



Innovative Solutions for *Listeria* Detection and Monitoring

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- **Challenges are increasing:**
 - Increase in susceptible persons with aging population
 - Increase in the amount of ready-to-eat foods
 - Increase in demand for organic foods
- **The economic impact of *Listeria* is enormous:**
 - In terms of human health (medical costs, loss of life)
 - Significant number of foods have been involved in international product recalls
 - Some manufacturers have suffered huge financial losses through recall, extra freight, product disposal, product reworking etc.

- The challenge for the food industry is to develop, implement, maintain and improve programs for *Listeria* monitoring and control
- The reference methods for *Listeria* in the EU are EN/ISO 11290-1 and EN/ISO 11290-2. The classical cultural methods for *Listeria* are time-intensive and laborious (4-7 days)

Solutions:

- Rapid analysis methods
- Reliable results to prove the absence of *Listeria*
- Fast confirmation of microbiological test results
- Long-term surveillance and outbreak studies

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Compared with conventional methods which can take several days, negative results are available with **BACGene** in less than 24 hours and with **BACSpec** in less than two days.

BACGene
real-time
PCR



BACSpec
ELISA



Cultural Method



- **Rapid analysis methods**
- **Reliable results to prove the absence of *Listeria***
- **Fast confirmation of microbiological test results**
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- **BACGene *Listeria* Multiplex (AFNOR certified)**
 - *Listeria* spp.
 - *Listeria monocytogenes*
- **BACGene *Listeria* spp. (AFNOR certified)**
- **BACGene *Listeria monocytogenes* (AFNOR certified)**
- **AOAC certification in preparation**



EGS 38/01-03/15
ALTERNATIVE ANALYTICAL
METHODS FOR AGRIBUSINESS
www.afnor-validation.com

2-in-1 detection of *Listeria monocytogenes* and *Listeria* spp. in one well improves *Listeria* testing!

Improving Time to Results

- **No secondary enrichment step necessary**
- **Simple lysis protocol**

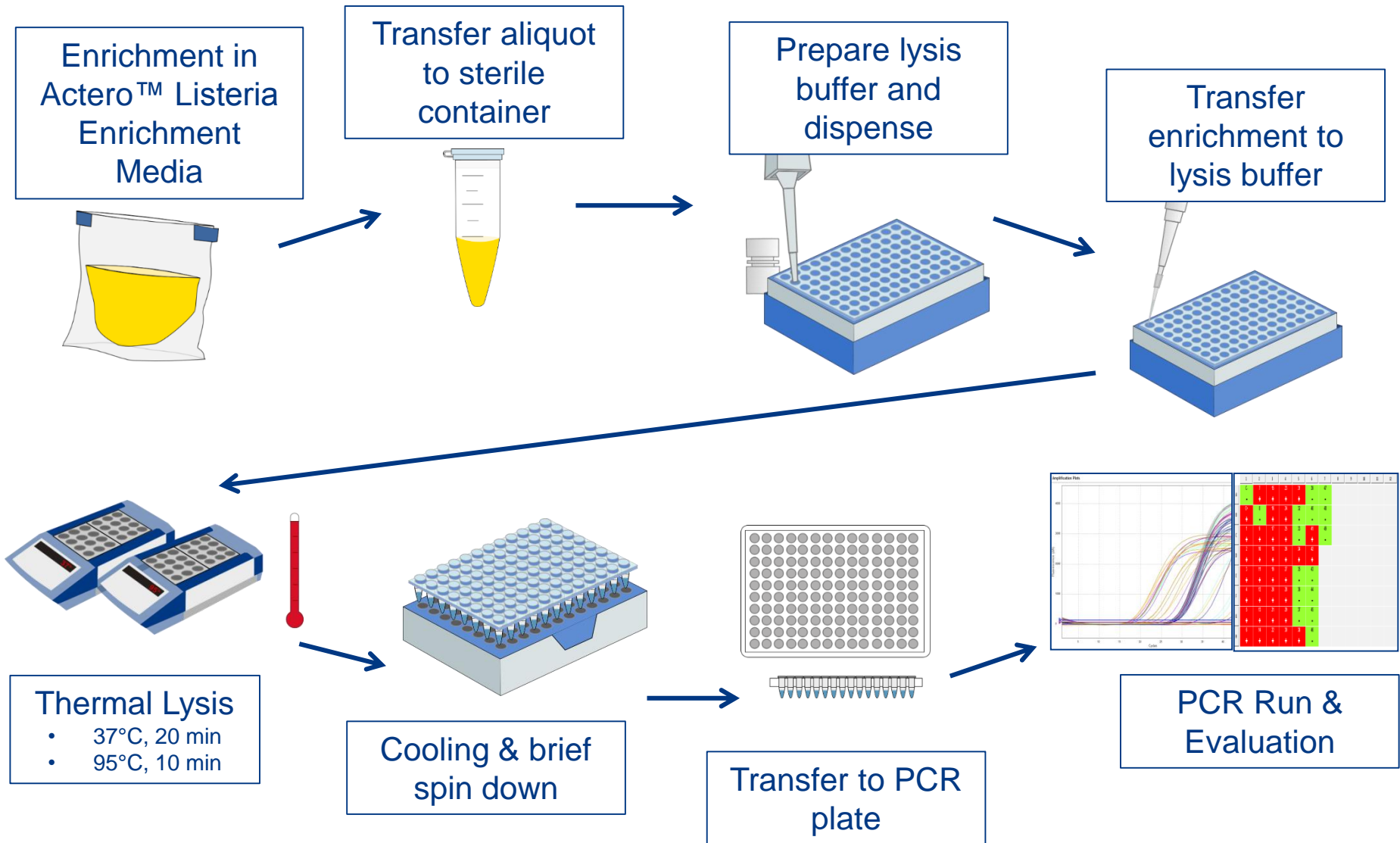
Minimizing Risk

- **Probe-based real-time PCR assay to guarantee excellent specificity**
- **Positive and Negative PCR controls to demonstrate PCR target detection and absence of contamination**
- **Internal Positive Control (IPC) for verification of PCR performance and confirmation of results**

Increasing Reliability

- **High quality kit with AFNOR Certification**
- **Automated data analysis based on validated evaluation algorithm**

Overview: BACGene Listeria



- **Rapid analysis methods**
- **Reliable results to prove the absence of Listeria**
- **Fast confirmation of microbiological test results**
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Confirmation of Presumptive Positives

Germ identification = Confirmation of microbiological test results

Traditional methods

- Microscopy
- Gram coloring
- Biochemical Galleries
- Antibodies

Defects of these methods

- Long time of answer
- High cost/price
- High technicality
- Difficulty of interpretation
- Sometimes unreliable

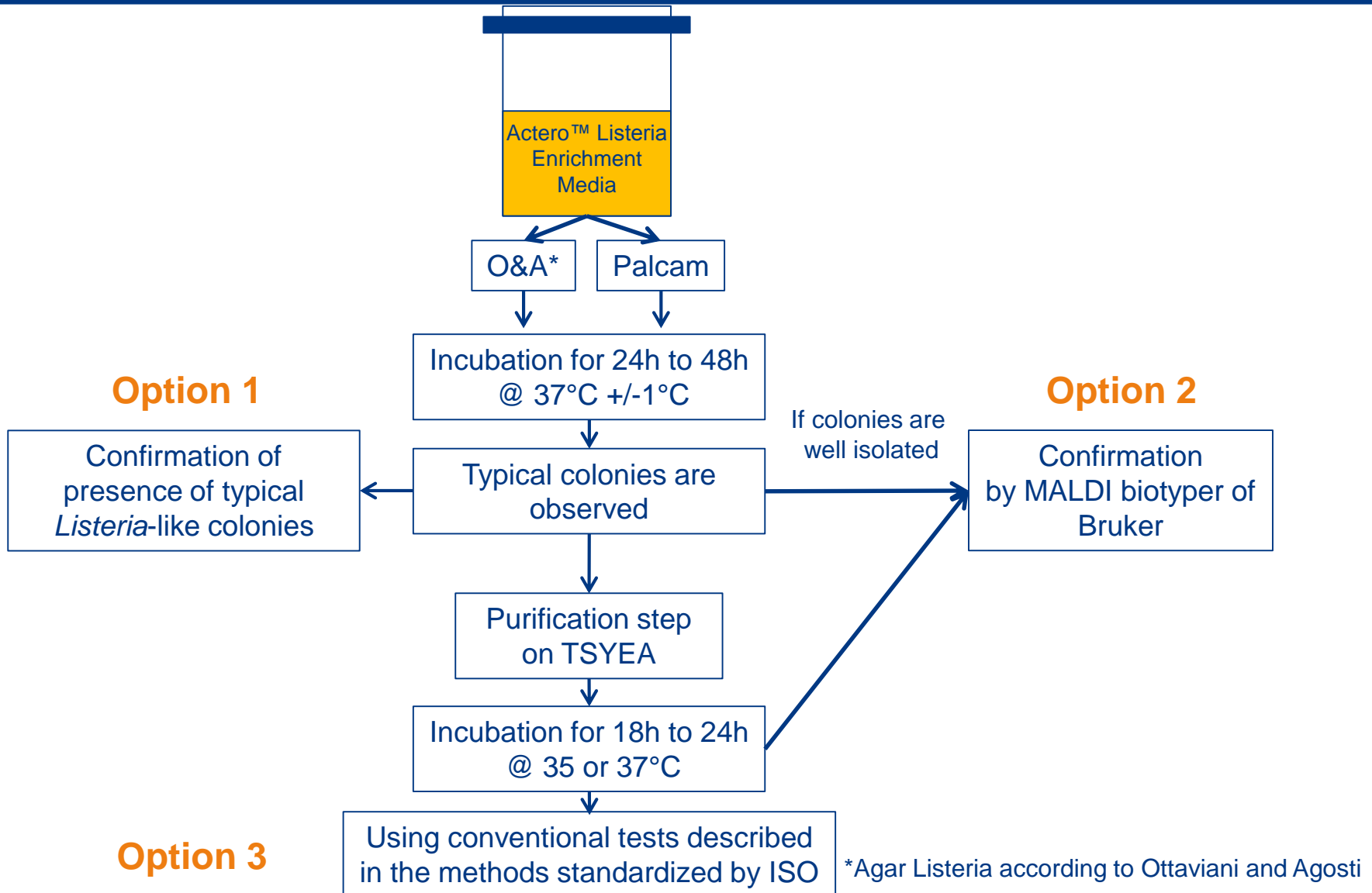


And regularly the result is not clear or unsatisfactory or arrive to late...

Need for a more reliable, faster and cheaper method



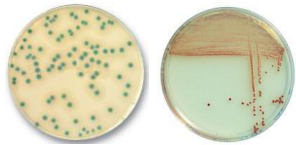
Options for Confirmation of Presumptive Positives



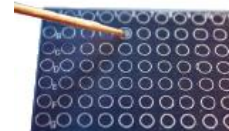
MALDI-TOF MS

Workflow of the method

1 Taking pure and fresh colonies



2 Deposit and covering sample with an adapted chemical matrix



5 Automatic Identification

Listeria monocytogenes	2.136
Listeria ivanovii	2.062
Salmonella sp	2.393
Campylobacter coli	2.102

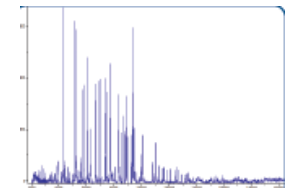


3 Sample Analysis:

Proteins are desorbed, charged with one or two protons and accelerated under a high vacuum ($5 \cdot 10^{-6}$ mBar) in an electric field up to a detector which measures their time of flight and thus their mass. All the analyzed proteins form a characteristic spectrum of each germ (of 2KDa to 20 KDa)



4 Processing of the spectra and comparison to the MALDI-TOF Data Base: more than 7000 reference spectra covering 2400 species and 500 genus of bacteria and yeast. Continuously updated by Eurofins LMO R&D Team



Time-to-result gains:

Listeria (AFNOR certified)

- **ISO 11290-1** **14-50%**
- **BACGene *Listeria* PCR** **~ 25%**

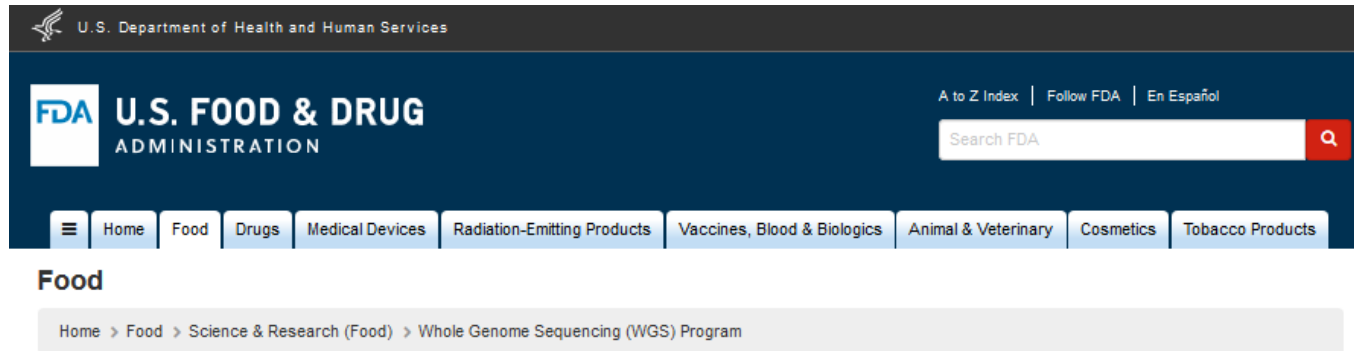
Salmonella

- **ISO6579** **20-50%**
- **BACGene *Salmonella* PCR** **~ 25%**

- **Rapid analysis methods**
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What Benefits?

Whole Genome Sequencing is used and recommended by the FDA:



The screenshot shows the FDA website header with the U.S. Department of Health and Human Services logo, the FDA logo, and the text 'U.S. FOOD & DRUG ADMINISTRATION'. Navigation links include 'A to Z Index', 'Follow FDA', and 'En Español'. A search bar is labeled 'Search FDA'. A menu bar contains links for 'Home', 'Food', 'Drugs', 'Medical Devices', 'Radiation-Emitting Products', 'Vaccines, Blood & Biologics', 'Animal & Veterinary', 'Cosmetics', and 'Tobacco Products'. Below the menu, the 'Food' section is highlighted, and a breadcrumb trail reads: 'Home > Food > Science & Research (Food) > Whole Genome Sequencing (WGS) Program'.

To Provide Benefits to the Food Industry

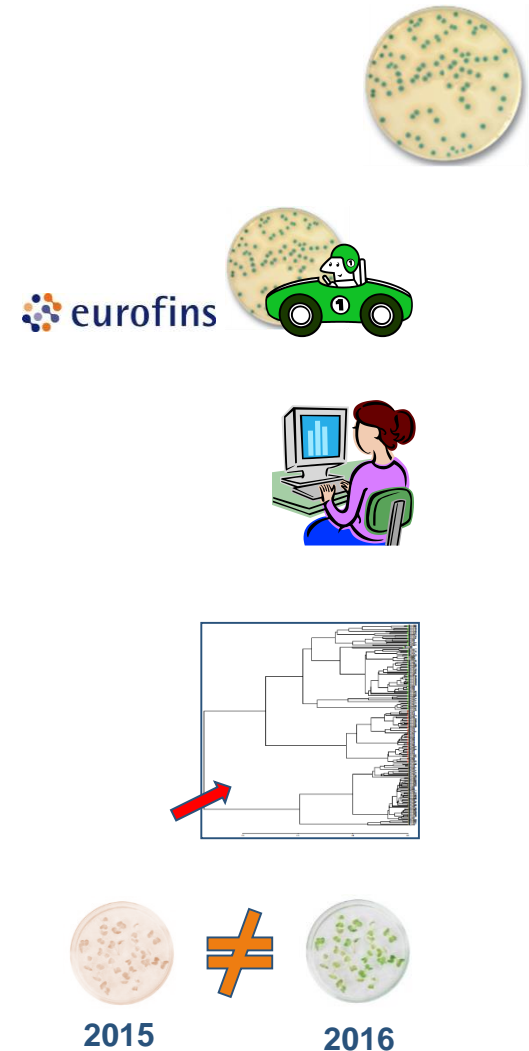
The benefits that come from leveraging the information whole genome sequencing provides aren't just limited to public health officials. Members of the food industry and diagnostic test developers can use it as well.

FDA regulated food manufacturers are required to ensure that their products are safe for consumption. Part of meeting this requirement is making sure that the ingredients they use are free from bacterial pathogens and if a bacterial pathogen is present, that the processing of the food include steps that are sufficient to kill the pathogen. Accordingly many food companies periodically test ingredients received from their suppliers as additional verification of the ingredients' safety. Similarly, companies may conduct environmental sampling in their processing plants to verify the effectiveness of their processing and sanitation controls. By conducting whole genome sequencing on any bacteria isolated from their sampling efforts companies will have detailed information about the pathogen(s) detected. They can then compare this to the genomic information publicly available in GenomeTrakr. In the case of an incoming ingredient, it may reveal information about the route source of the ingredient contamination in their supply chain. It may also identify which preventive or sanitary controls may have failed and need to be corrected.

The genomic information in the publicly accessible GenomeTrakr database can also be drawn on by the food industry, researchers, and instrument makers to develop rapid method and culture independent tests that can be utilized for screening for known pathogen strains.

How does it work?

- You have a **positive pathogen finding** (e.g. Listeria)
- The isolate is picked up and sent to us
- The genome of the isolate is fully sequenced (**Whole Genome Sequencing = WGS**)
- **Sequence Comparison** with
 - your **Private Pathogen Library** and/or
 - **GenomeTrakr Entries from the FDA**
- **Output: Same Strain? yes/no**



Rapid Detection

- **Fast enrichment of *Listeria***
- **Innovative 2-1 multiplex PCR**
 - *Listeria spp.*
 - *Listeria monocytogenes*

Rapid Confirmation

- **MALDI-TOF reduces confirmation time**

Long-Term Surveillance

- **Whole Genome Sequencing**

Laboratories involved in this presentation:

Eurofins GeneScan
Freiburg, Germany

**Eurofins
Laboratoires de
Microbiologie Ouest**
Nantes, France

Eurofins Genomics
Ebersberg, Germany



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Whole Genome Sequencing (WGS)



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*BAC Gene Listeria Multiplex,
MALDI-TOF*



Thank you for your attention!

Questions?