

Detection of *Staphylococcus aureus* and *Listeria monocytogenes* DNA in artificially contaminated spices

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Location: Berlin, BfR



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- BIOR – supporting partner in task 3.5 of work package 3 **Development and evaluation of rapid qualitative on-site detection/screening methods for biological contaminants** leading by VUP



- Tasks:
 - ✓ Contamination of spices and herbs samples with known concentration of bacterial pathogen (*Staphylococcus aureus*, *Listeria monocytogenes*)
 - ✓ Direct bacterial DNA extraction from spices and herbs using CTAB (cetyltrimethylammoniumbromide) protocol
 - ✓ Detection of bacterial DNA presence in samples by PCR



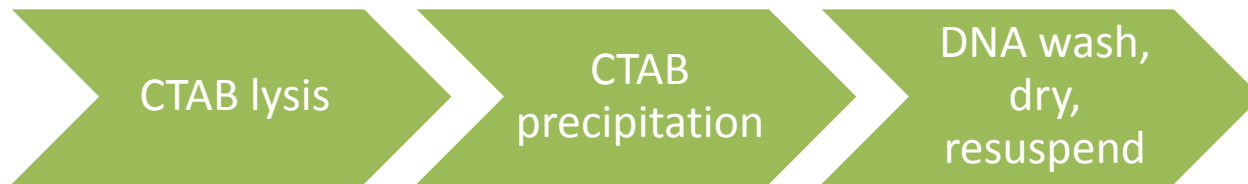
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CTAB DNA extraction

Type of samples:

plant, bacteria,
protozoa, fungi,
insects, animal



Amount of starting material:

μgs
– several grams

+ t°C
+ Proteinase K
+ Rnase
+ β-mercapthoethanol
+ Polyvinylpirrolidone (PVP)

CTAB + low salt conditions

Advantages: low costs, adjustable for a broad type of samples

Disadvantages: laborious, time consuming



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SPICED
spices & herbs

Staphylococcus aureus



- Gram (+), facultative anaerobe, non motile, non sporulate;
- Resistant to high salt conditions and desiccation;
- Common inhabitant of the skin and nasopharynx of humans and animals;
- Produces heat stable enterotoxins;
- *S. aureus* start to produce enterotoxins at concentration of $>10^5$ CFU/g;
- EU Regulation No 2073/2005 lays down food safety criteria for staphylococcal enterotoxins in dairy products:
 - Dairy products must be tested for enterotoxins if coagulase-positive staphylococci are detected at levels $>10^5$ CFU/g.



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Detection of *Staphylococcus aureus*

- Each matrix sample (parsley, basil, cinnamon, vanilla) was contaminated with different concentration (10^2 to 10^6 CFU per gramm) of *S. aureus*;
- DNA extracted with CTAB protocol from sample amount 0.5g;
- A protocol for **16S rRNA** gene specific for *S. aureus* was used
MASON W.J, BLEVINS J.S, BEENKEN K., WIBOWO N., OJHA N., SMELTZER M.S **Multiplex PCR Protocol for the Diagnosis of Staphylococcal Infection.** JOURNAL OF CLINICAL MICROBIOLOGY, Sept. 2001, p. 3332–3338

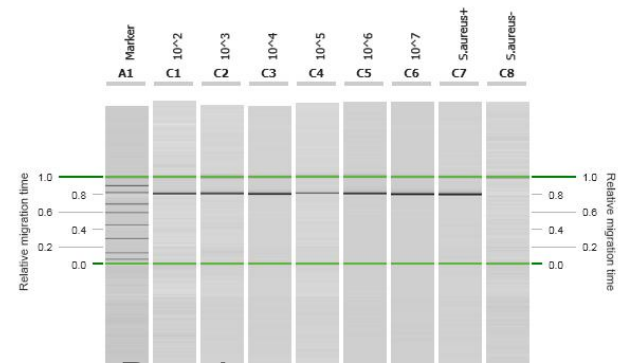


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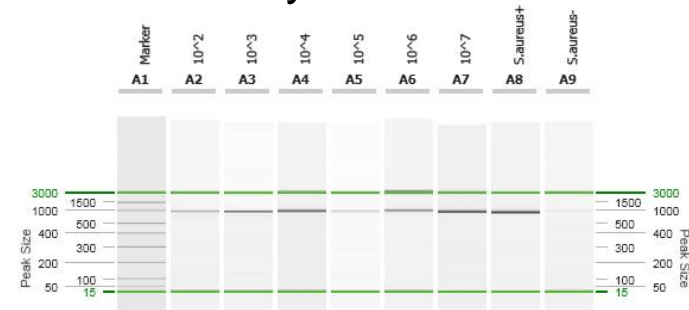


16S rRNA PCR results *S.aureus*

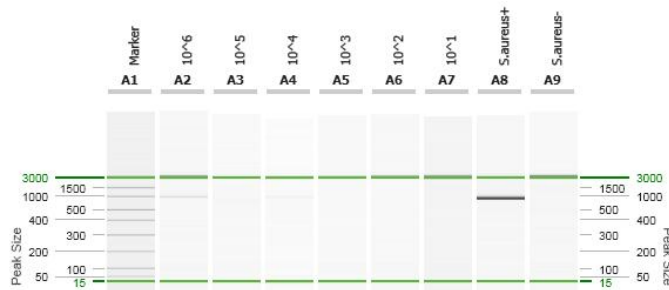
- *S.aureus* DNA was detectable down to 10^2 CFU/g in three matrices – parsley, basil, vanilla
- In cinnamon *S.aureus* DNA was detectable down to 10^4 CFU/g



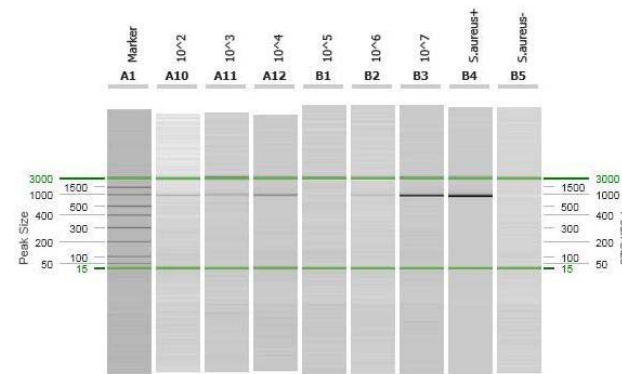
Parsley



Basil



Cinnamon

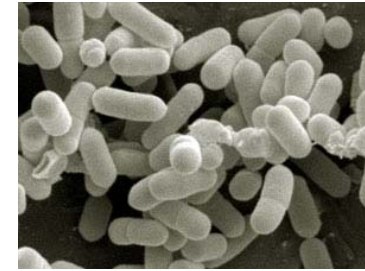


Vanilla



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Listeria monocytogenes



- Ubiquitous food-borne pathogen (soil, vegetation, water ...), can be found in unprocessed foods of animal origin and ready-to-eat (RTE) foods by post-processing contamination;
- Gram (+), facultative anaerobe, can grow at refrigerator temperatures, resistant for low pH
- EU Regulation No 2073/2005 lays down food safety criteria for *L. monocytogenes* in RTE foods:
 - In RTE food for infants and special medical purposes *L. monocytogenes* must not be present in 25 g,
 - *L. monocytogenes* must not be present in levels above 100 cfu/g during the shelf life of other RTE products.



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Detection of *Listeria monocytogenes*

- Each matrix sample (black pepper, paprika/chili, parsley, oregano, basil, cinnamon, nutmeg, allspice) was contaminated with different concentration (10^2 to 10^6 cfu per gramm) of *L. monocytogenes*;
- DNA extracted with CTAB protocol and CTAB + polyvinylpyrrolidone from sample amount 0.2g in duplicates;
- A universal bacterial primers for 16S rRNA gene
S. DORN-IN, R.T BASSITTA, K.SCHWAIGER, J. BAUER, C. S. HÖLZEL. **Specific amplification of bacterial DNA by optimized so-called universal bacterial primers in samples rich of plant DNA.** Journal of Microbiological Methods 113 (2015) 50–56
- Listeria monocytogenes prfA gene specific primers
A. KÉROUANTON, M.MARAULT, L.PETIT, J.GROUT, T. T. DAO, A. BRISABOIS. **Evaluation of a multiplex PCR assay as an alternative method for Listeria monocytogenes serotyping,** Journal of Microbiological Methods 80 (2010) 134–137
- Conventional and real time PCR detection for **prfA** gene in two replicates for each extraction was performed.



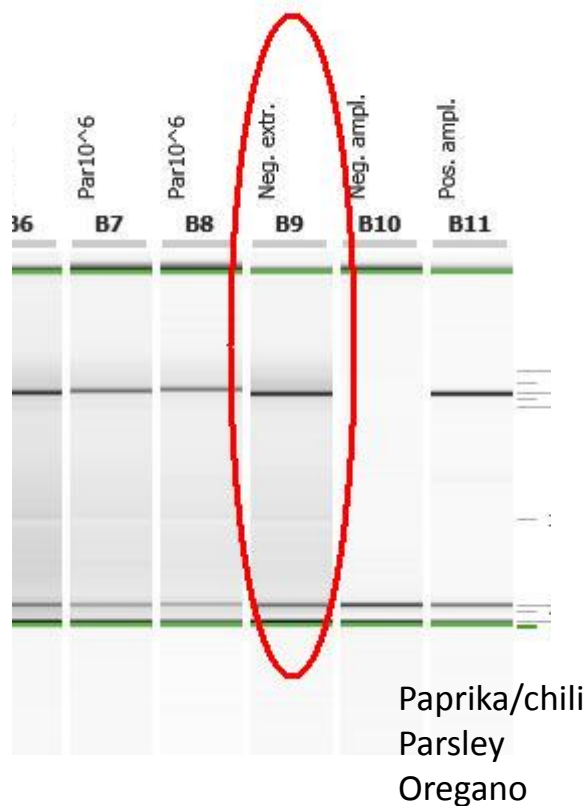
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Detection of *Listeria monocytogenes*

• A universal bacterial primers for 16S rRNA gene

S. DORN-IN, R.T BASSITTA, K.SCHWAIGER, J. BAUER, C. S. HÖLZEL. **Specific amplification of bacterial DNA by optimized so-called universal bacterial primers in samples rich of plant DNA.** Journal of Microbiological Methods 113 (2015) 50–56



NCBI BLAST/blastn suite/ Formatting Results - 57VCWCZ015

Oregano_negative

RID 57VCWCZ015 (Expires on 11-25 11:54 am)

Query ID Id|Query_46519
Description Oregano_negative
Molecule type nucleic acid
Query Length 563

Database Name nr
Description Nucleotide collection (nt)
Program BLASTN 2.3.0+ > Citation

Other reports: > Search Summary [Taxonomy reports] [Distance tree of results]

Graphic Summary
Descriptions

Sequences producing significant alignments:

Select: All None Selected 0

Alignments	Description	Max score	Total score	Query cover	E value	Ident	Accession
<input type="checkbox"/>	Ostracum vulgare subsp. vulgare chloroplast complete genome	1033	2066	100%	0.0	99%	J080002.1
<input type="checkbox"/>	Scutellaria insularis chloroplast complete genome	1027	2055	100%	0.0	99%	KT750009.1
<input type="checkbox"/>	Scutellaria baicalensis chloroplast complete genome	1027	2055	100%	0.0	99%	KR233163.2

Parsley_negative

RID 57VNC585015 (Expires on 11-25 11:59 am)

Query ID Id|Query_134041
Description Parsley_negative
Molecule type nucleic acid
Query Length 929

Database Name nr
Description Nucleotide collection (nt)
Program BLASTN 2.3.0+ > Citation

Other reports: > Search Summary [Taxonomy reports] [Distance tree of results]

Graphic Summary
Descriptions

Sequences producing significant alignments:

Select: All None Selected 0

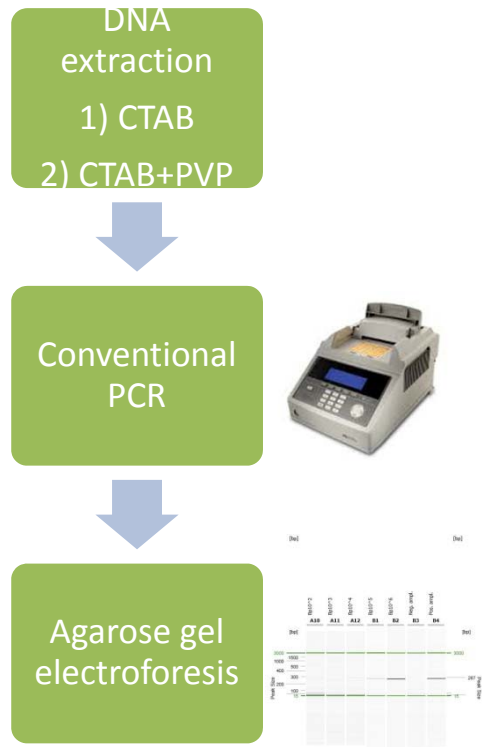
Alignments	Description	Max score	Total score	Query cover	E value	Ident	Accession
<input type="checkbox"/>	Anopelia deursiva voucher 170-92-1 chloroplast complete genome	1707	3414	99%	0.0	99%	KT781591.1
<input type="checkbox"/>	Heuchera paniculata var. saurensis voucher F-alk 97 (QS) chloroplast complete genome	1707	3414	99%	0.0	99%	KR473645.1
<input type="checkbox"/>	Heuchera paniculata var. saurensis voucher F-alk 97 (QS) complete genome	1707	2005	99%	0.0	99%	KR556021.1
<input type="checkbox"/>	Panax quinquefolius chloroplast complete genome	1707	3414	99%	0.0	99%	KT028714.1
<input type="checkbox"/>	Bupleurum falcatum chloroplast complete genome	1707	3414	99%	0.0	99%	KM207876.1



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Detection of *Listeria monocytogenes*



•*Listeria monocytogenes* **prfA** gene specific primers conventional PCR

Limit of detection of CTAB and CTAB+PVP extraction protocols combined with conventional PCR targeting *L.monocytogenes* *prfA* gene

Matrix	LOD CFU/g	
	CTAB	CTAB+PVP ¹⁾
Black pepper	>10 ⁵ ²⁾	10 ⁵
Paprika/chili	10 ⁵	10 ⁵
Oregano	10 ⁵	10 ⁶
Parsley	10 ⁶	10 ⁵
Basil	>10 ⁶	>10 ⁶
Nutmeg	>10 ⁶	10 ⁶
Allspice	>10 ⁶	>10 ⁶
Cinnamon	>10 ⁶	>10 ⁶

- One extraction was performed for each matrix
- Two of four tests were positive for this contamination level

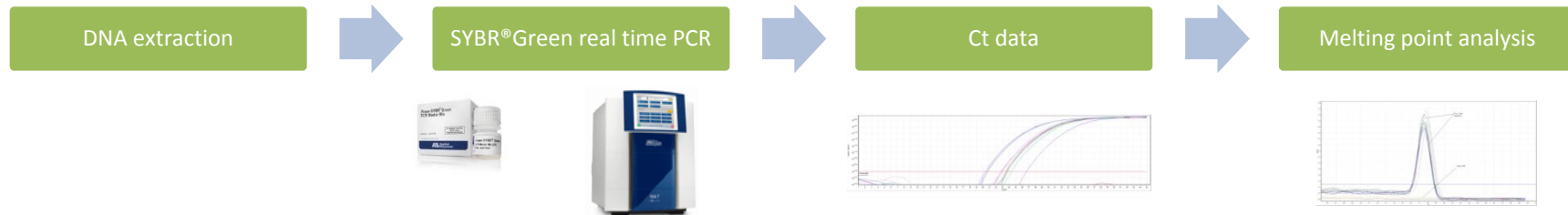


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Detection of *Listeria monocytogenes*

• *Listeria monocytogenes* prfA gene specific primers SYBR®Green real time PCR



Limit of detection and mean Ct values at level of contamination 10⁶ CFU/g of CTAB and CTAB+PVP extraction protocols combined with SYBR Green real-time PCR targeting *L.monocytogenes prfA* gene.

Matrix	CTAB		CTAB+PVP	
	LOD CFU/g	Ct at 10 ⁶ CFU/g	LOD CFU/g	Ct at 10 ⁶ CFU/g
Black pepper	10 ⁵	26.65	10 ⁴	25.00
Paprika/chili	10 ³	25.90	10 ³	25.35
Oregano	10 ⁵	27.03	10 ⁴	26.70
Parsley	10 ⁵	28.55	10 ⁴	27.20
Basil	10 ⁵	29.57	10 ³	20.52
Nutmeg	>10 ⁶		10 ⁵	30.38
Allspice	>10 ⁶		10 ⁶	30.71
Cinnamon	>10 ⁶		10 ⁶	30.10

Two extractions were performed for each matrix with two PCR replicates for each extraction.



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Conclusions

- Use of CTAB method for direct bacterial DNA isolation was successful for all spices and herbs
- Use of PVP in CTAB DNA extraction procedure may improve DNA recovery and PCR performance for some matrices (basil, nutmeg, allspice, cinnamon)
- Additional steps might be needed for difficult matrices like nutmeg, allspice and cinnamon to improve DNA recovery and PCR performance
- The sensitivity 10^3 - 10^4 CFU/g achieved for *Listeria monocytogenes* is not sufficient for routinely testing
- The sensitivity 10^2 - 10^4 CFU/g achieved for *Staphylococcus aureus* is sufficient for routinely testing



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**Thanks for your attention.
Questions and comments.**

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