

Evaluation Report

proficiency test

DLA ptALS2 (2020)

Allergen-Screening II:

Crustaceae, Egg, Fish, Milk, Molluscs, Mustard and Soybean

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| Status des EP-Bericht Status of PT-Report | Abschlussbericht / Final report (26 October 2020) Gültig ist die jeweils letzte Version/Korrektur des Berichts. Sie ersetzt alle vorangegangenen Versionen. Only the latest version/correction of the report is valid. It replaces all preceding versions. | | | | |
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| Unteraufträge Subcontractors | Im Rahmen dieser Eignungsprüfung wurden nachstehende Leistungen im Unterauftrag vergeben: Proteinbestimmung As part of the present proficency test the following services were subcontracted: protein determination | | | | |
| Vertraulichkeit Confidentiality | Die Teilnehmerergebnisse sind im EP-Bericht in anonymisierter Form mit Auswertenummern benannt. Daten einzelner Teilnehmer werden ausschließlich nach vorheriger Zustimmung des Teilnehmers an Dritte weitergegeben. Participant result are named anonymously with evaluation numbers in the PT report. Data of individual participants will be passed on to third parties only with prior consent of the participant. | | | | |

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1. Introduction

The participation in proficiency testing schemes is an essential element of the quality-management-system of every laboratory testing food and feed, cosmetics and food contact materials. The implementation of proficiency tests enables the participating laboratories to prove their own analytical competence under realistic conditions. At the same time they receive valuable data regarding the verification and/or validation of the particular testing method [1, 5].

The purpose of DLA is to offer proficiency tests for selected parameters in concentrations with practical relevance.

Realisation and evaluation of the present proficiency test follows the technical requirements of DIN EN ISO/IEC 17043 (2010) and DIN ISO 13528:2009 / ISO 13528:2015 [2, 3].

2. Realisation

2.1 Test material

Four PT-samples were provided for the qualitative detection of allergens in mg/kg range. To prepare the samples premixes were used at levels of about 5-10% of the allergenic ingredients concerned.

The respective raw materials for the allergens used were commercial egg powder, milk powder and soy flour and premixes produced by DLA from commercial mustard seeds as well as frozen king prawns, cod and mussels (s. Tab. 2). The mustard seeds were crushed, ground with addition of carrier substances and sieved (mesh 400 $\mu m)$. The frozen products were crushed, freeze dried and ground with addition of carrier substances and sieved by means of a centrifugal mill (mesh 250 $\mu m)$.

The composition of the allergen-premixes is given in table 1. The premixes were used for spiking of the PT-samples 1 to 4 (see Tab. 2).

After homogenisation the samples were portioned to approximately 20 g into metallised PET film bags.

<u>Table 1:</u> Composition of DLA-Samples

| Ingredients | Samples 1 - 4 |
|---|---------------|
| Potato powder (Ingredients: Potatoes, E471, E304, E223, E100) | 74,6 - 74,8 % |
| Maltodextrin | 24,8 - 25,0 % |
| Allergen-Premixes | 0,30 - 0,55 % |
| <pre>Ingredients:</pre> | |
| - Maltodextrin (30% - 88%) | |
| - Titanium dioxide (0,0% - 40%) | |
| - Sodium sulfate (0,0% - 7,7%) | |
| - Silicon dioxide (1,0% - 2,2%) | |
| - Allergens (5,0% - 10% each) | |

 $\underline{\text{Table 2:}}$ Added amounts of allergenic ingredients positive in mg/kg ranges** given as food item

| Ingredients * | Sample 1 | Sample 2 | Sample 3 | Sample 4 |
|---|------------------------|------------------------|------------------------|-------------------------|
| Crustaceae: King Prawns (Litopenaeus vannamei), freeze-dried (Protein 87%) | positive (75 - 150) | negative | positive (25 - 75) | negative |
| Egg: Whole egg powder (Protein 47%) | positive (75 - 150) | positive (25 - 75) | negative | negative |
| Fish: Cod (Gadus morhua), freez-dried (Protein 88%) | negative | positive (75 - 150) | positive (25 - 75) | negative |
| Milk: Skimmed milk powder (Protein 37%) | positive (25 - 75) | negative | negative | positive (100 - 225) |
| Molluscs: Yesso Scallop (Mizuhopecten yessoen-sis), freez-dried (Protein 76%) | negative | negative | positive (25 - 75) | positive (100 - 225) |
| Mustard, yellow: Sinapis alba (Protein 31%) | negative | positive (50 - 100) | negative | positive (50 - 100) |
| Mustard, brown: Brassica juncea (Protein 28%) | negative | negative | positive (50 - 100) | negative |
| Mustard, black: Brassica nigra (Protein 27%) | negative | positive (50 - 100) | negative | negative |
| Soya: Soyflour, not toasted (Protein 37%) | positive (75 - 150) | negative | negative | positive (25 - 75) |

^{*} Protein contents according to laboratory analysis (total nitrogen, Kjeldahl general factor F=6,25

Note: The metrological traceability of temperature, mass and volume during production of the PT samples is ensured by DAkkS calibrated reference materials.

The detectability or absence of the allergens was tested by DLA using lateral flow assays. The results are in agreement with the spiking of the PT samples 1-4 (see Table 3).

<u>Table 3:</u> Verification of detectability of the added allergens by lateral flow assays (AgraStrip $^{\circ}$ LFD, Romer Labs $^{\circ}$)

| Lateral Flow Device (LFD)* | Sample 1 | Sample 2 | Sample 3 | Sample 4 |
|----------------------------|----------|----------|----------|----------|
| AgraStrip® Crustaceae | positive | negative | positive | negative |
| AgraStrip® Egg | positive | positive | negative | negative |
| AgraStrip® Casein | positive | negative | negative | positive |
| AgraStrip® Soy | positive | negative | negative | positive |
| AgraStrip® Mustard | negative | positive | positive | positive |

^{*} Nachweisgrenze (NWG) jeweils 2-10 mg/kg / Limit of detection (LOD) 2-10 mg/kg each

^{**}Allergen contents of "food item" as indicated in the column of ingredients according gravimetric mixing

2.1.1 Homogeneity

The mixture homogeneity before bottling was examined 8-fold by microtracer analysis. It is a standardized method that is part of the international GMP certification system for feed [14].

Before mixing dye coated iron particles of μm size are added to the sample and the number of particles is determined after homogenization in taken aliquots. The evaluation of the mixture homogeneity is based on the Poisson distribution using the chi-square test. A probability of \geq 5 % is equivalent to a good homogeneous mixture and of \geq 25% to an excellent mixture [14, 15].

The microtracer analysis of the present PT samples 1-4 showed probabilities of 99%, 89%, 98% and 99%, respectively. Additionally particle number results were converted into concentrations, statistically evaluated according to normal distribution and compared to the standard deviation according to Horwitz. For the assessment HorRat values between 0,3 and 1,3 are to be accepted under repeat conditions (measurements within the laboratory) [17]. This gave HorRat values of 0,6, 0,8, 0,6 and 0,6, respectively. The results of microtracer analysis are given in the documentation.

2.1.2 Stability

A water activity (a_W) of < 0,5 is an important factor to ensure the stability of dry or dried products during storage. Optimum conditions for storage is the a_W value range of 0,15-0,3. In this range the lowest possible degradation rate is to be expected [16].

The experience with various DLA test materials showed good storage stability with respect to the durability of the sample (spoilage) and the content of the PT parameters for comparable food matrices and water activity (a_W value <0,5).

The a_W value of the PT samples was approx. 0,40 and 0,36 (21-22°C). The stability of the sample material was thus ensured during the investigation period under the specified storage conditions.

2.2 Sample shipment and information to the test

The portions of the test materials (sample 1 to 4) were sent to every participating laboratory in the $25^{\rm th}$ week of 2020. The testing method was optional. The tests should be finished at August $28^{\rm th}$ 2020 the latest (extended).

With the cover letter along with the sample shipment the following information was given to participants:

There are 4 different samples possibly containing the allergenic ingredients Crustaceae, Egg, Fish, Milk, Molluscs, Mustard (yellow/white, brown and black) and/or Soybean in a simple carrier matrix The evaluation of results is strictly qualitative (positive / negative).

The following analysis methods can be used:

- a) ELISA and Lateral Flow
- b) PCR
- C) LC/MS

Please note the attached information on the proficiency test. (see documentation, section 5.3 Information on the PT)

2.3 Submission of results

The participants submitted their results in standard forms, which have been sent by email or were available on our website. The results given as positive/negative were evaluated.

Queried and documented were the indicated results and details of the test methods like specificities, test kit manufacturer and hints about the procedure.

In case participants submitted several results for the same parameter obtained by different methods these results were evaluated with the same evaluation number with a letter as a suffix and indication of the related method.

All 29 participants submitted at least one result in time.

3. Evaluation

Different ELISA- and PCR-methods for the determination of allergens in foods are eventually using different antibodies and target-DNA, are usually calibrated with different reference materials and may utilize differing extraction methods. Among others this can induce different valuation of the presence and/or content of the analyte [25, 26, 27, 28]. Furthermore matrix- and/or processing of samples can have strong impact on the detectability of allergens by ELISA and PCR methods.

Therefore in the present PT the allergenic ingredients were provided for analysis in a simple matrix without further processing.

3.1 Agreement with consensus values from participants

The qualitative evaluation of the ELISA and PCR results of each participant was based on the agreement of the indicated results (positive or negative) with the **consensus values from participants**. A consensus value is determined if \geq 75% positive or negative results are available for a parameter.

The assessment will be in the form that the number of matching results followed by the number of samples for which a consensus value was obtained is indicated. Behind that the agreement is expressed as the percentage in parentheses.

3.2 Agreement with spiking of samples

The qualitative evaluation of the ELISA and PCR results of each participant was based on the agreement of the indicated results (positive or negative) with the **spiking of the four PT-samples**.

The assessment will be in the form that the number of matching results followed by the number of samples is indicated. Behind that the agreement is expressed as the percentage in parentheses.

4. Results

All following tables are anonymized. With the delivering of the evaluation-report the participants are informed about their individual evaluation-number.

The qualitative evaluation is carried out for each parameter for ELISA and PCR methods separately. Results of lateral flow methods were valuated together with ELISA methods, because they are usually based on antibody detection.

The participant results and evaluation are tabulated as follows:

| Evaluation number | Sample 1 | Sample 2 | Sample 3 | Sample 4 | Qualitative Valuation | Qualitative Valuation | Method | Remarks |
|-------------------|----------|----------|----------|----------|-----------------------------------|--------------------------------------|--------|---------|
| | pos/neg | pos/neg | pos/neg | pos/neg | Agreement with consensus value | Agreement with spiking of samples | | |

| | Sample 1 | Sample 2 | Sample 3 | Sample 4 |
|------------------|----------|----------|----------|----------|
| Number positive | | | | |
| Number negative | | | | |
| Percent positive | | | | |
| Percent negative | | | | |
| Consensus value | | | | |
| Spiking | | | | |

4.1 Proficiency Test Crustaceae

4.1.1 ELISA-Results: Crustaceae (King Prawns)

Qualitative valuation of results

| Evaluation number | Sample 1 | Sample 2 | Sample 3 | Sample 4 | Qualitative Valuation | Qualitative Valuation | Method | Remarks |
|-------------------|----------|----------|----------|----------|--------------------------------|--------------------------------------|--------|---------|
| | pos/neg | pos/neg | pos/neg | pos/neg | Agreement with consensus value | Agreement with spiking of samples | | |
| 8 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | AQ | |
| 18 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | AQ | |
| 5 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | BF | |
| 28 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | BF | |
| 17 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | IL | |
| 7 | positive | negative | positive | positive | 3/4 (75%) | 3/4 (75%) | RS-F | |
| 9 | positive | positive | positive | positive | 2/4 (50%) | 2/4 (50%) | RS-F | |
| 22 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 27 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 4 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | SP | |
| 12 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | SP | |
| 19 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | SP | |
| 21 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | SP | |

| | Sample 1 | Sample 2 | Sample 3 | Sample 4 |
|------------------|----------|----------|----------|----------|
| Number positive | 13 | 1 | 13 | 2 |
| Number negative | 0 | 12 | 0 | 11 |
| Percent positive | 100 | 8 | 100 | 15 |
| Percent negative | 0 | 92 | 0 | 85 |
| Consensus value | positive | negative | positive | negative |
| Spiking | positive | negative | positive | negative |

Methods:

AQ = AgraQuant, RomerLabs

BF = MonoTrace ELISA, BioFront Technologies

IL = Immunolab

RS-F= Ridascreen® Fast, R-Biopharm

SP = SensiSpec ELISA Kit, Eurofins

Comments:

The consensus values of results are in qualitative agreement with the spiking of samples.

Participant 8 has pointed to a possible cross-reactivity to molluscs for the used ELISA method AQ (see documentation).

Possible cross-reactivities should be documented in the manufacturer's test kit information.

4.1.2 PCR-Results: Crustaceae (King Prawns)

Qualitative valuation of results

| Evaluation number | Sample 1 | Sample 2 | Sample 3 | Sample 4 | Qualitative Valuation | Qualitative Valuation | Method | Remarks |
|-------------------|----------|----------|----------|----------|--------------------------------|--------------------------------------|--------|---------|
| | pos/neg | pos/neg | pos/neg | pos/neg | Agreement with consensus value | Agreement with spiking of samples | | |
| 15 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | ASU | |
| 27 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | ASU | |
| 1 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 2 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 5 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 7 | positive | negative | positive | positive | 3/4 (75%) | 3/4 (75%) | SFA | |
| 10 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 20 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 21 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 23 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 25 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 16 | positive | negative | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA-ID | |

| | Sample 1 | Sample 2 | Sample 3 | Sample 4 |
|------------------|----------|----------|----------|----------|
| Number positive | 12 | 0 | 12 | 1 |
| Number negative | 0 | 12 | 0 | 11 |
| Percent positive | 100 | 0 | 100 | 8 |
| Percent negative | 0 | 100 | 0 | 92 |
| Consensus value | positive | negative | positive | negative |
| Spiking | positive | negative | positive | negative |

Methods:

ASU = ASU §64 Methode/method

SFA = Sure Food ALLERGEN, R-Biopharm / Congen

SFA-ID = Sure Food Allergen ID, R-Biopharm / Congen

Comments:

The consensus values of results are in qualitative agreement with the spiking of samples.

4.2 Proficiency Test Egg

4.2.1 ELISA-Results: Egg (whole egg powder)

Qualitative valuation of results

| Evaluation number | Sample 1 | Sample 2 | Sample 3 | Sample 4 | Qualitative Valuation | Qualitative Valuation | Method | Remarks |
|-------------------|----------|----------|----------|----------|--------------------------------|-----------------------------------|--------|--------------|
| | pos/neg | pos/neg | pos/neg | pos/neg | Agreement with consensus value | Agreement with spiking of samples | | |
| 11 | positive | positive | negative | negative | 4/4 (100%) | 4/4 (100%) | AS | Lateral Flow |
| 28 | positive | positive | negative | negative | 4/4 (100%) | 4/4 (100%) | BF | |
| 14 | positive | positive | negative | negative | 4/4 (100%) | 4/4 (100%) | ES | |
| 17 | positive | positive | negative | negative | 4/4 (100%) | 4/4 (100%) | IL | |
| 3 | positive | positive | negative | negative | 4/4 (100%) | 4/4 (100%) | MI-II | |
| 4 | positive | positive | negative | negative | 4/4 (100%) | 4/4 (100%) | MI-II | |
| 8 | positive | positive | negative | negative | 4/4 (100%) | 4/4 (100%) | MI-II | |
| 12 | positive | positive | negative | negative | 4/4 (100%) | 4/4 (100%) | MI-II | |
| 7 | positive | positive | negative | negative | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 9 | positive | positive | negative | negative | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 20 | positive | positive | negative | negative | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 22 | positive | positive | negative | negative | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 19 | positive | positive | negative | negative | 4/4 (100%) | 4/4 (100%) | SP | |
| 21 | positive | positive | negative | negative | 4/4 (100%) | 4/4 (100%) | SP | |

| | Sample 1 | Sample 2 | Sample 3 | Sample 4 | |
|------------------|----------|------------------------|----------|----------|--|
| Number positive | 14 | 14 | 0 | 0 | |
| Number negative | 0 0 | | 14 | 14 | |
| Percent positive | 100 | 100 | 0 | 0 | |
| Percent negative | 0 | 0 | 100 | 100 | |
| Consensus value | positive | tive positive negative | | negative | |
| Spiking | positive | positive | negative | negative | |

Methods:

AS = AgraStrip (Lateral Flow), RomerLabs

BF = MonoTrace ELISA, BioFront Technologies

ES = ELISA-Systems

IL = Immunolab

MI-II = Morinaga Institute ELISA Kit II

RS-F= Ridascreen® Fast, R-Biopharm

SP = SensiSpec ELISA Kit, Eurofins

Comments:

The consensus values of results are in qualitative agreement with the spiking of samples.

4.2.2 PCR-Results: Egg (whole egg powder)

PCR methods were not applied by the participants.

4.3 Proficiency Test Fish

4.3.1 ELISA-Results: Fish (cod)

Qualitative valuation of results

| Evaluation number | Sample 1 | Sample 2 | Sample 3 | Sample 4 | Qualitative Valuation | Qualitative Valuation | Method | Remarks |
|-------------------|----------|----------|----------|----------|--------------------------------|--------------------------------------|--------|---------|
| | pos/neg | pos/neg | pos/neg | pos/neg | Agreement with consensus value | Agreement with spiking of samples | | |
| 8 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | AQ | |
| 26 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | ВС | |
| 28 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | BF | |
| 17 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | IL | |
| 19 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | SP | |
| 21 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | SP | |

| | Sample 1 | Sample 2 | Sample 3 | Sample 4 |
|------------------|--------------------------------|----------|----------|----------|
| Number positive | 0 | 6 | 6 | 0 |
| Number negative | 6 | 0 | 0 | 6 |
| Percent positive | 0 | 100 | 100 | 0 |
| Percent negative | 100 | 0 | 0 | 100 |
| Consensus value | e negative positive positive r | | negative | |
| Spiking | negative | positive | positive | negative |

Methods:

AQ = AgraQuant, RomerLabs

BC = BioCheck ELISA

BF = MonoTrace ELISA, BioFront Technologies

IL = Immunolab

SP = SensiSpec ELISA Kit, Eurofins

Comments:

The consensus values of results are in qualitative agreement with the spiking of samples.

4.3.2 PCR-Results: Fish (cod)

Qualitative valuation of results

| Evaluation number | Sample 1 | Sample 2 | Sample 3 | Sample 4 | Qualitative Valuation | Qualitative Valuation | Method | Remarks |
|-------------------|----------|----------|----------|----------|--------------------------------|-----------------------------------|--------|---------|
| | pos/neg | pos/neg | pos/neg | pos/neg | Agreement with consensus value | Agreement with spiking of samples | | |
| 15 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | Gl | |
| 12 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | GS | |
| 13 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | IM | |
| 3 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | MS | |
| 1 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 2 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 5 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 7 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 10 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 16 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 20 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 21 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 23 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 24 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 25 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | SFA | |
| 26 | negative | negative | positive | positive | 2/4 (50%) | 2/4 (50%) | SFA | |
| 4 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | div | |
| 9 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | div | |

| | Sample 1 | Sample 2 | Sample 3 | Sample 4 |
|------------------|------------------------------|----------|----------|----------|
| Number positive | 0 | 17 | 18 | 1 |
| Number negative | 18 | 1 | 0 | 17 |
| Percent positive | 0 | 94 | 100 | 6 |
| Percent negative | 100 | 6 | 0 | 94 |
| Consensus value | ralue negative positive posi | | positive | negative |
| Spiking | negative | positive | positive | negative |

Methods:

GI = GEN-IAL First Allergen

GS = Eurofins Genescan DNA nimal screen

IM = Imegen

MS = Microsynth

SFA = Sure Food ALLERGEN, R-Biopharm / Congen

div = keine genaue Angabe / andere Methode

div = not indicated / other method

Comments:

The consensus values of results are in qualitative agreement with the spiking of samples.

4.4 Proficiency Test Milk

4.4.1 ELISA-Results: Milk, Casein, β-Lactoglobulin

Qualitative valuation of results

| Evaluation number | Sample 1 | Sample 2 | Sample 3 | Sample 4 | Qualitative Valuation | Qualitative Valuation | Method | Remarks |
|-------------------|----------|----------|----------|----------|--------------------------------|--------------------------------------|--------|------------------------------|
| | pos/neg | pos/neg | pos/neg | pos/neg | Agreement with consensus value | Agreement with spiking of samples | | |
| 28 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | BF | |
| 14a | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | ES | Casein |
| 14b | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | ES | β-Lactoglobulin |
| 17 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | IL | |
| 4 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | MI-II | Casein |
| 8a | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | MI-II | Casein |
| 8b | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | MI-II | β-Lactoglobulin |
| 1 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | RS div | R-Biopharm Kit not specified |
| 3 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 7 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | RS-F | Casein |
| 9 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 22 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | RS-F | β-Lactoglobulin |
| 20a | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | RS-F | Casein |
| 20b | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | RS-F | β-Lactoglobulin |
| 12 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | SP | Casein |
| 19 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | SP | |
| 21 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | SP | |

| | Sample 1 | Sample 2 | Sample 3 | Sample 4 |
|------------------|-------------------------------|----------|----------|----------|
| Number positive | 17 | 0 | 0 | 17 |
| Number negative | 0 | 17 | 17 | 0 |
| Percent positive | 100 | 0 | 0 | 100 |
| Percent negative | 0 | 100 | 100 | 0 |
| Consensus value | ue positive negative negative | | positive | |
| Spiking | positive | negative | negative | positive |

Methods:

BF = MonoTrace ELISA, BioFront Technologies

ES = ELISA-Systems

IL = Immunolab

MI-II = Morinaga Institute ELISA Kit II

RS div= R-Biopharm

 ${\sf RS\text{-}F\text{=}Ridascreen} \& {\sf Fast}, \, {\sf R\text{-}Biopharm}$

SP = SensiSpec ELISA Kit, Eurofins

Comments:

The consensus values of results are in qualitative agreement with the spiking of samples.

4.4.2 PCR-Results: Milk (skimmed milk powder)

PCR methods were not applied by the participants.

4.5 Proficiency Test Molluscs

4.5.1 ELISA-Results: Molluscs (yesso scallop)

Qualitative valuation of results

| Evaluation number | Sample 1 | Sample 2 | Sample 3 | Sample 4 | Qualitative Valuation | Qualitative Valuation | Method | Remarks |
|-------------------|----------|----------|----------|----------|--------------------------------|--------------------------------------|--------|--|
| | pos/neg | pos/neg | pos/neg | pos/neg | Agreement with consensus value | Agreement with spiking of samples | | |
| 26 | positive | negative | positive | positive | 3/3 (100%) | 3/4 (75%) | DE | |
| 8 | positive | negative | positive | positive | 3/3 (100%) | 3/4 (75%) | ET | Sample 1 positive due to cross reactivity to crustacea suspected |
| 17 | negative | negative | positive | positive | 3/3 (100%) | 4/4 (100%) | IL | |
| 4 | - | negative | positive | positive | 3/3 (100%) | 3/3 (100%) | SP | Sample 1 traces at limit of detection |
| 19 | positive | negative | positive | positive | 3/3 (100%) | 3/4 (75%) | SP | Sample 1 and 3 positive due to cross reactivity to crustacea |
| 21 | negative | negative | positive | positive | 3/3 (100%) | 4/4 (100%) | SP | |

| | Sample 1 | Sample 2 | Sample 3 | Sample 4 |
|------------------|----------|----------|----------|----------|
| Number positive | 3 | 0 | 6 | 6 |
| Number negative | 2 | 6 | 0 | 0 |
| Percent positive | 60 | 0 | 100 | 100 |
| Percent negative | 40 | 100 | 0 | 0 |
| Consensus value | none | negative | positive | positive |
| Spiking | negative | negative | positive | positive |

Methods:

DE = Demeditec ELISA

ET = Elution Technologies ELISA Kit

IL = Immunolab

SP = SensiSpec ELISA Kit, Eurofins

Comments:

The consensus values of sample 2, 3 and 4 are in qualitative agreement with the spiking of samples. For sample 1 (without addition of molluscs) no consensus value with $\geq 75\%$ positive or negative results was obtained. Two participants have pointed to a possible cross-reactivity to Crustaceae (methods ET and SP). Samples 1 and 3 contain king prawns. Possible cross-reactivities should be documented in the manufacturer's test kit information.

4.5.2 PCR-Results: Molluscs (yesso scallop)

Qualitative valuation of results

| Evaluation number | Sample 1 | Sample 2 | Sample 3 | Sample 4 | Qualitative Valuation | Qualitative Valuation | Method | Remarks |
|-------------------|----------|----------|----------|----------|--------------------------------|--------------------------------------|--------|---------|
| | pos/neg | pos/neg | pos/neg | pos/neg | Agreement with consensus value | Agreement with spiking of samples | | |
| 22 | negative | negative | positive | positive | 4/4 (100%) | 4/4 (100%) | 4L | |
| 1 | negative | negative | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 2 | negative | negative | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 5 | negative | negative | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 7 | negative | negative | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 10 | negative | negative | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 13 | negative | negative | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 18 | negative | negative | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 20 | negative | negative | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 21 | negative | negative | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 25 | negative | negative | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 26 | negative | negative | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 9 | negative | negative | positive | positive | 4/4 (100%) | 4/4 (100%) | div | |

| | Sample 1 | Sample 2 | Sample 3 | Sample 4 |
|------------------|----------------------------|----------|----------|----------|
| Number positive | 0 | 0 | 13 | 13 |
| Number negative | 13 | 13 | 0 | 0 |
| Percent positive | 0 | 0 | 100 | 100 |
| Percent negative | 100 | 100 | 0 | 0 |
| Consensus value | negative negative positive | | positive | |
| Spiking | negative | negative | positive | positive |

Methods:

4L = 4LAB Diagnostics

SFA = Sure Food ALLERGEN, R-Biopharm / Congen

div = keine genaue Angabe / andere Methode

div = not indicated / other method

Comments:

The consensus values of results are in qualitative agreement with the $spiking\ of\ samples.$

4.6 Proficiency Test Mustard

4.6.1 ELISA-Results: Mustard, in general

Qualitative valuation of results

| Evaluation number | Sample 1 | Sample 2 | Sample 3 | Sample 4 | Qualitative Valuation | Qualitative Valuation | Method | Remarks |
|-------------------|----------|----------|----------|----------|--------------------------------|--------------------------------------|--------|--------------|
| | pos/neg | pos/neg | pos/neg | pos/neg | Agreement with consensus value | Agreement with spiking of samples | | |
| 11 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | AS | Lateral Flow |
| 28 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | BF | |
| 17 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | IL | |
| 7 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 10 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 18 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 20 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 12 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | SP | |
| 19 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | SP | |
| 21 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | SP | |
| 4 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | VT | |
| 8 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | VT | |

| | Sample 1 | Sample 2 | Sample 3 | Sample 4 |
|------------------|----------|----------|----------|----------|
| Number positive | 0 | 12 | 12 | 12 |
| Number negative | 12 | 0 | 0 | 0 |
| Percent positive | 0 | 100 | 100 | 100 |
| Percent negative | 100 | 0 | 0 | 0 |
| Consensus value | negative | positive | positive | positive |
| Spiking | negative | positive | positive | positive |

Methods:

AS = AgraStrip (Lateral Flow), RomerLabs
BF = MonoTrace ELISA, BioFront Technologies

IL = Immunolab

RS-F= Ridascreen® Fast, R-Biopharm

SP = SensiSpec ELISA Kit, Eurofins

VT = Veratox, Neogen

Comments:

The consensus values of results are in qualitative agreement with the spiking of samples (sample 2 and 4 yellow mustard, sample 2 black mustard and sample 3 brown mustard).

4.6.2 PCR-Results: Mustard

Qualitative valuation of results

4.6.2.1 Mustard, in general

| Evaluation number | Sample 1 | Sample 2 | Sample 3 | Sample 4 | Qualitative Valuation | Qualitative Valuation | Method | Remarks |
|-------------------|----------|----------|----------|----------|--------------------------------|--------------------------------------|--------|---------|
| | pos/neg | pos/neg | pos/neg | pos/neg | Agreement with consensus value | Agreement with spiking of samples | | |
| 4 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | ASU | |
| 1 | positive | positive | positive | positive | 3/4 (75%) | 3/4 (75%) | SFA | |
| 2 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 5 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 8 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 16 | negative | - | - | - | 1/1 (100%) | 1/1 (100%) | SFA | |
| 20 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 21 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 29 | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | SFA | |

| | Sample 1 | Sample 2 | Sample 3 | Sample 4 |
|------------------|----------------|-------------------|----------|----------|
| Number positive | 1 | 8 | 8 | 8 |
| Number negative | per negative 8 | | 0 | 0 |
| Percent positive | 11 | 100 | 100 | 100 |
| Percent negative | 89 | 0 | 0 | 0 |
| Consensus value | negative | positive positive | | positive |
| Spiking | negative | positive | positive | positive |

Methods:

ASU = ASU §64 Methode/method SFA = Sure Food ALLERGEN, R-Biopharm / Congen

Comments:

Some participants used PCR methods for the detection of mustard without differentiating the varieties.

The consensus values of results are in qualitative agreement with the spiking of samples (sample 2 and 4 yellow mustard, sample 2 black mustard and sample 3 brown mustard).

4.6.2.2 Mustard, yellow (Sinapis alba)

| Evaluation number | Sample 1 | Sample 2 | Sample 3 | Sample 4 | Qualitative Valuation | Qualitative Valuation | Method | Remarks |
|-------------------|----------|----------|----------|----------|--------------------------------|--------------------------------------|--------|---------|
| | pos/neg | pos/neg | pos/neg | pos/neg | Agreement with consensus value | Agreement with spiking of samples | | |
| 6 | negative | positive | negative | positive | 4/4 (100%) | 4/4 (100%) | ASU | |
| 27 | negative | positive | negative | positive | 4/4 (100%) | 4/4 (100%) | ASU | |
| 15 | negative | positive | negative | positive | 4/4 (100%) | 4/4 (100%) | Gl | |
| 3 | negative | positive | negative | positive | 4/4 (100%) | 4/4 (100%) | MS | |
| 9 | negative | positive | negative | positive | 4/4 (100%) | 4/4 (100%) | div | |

| | Sample 1 | Sample 2 | Sample 3 | Sample 4 |
|------------------|----------|----------|----------|----------|
| Number positive | 0 | 5 | 0 | 5 |
| Number negative | 5 | 0 | 5 | 0 |
| Percent positive | 0 | 100 | 0 | 100 |
| Percent negative | 100 | 0 | 100 | 0 |
| Consensus value | negative | positive | negative | positive |
| Spiking | negative | positive | negative | positive |

Methods:

ASU = ASU §64 Methode/method

GI = GEN-IAL First Allergen

MS = Microsynth

div = keine genaue Angabe / andere Methode

div = not indicated / other method

Comments:

Five participants tested for mustard species by PCR. Yellow mustard (Sinapis alba) was detected in sample 2 and 4 by all of them. The consensus values of results are in qualitative agreement with the spiking of samples.

4.6.2.3 Mustard, brown and black (Brassica juncea / nigra)

| Evaluation number | Sample 1 | Sample 2 | Sample 3 | Sample 4 | Qualitative Valuation | Qualitative Valuation | Method | Remarks |
|-------------------|----------|----------|----------|----------|--------------------------------|--------------------------------------|--------|---------|
| | pos/neg | pos/neg | pos/neg | pos/neg | Agreement with consensus value | Agreement with spiking of samples | | |
| 27 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | ASU | |
| 15 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | GI | |
| 3a | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | MS | |
| 3b | negative | positive | positive | positive | 4/4 (100%) | 4/4 (100%) | MS | |
| 9 | negative | positive | positive | negative | 4/4 (100%) | 4/4 (100%) | div | |

| | Sample 1 | Sample 2 | Sample 3 | Sample 4 |
|------------------|----------|----------|----------|----------|
| Number positive | 0 | 5 | 5 | 1 |
| Number negative | 5 | 0 | 0 | 4 |
| Percent positive | 0 | 100 | 100 | 20 |
| Percent negative | 100 | 0 | 0 | 80 |
| Consensus value | negative | positive | positive | negative |
| Spiking | negative | positive | positive | negative |

Methods:

ASU = ASU §64 Methode/method

GI = GEN-IAL First Allergen

MS = Microsynth

div = keine genaue Angabe / andere Methode

div = not indicated / other method

<u>Comments:</u>

Moreover four participants detected Brassica species in sample 2 (containing black mustard, Brassica nigra) and sample 3 (containing brown mustard, Brassica juncea). One participant also obtained a positive result for sample 4.

The consensus values of results are in qualitative agreement with the spiking of samples.

4.7 Proficiency Test Soya

4.7.1 ELISA-Results: Soya (soyflour)

Qualitative valuation of results

| Evaluation number | Sample 1 | Sample 2 | Sample 3 | Sample 4 | Qualitative Valuation | Qualitative Valuation | Method | Remarks |
|-------------------|----------|----------|----------|----------|--------------------------------|--------------------------------------|--------|--------------|
| | pos/neg | pos/neg | pos/neg | pos/neg | Agreement with consensus value | Agreement with spiking of samples | | |
| 11 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | AS | Lateral Flow |
| 28 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | BF | |
| 14 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | ES | |
| 17 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | IL | |
| 19 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | IL | |
| 4 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | MI-II | |
| 8 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | MI-II | |
| 7 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 9 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 12 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 20 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 27 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | RS-F | |
| 21 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | SP | |

| | Sample 1 | Sample 2 | Sample 3 | Sample 4 |
|------------------|----------|----------|----------|----------|
| Number positive | 13 | 0 | 0 | 13 |
| Number negative | 0 | 13 | 13 | 0 |
| Percent positive | 100 | 0 | 0 | 100 |
| Percent negative | 0 | 100 | 100 | 0 |
| Consensus value | positive | negative | negative | positive |
| Spiking | positive | negative | negative | positive |

Methods:

AS = AgraStrip (Lateral Flow), RomerLabs

BF = MonoTrace ELISA, BioFront Technologies

ES = ELISA-Systems

IL = Immunolab

MI-II = Morinaga Institute ELISA Kit II

RS-F= Ridascreen® Fast, R-Biopharm

SP = SensiSpec ELISA Kit, Eurofins

<u>Comments:</u>

The consensus values of results are in qualitative agreement with the spiking of samples.

4.7.2 PCR-Results: Soya (soyflour)

Qualitative valuation of results

| Evaluation number | Sample 1 | Sample 2 | Sample 3 | Sample 4 | Qualitative Valuation | Qualitative Valuation | Method | Remarks |
|-------------------|----------|----------|----------|----------|--------------------------------|--------------------------------------|--------|---------|
| | pos/neg | pos/neg | pos/neg | pos/neg | Agreement with consensus value | Agreement with spiking of samples | | |
| 6 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | ASU | |
| 27 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | ASU | |
| 15 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | GI | |
| 3 | negative | negative | negative | positive | 3/4 (75%) | 3/4 (75%) | MS | |
| 1 | positive | positive | positive | positive | 2/4 (50%) | 2/4 (50%) | SFA | |
| 2 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 5 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 16 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 20 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 21 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | SFA | |
| 4 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | div | |
| 9 | positive | negative | negative | positive | 4/4 (100%) | 4/4 (100%) | div | |

| | Sample 1 | Sample 2 | Sample 3 | Sample 4 |
|------------------|----------|----------|----------|----------|
| Number positive | 11 | 1 | 1 | 12 |
| Number negative | 1 | 11 | 11 | 0 |
| Percent positive | 92 | 8 | 8 | 100 |
| Percent negative | 8 | 92 | 92 | 0 |
| Consensus value | positive | negative | negative | positive |
| Spiking | positive | negative | negative | positive |

Methods:

ASU = ASU §64 Methode/method GI = GEN-IAL First Allergen

MS = Microsynth

SFA = Sure Food ALLERGEN, R-Biopharm / Congen

div = keine genaue Angabe / andere Methode

div = not indicated / other method

<u>Comments:</u>

The consensus values of results are in qualitative agreement with the $spiking\ of\ samples.$

5. Documentation

5.1 Details by the participants

 $\underline{\text{Note:}}$ Information given in German was translated by DLA to the best of our knowledge (without guarantee of correctness).

5.1.1 ELISA: Crustaceae

Primary data

| Meth. Abbr. | Evaluation number | Date of analysis | Result Sample 1 | Result Sample 2 | Result Sample 3 | Result Sample 4 | Limit of detection | Limit of detection given as | Method |
|----------------|-------------------|------------------|------------------------|------------------------|------------------------|------------------------|--------------------|-----------------------------|--|
| | | Day/ Month | positive / negative | positive / negative | positive / negative | positive / negative | mg/kg | e.g. food/ protein | Test-Kit + Manufacturer |
| AQ | 8 | 15.07.20 | positive | negative | positive | negative | 0,02 | tropomyosin | AQ = AgraQuant, RomerLabs |
| AQ | 18 | Aug | pos | neg | pos | neg | 0,1 | Food item, total | AQ = AgraQuant, RomerLabs |
| BF | 5 | | positive | negative | positive | negative | 1 | Food item, total | BF = MonoTrace ELISA, BioFront Technologies |
| BF | 28 | 28/8 | positive | negative | positive | negative | 0,07 | Food item, total | BF = MonoTrace ELISA, BioFront Technologies |
| IL | 17 | | positive | negative | positive | negative | | Food item, total | IL = Immunolab |
| RS-F | 7 | 30.07.20 | positive | negative | positive | positive | 20 | Food item, total | RS-F= Ridascreen® Fast, R- Biopharm |
| RS-F | 9 | | positive | positive | positive | positive | 2 | Protein | R-BIOPHARM R7312 |
| RS-F | 22 | 08.07.20 | positive | negative | positive | negative | 20 | Food item, total | RS-F= Ridascreen® Fast, R- Biopharm |
| RS-F | 27 | 29.06.20 | positive | negative | positive | negative | 2 | Please select! | RS-F= Ridascreen® Fast, R- Biopharm |
| SP | 4 | 30.06. | positive | negative | positive | negative | 0,02 | Please select! | SP = SensiSpec, Eurofins Technologies |
| SP | 12 | | positive | negative | positive | negative | 0,02 | Protein (tropomyosin) | SP = SensiSpec, Eurofins Technologies |
| SP | 19 | 23.06.20 | positive | negative | positive | negative | 0,009 | Shrimp Tropomyosin | SP = SensiSpec, Eurofins Technologies |
| SP | 21 | 10.08.20 | positive | negative | positive | negative | 0,01 | Food item, total | SP = SensiSpec, Eurofins Technologies |

| Meth. Abbr. | Evaluation number | Method-No. / Test-Kit No. | Specifity | Remarks to the Method (Extraction and Determination) | Further Remarks |
|----------------|-------------------|---|---|---|--|
| | | Article-No. / ASU-No. | Antibody | e.g. Extractionbuffer / Time / Temperature | |
| AQ | 8 | | | as stipulated in kit insert | Kit present weak cross reactivity to molluscs |
| AQ | 18 | 10002076 (COKAL2248) | Unknown | | Reported as 'Crustacea' |
| BF | 5 | | | | |
| BF | 28 | | Monoclonal antibodies | 1:10 extractraction ratio, 1 hour at 42C | no |
| IL | 17 | | | | |
| RS-F | 7 | RIDASCREEN® FAST Crustacean (2nd generation) Art. No. R7312 / 14139 | The antibody specifically detects crustacean proteins such as tropomyosin | As per kit instructions | no |
| RS-F | 9 | | | | Sample 1 and Sample 3 are out of range |
| RS-F | 22 | R 7312 | ANTI-TROPOMIOSIN | EXTRACTION: BUFFER 10 MINUTI / 60°C DETERMINATION 30 MINUTI / 20-25°C | |
| RS-F | 27 | R7312 | Tropomyosin | As per kit instructions | reported as crustaceen |
| SP | 4 | HU0030006 | recognizes the crustacean tropomyosin | According to manufacturer information | Tropomyosin crustaceans |
| SP | 12 | HU0030006 | | | Reported as ug/Kg tropomyosin from crustaceans |
| SP | 19 | | | | |
| SP | 21 | | | | |

5.1.2 ELISA: Egg

Primary data

| Meth. Abbr. | Evaluation number | Date of analysis | Result Sample 1 | Result Sample 2 | Result Sample 3 | Result Sample 4 | Limit of detection | Limit of detection given as | Method |
|----------------|-------------------|------------------|------------------------|------------------------|------------------------|------------------------|--------------------|-----------------------------|--|
| | | Day/ Month | positive / negative | positive / negative | positive / negative | positive / negative | mg/kg | e.g. food/ protein | Test-Kit + Manufacturer |
| AS | 11 | 10.07.20 | positive | positive | negative | negative | 2 | | AgraStrip Egg/Romer Labs |
| BF | 28 | 28/8 | positive | positive | negative | negative | 0,3 | Whole egg powder | BF = MonoTrace ELISA, BioFront Technologies |
| ES | 14 | | positive | positive | negative | negative | 5 ppm | Egg white powder | ES = ELISA-Systems |
| IL | 17 | | positive | positive | negative | negative | | Please select! | IL = Immunolab |
| MI-II | 3 | 07.07.20 | positive | positive | negative | negative | 10 | Food item, total | MI-II = Morinaga Institute ELISA II |
| MI-II | 4 | 29.06. | positive | positive | negative | negative | 0,31 | Please select! | MI-II = Morinaga Institute ELISA II |
| MI-II | 8 | 20.07.20 | positive | positive | negative | negative | 0,312 | egg protein | MI-II = Morinaga Institute ELISA II |
| MI-II | 12 | | positive | positive | negative | negative | 0,31 | Whole Egg Protein | MI-II = Morinaga Institute ELISA II |
| RS-F | 7 | 07.08.20 | positive | positive | negative | negative | 0,5 | Whole egg powder | RS-F= Ridascreen® Fast, R- Biopharm |
| RS-F | 9 | | positive | positive | negative | negative | 0,1 | protein | R-BIOPHARM 6402 |
| RS-F | 20 | 14.08.20 | positive | positive | negative | negative | 0,5 | Whole egg powder | RS-F= Ridascreen® Fast, R- Biopharm |
| RS-F | 22 | 07.07.20 | positive | positive | negative | negative | 0,5 | Whole egg powder | RS-F= Ridascreen® Fast, R- Biopharm |
| SP | 19 | 22.06.20 | positive | positive | negative | negative | 0.05 | Egg white protein | SP = SensiSpec, Eurofins Technologies |
| SP | 21 | 10.08.20 | positive | positive | negative | negative | 0,05 | Food item, total | SP = SensiSpec, Eurofins Technologies |

| Meth. Abbr. | Evaluation number | Method-No. / Test-Kit No. | Specifity | Remarks to the Method (Extraction and Determination) | Further Remarks |
|----------------|-------------------|--|---|--|--|
| | | Article-No./ ASU-No. | Antibody | e.g. Extractionbuffer / Time / Temperature | |
| AS | 11 | 1000003564 | | | |
| BF | 28 | | Monoclonal antibodies | 1:20 extractraction ratio, 1 hour at 60C | no |
| ES | 14 | ES-6020, Transia | polyclonal, anti ovomucoid/ ovalbumin polyclonal, anti ovomucoid/ ovalbumin poly- clonal, anti ovomucoid/ oval- bumin | Extractions buffer, 3x10min, room temperature | LOD 0,5 ppm |
| IL | 17 | | | | |
| MI-II | 3 | | | | |
| MI-II | 4 | M2111 | recognizes the egg white protein ovalbumin | According to manufacturer information | whole egg protein |
| MI-II | 8 | | | as stipulated in kit insert | |
| MI-II | 12 | M2111 | | | Reported as whole egg protein mg/Kg |
| RS-F | 7 | RIDASCREEN® FAST Ei / Egg Protein (ART. No R6402) / 15339 | The antibodies specifically detect the antigens ovalbumin and ovomucoid of hen's egg | As per kit instructions | no |
| RS-F | 9 | | | | Sample 1 and Sample 2 are out of range |
| RS-F | 20 | RIDASCREEN® FAST Ei / Egg Protein (Art. Nr.: R6402) | The specific antibodies detect the egg white proteins ovalbumin and ovomucoid. | Preparation of the sample and test implementation following the instruction of RIDASCREEN® FAST Ei / Egg Protein (Art. Nr.: R6402) Lot 15339 - extraction with diluted Allergen Extraction buffer 10 min at 60°C | |
| RS-F | 22 | R 6402 | ANTI- OVOALBUMIN ANTI- OVOMUCOID | EXTRACTION: BUFFER 10 MINUTI / 60°C DETERMINATION 30 MINUTI / 20-25°C | |
| SP | 19 | | | | |
| SP | 21 | | | | |

5.1.3 ELISA: Fish

Primary data

| Meth. Abbr. | Evaluation number | Date of analysis | Result Sample 1 | Result Sample 2 | Result Sample 3 | Result Sample 4 | Limit of detection | Limit of detection given as | Method |
|----------------|-------------------|------------------|------------------------|------------------------|------------------------|------------------------|--------------------|-----------------------------|--|
| | | Day/ Month | positive / negative | positive / negative | positive / negative | positive / negative | mg/kg | e.g. food/ protein | Test-Kit + Manufacturer |
| AQ | 8 | 06.07.20 | negative | positive | positive | negative | 4 | Food item (cod) | AQ = AgraQuant, RomerLabs |
| ВС | 26 | 11.07.20 | negative | positive | positive | negative | 5 | Food item, total | BC = BioCheck ELISA |
| BF | 28 | 28/8 | negative | positive | positive | negative | 0,3 | Food item, total | BF = MonoTrace ELISA, BioFront Technologies |
| IL | 17 | | negative | positive | positive | negative | | Food item, total | IL = Immunolab |
| SP | 19 | 22.06.20 | negative | positive | positive | negative | "1,4" | Food item, fresh | SP = SensiSpec, Eurofins Technologies |
| SP | 21 | 10.08.20 | negative | positive | positive | negative | 1,4 | Food item, total | SP = SensiSpec, Eurofins Technologies |

| Meth. Abbr. | Evaluation number | Method-No. / Test-Kit No. | Specifity | Remarks to the Method (Extraction and Determination) | Further Remarks |
|----------------|-------------------|------------------------------|-------------------------|--|-----------------|
| | | Article-No. / ASU-No. | Antibody | e.g. Extractionbuffer / Time / Temperature | |
| AQ | 8 | | | as stipulated in kit insert | |
| ВС | 26 | As Per Kit Instructions | As Per Kit Instructions | As Per Kit Instructions | |
| BF | 28 | | Monoclonal antibodies | 1:10 extractraction ratio, 1 hour boiling | no |
| IL | 17 | | | | |
| SP | 19 | | | | |
| SP | 21 | | | | |

5.1.4 ELISA: Milk

Primary data

| Meth. Abbr. | Evaluation number | Date of analysis | Result Sample 1 | Result Sample 2 | Result Sample 3 | Result Sample 4 | Limit of detection | Limit of detection given as | Method |
|----------------|-------------------|------------------|------------------------|------------------------|------------------------|------------------------|--|---------------------------------------|--|
| | | Day/ Month | positive / negative | positive / negative | positive / negative | positive / negative | mg/kg | e.g. food/ protein | Test-Kit + Manufacturer |
| BF | 28 | 28/8 | positive | negative | negative | positive | 0,48 | Milk powder | BF = MonoTrace ELISA, BioFront Technologies |
| ES | 14a | | positive | negative | negative | positive | 10 ppm Casein | Skimmed milk powder equivalents | ES = ELISA-Systems |
| ES | 14b | | positive | negative | negative | positive | 1 ppm ß- Lactglobulin | ß-Lactglobulin | ES = ELISA-Systems |
| IL | 17 | | positive | negative | negative | positive | | Milk powder | IL = lmmunolab |
| MI-II | 4 | 26.06. | positive | negative | negative | positive | 0,31 | Please select! | MI-II = Morinaga Institute ELISA II |
| MI-II | 8a | 29.07.20 | positive | negative | negative | positive | 0,312 | Milk powder | MI-II = Morinaga Institute ELISA II |
| MI-II | 8b | 29.07.20 | positive | negative | negative | positive | 0,312 | Milk powder | MI-II = Morinaga Institute ELISA II |
| RS div | 1 | | positive | negative | negative | positive | | Please select! | Selection ELISA-Kits: |
| RS-F | 3 | 07.07.20 | positive | negative | negative | positive | 10 | Food item, total | RS-F= Ridascreen® Fast, R- Biopharm |
| RS-F | 7 | 08.04.20 | positive | negative | negative | positive | 0,5 | Casein | RS-F= Ridascreen® Fast, R- Biopharm |
| RS-F | 9 | | positive | negative | negative | positive | 0,7 | protein | R-BIOPHARM 4652 |
| RS-F | 22 | 09.07.20 | positive | negative | negative | positive | 0,167 | Food item, total | RS-F= Ridascreen® Fast, R- Biopharm |
| RS-F | 20a | 04.08.20 | positive | negative | negative | positive | 2,5 mg/kg (ppm) casein | Food item, total | RS-F= Ridascreen® Fast, R- Biopharm |
| RS-F | 20b | 04.08.20 | positive | negative | negative | positive | 0,167 mg/kg (ppm) β- lactoglobulin | Food item, total | RS-F= Ridascreen® Fast, R- Biopharm |
| SP | 12 | | positive | negative | negative | positive | 0,2 | Protein (casein) | SP = SensiSpec, Eurofins Technologies |
| SP | 19 | 22.06.20 | positive | negative | negative | positive | 0.05 | Casein+BLG | SP = SensiSpec, Eurofins Technologies |
| SP | 21 | 10.08.20 | positive | negative | negative | positive | 0,05 | Food item, total | SP = SensiSpec, Eurofins Technologies |

| Meth. Abbr. | Evaluation number | Method-No. / Test-Kit No. | Specifity | Remarks to the Method (Extraction and Determination) | Further Remarks |
|----------------|-------------------|---|---|---|--|
| | | Article-No. / ASU-No. | Antibody | e.g. Extractionbuffer / Time / Temperature | |
| BF | 28 | | Monoclonal antibodies | 1:10 extractraction ratio, 1 hour at 60C | no |
| ES | 14a | ES-6030 Transia | polyclonal, anti bovine alpha- Casein | extraction buffer for both ELISAs, 2x15min and 1x10min, room temperature | we use 2 ELISAs: LOD 1 ppm skimmed milk powder equi- valents or LOD 0.1 ppm ß-lac- toglobulin |
| ES | 14b | ES-6034 Transia | polyclonal, anti ß- Lactglobulin | extraction buffer for both ELISAs, 2x15min and 1x10min, room temperature | we use 2 ELISAs: LOD 1 ppm skimmed milk powder equi- valents or LOD 0.1 ppm ß-lac- toglobulin |
| IL | 17 | | | | |
| MI-II | 4 | M2113 Casein | recognizes cow's milk casein | According to manufacturer information | milk protein |
| MI-II | 8a | | CASEIN | as stipulated in kit insert | |
| MI-II | 8b | | BLG | as stipulated in kit insert | |
| RS div | 1 | | | R biopharm | |
| RS-F | 3 | | | | |
| RS-F | 7 | RIDASCREEN® FAST casein Art. N° R4612 / 22060 | The antibodies specifically detect Casein | As per kit instructions | no |
| RS-F | 9 | | | | |
| RS-F | 22 | R 4912 | ANTI-COW BETALACTOGLOBULIN | EXTRACTION: BUFFER1 10 MIN/ 100°C BUFFER 2 10 MIN/60°C DETERMINATION 30 MINUTI / 20-25°C | |
| RS-F | | RIDASCREEN® FAST Casein (R4612) | The used antibodies specifically detect caseins of cow's milk. | Preparation of the sample and test implementation following the instruction of RIDASCREEN® FAST Casein (R4612), Lot 22060 casein - extraction with Extractor 2 cook it for 10 min at 100 °C in a water bath and then adding Allergen extraction buffer containing Additive 1 (A-AEP) and extract for 10 min at 60 °C in a water bath | |
| RS-F | 20b | RIDASCREEN® FAST β- Lactoglobulin (Art. No. R4912) | The antibodies specifically detect β-lactoglobulin of cow's milk. | Preparation of the sample and test implementation following the instruction of RIDASCREEN® FAST β-Lactoglobulin (Art. No. R4912), Lot 24090 β-Lactoglobulin - extraction with Extractor 2 cook it for 10 min at 100 °C in a water bath and then adding Allergen extraction buffer containing Additive 1 (A-AEP) and extract for 10 min at 60 °C in a water bath | |
| SP | 12 | HU0030003 | | | Milk detected as casein mg/Kg |
| SP | 19 | | | | |
| SP | 21 | | | | |

5.1.5 ELISA: Molluscs

Primary data

| Meth. Abbr. | Evaluation number | Date of analysis | Result Sample 1 | Result Sample 2 | Result Sample 3 | Result Sample 4 | Limit of detection | Limit of detection given as | Method |
|----------------|-------------------|------------------|---------------------|------------------------|------------------------|------------------------|--------------------|-----------------------------|--|
| | | Day/ Month | positive / negative | positive / negative | positive / negative | positive / negative | mg/kg | e.g. food/ protein | Test-Kit + Manufacturer |
| DE | 26 | 24.07.20 | positive | negative | positive | positive | 0,01 | Food item, total | other: please fill in! |
| ET | 8 | 06.07.20 | positive | negative | positive | positive | 1 | molluscs protein | ET = Elution Technologies ELISA Kit |
| IL | 17 | | negative | negative | positive | positive | | Food item, total | IL = Immunolab |
| SP | 4 | 30.06. | - | negative | positive | positive | 0,03 | Please select! | SP = SensiSpec, Eurofins Technologies |
| SP | 19 | 23.06.20 | positive* | negative | positive* | positive | 0,017 | Garden Snail Tropomyosin | SP = SensiSpec, Eurofins Technologies |
| SP | 21 | 10.08.20 | negative | negative | positive | positive | 0,0017 | Food item, total | SP = SensiSpec, Eurofins Technologies |

| Meth. Abbr. | Evaluation number | Method-No. / Test-Kit No. | Specifity | Remarks to the Method (Extraction and Determination) | Further Remarks |
|----------------|-------------------|------------------------------|---------------------------------------|---|---|
| | | Article-No. / ASU-No. | Antibody | e.g. Extractionbuffer / Time / Temperature | |
| DE | 76 | As Per Kit Instructions | As Per Kit Instructions | As Per Kit Instructions | DeMediTec GmBH Test Kit |
| ET | 8 | | | as stipulated in kit insert | Use kit 3M with cross reactivity to crustacea. Result #1 = Cross reactivity to crustacea suspected. |
| IL | 17 | | | | |
| SP | 1 1 | HU0030015/00300 39 | recognizes the mollusc tropomyosin | According to manufacturer information | Tropomyosin molluscs; Sample 1: traces at the limit of detection (positive <0.03mg/kg) |
| SP | 19 | | | | *positive due to cross-reaction of crustaceans |
| SP | 21 | | | | |

5.1.6 ELISA: Mustard

Primary data

| Meth. Abbr. | Evaluation number | Date of analysis | Result Sample 1 | Result Sample 2 | Result Sample 3 | Result Sample 4 | Limit of detection | Limit of detection given as | Method |
|----------------|-------------------|------------------|------------------------|------------------------|------------------------|------------------------|--------------------|-----------------------------|--|
| | | Day/ Month | positive / negative | positive / negative | positive / negative | positive / negative | mg/kg | e.g. food/ protein | Test-Kit + Manufacturer |
| AS | 11 | 10.07.20 | negative | positive | positive | positive | 2 | | AgraStrip Mustard / Romer Labs |
| BF | 28 | 28/8 | negative | positive | positive | positive | 0,13 | Food item, total | BF = MonoTrace ELISA, BioFront Technologies |
| IL | 17 | | negative | positive | positive | positive | | Food item, total | IL = Immunolab |
| RS-F | 7 | 06.08.20 | negative | positive | positive | positive | 2,5 | Food item, total | RS-F= Ridascreen® Fast, R- Biopharm |
| RS-F | 10 | | negative | positive | positive | positive | 0,5 | Food item, total | RS-F= Ridascreen® Fast, R- Biopharm |
| RS-F | 18 | Aug | neg | pos | pos | pos | 0,5 | Food item, total | RS-F= Ridas creen® Fast, R- Biopharm |
| RS-F | 20 | 14.08.20 | negative | positive | positive | positive | 0,5 | Food item, total | RS-F= Ridas creen® Fast, R- Biopharm |
| SP | 12 | | negative | positive | positive | positive | 2 | Food item, total | SP = SensiSpec, Eurofins Technologies |
| SP | 19 | 22.06.20 | negative | positive | positive | positive | 1 | food item, dried | SP = SensiSpec, Eurofins Technologies |
| SP | 21 | 10.08.20 | negative | positive | positive | positive | 1 | Food item, total | SP = SensiSpec, Eurofins Technologies |
| VT | 4 | 26.06. | negative | positive | positive | positive | 2,5 | Mustard | Neogen Veratox Senf |
| VT | 8 | 15.07.20 | negative | positive | positive | positive | 2,5 | Food item, total | VT = Veratox, Neogen |

| Meth. Abbr. | Evaluation number | Method-No. / Test-Kit No. | Specifity | Remarks to the Method (Extraction and Determination) | Further Remarks |
|----------------|-------------------|---|---|---|---|
| | | Article-No. / ASU-No. | Antibody | e.g. Extractionbuffer / Time / Temperature | |
| AS | 11 | 1000000977 | | | |
| BF | 28 | | Monoclonal antibodies | 1:20 extractraction ratio, 1 hour at 60C | assay detects yellow/white, brown, black mustard |
| IL | 17 | | | | Cross reactivity to brown mustard and black mustard |
| RS-F | 7 | Ridascreen® FAST Mustard R6152, R- Biopharm / 14489 | The antibody specifically detects white, yellow, brown and black mustard. | As per kit instructions. Kit uses general mustard screening. Yellow, brown and black mustard cannot be differentiated | no |
| RS-F | 10 | | | mustard extraction buffer, 10 min, 60°C | |
| RS-F | 18 | R6152 | Unkown | | yellow, brown and black reported in total as 'mustard' |
| RS-F | 20 | RIDASCREEN® FAST Senf/Mustard (Art. Nr.: R6152) | | Preparation of the sample and test implementation following the instruction of RIDASCREEN® FAST Senf/Mustard (Art. Nr.: R6152), Lot 14489 - extraction with diluted Allergen Extraction buffer 10 min at 60°C | |
| SP | 12 | HU0030016 | | | Test does NOT separate out species e.g. black, yellow ets. Reported as mg/Kg mustard. |
| SP | 19 | | | | Cross-reactivty: Yellow 100%, brown 59%, black 50% |
| SP | 21 | | | | |
| VT | 4 | 8400 | recognizes mustard protein from seeds of white mustard (Sinapis alba), black mustard (Brassica nigra) and brown mustard (Brassica juncea) | According to manufacturer information | |
| VT | 8 | | | as stipulated in kit insert | |

5.1.7 ELISA: Soya

Primary data

| Meth. Abbr. | Evaluation number | Date of analysis | Result Sample 1 | Result Sample 2 | Result Sample 3 | Result Sample 4 | Limit of detection | Limit of detection given as | Method |
|----------------|-------------------|------------------|------------------------|------------------------|------------------------|------------------------|--------------------|-----------------------------|--|
| | | Day/ Month | positive / negative | positive / negative | positive / negative | positive / negative | mg/kg | e.g. food/ protein | Test-Kit + Manufacturer |
| AS | 11 | 10.07.20 | positive | negative | negative | positive | 2 | | AgraStrip Soy / Romer Labs |
| BF | 28 | 28/8 | positive | negative | negative | positive | 0,16 | Food item, total | BF = MonoTrace ELISA, BioFront Technologies |
| ES | 14 | | positive | negative | negative | positive | 25 ppm | Soya protein | ES = ELISA-Systems |
| IL | 17 | | positive | negative | negative | positive | | Please select! | IL = Immunolab |
| IL | 19 | 23.06.20 | positive | negative | negative | positive | 0,2 | Total protein | IL = Immunolab |
| MI-II | 4 | 29.06. | positive | negative | negative | positive | 0,31 | Please select! | MI-II = Morinaga Institute ELISA II |
| MI-II | 8 | 21.07.20 | positive | negative | negative | positive | 0,312 | soya protein | MI-II = Morinaga Institute ELISA II |
| RS-F | 7 | 03.08.20 | positive | negative | negative | positive | 2,5 | Food item, total | RS-F= Ridascreen® Fast, R- Biopharm |
| RS-F | 9 | | positive | negative | negative | positive | 0,24 | protein | R-BIOPHARM 7102 |
| RS-F | 12 | | positive | negative | negative | positive | 2,5 | Protein | RS-F= Ridascreen® Fast, R- Biopharm |
| RS-F | 20 | 04.08.20 | positive | negative | negative | positive | 2,5 | Food item, total | RS-F= Ridascreen® Fast, R- Biopharm |
| RS-F | 27 | 23.06.20 | positive | negative | negative | positive | 0,24 | Please select! | RS-F= Ridascreen® Fast, R- Biopharm |
| SP | 21 | 10.08.20 | positive | negative | negative | positive | 0,016 | Please select! | SP = SensiSpec, Eurofins Technologies |

| Meth. Abbr. | Evaluation number | Method-No. / Test-Kit No. | Specifity | Remarks to the Method (Extraction and Determination) | Further Remarks |
|----------------|-------------------|--|---|--|--|
| | | Article-No. / ASU-No. | Antibody | e.g. Extractionbuffer / Time / Temperature | |
| AS | 11 | 1000002188 | | | |
| BF | 28 | | Monoclonal antibodies | 1:20 extractraction ratio, 1 hour boiling | no |
| ES | 14 | ES-6012, Transia | polyclonal, anti-soy trypsin inhibitor and anti-soy flour protein | Extraction buffer, 2x30min and 1x15 min, room temperature | LOD 2,5 ppm |
| IL | 17 | | | | |
| IL | 19 | | | | |
| MI-II | 4 | M2117 | recognizes the soy protein beta-conglycinin | According to manufacturer information | soy protein |
| MI-II | 8 | | | as stipulated in kit insert | |
| RS-F | 7 | Ridas creen® FAST Soy R7102, R- Biopharm / 24180 | Against Heat processed soya proteins. (Glycinin (408%, beta-conglycinin 7.3%, tripsin inhibitor 0.46%) | As per kit instructions | no |
| RS-F | 9 | | | | Sample 1 and Sample 4 are out of range |
| RS-F | 12 | R7102 | | | Reported as soya protein mg/Kg |
| RS-F | 20 | RIDASCREEN® FAST Soya (Art. No. R7102) | The antibodies specifically detect heated soya proteins | Preparation of the sample and test implementation following the instruction of RIDASCREEN® FAST Soya (Art. No. R7102), Lot 13339 - extraction with Extractor 3 and diluted Allergen Extraction Buffer for 10 min at 100 °C | |
| RS-F | 27 | R7102 | heated soyproteins | according to test kit instructions | Reported as soyprotein |
| SP | 21 | | | | |

5.1.8 PCR: Crustaceae

Primary data

| Meth. Abbr. | Evaluation number | Date of analysis | Result Sample 1 | Result Sample 2 | Result Sample 3 | Result Sample 4 | Limit of detection | Limit of detection given as | Method |
|----------------|-------------------|------------------|------------------------|------------------------|------------------------|------------------------|--------------------|-----------------------------|--|
| | | Day/ Month | positive / negative | positive / negative | positive / negative | positive / negative | mg/kg | e.g. food/ protein | Test-Kit + Manufacturer |
| ASU | 15 | 18.08.20 | positive | negative | positive | negative | | Please select! | ASU = ASU §64 Methode/method |
| ASU | 27 | 01.07.20 | positive | negative | positive | negative | | Please select! | ASU = ASU §64 Methode/method |
| SFA | 1 | | positive | negative | positive | negative | | Please select! | Selection PCR-Methods |
| SFA | 2 | 23.06.20 | positive | negative | positive | negative | 0,4 | Food item, total | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 5 | | positive | negative | positive | negative | 0,4 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 7 | 07.08.20 | positive | negative | positive | positive | 2,5 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 10 | | positive | negative | positive | negative | 2 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 20 | 13.08.20 | positive | negative | positive | negative | 0,4 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 21 | 10.08.20 | positive | negative | positive | negative | 0,4 | Food item, total | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 23 | | positive | negative | positive | negative | 0,4 | Food item, total | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 25 | 20.08.20 | positive | negative | positive | negative | 100 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA-ID | 16 | | positive | negative | positive | negative | <0.4 mg/kg | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |

| Meth. Abbr. | Evaluation number | Method-No. / Test-Kit No. | Specifity | Remarks to the Method (Extraction and Determination) | Further Remarks |
|----------------|-------------------|--|---|---|--|
| | | Article-No. / ASU-No. | Target-Sequence / -DNA | e.g. Extraction / Enzymes / Clean-Up / Real Time PCR / Gel electrophoresis / Cycles | |
| ASU | 15 | L12.01-3 ; 07/2012 | | Simplex EasySpinFood DNA Kit / GEN-IAL, endpoint PCR with subsequent sequencing | |
| ASU | 27 | L12.01-3 | | according to ASU method | |
| SFA | 1 | | | CTAB extraction + PCR CONGEN | |
| SFA | 2 | S3612 | Crustacea | Extraction with SureFood® Prep Advanced protocol 1 (S1053) | K01, QE to abalone (Haliotis) 100 % |
| SFA | 5 | | | | |
| SFA | 7 | SureFood® ALLERGEN Crustaceans Art. No. S3612 / 20150 | Not specified in kit | As per kit instructions | no |
| SFA | 10 | | | prep advance surefood/taq polymerase/ RT PCR/45 cycles | |
| SFA | 20 | SureFood® ALLERGEN Crustaceans - Art. No. S3612 | The real-time PCR test detects DNA of crustaceans (Crustacea) | DNA preparation with SureFood® PREP Advanced (Principle according to protocol 2: Lysis at 65°C - Pre-filtration and setting of optimal binding conditions - Binding of the nucleic acids on a Spin Filter - Purification of the bound nucleic acids - Drying of the Spin Filter - First Elution of nucleic acids from the Spin Filter - Repeated setting of optimal binding conditions - Second binding of the nucleic acids on a Spin Filter - Second purification of the bound nucleic acids - Drying of the Spin Filter - Elution of nucleic acids from the Spin Filter for analysis) and real-time PCR (45 cycles following kit setup instructions) with Bio-Rad CFX96, Lot 11349 | |
| SFA | 21 | | | , | |
| SFA | 23 | S3612/11349 | | Extraction= Sure Food PREP Advanced Determination = real tim | |
| SFA | 25 | S3612 | | qiagen dneasy kit/real time PCR/45 cycles | |
| SFA-ID | 16 | S3112 | | real time PCR | |

5.1.9 PCR: Fish

Primary data

| Meth. Abbr. | Evaluation number | Date of analysis | Result Sample 1 | Result Sample 2 | Result Sample 3 | Result Sample 4 | Limit of detection | Limit of detection given as | Method |
|----------------|-------------------|------------------|------------------------|------------------------|------------------------|------------------------|--------------------|-----------------------------|--|
| | | Day/ Month | positive / negative | positive / negative | positive / negative | positive / negative | mg/kg | e.g. food/ protein | Test-Kit + Manufacturer |
| GI | 15 | 12.08.20 | negative | positive | positive | negative | 15 | Allergen-DNA | First-Fish Kit / GEN-IAL |
| GS | 12 | | negative | positive | positive | negative | 0,001 | Food item, total | Eurofins Genescan DNAnimal screen fish |
| IM | 13 | 12.08.20 | negative | positive | positive | negative | 4 | Please select! | other: IMEGEN |
| MS | 3 | 13.07.20 | negative | positive | positive | negative | 10 | Allergen-DNA | MS = Microsynth |
| SFA | 1 | | negative | positive | positive | negative | | Please select! | Selection PCR-Methods |
| SFA | 2 | 23.06.20 | negative | positive | positive | negative | 1 | Food item, total | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 5 | | negative | positive | positive | negative | 0,4 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 7 | 31.07.20 | negative | positive | positive | negative | 2,5 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 10 | | negative | positive | positive | negative | 5 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 16 | | negative | positive | positive | negative | <1 mg/kg | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 20 | 13.08.20 | negative | positive | positive | negative | 0,4 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 21 | 10.08.20 | negative | positive | positive | negative | 1 | Food item, total | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 23 | | negative | positive | positive | negative | 1 | Food item, total | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 24 | | negative | positive | positive | negative | 1 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 25 | 20.08.20 | negative | positive | positive | negative | 100 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 26 | 22.08.20 | negative | negative | positive | positive | 1 | Food item, total | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| div | 4 | 13.07. | negative | positive | positive | negative | 20 | Allergen-DNA | Selection PCR methods |
| div | 9 | | negative | positive | positive | negative | 0,008 | | in-house method |

| Meth. Abbr. | Evaluation number | Method-No. / Test-Kit No. | Specifity | Remarks to the Method (Extraction and Determination) | Further Remarks |
|----------------|-------------------|--|--|---|---|
| | | Article-No. / ASU-No. | Target-Sequence / -DNA | e.g. Extraction / Enzymes / Clean-Up / Real Time PCR / Gel electrophoresis / Cycles | |
| GI | 15 | PHF0050, L10.00- 12 | | Simplex EasySpinFood DNA Kit/GEN-IAL, RealTime PCR | |
| GS | 12 | 5422211310 | | | |
| IM | 13 | | | CTAB/ kit /PCR real time | |
| MS | 3 | | | Wizard Extraktion, Real Time PCR | |
| SFA | 1 | | | CTAB extraction + PCR CONGEN | |
| SFA | 2 | S3610 | Osteichthyes (bony fish) | Extraction with SureFood® Prep Advanced protocol 1 (S1053) | K01, QE to muscovy duck (Cairina moschata) 100 % |
| SFA | 5 | | | | |
| SFA | 7 | SureFood® ALLERGEN fish Art. No. S3610/20150 | Not specified in kit | As per kit instructions | no |
| SFA | 10 | | | prep advance surefood/taq polymerase/ RT PCR/45 cycles | |
| SFA | 16 | S3610 | | real time PCR | |
| SFA | 20 | SureFood® ALLERGEN Fish - Art. No. S3610 | The real-time PCR test detects DNA of fish | DNA preparation with SureFood® PREP Advanced (Principle according to protocol 2: Lysis at 65°C - Pre-filtration and setting of optimal binding conditions - Binding of the nucleic acids on a Spin Filter - Purification of the bound nucleic acids - Drying of the Spin Filter - First Elution of nucleic acids from the Spin Filter - Repeated setting of optimal binding conditions - Second binding of the nucleic acids on a Spin Filter - Second purification of the bound nucleic acids - Drying of the Spin Filter - Elution of nucleic acids from the Spin Filter for analysis) and real-time PCR (45 cycles following kit setup instructions) with Bio-Rad CFX96, Lot 14309 | |
| SFA | 21 | | | | |
| SFA | 23 | S3610/14309 | | Extraction= Sure Food PREP Advanced Determination = real time | |
| SFA | 24 | S3610 | DNA fragment present solely in fish | CTAB DNA extraction/Real time PCR | Analyst: LP/AP |
| SFA | 25 | S3610 | | | |
| SFA | 26 | As Per Kit Instructions | As Per Kit Instructions | As Per Kit Instructions | |
| div | 4 | internal method | | CTAB / Proteinase K / Promega Wizard DNA CleanUp / Real-time PCR 45 cycles | |
| div | 9 | | | | |

5.1.10 PCR: Molluscs

Primary data

| Meth. Abbr. | Evaluation number | Date of analysis | Result Sample 1 | Result Sample 2 | Result Sample 3 | Result Sample 4 | Limit of detection | Limit of detection given as | Method |
|----------------|-------------------|------------------|------------------------|------------------------|------------------------|------------------------|--------------------------------|-----------------------------|--|
| | | Day/ Month | positive / negative | positive / negative | positive / negative | positive / negative | mg/kg | e.g. food/ protein | Test-Kit + Manufacturer |
| 4L | 22 | 23.07.20 | negative | negative | positive | positive | A COPY OF HAPLOID GENOME | Allergen DNA | 4L = 4LAB Diagnostics |
| SFA | 1 | | negative | negative | positive | positive | | Please select! | Selection PCR-Methods |
| SFA | 2 | 23.06.20 | negative | negative | positive | positive | 0,4 | Food item, total | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 5 | | negative | negative | positive | positive | 0,4 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 7 | 31.07.20 | negative | negative | positive | positive | | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 10 | | negative | negative | positive | positive | 2 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 13 | 07.07.20 | negative | negative | positive | positive | 100 | Please select! | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 18 | Aug | neg | neg | pos | pos | 0,4 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 20 | 13.08.20 | negative | negative | positive | positive | 0,4 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 21 | 10.08.20 | negative | negative | positive | positive | 0,4 | Food item, total | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 25 | 20.08.20 | negative | negative | positive | positive | 40 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 26 | 22.08.20 | negative | negative | positive | positive | 1 | Food item, total | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| div | 9 | | negative | negative | positive | positive | 0,08 | | in-house method |

| Meth. Abbr. | Evaluation number | Method-No. / Test-Kit No. | Specifity | Remarks to the Method (Extraction and Determination) | Further Remarks |
|----------------|-------------------|--|--|---|-----------------|
| | | Article-No. / ASU-No. | Target-Sequence / -DNA | e.g. Extraction / Enzymes / Clean-Up / Real Time PCR / Gel electrophoresis / Cycles | |
| 4L | 22 | IC-02-1008 | MOLLUSC DNA | EXTRACTION WITH GREES DAN FOOD KIT KIT IC-02-0095 | |
| SFA | 1 | | | CTAB extraction + PCR CONGEN | |
| SFA | 2 | S3613 | Gastropods, Decabrachia, Bivalvia | Extraction with SureFood® Prep Advanced protocol 1 (S1053) | K02 |
| SFA | 5 | | | | |
| SFA | 7 | SureFood® ALLERGEN mollusc Art. No. S3613 / 23040 | Not specified in kit | As per kit instructions | no |
| SFA | 10 | | | prep advance surefood/taq polymerase/ RT PCR/45 cycles | |
| SFA | 13 | | | CTAB/ kit /PCR real time | |
| SFA | 18 | S3613 | Uknown | Tris extraction with column clean-up, real-time PCR detection | |
| SFA | 20 | SureFood® ALLERGEN Molluscs - Art. No. S3613 | The real-time PCR test detects DNA of molluscs | DNA preparation with SureFood® PREP Advanced (Principle according to protocol 2: Lysis at 65°C - Pre-filtration and setting of optimal binding conditions - Binding of the nucleic acids on a Spin Filter - Purification of the bound nucleic acids - Drying of the Spin Filter - First Elution of nucleic acids from the Spin Filter - Repeated setting of optimal binding conditions - Second binding of the nucleic acids on a Spin Filter - Second purification of the bound nucleic acids - Drying of the Spin Filter - Elution of nucleic acids from the Spin Filter for analysis) and real-time PCR (45 cycles following kit setup instructions) with Bio-Rad CFV96, Lot 13089 | |
| SFA | 21 | | | | |
| SFA | 25 | S3613 | | | |
| SFA | 1 76 | As Per Kit Instructions | As Per Kit Instructions | As Per Kit Instructions | |
| div | 9 | | | | |

5.1.11 PCR: Mustard, in general

Primary data

| Meth. Abbr. | Evaluation number | Date of analysis | Result Sample 1 | Result Sample 2 | Result Sample 3 | Result Sample 4 | Limit of detection | Limit of detection given as | Method |
|----------------|-------------------|------------------|------------------------|------------------------|------------------------|------------------------|--------------------|-----------------------------|--|
| | | Day/ Month | positive / negative | positive / negative | positive / negative | positive / negative | mg/kg | e.g. food/ protein | Test-Kit + Manufacturer |
| ASU | 4 | 13.07. | negative | positive | positive | positive | 5 | Allergen-DNA | ASU |
| SFA | 1 | | positive | positive | positive | positive | | Please select! | Selection PCR-Methods |
| SFA | 2 | 23.06.20 | negative | positive | positive | positive | 0,4 | Food item, total | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 5 | | negative | positive | positive | positive | 0,4 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 8 | 16.07.20 | negative | positive | positive | positive | | Allergen DNA | SFA-Q = Sure Food Allergen Quant, R-Biopharm / Congen |
| SFA | 16 | | negative | • | • | - | <0.4 mg/kg | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 20 | 13.08.20 | negative | positive | positive | positive | 0,4 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 21 | 10.08.20 | negative | positive | positive | positive | 0,4 | Food item, total | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 29 | | negative | positive | positive | positive | 0,4 | other: In general | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |

| Meth. Abbr. | Evaluation number | Method-No. / Test-Kit No. | Specifity | Remarks to the Method (Extraction and Determination) | Further Remarks |
|----------------|-------------------|------------------------------|---|---|--|
| | | Article-No. / ASU-No. | Target-Sequence / -DNA | e.g. Extraction / Enzymes / Clean-Up / Real Time PCR / Gel electrophoresis / Cycles | |
| ASU | 4 | L 08.00-65:2017-10 | | CTAB / Proteinase K / Promega Wizard DNA CleanUp / Real- time PCR 45 cycles | |
| SFA | 1 | | | CTAB extraction + PCR CONGEN (mustard screening) | |
| SFA | 2 | S3609 | yellow mustard (Sinapis alba), brown mustard (Brassica juncea), black mustard (Brassica nigra), ethiopian mustard (Brassica carinata), field mustard (Sinapis arvensis) | Extraction with SureFood® Prep Advanced protocol 1 (S1053) | K02, no differentiation between yellow, brown and black mustard |
| SFA | 5 | | | | The kit used for mustard's determination detects all three species listed without distinction. |
| SFA | 8 | S3609 | | cleaning using SureFood Prep Advanced S1053, real time PCR, 45 cycles | |
| SFA | 16 | S3609 | | real time PCR | Test cannot distinguish between different types of mustard |
| SFA | | Mustard - Art. No. S3609 | The test detects DNA of white mustard (Sinapis alba), indian mustard (Brassica juncea) und black mustard (Brassica nigra). The results are for mustard, in general | DNA preparation with SureFood® PREP Advanced (Principle according to protocol 2: Lysis at 65°C - Pre-filtration and setting of optimal binding conditions - Binding of the nucleic acids on a Spin Filter - Purification of the bound nucleic acids - Drying of the Spin Filter - First Elution of nucleic acids from the Spin Filter - Repeated setting of optimal binding conditions - Second binding of the nucleic acids on a Spin Filter - Second purification of the bound nucleic acids - Drying of the Spin Filter - Elution of nucleic acids from the Spin Filter for analysis) and real-time PCR (45 cycles following kit setup instructions) with Bio-Rad CFX96, Lot 13059 | The results are for mustard, in general. |
| SFA | 21 | | | | |
| SFA | 29 | | | R-Biopharm Kit for extraction DNA. We used a real time PCR with 45 cycles.R-Biopharm Kit for extraction DNA. We used a real time PCR with 45 cycles. | |

5.1.12 PCR: Mustard, Sinapis alba

Primary data

| Meth. Abbr. | Evaluation number | Date of analysis | Result Sample 1 | Result Sample 2 | Result Sample 3 | Result Sample 4 | Limit of detection | Limit of detection given as | Method |
|----------------|-------------------|------------------|------------------------|------------------------|------------------------|------------------------|--------------------|-----------------------------|---------------------------------|
| | | Day/ Month | positive / negative | positive / negative | positive / negative | positive / negative | mg/kg | e.g. food/ protein | Test-Kit + Manufacturer |
| ASU | 6 | 13.07.20 | negative | positive | negative | positive | 10 | Food item, total | ASU = ASU §64 Methode/method |
| ASU | 27 | 01.07.20 | negative | positive | negative | positive | | Please select! | ASU = ASU §64 Methode/method |
| GI | 15 | 12.08.20 | negative | positive | negative | positive | 10 | Allergen-DNA | GI = GEN-IAL First Allergen |
| MS | 3 | 13.07.20 | negative | positive | negative | positive | 10 | Allergen-DNA | MS = Microsynth |
| div | 9 | | negative | positive | negative | positive | 0,008 | | in-house method |

Other details to the Methods

| Meth. Abbr. | Evaluation number | Method-No. / Test-Kit No. | Specifity | Remarks to the Method (Extraction and Determination) | Further Remarks |
|----------------|-------------------|------------------------------|------------------------|--|-----------------|
| | | Article-No. / ASU-No. | Target-Sequence / -DNA | e.g. Extraction / Enzymes / Clean-Up / Real Time PCR / Gel electrophoresis / Cycles | |
| ASU | 6 | L08.00-59 | MADSD-F, MADSD-R | CTAB | |
| ASU | 27 | L08.00-65 | | according to ASU method | |
| GI | 15 | PMUS0050, L08.00- 64 | | Simplex EasySpinFood DNA Kit/GEN-IAL, RealTime PCR | |
| MS | 3 | | | Wizard extraction, Real Time PCR | |
| div | 9 | | | | |

5.1.13 PCR: Mustard, Brassica juncea/ Brassica nigra

Primary data

| Meth. Abbr. | Evaluation number | Date of analysis | Result Sample 1 | Result Sample 2 | Result Sample 3 | Result Sample 4 | Limit of detection | Limit of detection given as | Method |
|----------------|-------------------|------------------|------------------------|------------------------|------------------------|------------------------|--------------------|-----------------------------|---------------------------------|
| | | Day/ Month | positive / negative | positive / negative | positive / negative | positive / negative | mg/kg | e.g. food/ protein | Test-Kit + Manufacturer |
| ASU | 27 | 01.07.20 | negative | positive | positive | negative | | Please select! | ASU = ASU §64 Methode/method |
| GI | 15 | 12.08.20 | negative | positive | positive | negative | 5 | Allergen-DNA | GI = GEN-IAL First Allergen |
| MS | 3a | 13.07.20 | negative | positive | positive | negative | 10 | Allergen-DNA | MS = Microsynth |
| MS | 3b | 13.07.20 | negative | positive | positive | positive | 10 | Allergen-DNA | MS = Microsynth |
| div | 9 | | negative | positive | positive | negative | 0,008 | | in-house method |

| Meth. Abbr. | Evaluation number | Method-No. / Test-Kit No. | Specifity | Remarks to the Method (Extraction and Determination) | Further Remarks |
|----------------|-------------------|------------------------------|------------------------|--|---|
| | | Article-No. / ASU-No. | Target-Sequence / -DNA | e.g. Extraction / Enzymes / Clean-Up / Real Time PCR / Gel electrophoresis / Cycles | |
| ASU | 27 | L08.00-65 | | according to ASU method | brown and black mustard |
| GI | 15 | PMUS0050, L08.00- 64 | | ISIMPLEY EASYSPINEOOD DNA KIT/GEN-IAL REALTIME PCR | brown mustard is detected together with black mustard |
| MS | 3a | | | Wizard Extraktion, Real Time PCR | brown mustard |
| MS | 3b | | | Wizard Extraktion, Real Time PCR | black mustard |
| div | 9 | | | | black mustard |

5.1.14 PCR: Soya

Primary data

| Meth. Abbr. | Evaluation number | Date of analysis | Result Sample 1 | Result Sample 2 | Result Sample 3 | Result Sample 4 | Limit of detection | Limit of detection given as | Method |
|----------------|-------------------|------------------|------------------------|------------------------|------------------------|------------------------|--------------------|-----------------------------|--|
| | | Day/ Month | positive / negative | positive / negative | positive / negative | positive / negative | mg/kg | e.g. food/ protein | Test-Kit + Manufacturer |
| ASU | 6 | 13.07.20 | positive | negative | negative | positive | 10 | Food item, total | ASU = ASU §64 Methode/method |
| ASU | 27 | 01.07.20 | positive | negative | negative | positive | | Please select! | ASU = ASU §64 Methode/method |
| Gl | 15 | 12.08.20 | positive | negative | negative | positive | 10 | Allergen-DNA | GI = GEN-IAL First Allergen |
| MS | 3 | 13.07.20 | negative | negative | negative | positive | 10 | Allergen-DNA | MS = Microsynth |
| SFA | 1 | | positive | positive | positive | positive | | Please select! | Selection PCR-Methods |
| SFA | 2 | 23.06.20 | positive | negative | negative | positive | 0,4 | Food item, total | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 5 | | positive | negative | negative | positive | 0,4 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 16 | | positive | negative | negative | positive | <0.4 mg/kg | Allergen DNA | SFA-Q = Sure Food Allergen Quant, R-Biopharm / Congen |
| SFA | 20 | 13.08.20 | positive | negative | negative | positive | 0,4 | Allergen DNA | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| SFA | 21 | 10.08.20 | positive | negative | negative | positive | 0,4 | Food item, total | SFA = Sure Food ALLERGEN, R-Biopharm / Congen |
| div | 4 | 13.07. | positive | negative | negative | positive | 5 | Allergen-DNA | Selection PCR methods |
| div | 9 | | positive | negative | negative | positive | 0,02 | | in-house method |

| Meth. Abbr. | Evaluation number | Method-No. / Test-Kit No. | Specifity | Remarks to the Method (Extraction and Determination) | Further Remarks |
|----------------|-------------------|--|---|---|-----------------|
| | | Article-No. / ASU-No. | Target-Sequence / -DNA | e.g. Extraction / Enzymes / Clean-Up / Real Time PCR / Gel electrophoresis / Cycles | |
| ASU | 6 | L08.00-59 | Lectin-F; Lectin-R | CTAB | |
| ASU | 27 | L08.00-59 und L08.00-65 | | according to ASU methods | |
| Gl | 15 | PSOY 0050, L08.00-65 | | Simplex EasySpinFood DNA Kit/GEN-IAL, RealTime PCR | |
| MS | 3 | | | Wizard extraction, Real Time PCR | |
| SFA | 1 | | | CTAB extraction + PCR CONGEN | |
| SFA | 2 | S3601 | Glycine max | Extraction with SureFood® Prep Advanced protocol 1 (S1053) | K02 |
| SFA | 5 | | | | |
| SFA | 16 | S3601 | | real time PCR | |
| SFA | 20 | SureFood® ALLERGEN Soya - Art. No. S3601 | The real-time PCR test detects soya DNA (Glycine max) | DNA preparation with SureFood® PREP Advanced (Principle according to protocol 2: Lysis at 65°C - Pre-filtration and setting of optimal binding conditions - Binding of the nucleic acids on a Spin Filter - Purification of the bound nucleic acids - Drying of the Spin Filter - First Elution of nucleic acids from the Spin Filter - Repeated setting of optimal binding conditions - Second binding of the nucleic acids on a Spin Filter - Second purification of the bound nucleic acids - Drying of the Spin Filter - Elution of nucleic acids from the Spin Filter for analysis) and real-time PCR (45 cycles following kit setup instructions) with Bio-Rad CFX96, Lot 24060 | |
| SFA | 21 | | | , | |
| div | 4 | internal method | | CTAB / Proteinase K / Promega Wizard DNA CleanUp / Real-time PCR 45 cycles | |
| div | 9 | | | | |

5.2 Homogeneity

5.2.1 Mixture homogeneity before bottling

Microtracer Homogeneity Test

DLA ptALS2 Sample 1

Weight whole sample 1,01 kg Microtracer FSS-rot lake Particle size 75 – 300 μm Weight per particle 2,0 Addition of tracer 20,6 mg/kg

Result of analysis

| Sample | Weight [g] | Particle number | Particles [mg/kg] |
|--------|------------|--------------------|----------------------|
| 1 | 4,97 | 43 | 17,3 |
| I | 4,97 | 43 | 5, ۱۲ |
| 2 | 5,02 | 44 | 17,5 |
| 3 | 5,01 | 50 | 20,0 |
| 4 | 4,98 | 44 | 17,7 |
| 5 | 4,98 | 41 | 16,5 |
| 6 | 5,00 | 47 | 18,8 |
| 7 | 5,03 | 47 | 18,7 |
| 8 | 5,02 | 48 | 19,1 |

| Poisson distribution | | |
|----------------------|------|-----------|
| Number of samples | 8 | |
| Degree of freedom | 7 | |
| Mean | 45,5 | Particles |
| Standard deviation | 2,85 | Particles |
| χ² (CHI-Quadrat) | 1,25 | |
| Probability | 99 | % |
| Recovery rate | 88 | % |

| Normal distribution | | |
|----------------------------|------|-------|
| Number of samples | 8 | |
| Mean | 18,2 | mg/kg |
| Standard deviation | 1,14 | mg/kg |
| rel. Standard deviaton | 6,3 | % |
| Horwitz standard deviation | 10,3 | % |
| HorRat-value | 0,61 | |
| Recovery rate | 88 | % |

Microtracer Homogeneity Test DLA ptALS2 Sample 2

Weight whole sample 1,00 kg Microtracer FSS-rot lake 75 – 300 μm Particle size Weight per particle 2,0 μg Addition of tracer 29,3 mg/kg

Result of analysis

| Sample | Weight [g] | Particle number | Particles [mg/kg] |
|--------|------------|--------------------|----------------------|
| 1 | 5,03 | 67 | 26,6 |
| 2 | 4,97 | 75 | 30,2 |
| 3 | 5,03 | 72 | 28,6 |
| 4 | 5,02 | 74 | 29,5 |
| 5 | 5,02 | 77 | 30,7 |
| 6 | 5,03 | 65 | 25,8 |
| 7 | 4,99 | 78 | 31,3 |
| 8 | 4,96 | 64 | 25,8 |

| Poisson distribution | | |
|----------------------|------|-----------|
| Number of samples | 8 | |
| Degree of freedom | 7 | |
| Mean | 71,5 | Particles |
| Standard deviation | 5,51 | Particles |
| χ² (CHl-Quadrat) | 2,97 | |
| Probability | 89 | % |
| Recovery rate | 97 | % |

| Normal distribution | | |
|----------------------------|------|-------|
| Number of samples | 8 | |
| Mean | 28,6 | mg/kg |
| Standard deviation | 2,20 | mg/kg |
| rel. Standard deviaton | 7,7 | % |
| Horwitz standard deviation | 9,7 | % |
| HorRat-value | 0,80 | |
| Recovery rate | 97 | % |

Microtracer Homogeneity Test DLA ptALS2 Sample 3

Result of analysis

| Sample | Weight [g] | Particle | Particles |
|--------|------------|----------|-----------|
| Campic | | number | [mg/kg] |
| 1 | 5,05 | 65 | 25,7 |
| 2 | 5,04 | 71 | 28,2 |
| 3 | 5,01 | 63 | 25,1 |
| 4 | 4,98 | 57 | 22,9 |
| 5 | 4,97 | 66 | 26,6 |
| 6 | 5,05 | 66 | 26,1 |
| 7 | 4,96 | 68 | 27,4 |
| 8 | 5.01 | 65 | 25.9 |

| 8 | |
|------|--------------|
| - | |
| / | |
| 65,1 | Particles |
| 3,96 | Particles |
| 1,69 | |
| 98 | % |
| 05 | % |
| | 3,96 1,69 |

| Normal distribution | | |
|----------------------------|------|-------|
| Number of samples | 8 | |
| Mean | 26,0 | mg/kg |
| Standard deviation | 1,58 | mg/kg |
| rel. Standard deviaton | 6,1 | % |
| Horwitz standard deviation | 9,8 | % |
| HorRat-value | 0,62 | |
| Recovery rate | 95 | % |

Microtracer Homogeneity Test DLA ptALS2 Sample 4

Result of analysis

| Sample | Weight [g] | Particle number | Particles [mg/kg] |
|--------|------------|--------------------|----------------------|
| 1 | 5,05 | 49 | 19,4 |
| 2 | 5,00 | 45 | 18,0 |
| 3 | 5,00 | 49 | 19,6 |
| 4 | 4,99 | 46 | 18,4 |
| 5 | 4,98 | 48 | 19,3 |
| 6 | 5,01 | 41 | 16,4 |
| 7 | 5,00 | 48 | 19,2 |
| 8 | 4,98 | 50 | 20,1 |

| Poisson distribution | | |
|----------------------|------|-----------|
| Number of samples | 8 | |
| Degree of freedom | 7 | |
| Mean | 47,0 | Particles |
| Standard deviation | 2,95 | Particles |
| χ² (CHI-Quadrat) | 1,29 | |
| Probability | 99 | % |
| Recovery rate | 91 | % |

| Normal distribution | | |
|----------------------------|------|-------|
| Number of samples | 8 | |
| Mean | 18,8 | mg/kg |
| Standard deviation | 1,18 | mg/kg |
| rel. Standard deviaton | 6,3 | % |
| Horwitz standard deviation | 10,3 | % |
| HorRat-value | 0,61 | |
| Recovery rate | 91 | % |

5.3 Information on the Proficiency Test (PT)

Before the PT the participants received the following information in the sample cover letter:

| PT number | DLA ptALS2 (2020) | |
|--------------------------------------|--|--|
| PT name | Allergen-Screening II - 4 Samples qualitative: Crustaceae, Egg, Fish, Milk, Molluscs, Mustard (yellow/white, brown and black), Soybean | |
| Sample matrix | Samples 1-4: Carrier matrix / ingredients: potato powder (appr. 75%), maltodextrin (appr. 25%), other food additives and allergenic foods | |
| Number of samples and sample amount | 4 different Samples 1-4: 20 g each | |
| Storage | Samples A + B: room temperature (PT period), cooled 2 - 10°C (long term) | |
| Intentional use | Laboratory use only (quality control samples) | |
| Parameter | Qualitative: Crustaceae, Egg, Fish, Milk, Molluscs, Mustard (yellow/white, brown and black) and Soybean Samples 1-4: appr. 25 - 250 mg/kg | |
| Methods of analysis | The analytical methods ELISA (+ Lateral Flow), PCR and LC-MS can be applied for qualitative determinations. | |
| Notes to analysis | The analysis of PT samples should be performed like a routine laboratory analysis. In general we recommend to homogenize a representative sample amount before analysis according to good laboratory practice, especially in case of low sample weights. | |
| Result sheet | One result each should be determined for Samples 1-4. The results should be filled in the result submission file. | |
| Units | posititv / negativ (limit of detection mg/kg) | |
| Number of digits | at least 2 | |
| Result submission | The result submission file should be sent by e-mail to: pt@dla-lvu.de | |
| Last Deadline | the latest August 28th 2020 | |
| Evaluation report | The evaluation report is expected to be completed 6 weeks after deadline of result submission and sent as PDF file by e-mail. | |
| Coordinator and contact person of PT | Matthias Besler-Scharf PhD | |

^{*} Control of mixture homogeneity and qualitative testings are carried out by DLA. Any testing of the content, homogeneity and stability of PT parameters is subcontracted by DLA.

6. Index of participant laboratories

| Teilnehmer / Participant | Ort / Town | Land / Country |
|--------------------------|------------|----------------|
| | | SPAIN |
| | | SPAIN |
| | | USA |
| | | SPAIN |
| | | CANADA |
| | | ITALY |
| | | SPAIN |
| | | Germany |
| | | Germany |
| | | ITALY |
| | | Germany |
| | | FRANCE |
| | | ITALY |
| | | BRAZIL |
| | | GREAT BRITAIN |
| | | Germany |
| | | Germany |
| | | SWEDEN |
| | | SPAIN |
| | | SWITZERLAND |
| | | ITALY |
| | | Germany |
| | | Germany |
| | | Germany |
| | | GREAT BRITAIN |
| | | ITALY |
| | | FRANCE |
| | | GREAT BRITAIN |
| | | GREAT BRITAIN |

[Die Adressdaten der Teilnehmer wurden für die allgemeine Veröffentlichung des Auswerte-Berichts nicht angegeben.]

[The address data of the participants were deleted for publication of the evaluation report.]

7. Index of references

- 1. DIN EN ISO/IEC 17025:2005; Allgemeine Anforderungen an die Kompetenz von Prüf- und Kalibrierlaboratorien / General requirements for the competence of testing and calibration laboratories
- 2. DIN EN ISO/IEC 17043:2010; Konformitätsbewertung Allgemeine Anforderungen an Eignungsprüfungen / Conformity assessment General requirements for proficiency testing
- 3. ISO 13528:2015 & DIN ISO 13528:2009; Statistische Verfahren für Eignungsprüfungen durch Ringversuche / Statistical methods for use in proficiency testing by interlaboratory comparisons
- $4.~\mathrm{ASU}$ §64 LFGB: Planung und statistische Auswertung von Ringversuchen zur Methodenvalidierung / DIN ISO 5725 series part 1, 2 and 6 Accuracy (trueness and precision) of measurement methods and results
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