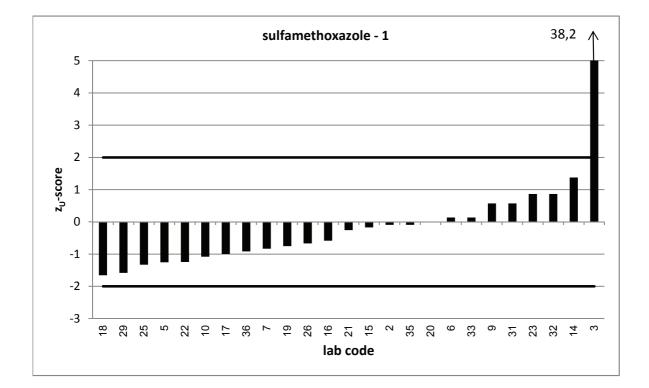


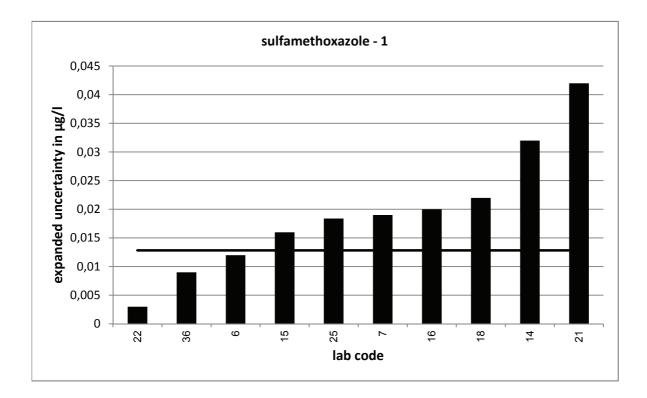


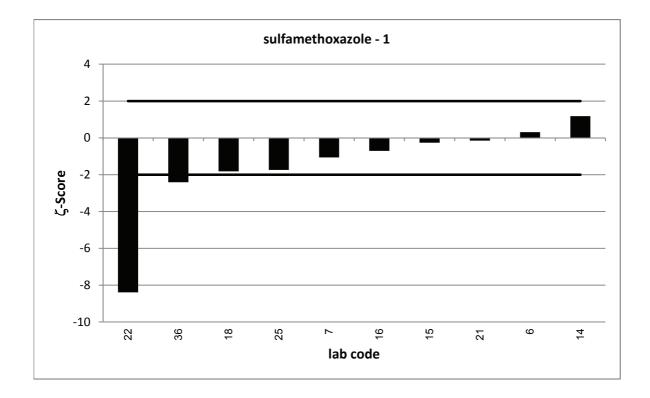


PT 5/16 - T	sulfamethoxazole - 1					
assigned value [µg/l]*			0,1081 ± 0,002			
upper tolera	ance limit [µg/l]		0,1356			
lower tolera	nce limit [µg/l]		0,08384			
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**	
2	0,107			-0,1	s	
3	0,632			38,2	u	
5	0,0929			-1,3	S	
6	0,11	0,012	0,3	0,1	S	
7	0,098	0,019	-1,1	-0,8	S	
9	0,116			0,6	S	
10	0,095			-1,1	S	
14	0,127	0,032	1,2	1,4	S	
15	0,106	0,016	-0,3	-0,2	S	
16	0,101	0,02	-0,7	-0,6	S	
17	0,096			-1,0	S	
18	0,088	0,022	-1,8	-1,7	S	
19	0,099			-0,8	S	
20	0,108			0,0	S	
21	0,105	0,042	-0,1	-0,3	S	
22	0,093	0,003	-8,4	-1,2	S	
23	0,12			0,9	S	
25	0,092	0,018	-1,7	-1,3	S	
26	0,1			-0,7	S	
29	0,0889			-1,6	S	
31	0,116			0,6	S	
32	0,12			0,9	S	
33	0,11			0,1	S	
35	0,107			-0,1	S	
36	0,097	0,009	-2,4	-0,9	S	

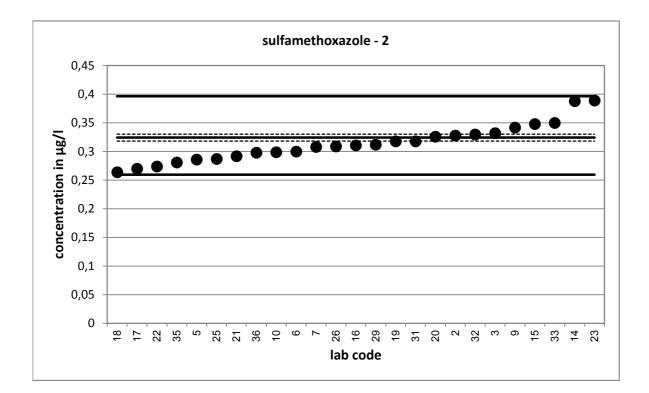


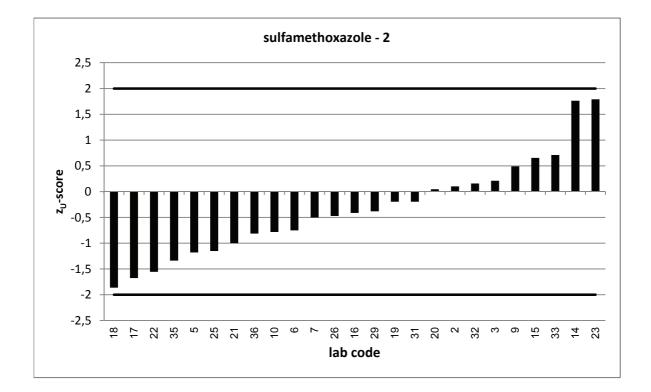


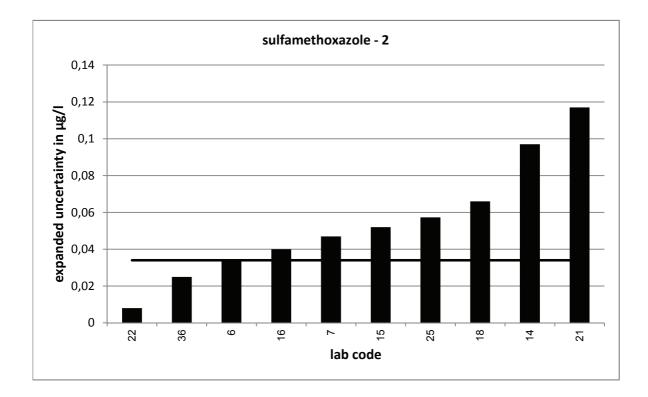


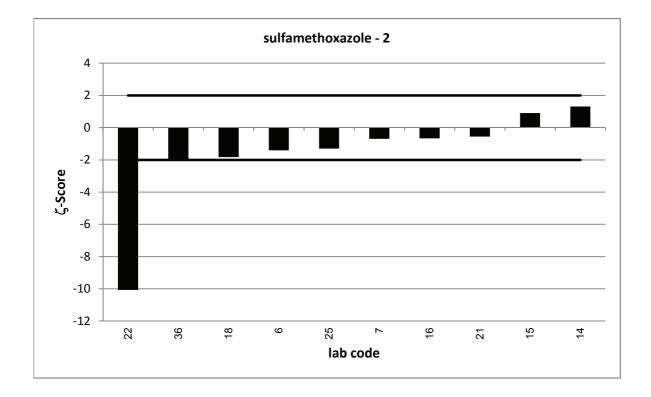


PT 5/16 - T	sulfamethoxazole - 2					
assigned value [µg/l]*			0,3243 ± 0,006			
upper tolera	nce limit [µg/l]		0,3965			
lower tolera	nce limit [µg/l]		0,2596			
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**	
2	0,328			0,1	S	
3	0,332			0,2	S	
5	0,2861			-1,2	S	
6	0,3	0,034	-1,4	-0,8	S	
7	0,308	0,047	-0,7	-0,5	S	
9	0,342			0,5	S	
10	0,299			-0,8	S	
14	0,388	0,097	1,3	1,8	S	
15	0,348	0,052	0,9	0,7	S	
16	0,311	0,04	-0,7	-0,4	S	
17	0,27			-1,7	S	
18	0,264	0,066	-1,8	-1,9	S	
19	0,318			-0,2	S	
20	0,326			0,0	S	
21	0,292	0,117	-0,6	-1,0	S	
22	0,274	0,008	-10,1	-1,6	S	
23	0,389			1,8	S	
25	0,287	0,057	-1,3	-1,2	S	
26	0,309			-0,5	S	
29	0,312			-0,4	S	
31	0,318			-0,2	S	
32	0,33			0,2	S	
33	0,35			0,7	S	
35	0,281			-1,3	S	
36	0,298	0,025	-2,0	-0,8	S	

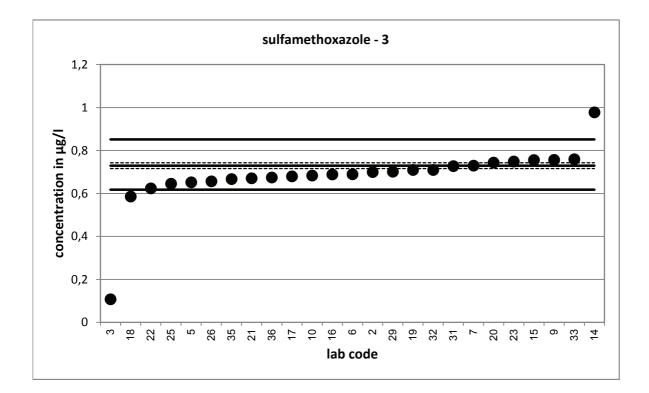


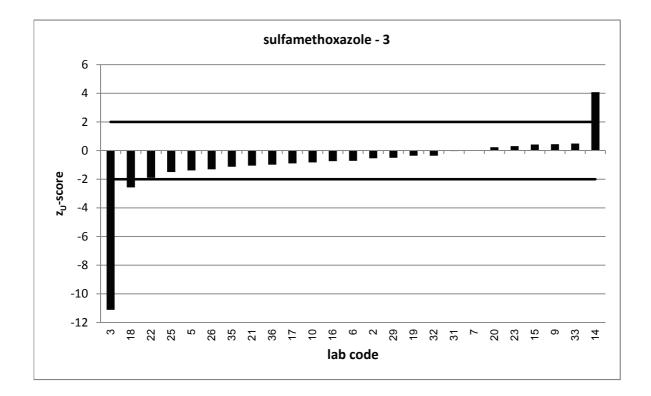


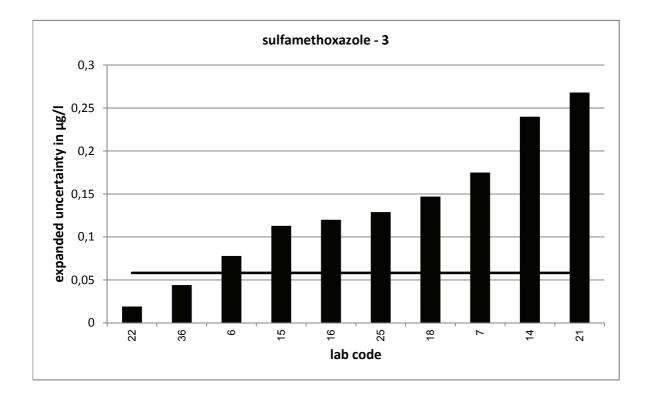


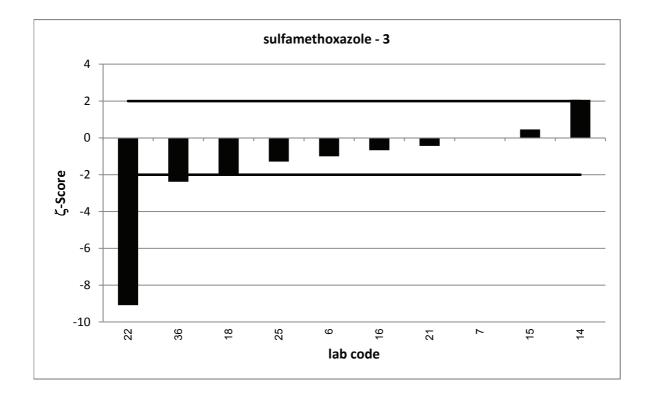


PT 5/16 - T	sulfamethoxazole - 3				
assigned va		0,7298	± 0,0135		
upper tolera	nce limit [µg/l]		0,8514		
lower tolera	nce limit [µg/l]		0,6178		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
2	0,7			-0,5	S
3	0,108			-11,1	u
5	0,6523			-1,4	S
6	0,69	0,078	-1,0	-0,7	S
7	0,73	0,175	0,0	0,0	S
9	0,757			0,4	S
10	0,684			-0,8	S
14	0,978	0,24	2,1	4,1	u
15	0,756	0,113	0,5	0,4	S
16	0,689	0,12	-0,7	-0,7	S
17	0,68			-0,9	S
18	0,586	0,147	-1,9	-2,6	q
19	0,71			-0,4	S
20	0,744			0,2	S
21	0,671	0,268	-0,4	-1,0	S
22	0,624	0,019	-9,1	-1,9	S
23	0,749			0,3	S
25	0,646	0,129	-1,3	-1,5	S
26	0,657			-1,3	S
29	0,702			-0,5	S
31	0,728			0,0	S
32	0,71			-0,4	S
33	0,76			0,5	S
35	0,667			-1,1	S
36	0,675	0,044	-2,4	-1,0	S

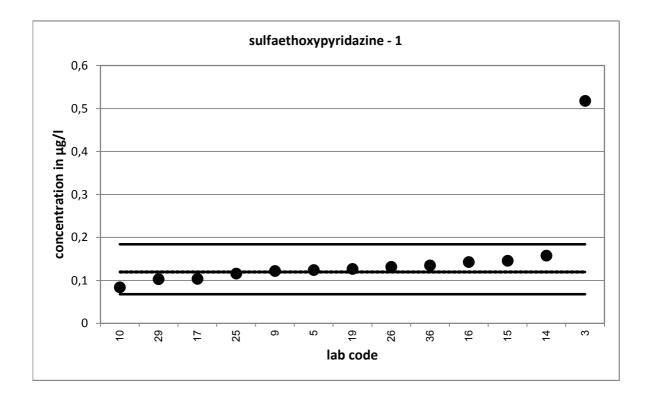


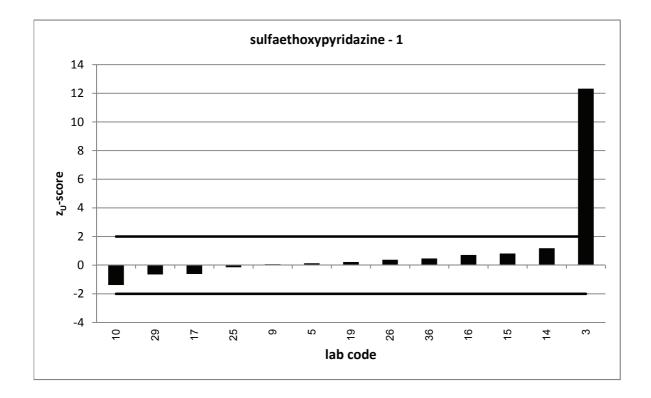


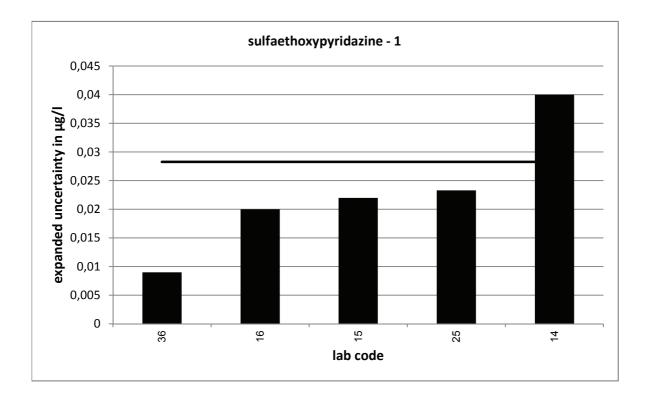


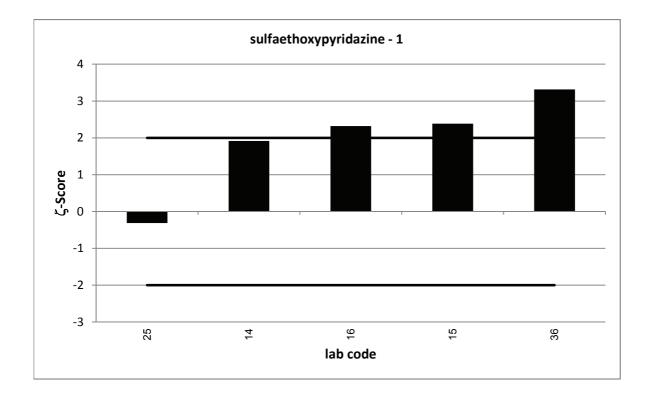


PT 5/16 - TW S2 sulfaethoxypyridazine - 1						
		suilaei				
assigned va			0,1196	± 0,0022		
upper tolera	ance limit [µg/l]		0,1843			
lower tolera	nce limit [µg/l]		0,06813			
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**	
3	0,518			12,3	u	
5	0,1242			0,1	S	
9	0,122			0,1	S	
10	0,084			-1,4	S	
14	0,158	0,04	1,9	1,2	S	
15	0,146	0,022	2,4	0,8	S	
16	0,143	0,02	2,3	0,7	S	
17	0,104			-0,6	S	
19	0,127			0,2	S	
25	0,116	0,023	-0,3	-0,1	S	
26	0,132			0,4	S	
29	0,103			-0,6	S	
36	0,135	0,009	3,3	0,5	S	

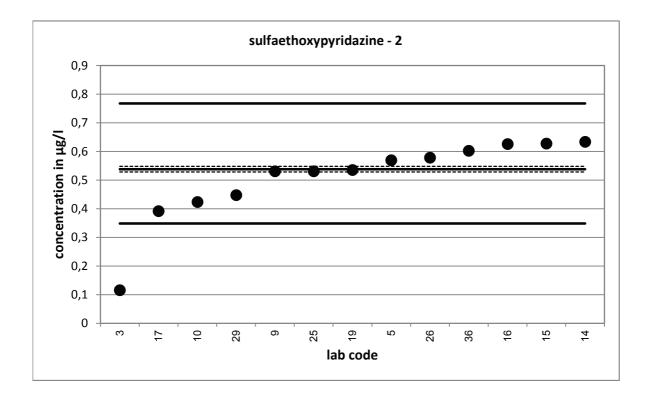


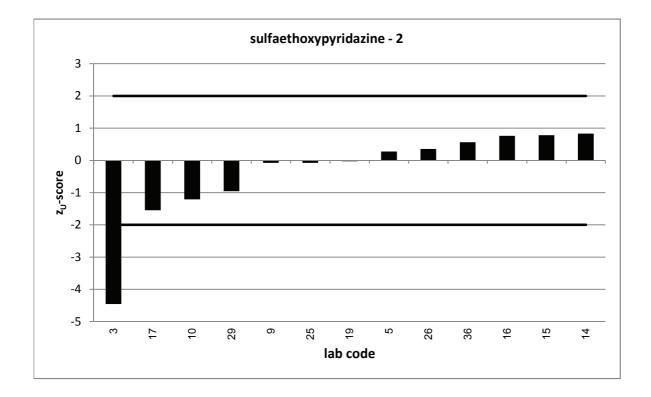


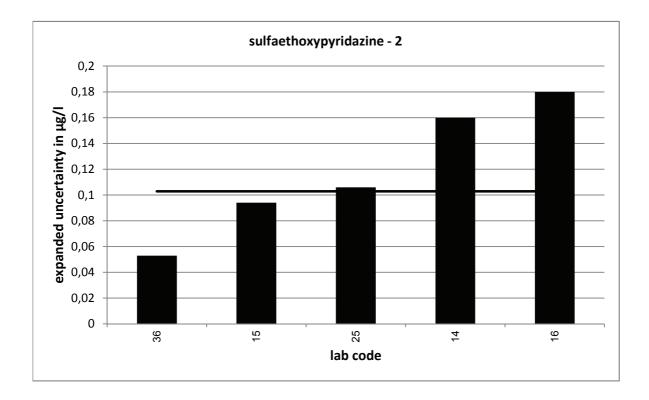


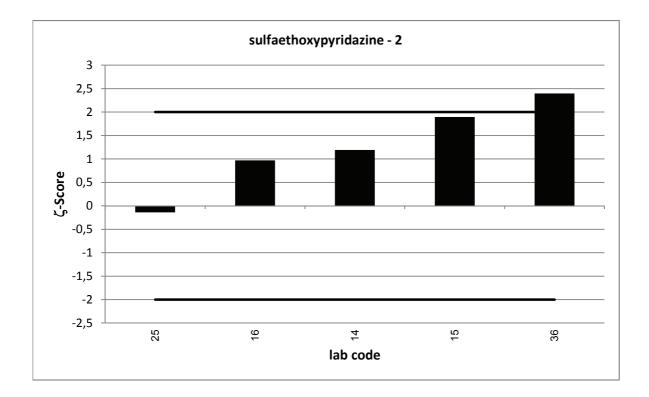


PT 5/16 - T	W S2	sulfaet	sulfaethoxypyridazine - 2		
assigned va	lue [µg/l]*		0,5384	± 0,0099	
upper tolera	nce limit [µg/l]		0,7676		
lower tolera	nce limit [µg/l]		0,3489		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
3	0,116			-4,5	u
5	0,57			0,3	S
9	0,531			-0,1	S
10	0,424			-1,2	S
14	0,634	0,16	1,2	0,8	S
15	0,628	0,094	1,9	0,8	S
16	0,626	0,18	1,0	0,8	S
17	0,392			-1,5	S
19	0,536			0,0	S
25	0,531	0,106	-0,1	-0,1	S
26	0,579			0,4	S
29	0,448			-1,0	S
36	0,603	0,053	2,4	0,6	S

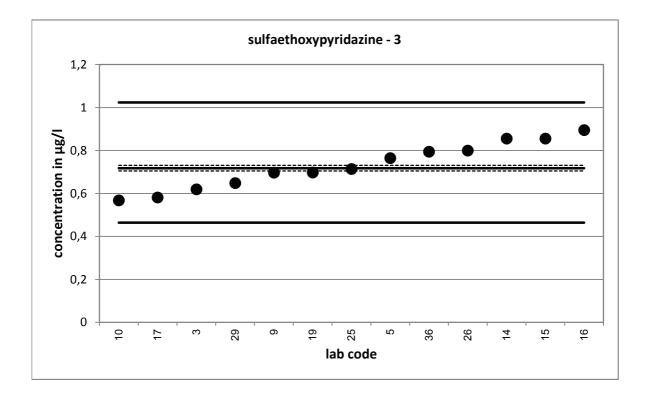


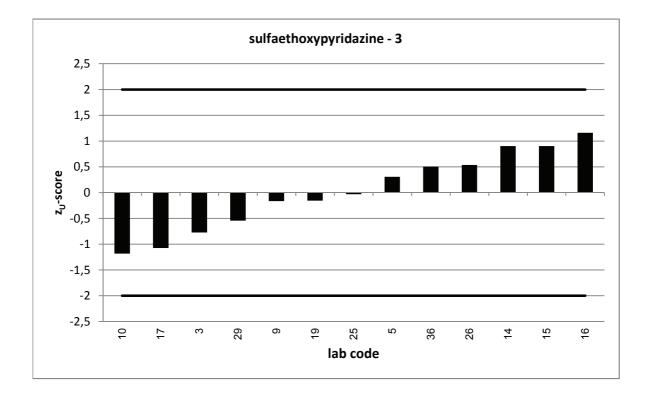


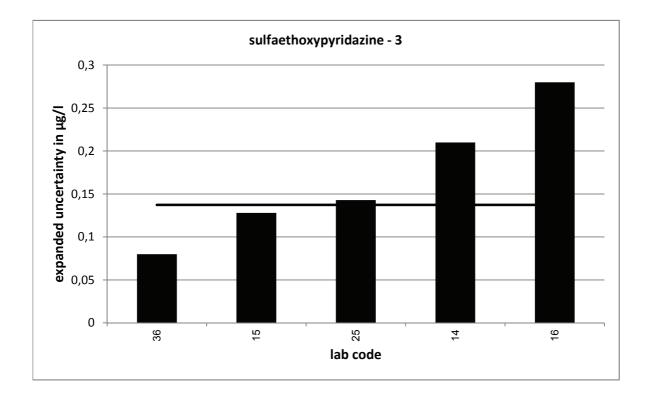


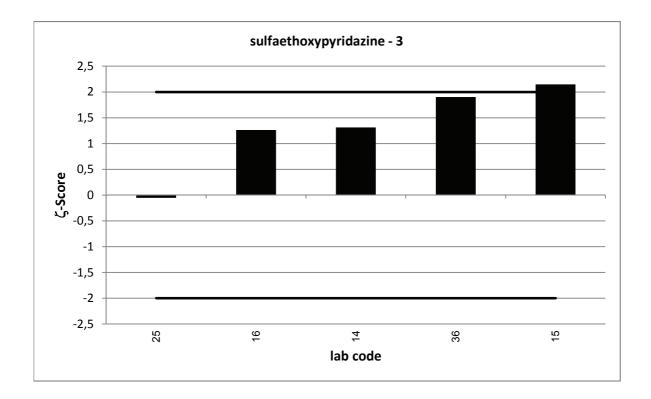


PT 5/16 - TW S2 sulfaethoxypyridazine - 3						
		suitaet				
assigned va	alue [µg/l]*		0,7179	± 0,0131		
upper tolera	ance limit [µg/l]		1,023			
lower tolera	nce limit [µg/l]		0,4645			
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**	
3	0,62			-0,8	S	
5	0,7647			0,3	S	
9	0,697			-0,2	S	
10	0,568			-1,2	S	
14	0,856	0,21	1,3	0,9	S	
15	0,856	0,128	2,1	0,9	S	
16	0,895	0,28	1,3	1,2	S	
17	0,582			-1,1	S	
19	0,698			-0,2	S	
25	0,714	0,143	-0,1	0,0	S	
26	0,8			0,5	S	
29	0,649			-0,5	S	
36	0,795	0,08	1,9	0,5	S	

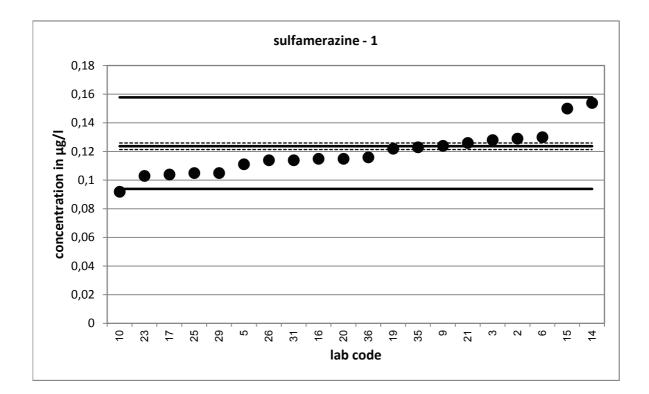


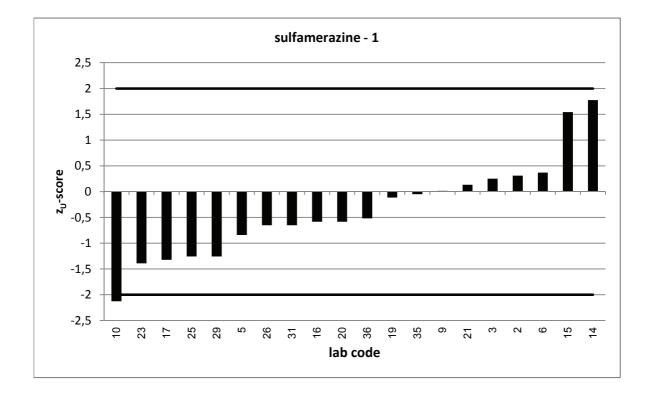


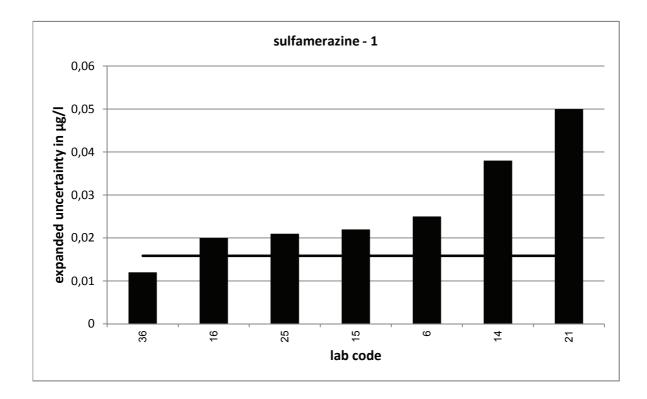


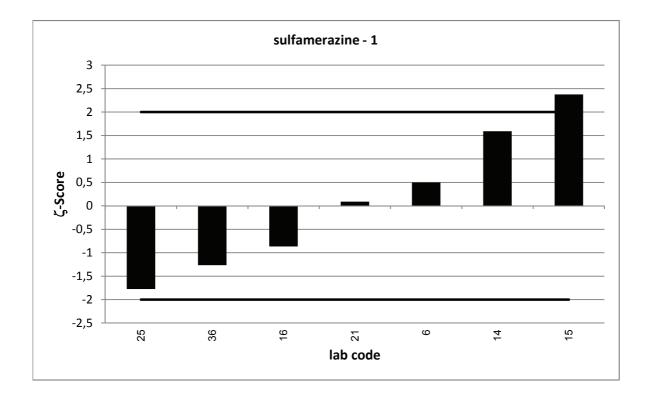


PT 5/16 - T	W S2	sulfamerazine - 1			
assigned va	lue [µg/l]*		0,1237	± 0,0023	
upper tolera	ance limit [µg/l]		0,1578		
lower tolera	nce limit [µg/l]		0,0939		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
2	0,129			0,3	s
3	0,128			0,3	S
5	0,1112			-0,8	S
6	0,13	0,025	0,5	0,4	S
9	0,124			0,0	S
10	0,092			-2,1	q
14	0,154	0,038	1,6	1,8	S
15	0,15	0,022	2,4	1,5	S
16	0,115	0,02	-0,9	-0,6	S
17	0,104			-1,3	S
19	0,122			-0,1	S
20	0,115			-0,6	S
21	0,126	0,05	0,1	0,1	S
23	0,103			-1,4	S
25	0,105	0,021	-1,8	-1,3	S
26	0,114			-0,7	S
29	0,105			-1,3	S
31	0,114			-0,7	S
35	0,123			0,0	S
36	0,116	0,012	-1,3	-0,5	S

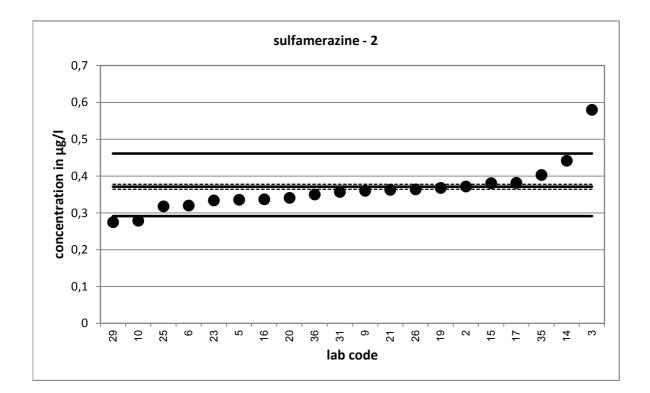


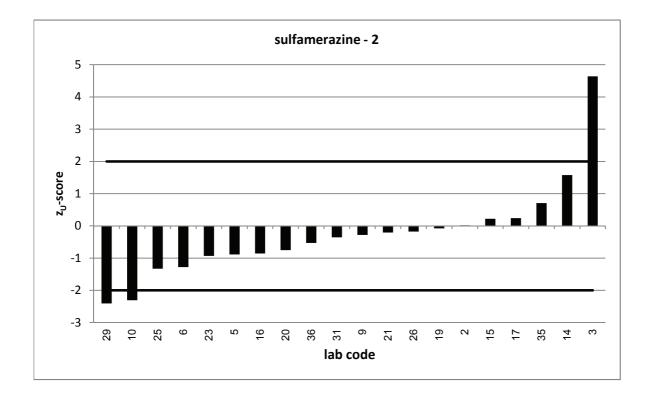


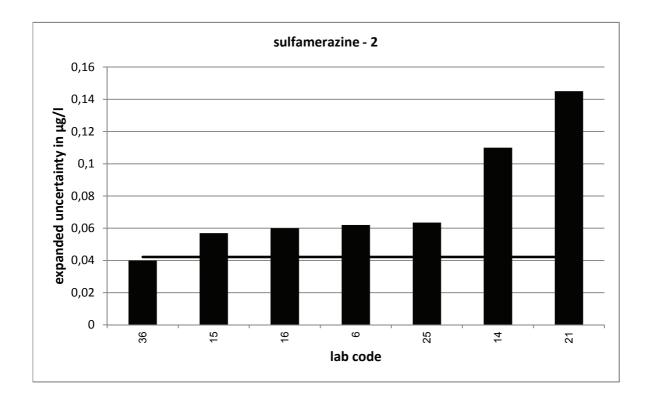


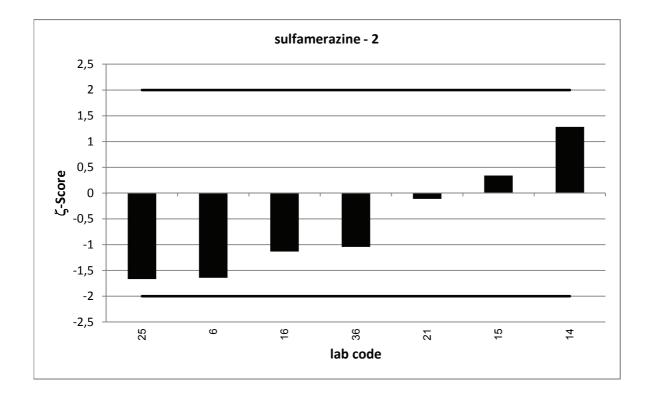


PT 5/16 - T	W S2	sulfamerazine - 2				
assigned va	lue [µg/l]*		0,3712	± 0,0068	± 0,0068	
upper tolera	nce limit [µg/l]		0,4612			
lower tolera	nce limit [µg/l]		0,2913			
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**	
2	0,372			0,0	s	
3	0,58			4,6	u	
5	0,3358			-0,9	S	
6	0,32	0,062	-1,6	-1,3	S	
9	0,36			-0,3	S	
10	0,279			-2,3	q	
14	0,442	0,11	1,3	1,6	S	
15	0,381	0,057	0,3	0,2	S	
16	0,337	0,06	-1,1	-0,9	S	
17	0,382			0,2	S	
19	0,368			-0,1	S	
20	0,341			-0,8	S	
21	0,363	0,145	-0,1	-0,2	S	
23	0,334			-0,9	S	
25	0,318	0,064	-1,7	-1,3	S	
26	0,364			-0,2	S	
29	0,275			-2,4	q	
31	0,357			-0,4	S	
35	0,403			0,7	S	
36	0,35	0,04	-1,0	-0,5	S	

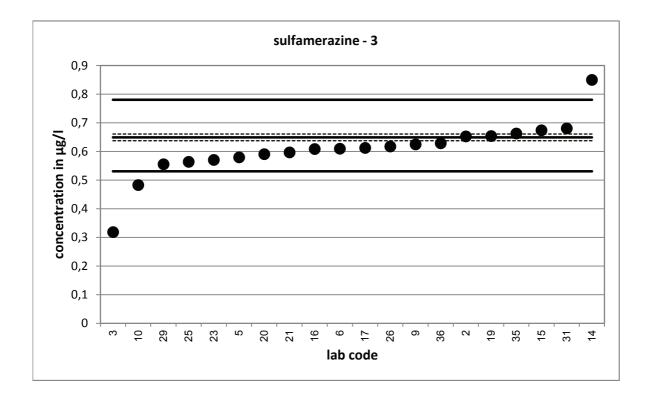


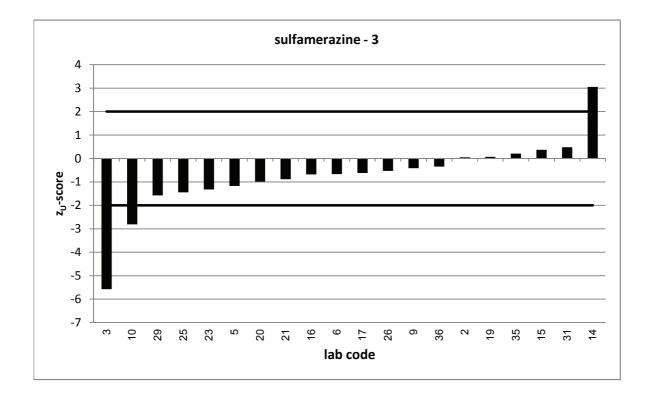


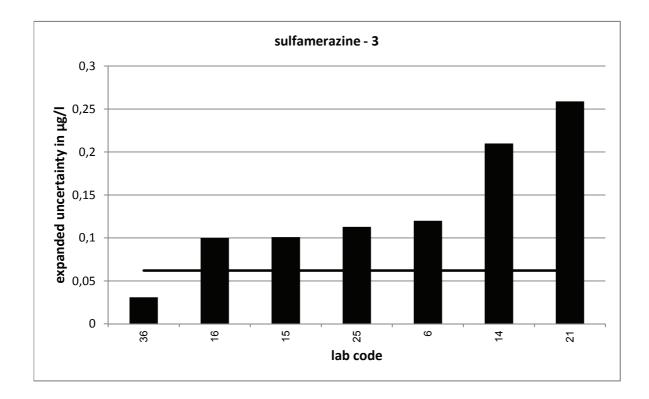


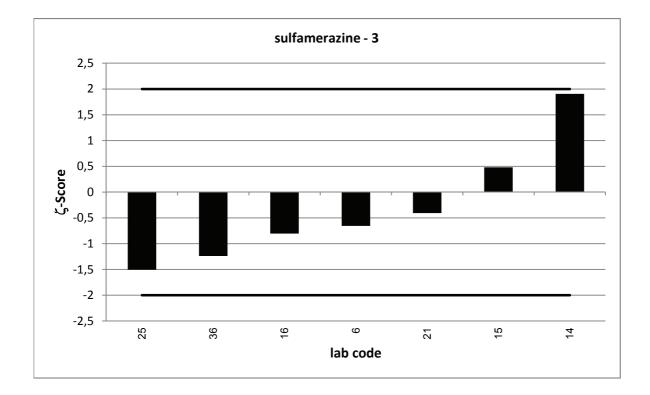


PT 5/16 - T	W S2	sulfamerazine - 3			
assigned va				± 0,0118	
	nce limit [µg/l]		0,7808		
	nce limit [µg/l]		0,5309		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
2	0,653			0,1	s
3	0,319			-5,6	u
5	0,5799			-1,2	S
6	0,61	0,12	-0,7	-0,7	S
9	0,625			-0,4	S
10	0,483			-2,8	q
14	0,85	0,21	1,9	3,1	u
15	0,674	0,101	0,5	0,4	S
16	0,609	0,1	-0,8	-0,7	S
17	0,613			-0,6	S
19	0,654			0,1	S
20	0,591			-1,0	S
21	0,597	0,259	-0,4	-0,9	S
23	0,571			-1,3	S
25	0,564	0,113	-1,5	-1,4	S
26	0,618			-0,5	S
29	0,556			-1,6	S
31	0,681			0,5	S
35	0,663			0,2	S
36	0,629	0,031	-1,2	-0,3	S

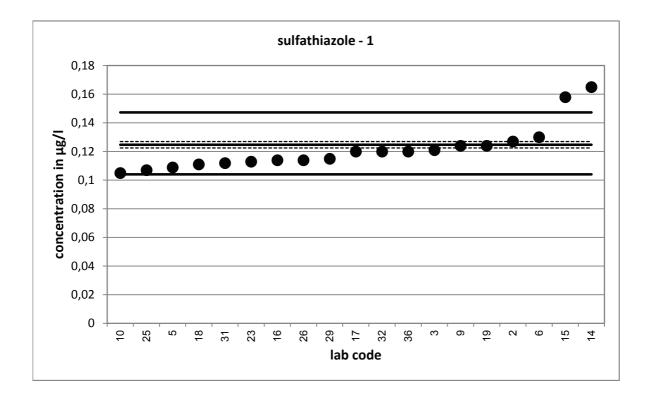


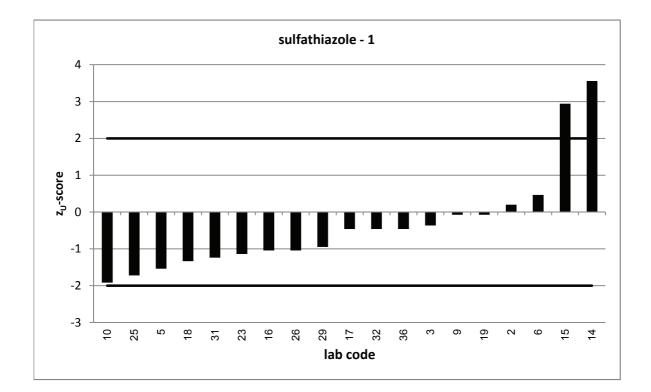


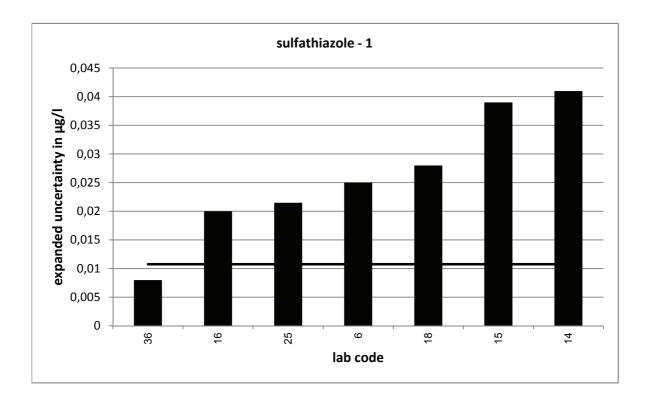


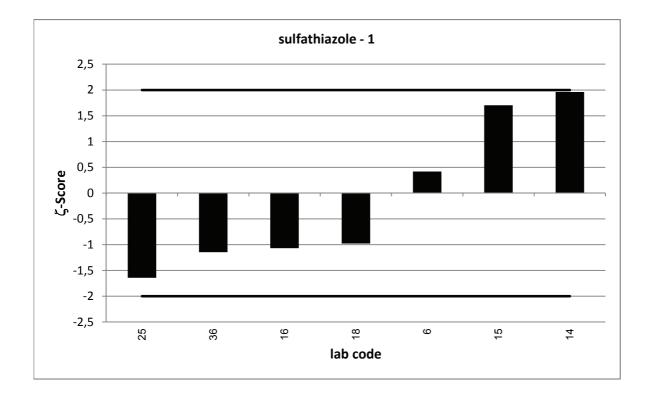


PT 5/16 - T	W S2	sulfath	iazole - 1		
assigned va	lue [µg/l]*		0,1248	± 0,0022	
upper tolera	nce limit [µg/l]		0,1474		
lower tolera	nce limit [µg/l]		0,1041		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
2	0,127			0,2	S
3	0,121			-0,4	S
5	0,1089			-1,5	S
6	0,13	0,025	0,4	0,5	S
9	0,124			-0,1	S
10	0,105			-1,9	S
14	0,165	0,041	2,0	3,6	u
15	0,158	0,039	1,7	2,9	q
16	0,114	0,02	-1,1	-1,0	S
17	0,12			-0,5	S
18	0,111	0,028	-1,0	-1,3	S
19	0,124			-0,1	S
23	0,113			-1,1	S
25	0,107	0,022	-1,6	-1,7	S
26	0,114			-1,0	S
29	0,115			-0,9	S
31	0,112			-1,2	S
32	0,12			-0,5	S
36	0,12	0,008	-1,1	-0,5	S

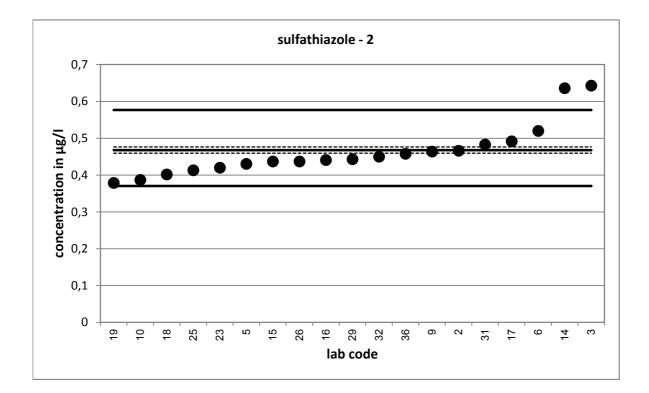


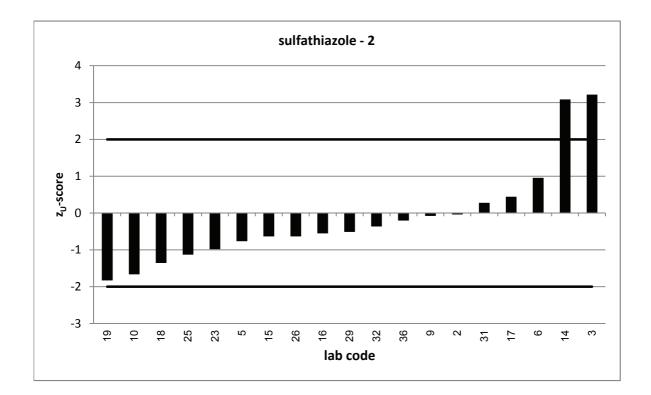






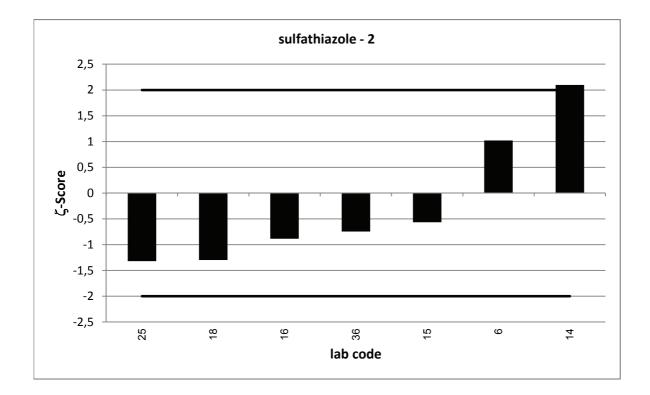
PT 5/16 - T	W S2	sulfath	iazole - 2	2	
assigned va	lue [µg/l]*		0,4678	± 0,0084	
upper tolera	nce limit [µg/l]		0,5768		
lower tolera	nce limit [µg/l]		0,3706		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
2	0,466			0,0	S
3	0,643			3,2	u
5	0,4307			-0,8	S
6	0,52	0,102	1,0	1,0	S
9	0,464			-0,1	S
10	0,387			-1,7	S
14	0,636	0,16	2,1	3,1	u
15	0,437	0,109	-0,6	-0,6	S
16	0,441	0,06	-0,9	-0,6	S
17	0,492			0,4	S
18	0,402	0,101	-1,3	-1,4	S
19	0,379			-1,8	S
23	0,42			-1,0	S
25	0,413	0,083	-1,3	-1,1	S
26	0,437			-0,6	S
29	0,443			-0,5	S
31	0,483			0,3	S
32	0,45			-0,4	S
36	0,458	0,025	-0,7	-0,2	S



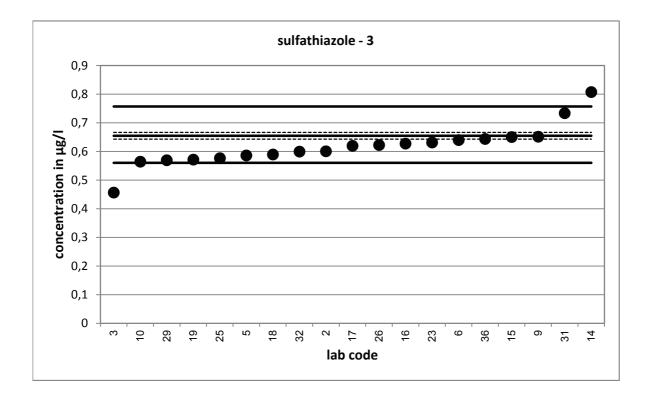


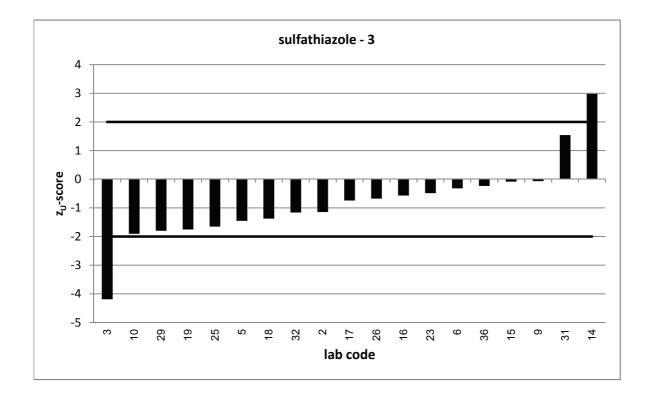
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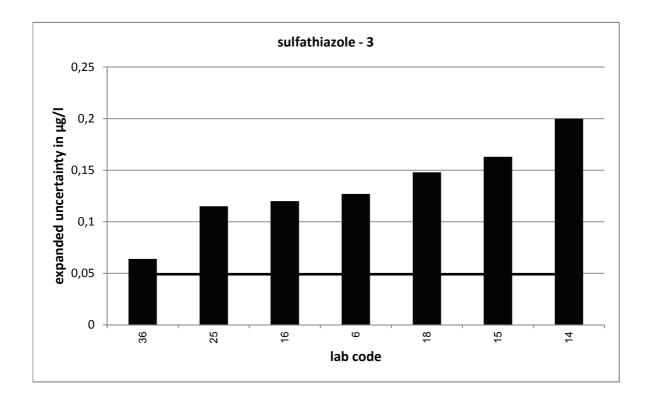


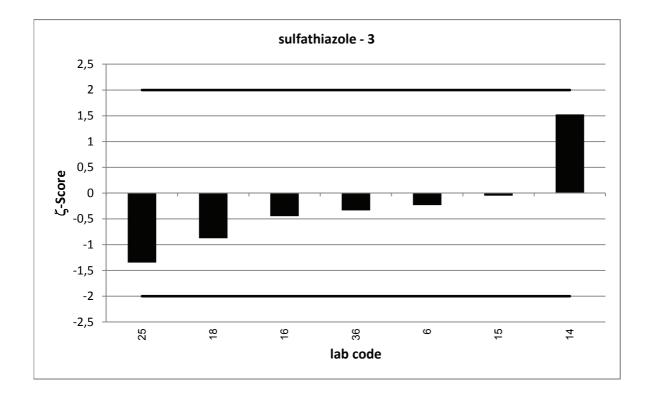


PT 5/16 - T	W S2	sulfath	iazole - 3			
assigned va	lue [µg/l]*			± 0,0117		
	ance limit [µg/l]		0,7574			
	nce limit [µg/l]		0,5604			
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**	
2	0,601			-1,1	S	
3	0,457			-4,2	u	
5	0,5863			-1,5	S	
6	0,64	0,127	-0,2	-0,3	S	
9	0,652			-0,1	S	
10	0,565			-1,9	S	
14	0,808	0,2	1,5	3,0	u	
15	0,651	0,163	0,0	-0,1	S	
16	0,628	0,12	-0,4	-0,6	S	
17	0,62			-0,7	S	
18	0,59	0,148	-0,9	-1,4	S	
19	0,572			-1,8	S	
23	0,632			-0,5	S	
25	0,577	0,115	-1,3	-1,6	S	
26	0,623			-0,7	S	
29	0,57			-1,8	S	
31	0,734			1,5	S	
32	0,6			-1,2	S	
36	0,644	0,064	-0,3	-0,2	S	

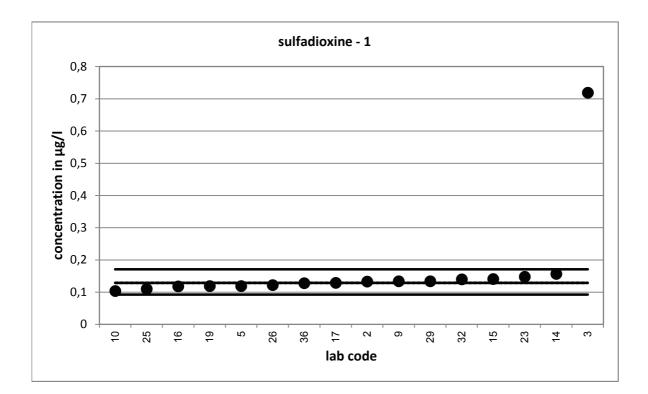


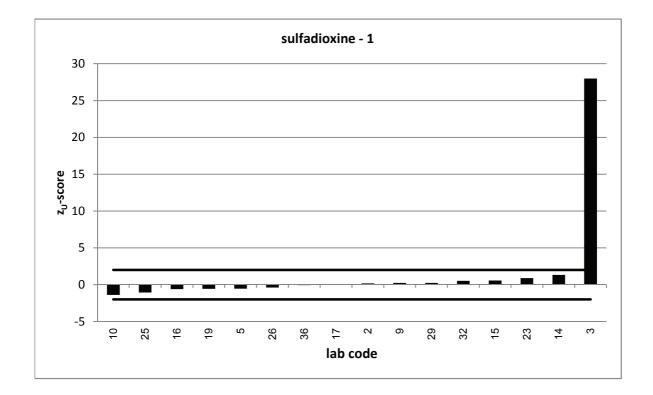


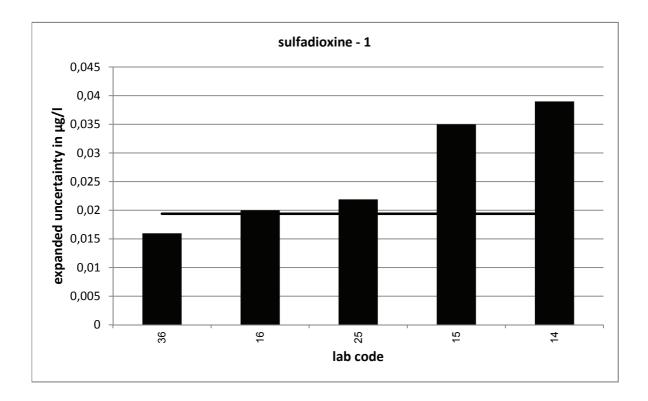


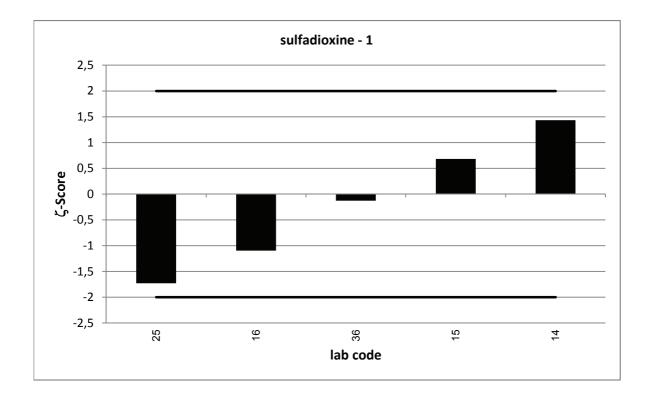


PT 5/16 - T	W S2	sulfadi	oxine - 1			
assigned va	llue [µg/l]*		0,129	± 0,0023		
upper tolera	nce limit [µg/l]		0,1712			
lower tolera	nce limit [µg/l]		0,09274			
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**	
2	0,133			0,2	S	
3	0,719			28,0	u	
5	0,1191			-0,5	S	
9	0,134			0,2	S	
10	0,104			-1,4	S	
14	0,157	0,039	1,4	1,3	S	
15	0,141	0,035	0,7	0,6	S	
16	0,118	0,02	-1,1	-0,6	S	
17	0,129			0,0	S	
19	0,119			-0,6	S	
23	0,148			0,9	S	
25	0,11	0,022	-1,7	-1,0	S	
26	0,122			-0,4	S	
29	0,134			0,2	S	
32	0,14			0,5	S	
36	0,128	0,016	-0,1	-0,1	S	

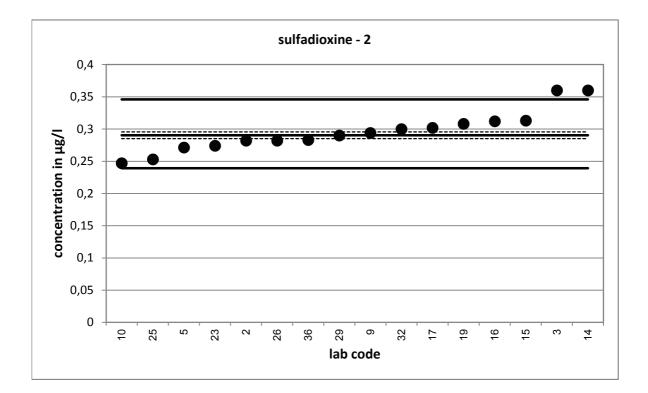


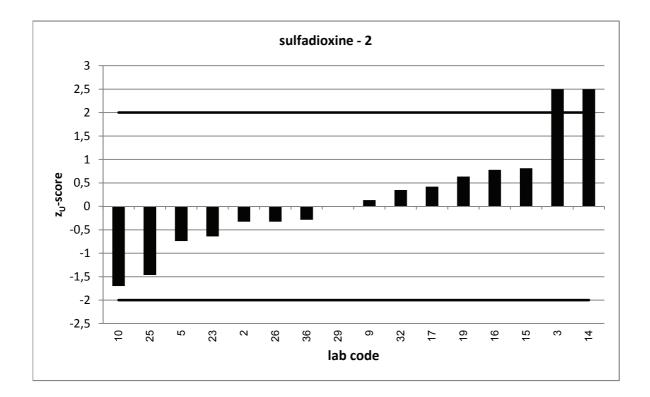


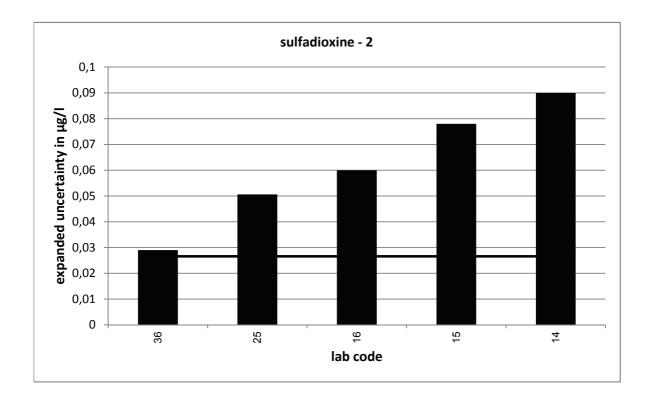


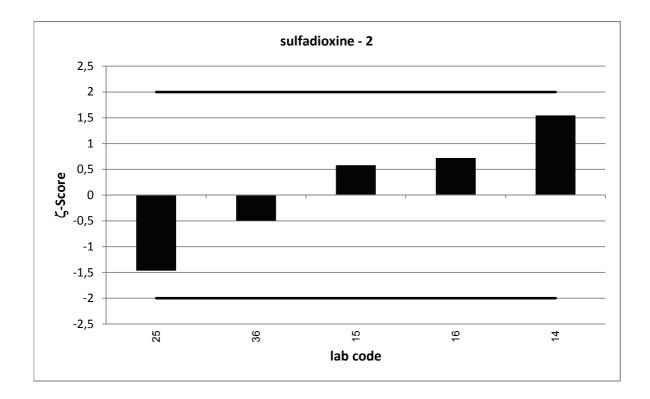


PT 5/16 - T	W S2	sulfadi	oxine - 2			
assigned va	lue [µg/l]*		0,2903	± 0,0052		
upper tolera	nce limit [µg/l]		0,3461			
lower tolera	nce limit [µg/l]		0,2393			
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**	
2	0,282			-0,3	S	
3	0,36			2,5	q	
5	0,2714			-0,7	S	
9	0,294			0,1	S	
10	0,247			-1,7	S	
14	0,36	0,09	1,5	2,5	q	
15	0,313	0,078	0,6	0,8	S	
16	0,312	0,06	0,7	0,8	S	
17	0,302			0,4	S	
19	0,308			0,6	S	
23	0,274			-0,6	S	
25	0,253	0,051	-1,5	-1,5	S	
26	0,282			-0,3	S	
29	0,29			0,0	S	
32	0,3			0,3	S	
36	0,283	0,029	-0,5	-0,3	S	

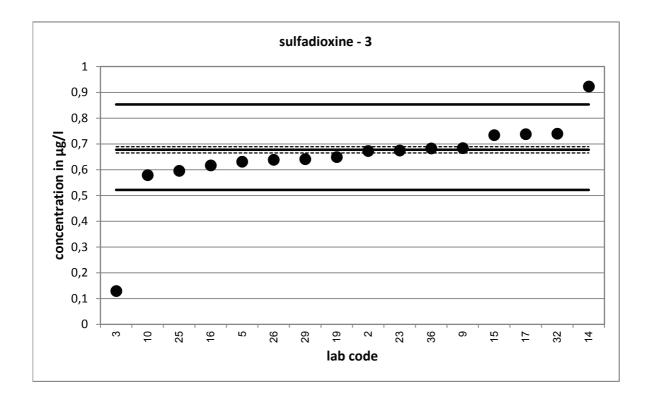


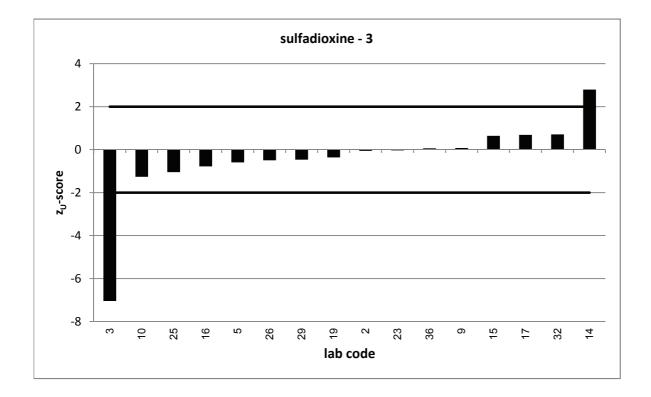


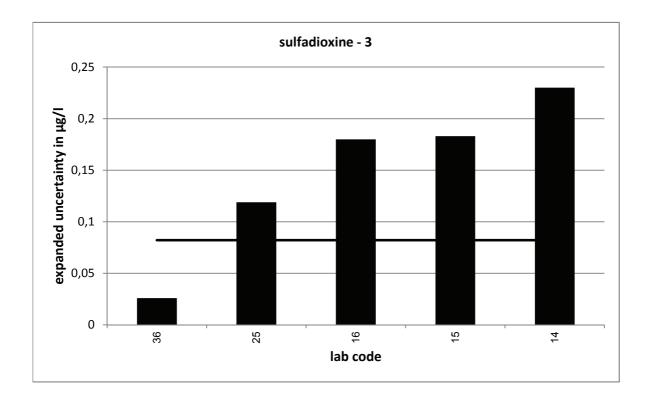


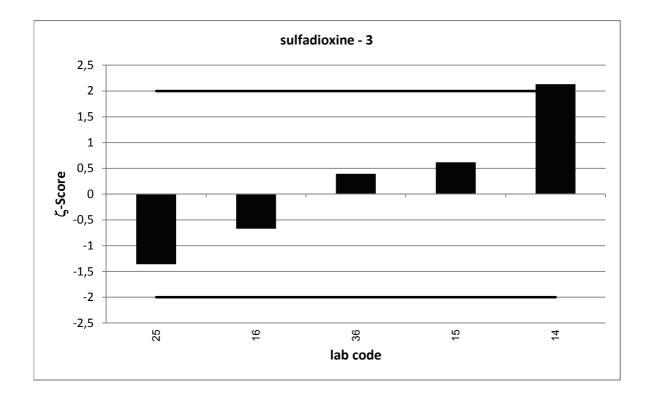


PT 5/16 - T	W S2	sulfadi	oxine - 3			
assigned va	llue [µg/l]*		0,6774	± 0,0121		
upper tolera	nce limit [µg/l]		0,8529			
lower tolera	nce limit [µg/l]		0,522			
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**	
2	0,673			-0,1	S	
3	0,13			-7,0	u	
5	0,6313			-0,6	S	
9	0,684			0,1	S	
10	0,579			-1,3	S	
14	0,923	0,23	2,1	2,8	q	
15	0,734	0,183	0,6	0,6	S	
16	0,617	0,18	-0,7	-0,8	S	
17	0,738			0,7	S	
19	0,649			-0,4	S	
23	0,675			0,0	S	
25	0,596	0,119	-1,4	-1,0	S	
26	0,639			-0,5	S	
29	0,641			-0,5	S	
32	0,74			0,7	S	
36	0,683	0,026	0,4	0,1	S	

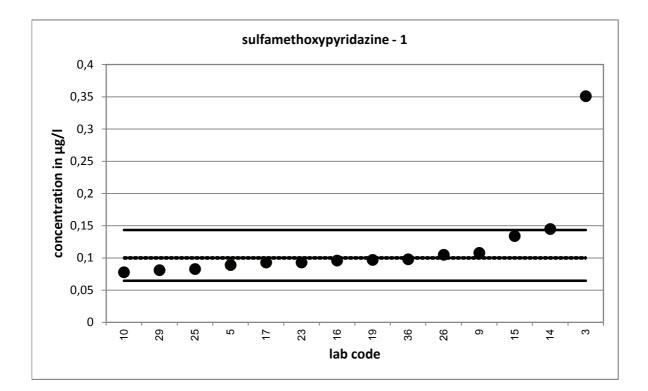


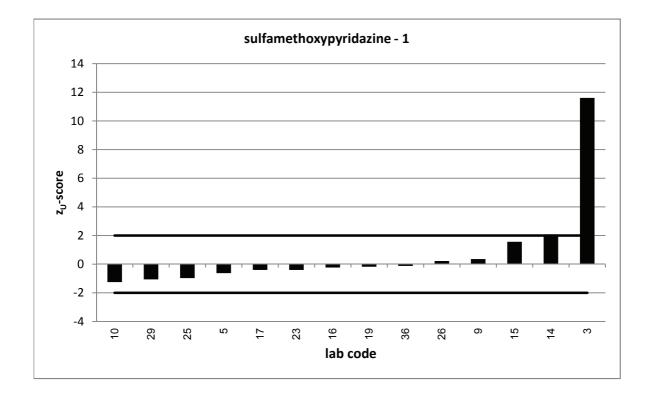


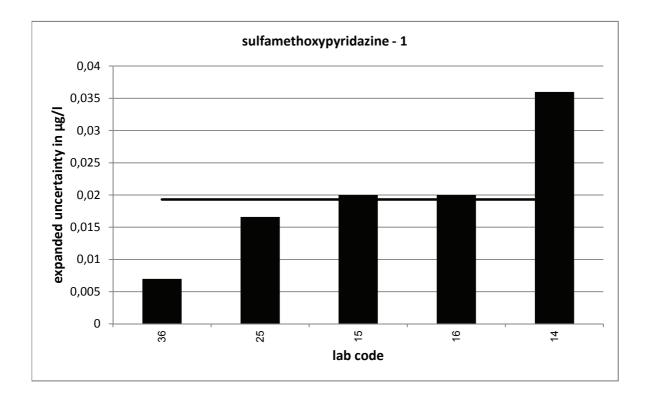


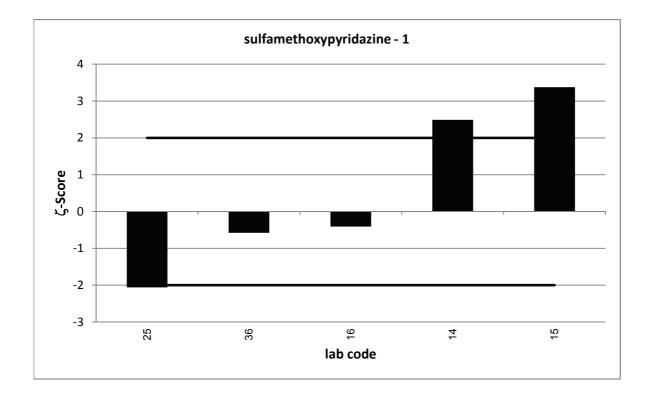


PT 5/16 - TW S2 su			sulfamethoxypyridazine - 1			
assigned va				± 0,0019		
upper tolera	ance limit [µg/l]		0,1433			
lower tolera	nce limit [µg/l]		0,06456			
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**	
3	0,351			11,6	u	
5	0,0891			-0,6	S	
9	0,108			0,4	S	
10	0,078			-1,2	S	
14	0,145	0,036	2,5	2,1	q	
15	0,134	0,02	3,4	1,6	S	
16	0,096	0,02	-0,4	-0,2	S	
17	0,093			-0,4	S	
19	0,097			-0,2	S	
23	0,093			-0,4	S	
25	0,0829	0,017	-2,1	-1,0	S	
26	0,105			0,2	S	
29	0,0813			-1,1	S	
36	0,098	0,007	-0,6	-0,1	S	

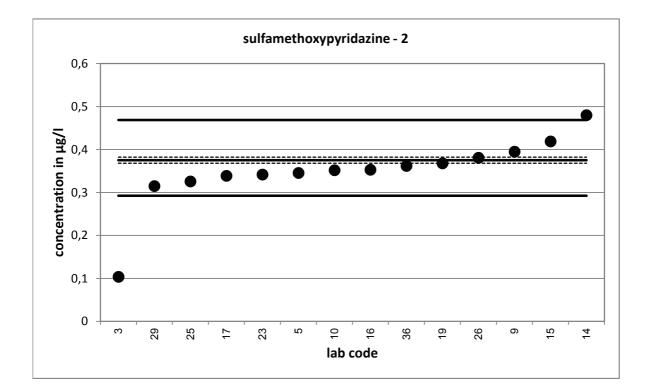


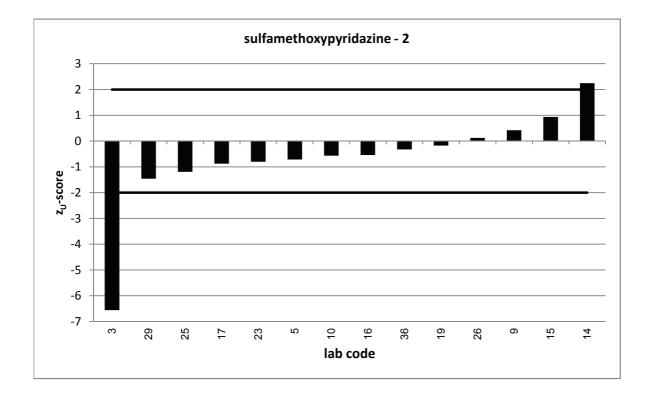


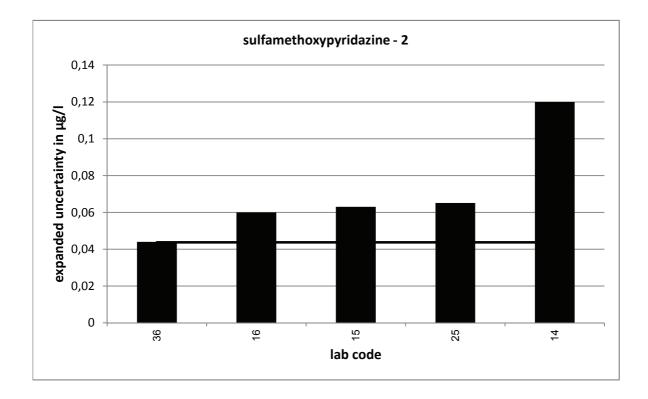


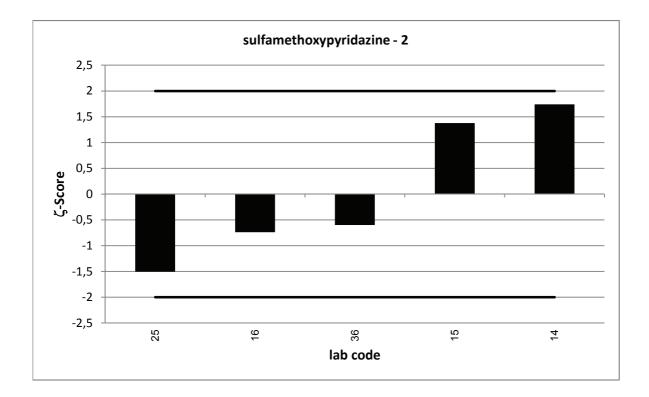


PT 5/16 - TW S2 sulfamethoxypy			ridazine -	2	
assigned va				± 0,0071	
	ance limit [µg/l]		0,4687		
lower tolera	nce limit [µg/l]		0,2925		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
3	0,104			-6,6	u
5	0,3456			-0,7	S
9	0,395			0,4	S
10	0,352			-0,6	S
14	0,48	0,12	1,7	2,2	q
15	0,419	0,063	1,4	0,9	S
16	0,353	0,06	-0,7	-0,5	S
17	0,339			-0,9	S
19	0,368			-0,2	S
23	0,342			-0,8	S
25	0,326	0,065	-1,5	-1,2	S
26	0,381			0,1	S
29	0,315			-1,5	S
36	0,362	0,044	-0,6	-0,3	S

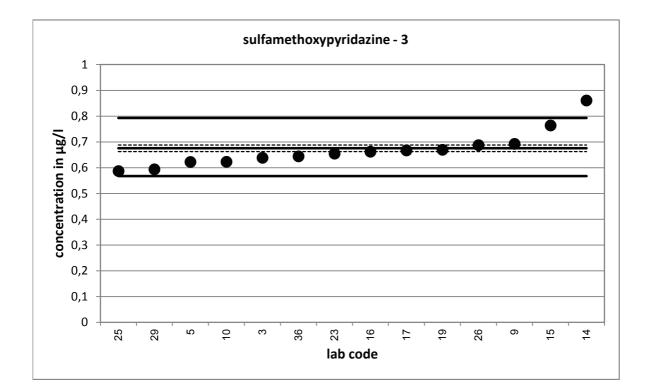


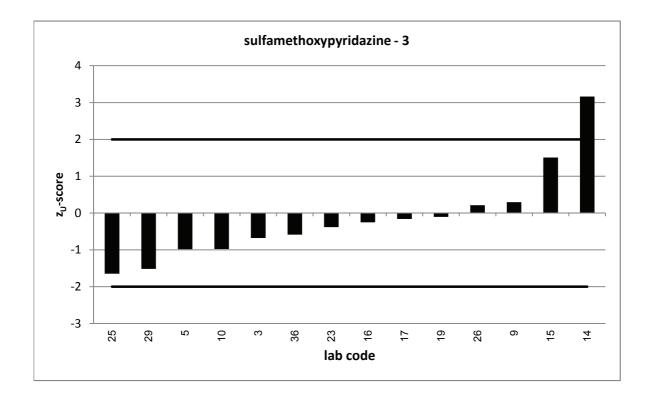


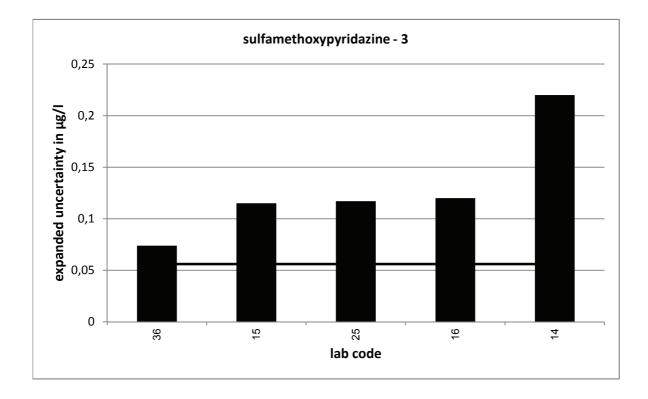


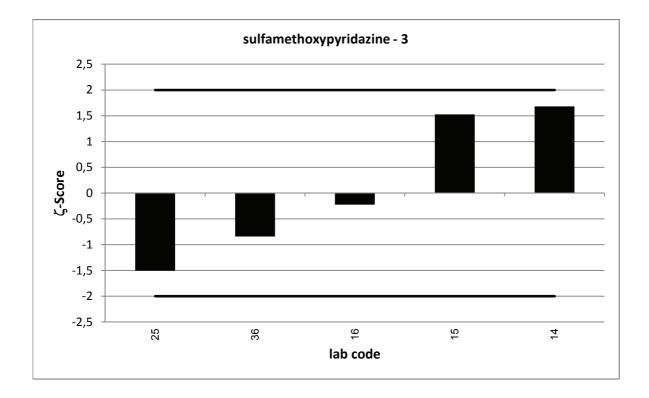


PT 5/16 - TW S2 sulfametho			ethoxvpv	ridazine -	3
assigned va				± 0,0127	_
	nce limit [µg/l]		0,7928		
	nce limit [µg/l]		0,5679		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
3	0,639			-0,7	s
5	0,6227			-1,0	s
9	0,693			0,3	S
10	0,623			-1,0	S
14	0,861	0,22	1,7	3,2	u
15	0,764	0,115	1,5	1,5	S
16	0,662	0,12	-0,2	-0,3	S
17	0,667			-0,2	S
19	0,67			-0,1	S
23	0,655			-0,4	S
25	0,587	0,117	-1,5	-1,6	S
26	0,688			0,2	S
29	0,594			-1,5	S
36	0,644	0,074	-0,8	-0,6	S

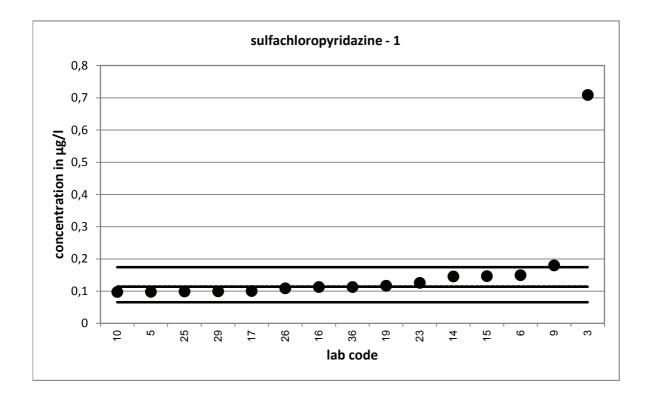


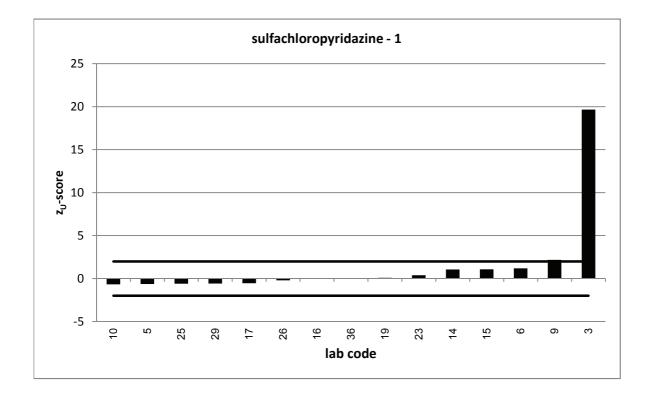


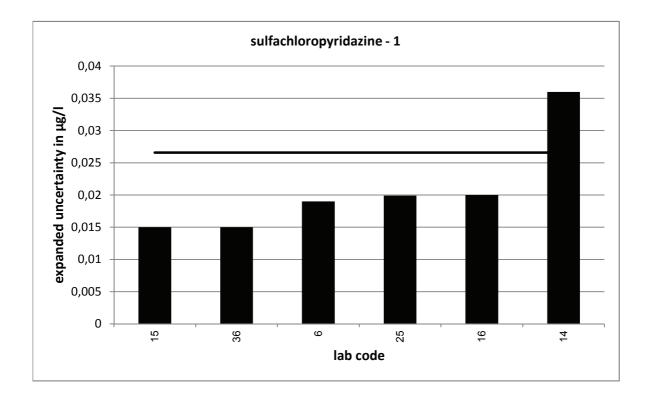


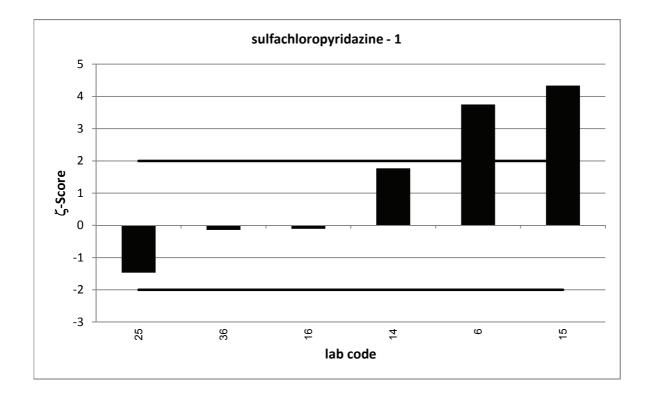


PT 5/16 - T	W S2	sulfact	loropyric	lazine - 1	
assigned va	lue [µg/l]*			± 0,0023	
upper tolera	nce limit [µg/l]		0,1746		
lower tolera	nce limit [µg/l]		0,0656		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
3	0,709			19,7	u
5	0,0985			-0,6	S
6	0,15	0,019	3,8	1,2	S
9	0,18			2,2	q
10	0,098			-0,7	S
14	0,146	0,036	1,8	1,1	S
15	0,147	0,015	4,3	1,1	S
16	0,113	0,02	-0,1	0,0	S
17	0,101			-0,5	S
19	0,117			0,1	S
23	0,126			0,4	S
25	0,0994	0,02	-1,5	-0,6	S
26	0,109			-0,2	S
29	0,1			-0,6	S
36	0,113	0,015	-0,1	0,0	S

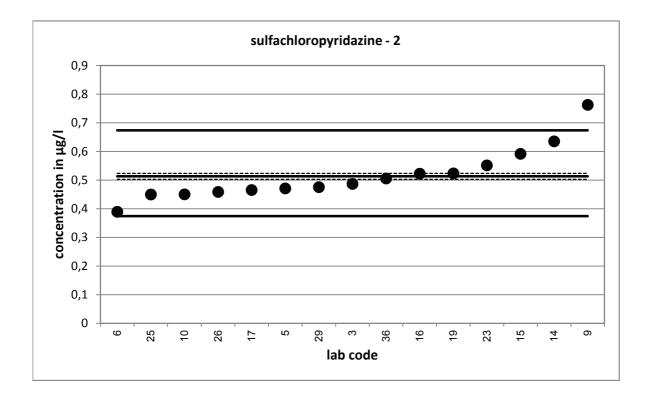


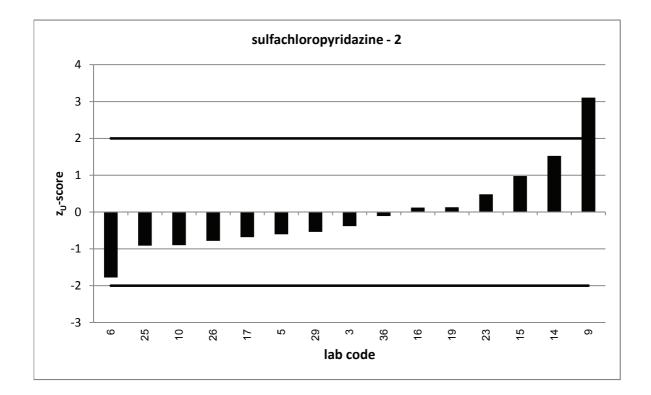


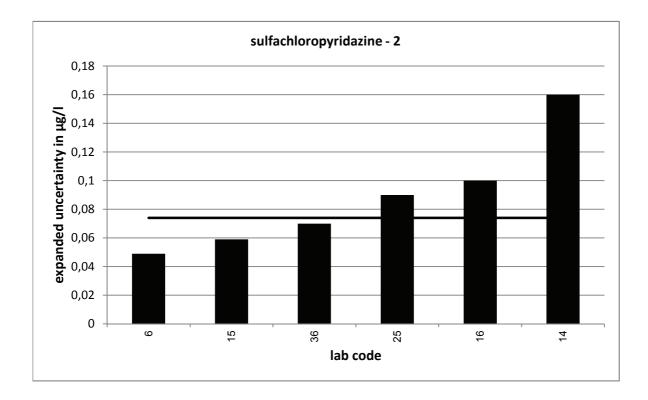


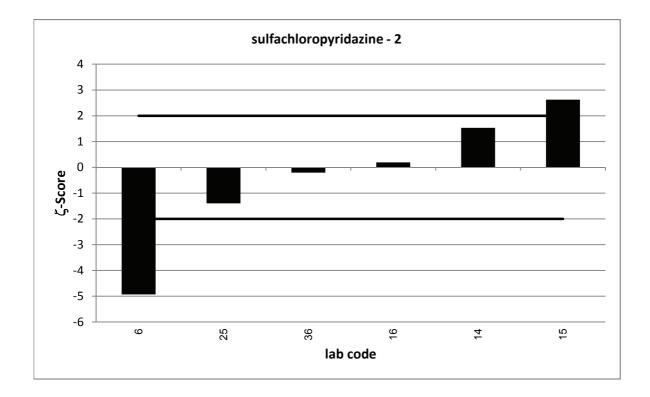


PT 5/16 - T	W S2	sulfach	loropyrid	lazine - 2	
assigned va	lue [µg/l]*			± 0,0104	
upper tolera	nce limit [µg/l]		0,6741		
lower tolera	nce limit [µg/l]		0,3746		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
3	0,487			-0,4	S
5	0,4715			-0,6	S
6	0,39	0,049	-4,9	-1,8	S
9	0,763			3,1	u
10	0,451			-0,9	S
14	0,636	0,16	1,5	1,5	S
15	0,592	0,059	2,6	1,0	S
16	0,523	0,1	0,2	0,1	S
17	0,466			-0,7	S
19	0,524			0,1	S
23	0,552			0,5	S
25	0,45	0,09	-1,4	-0,9	S
26	0,459			-0,8	S
29	0,476			-0,5	S
36	0,506	0,07	-0,2	-0,1	S

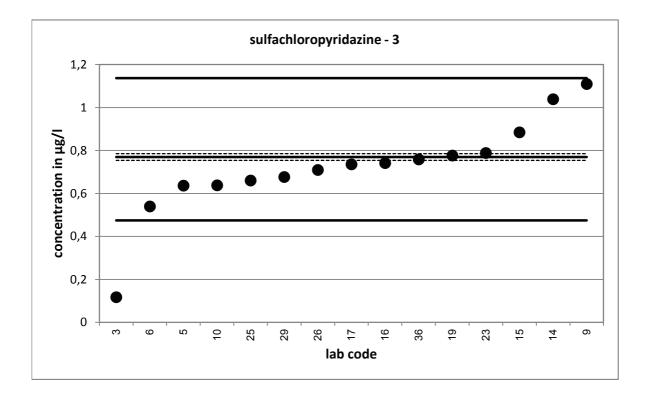


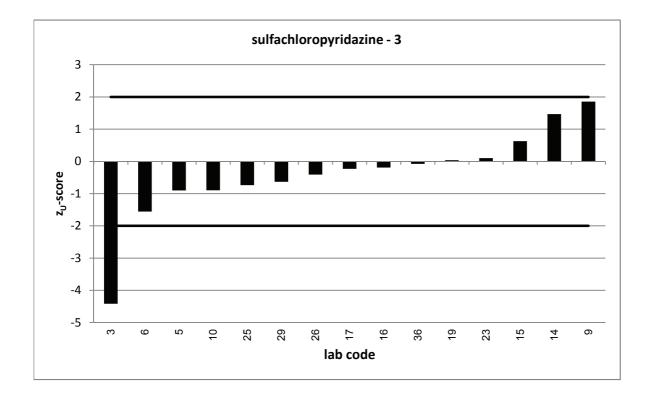


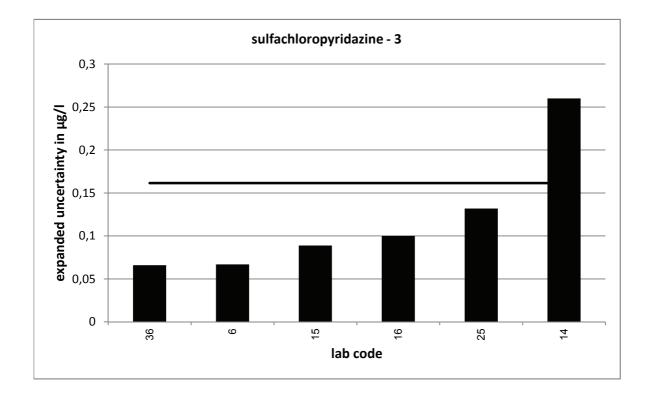


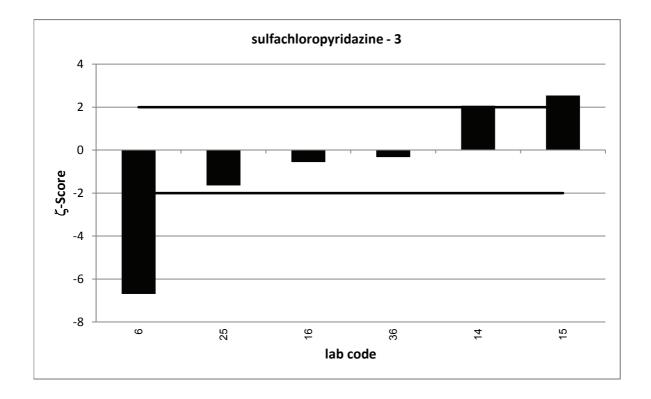


PT 5/16 - T	W S2	sulfach	loropyrid	azine - 3	
assigned va	lue [µg/l]*			± 0,0156	
upper tolera	nce limit [µg/l]		1,137		
lower tolera	nce limit [µg/l]		0,4748		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
3	0,118			-4,4	u
5	0,6369			-0,9	S
6	0,54	0,067	-6,7	-1,6	S
9	1,11			1,9	S
10	0,638			-0,9	S
14	1,039	0,26	2,1	1,5	S
15	0,885	0,089	2,5	0,6	S
16	0,742	0,1	-0,6	-0,2	S
17	0,736			-0,2	S
19	0,777			0,0	S
23	0,789			0,1	S
25	0,661	0,132	-1,6	-0,7	S
26	0,71			-0,4	S
29	0,677			-0,6	S
36	0,759	0,066	-0,3	-0,1	S

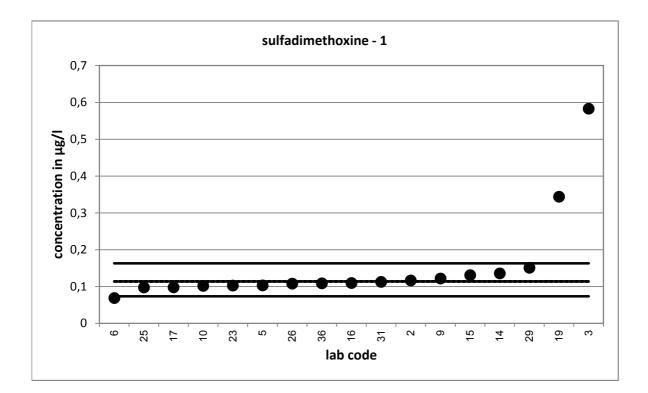


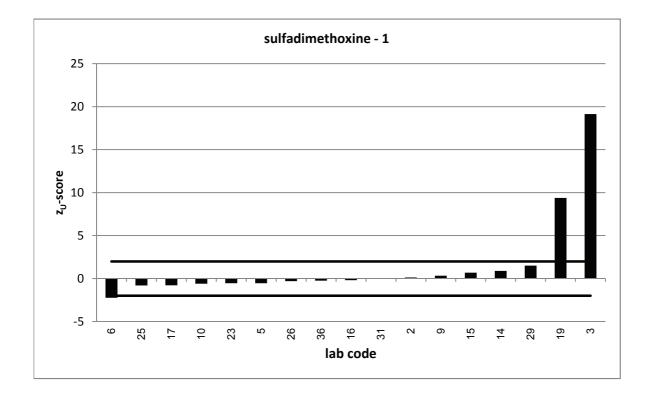


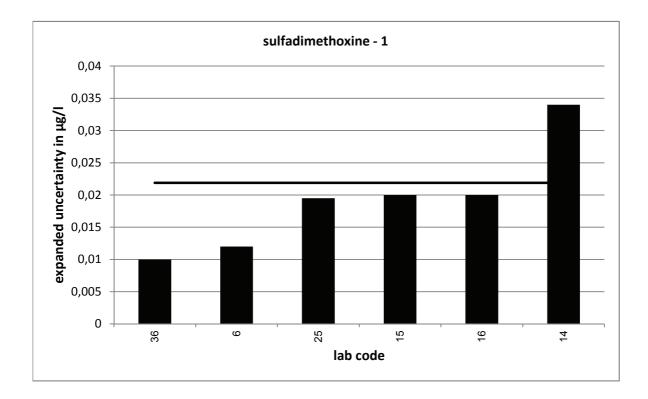


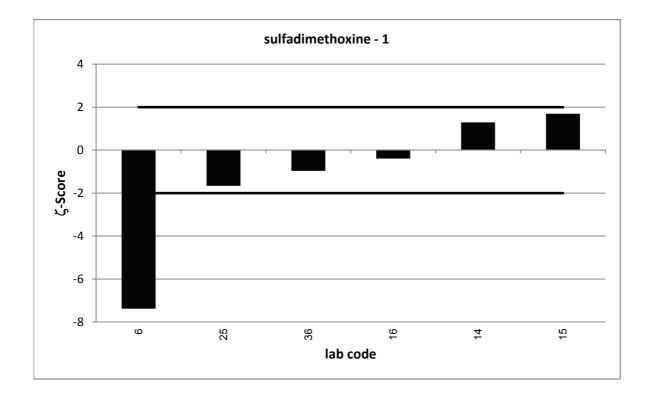


PT 5/16 - T	PT 5/16 - TW S2 sulfadimethox			e - 1	
assigned va	lue [µg/l]*		0,1139	± 0,0021	
upper tolera	nce limit [µg/l]		0,1629		
lower tolera	nce limit [µg/l]		0,07365		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
2	0,117			0,1	S
3	0,583			19,1	u
5	0,1032			-0,5	S
6	0,069	0,012	-7,4	-2,2	q
9	0,122			0,3	S
10	0,102			-0,6	S
14	0,136	0,034	1,3	0,9	S
15	0,131	0,02	1,7	0,7	S
16	0,11	0,02	-0,4	-0,2	S
17	0,098			-0,8	S
19	0,344			9,4	u
23	0,103			-0,5	S
25	0,0977	0,02	-1,7	-0,8	S
26	0,108			-0,3	S
29	0,151			1,5	S
31	0,113			0,0	S
36	0,109	0,01	-1,0	-0,2	S

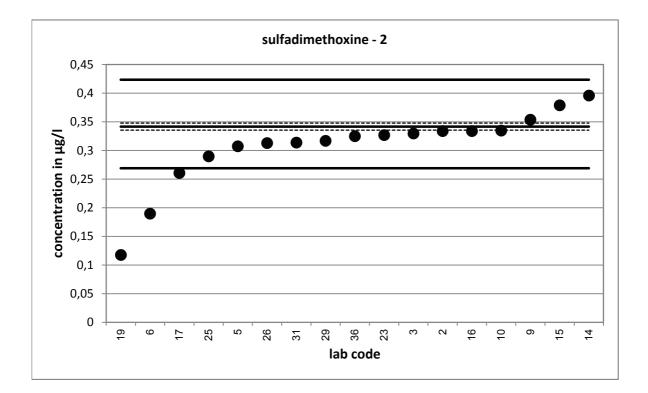


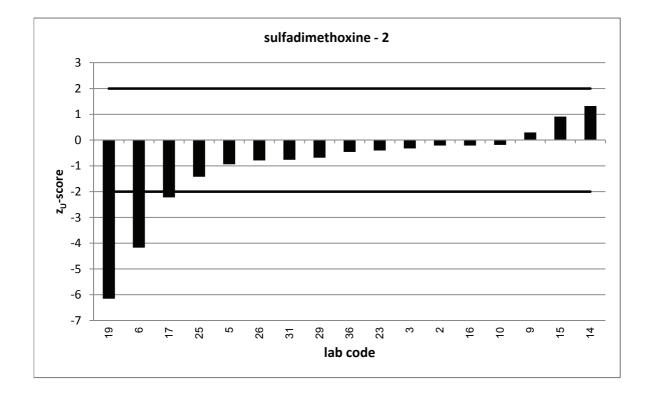


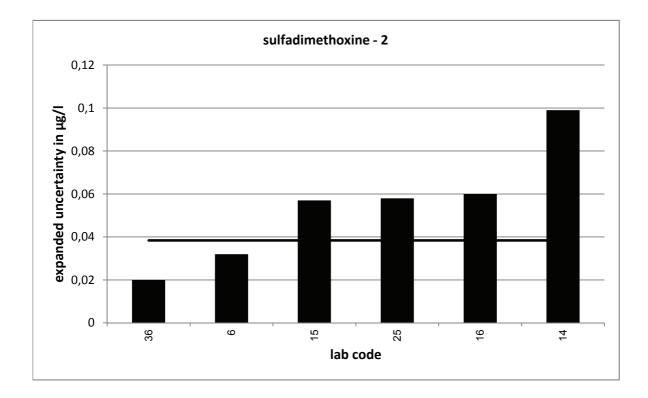


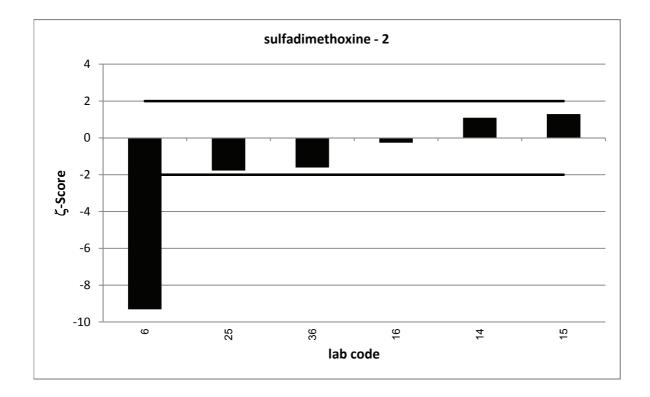


PT 5/16 - T	5/16 - TW S2 sulfadimethoxine			ie - 2	
assigned va	lue [µg/l]*		0,3418	± 0,0062	
upper tolera	nce limit [µg/l]		0,4237		
lower tolera	nce limit [µg/l]		0,269		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
2	0,334			-0,2	S
3	0,33			-0,3	S
5	0,3076			-0,9	S
6	0,19	0,032	-9,3	-4,2	u
9	0,354			0,3	S
10	0,335			-0,2	S
14	0,396	0,099	1,1	1,3	S
15	0,379	0,057	1,3	0,9	S
16	0,334	0,06	-0,3	-0,2	S
17	0,261			-2,2	q
19	0,118			-6,2	u
23	0,327			-0,4	S
25	0,29	0,058	-1,8	-1,4	S
26	0,313			-0,8	S
29	0,317			-0,7	S
31	0,314			-0,8	S
36	0,325	0,02	-1,6	-0,5	S

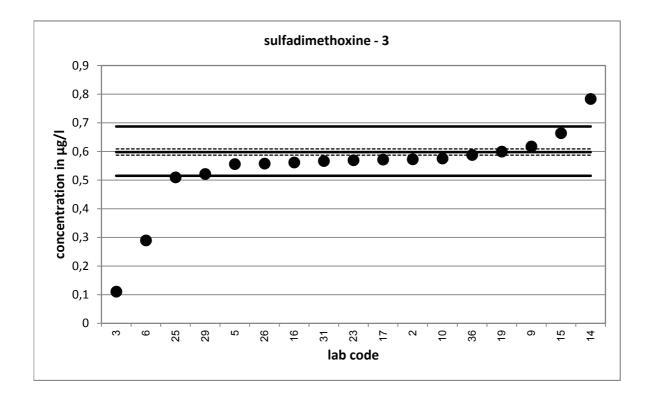


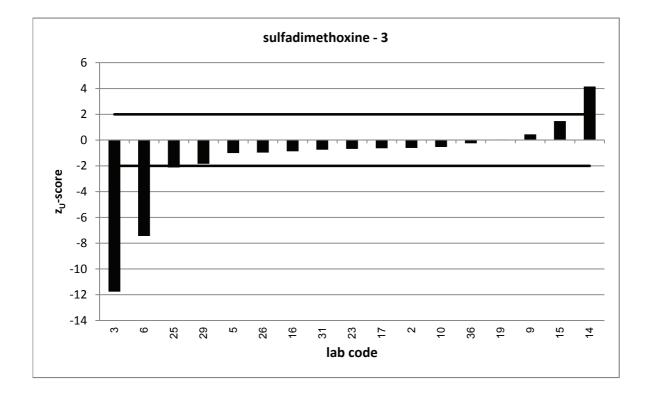


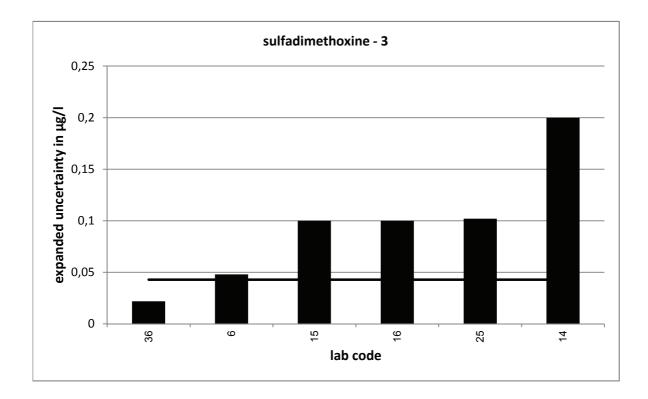


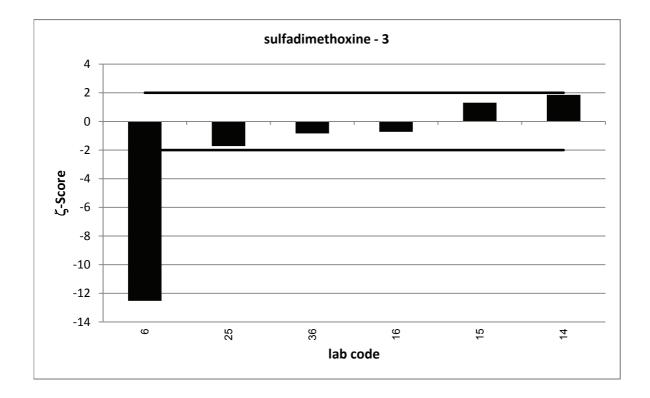


PT 5/16 - T	PT 5/16 - TW S2 sulfadimethoxir			ie - 3	
assigned va	llue [µg/l]*		0,5982	± 0,0109	
upper tolera	nce limit [µg/l]		0,6875		
lower tolera	nce limit [µg/l]		0,5153		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
2	0,573			-0,6	S
3	0,111			-11,8	u
5	0,5562			-1,0	S
6	0,29	0,048	-12,5	-7,4	u
9	0,618			0,4	S
10	0,576			-0,5	S
14	0,784	0,2	1,9	4,2	u
15	0,664	0,1	1,3	1,5	S
16	0,562	0,1	-0,7	-0,9	S
17	0,572			-0,6	S
19	0,6			0,0	S
23	0,57			-0,7	S
25	0,51	0,102	-1,7	-2,1	q
26	0,558			-1,0	S
29	0,522			-1,8	S
31	0,567			-0,8	S
36	0,588	0,022	-0,8	-0,2	S

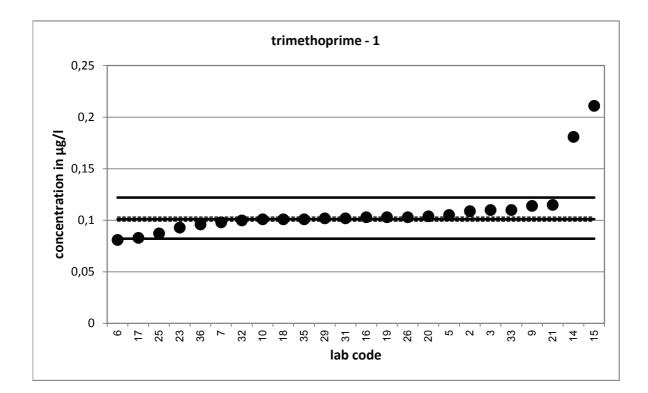


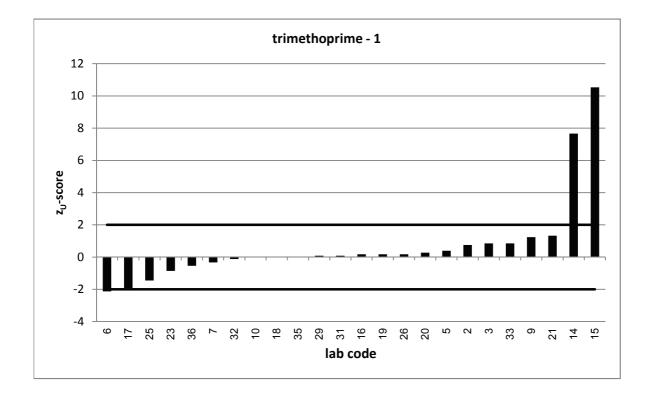


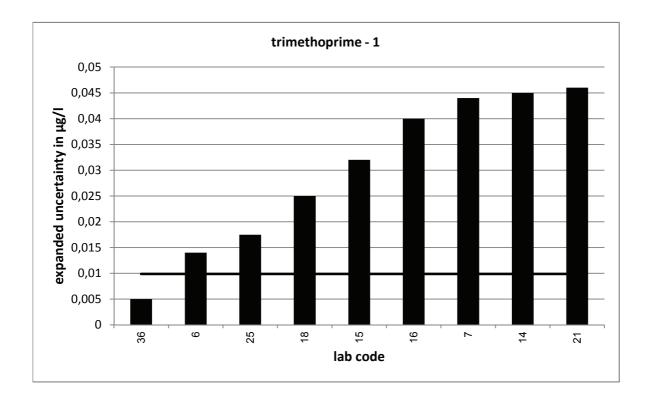




PT 5/16 - T	W S2	trimeth	oprime -	1		
assigned va	lue [µg/l]*		0,1011		± 0,0019	
upper tolera	nce limit [µg/l]		0,122			
	nce limit [µg/l]		0,08222			
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**	
2	0,109			0,8	S	
3	0,11			0,9	S	
5	0,1052			0,4	S	
6	0,081	0,014	-2,9	-2,1	q	
7	0,098	0,044	-0,1	-0,3	S	
9	0,114			1,2	S	
10	0,101			0,0	S	
14	0,181	0,045	3,5	7,7	u	
15	0,211	0,032	6,9	10,5	u	
16	0,103	0,04	0,1	0,2	S	
17	0,083			-1,9	S	
18	0,101	0,025	0,0	0,0	S	
19	0,103			0,2	S	
20	0,104			0,3	S	
21	0,115	0,046	0,6	1,3	S	
23	0,093			-0,9	S	
25	0,0874	0,018	-1,6	-1,5	S	
26	0,103			0,2	S	
29	0,102			0,1	S	
31	0,102			0,1	S	
32	0,1			-0,1	s	
33	0,11			0,9	S	
35	0,101			0,0	S	
36	0,096	0,005	-1,9	-0,5	S	

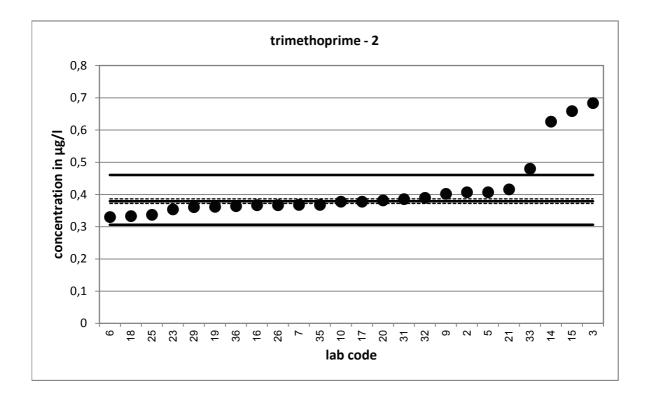


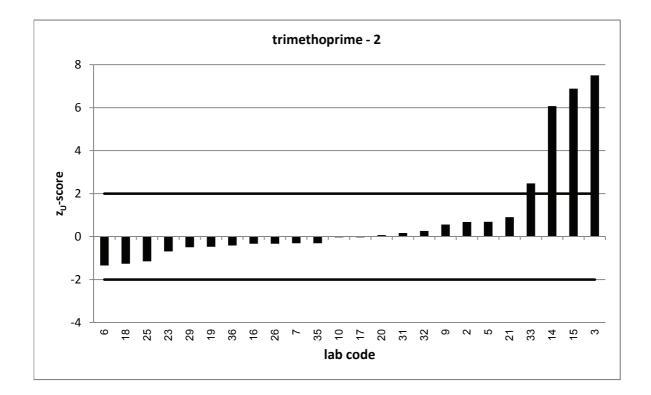


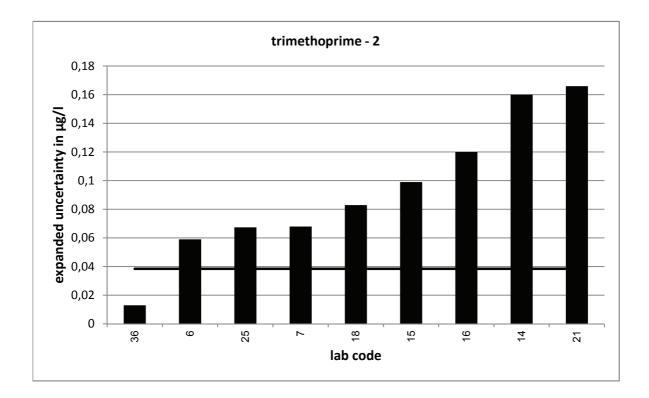


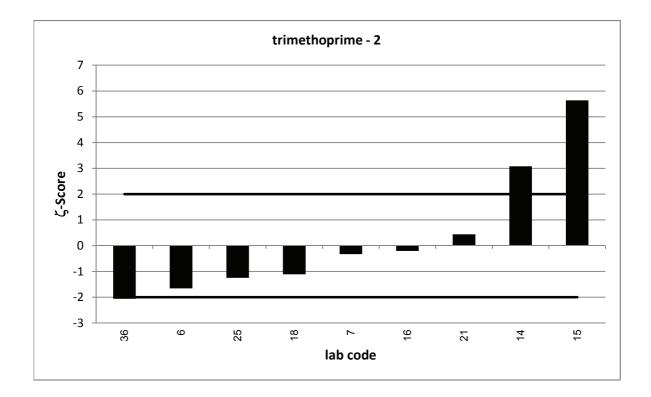


PT 5/16 - T	W S2	trimeth	oprime -	2	
assigned va	alue [µg/l]*	0,3793		± 0,0071	
	ance limit [µg/l]		0,4605		
	nce limit [µg/l]		0,3059		
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**
2	0,407			0,7	S
3	0,684			7,5	u
5	0,4073			0,7	S
6	0,33	0,059	-1,7	-1,3	S
7	0,368	0,068	-0,3	-0,3	S
9	0,402			0,6	S
10	0,378			0,0	S
14	0,626	0,16	3,1	6,1	u
15	0,659	0,099	5,6	6,9	u
16	0,367	0,12	-0,2	-0,3	S
17	0,378			0,0	S
18	0,333	0,083	-1,1	-1,3	S
19	0,362			-0,5	S
20	0,382			0,1	S
21	0,416	0,166	0,4	0,9	S
23	0,354			-0,7	S
25	0,337	0,067	-1,2	-1,2	S
26	0,367			-0,3	S
29	0,361			-0,5	S
31	0,386			0,2	S
32	0,39			0,3	S
33	0,48			2,5	q
35	0,368			-0,3	S
36	0,364	0,013	-2,1	-0,4	S









PT 5/16 - T	W S2	trimeth	oprime -	3	3	
assigned va	alue [µg/l]*	ug/l]* 0,6827		± 0,0128		
	ance limit [µg/l]		0,8509			
lower tolera	nce limit [µg/l]		0,5335			
lab code	result [µg/l]	±	ζ-score	z _u -score	assessm.**	
2	0,635			-0,6	s	
3	0,368			-4,2	u	
5	0,7383			0,7	S	
6	0,49	0,087	-4,4	-2,6	q	
7	0,705	0,194	0,2	0,3	S	
9	0,721			0,5	S	
10	0,669			-0,2	S	
14	1	0,25	2,5	3,8	u	
15	1,064	0,16	4,8	4,5	u	
16	0,664	0,22	-0,2	-0,3	S	
17	0,533			-2,0	S	
18	0,684	0,171	0,0	0,0	S	
19	0,649			-0,5	S	
20	0,699			0,2	S	
21	0,709	0,284	0,2	0,3	S	
23	0,621			-0,8	S	
25	0,603	0,121	-1,3	-1,1	S	
26	0,671			-0,2	S	
29	0,551			-1,8	S	
31	0,716			0,4	S	
32	0,65			-0,4	S	
33	0,72			0,4	S	
35	0,706			0,3	S	
36	0,646	0,034	-2,0	-0,5	S	

