



Detection of *Echinococcus spp.* and other taeniid species in lettuces and berries: two international multicentre studies from the MEmE project.

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MEmE:

Multi-centre study on *E. multilocularis* and *E. granulosus s.l.* in Europe:
development and harmonisation of diagnostic methods in the food chain

International multicentre collaborative project

20 European countries & 3 international external partners

Context:

- **Routes of human AE and CE infection** difficult to identify
 - Long asymptomatic period
- **Scarce data** about foodborne transmission
 - Which relative importance ?

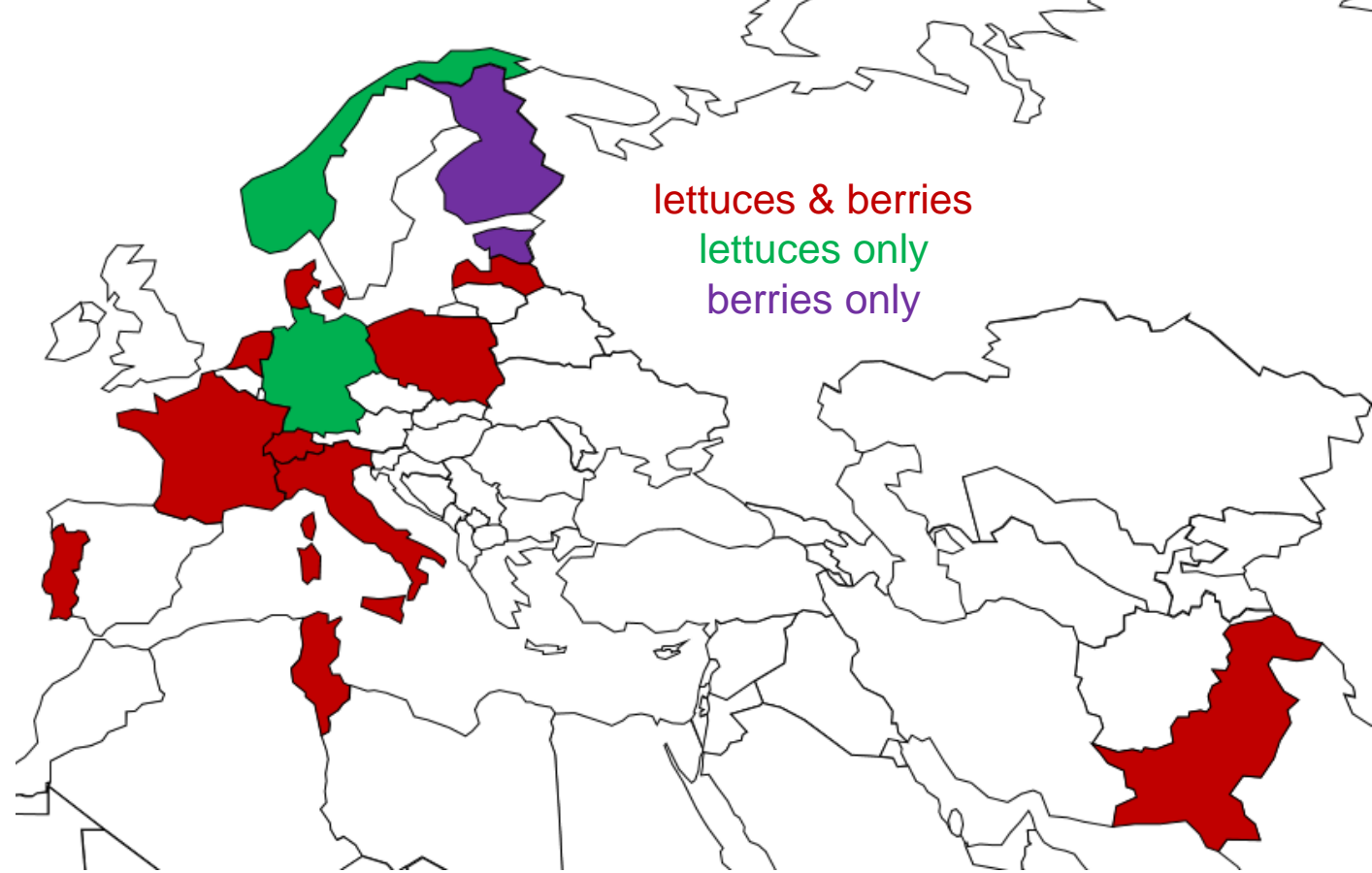
Aim:

**Provide data about the proportion of lettuces and berries with
DNA of Em, Eg and others taeniid species**



Sampling

15 European countries
(18 labs)
+ **Tunisia**
+ **Pakistan**



Summer 2021: 1,117 lettuces + 71 other vegetables
(50-100/lab)

Summer 2022: 300 batches of strawberries + 130 blueberries + 50 others
(20-30/lab)

Collected mainly from local markets (wild)
but also private kitchen garden, supermarkets



Methods

Washing

500ml Tween + hand shaking → sedimentation



Sequential sieving

105µm → 40µm → 20µm

(Guggisberg et al. 2020)



Molecular detection

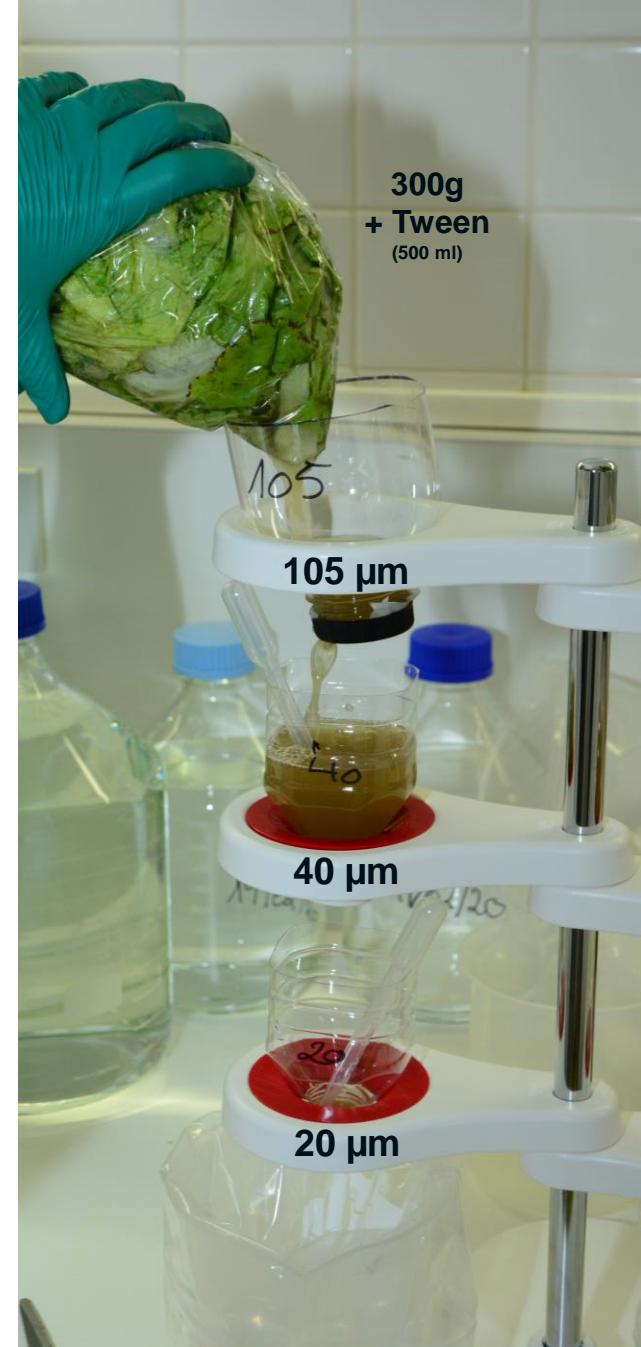
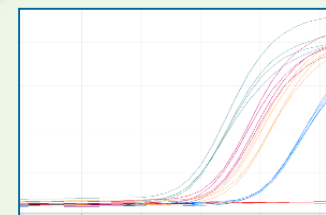
DNA extraction from pellet

Real-time PCR: **Em**

(Knapp et al. 2016)

Eg sl (Maksimov et al. 2020)

PCR/sequencing: **other taeniids** (Trachsel et al. 2007)





Limit of detection

Evaluation in Anses lab conditions

n=24



95%



75%



50%

- 2 eggs (5/5) in original method from Guggisberg et al. 2020

n=8



95%



88%



88%

- Not evaluated previously on berries

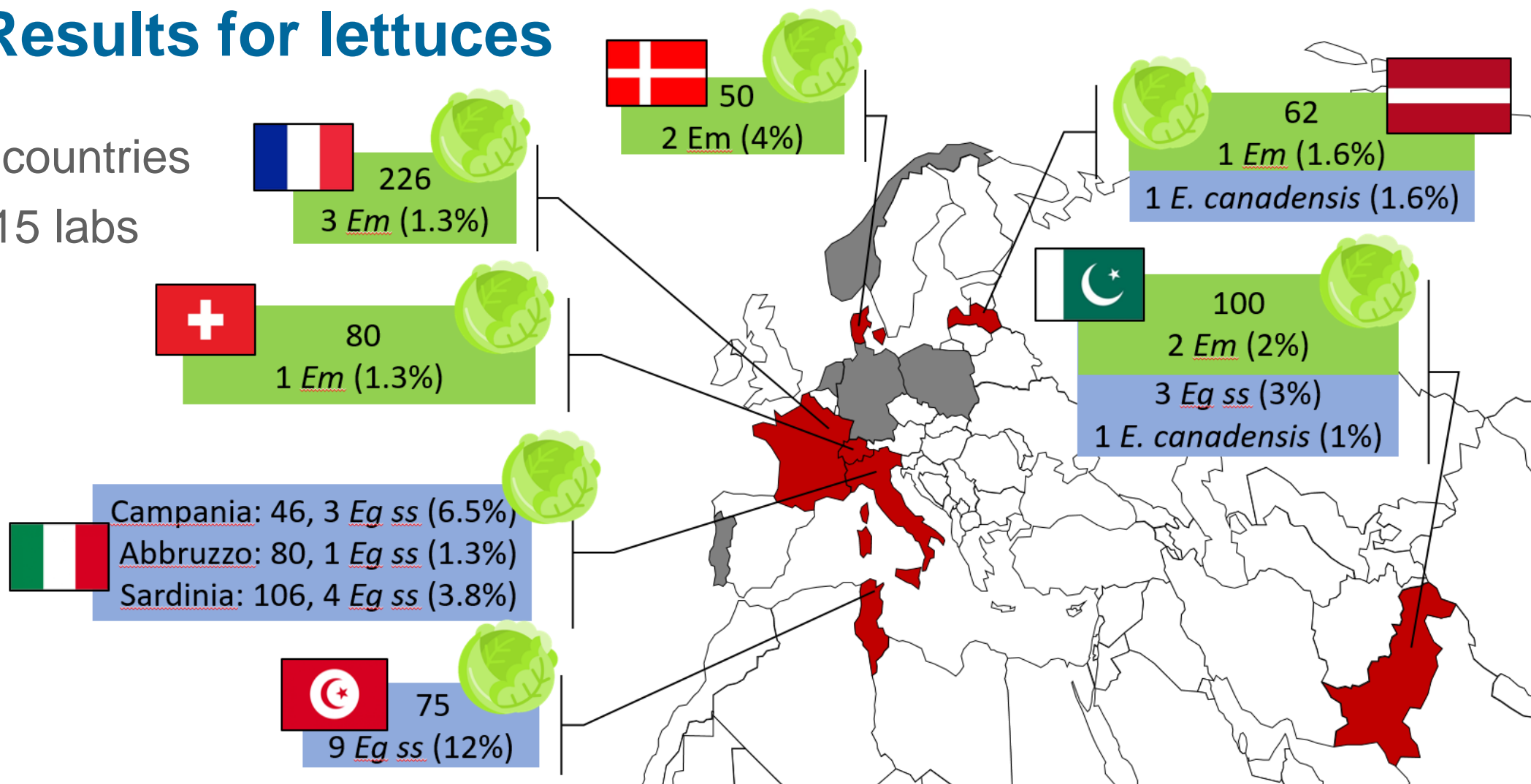
➔ Relevant limit of detection



Results for lettuces

12 countries

15 labs



Global results in European endemic areas:

Em: 1.2% (0,5-2,5)

Eg sl: 1.3% (0,6-2,4)

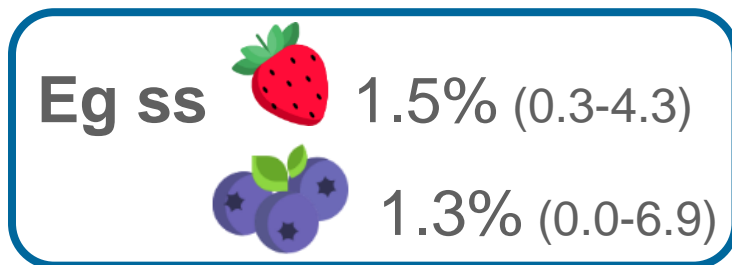
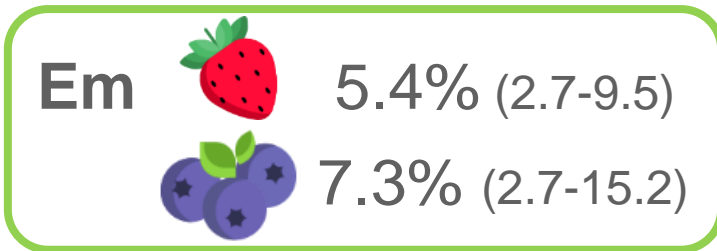
Other taeniid: 1.7% (1-2,7)



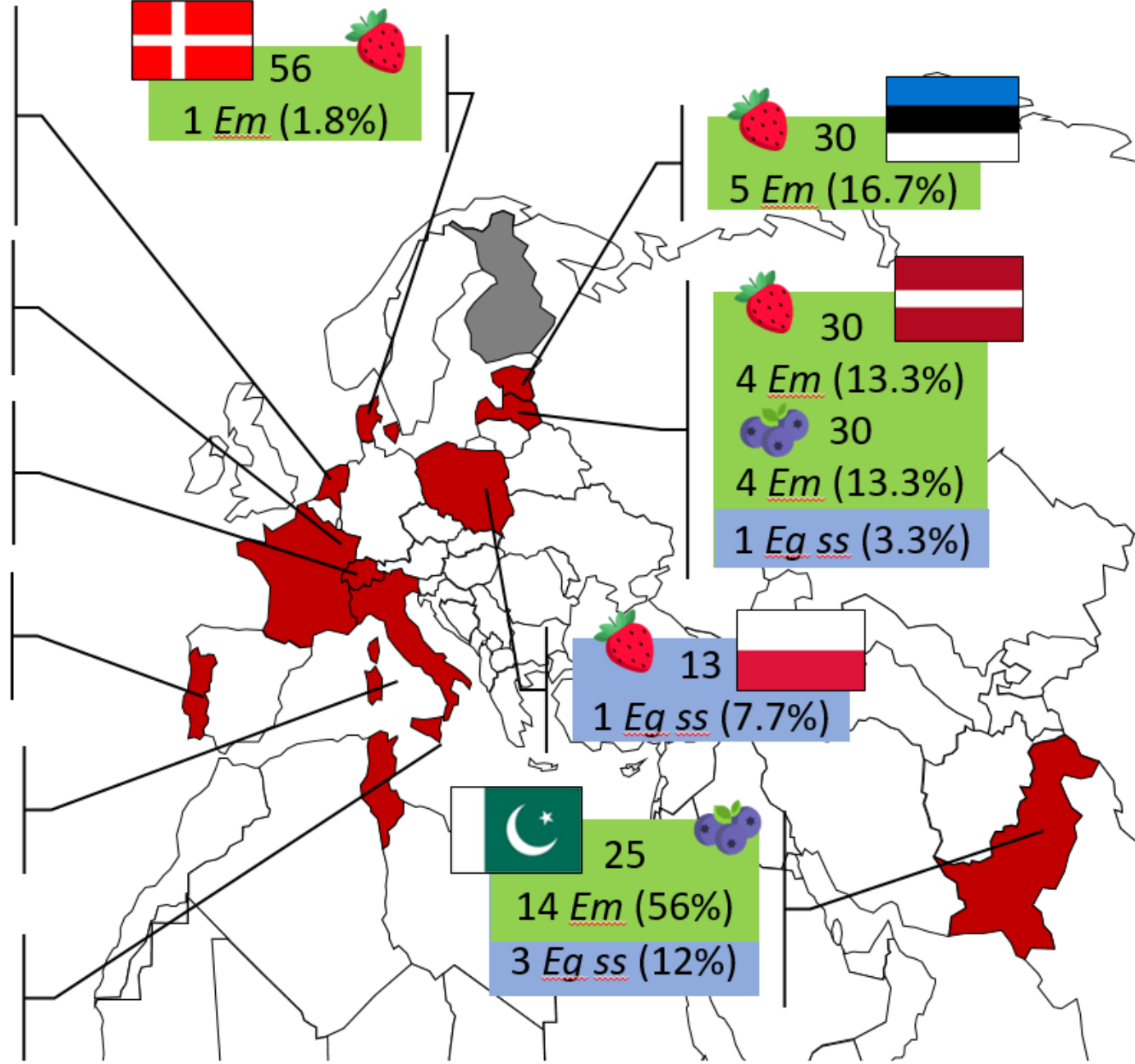
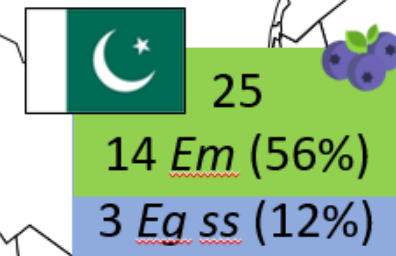
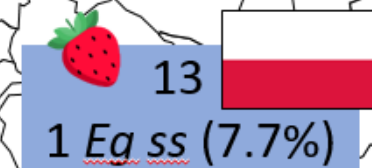
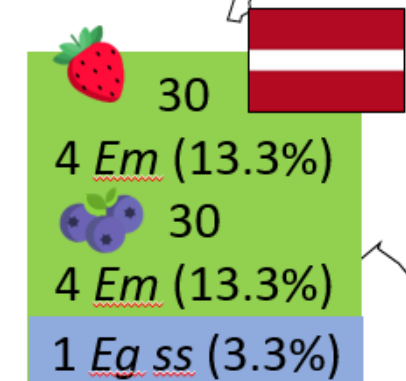
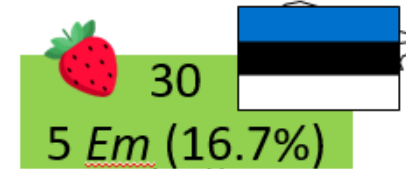
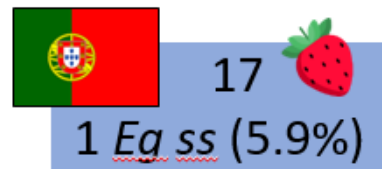
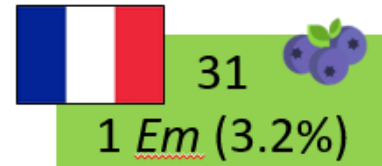
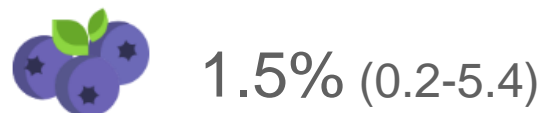
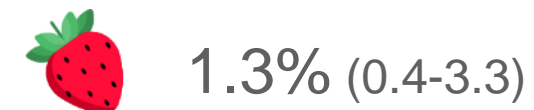
Results for berries

12 countries, 12 labs

European endemic areas



Other taeniids (all areas)





Complementary results

- **Vegetables others than lettuces** (n=69): chard, parsley, basil, sorrel, endive, spinach, beet and carrot leaves ...
 - ✓ **1 chard** from North Germany positive for *E. multilocularis*
- **Others berries** (n=50): raspberries, blackberries, currant, ...
 - ✓ **1 raspberry** from Switzerland positive for *E. multilocularis*
- **Other taeniid species detected**
 - ✓ mainly *Hydatigera* sp. and *T. hydatigena*
- **Global proportion of DNA from all Taenidae species** (Em, Eg, taeniid)
 - ✓ **5.4%** in lettuces
 - ✓ **11.9%** in berries



✓ 1.7%



✓ 2.5%

**Transfer of taeniid eggs from feces to food
is not a rare event**



Discussion

- **One step more in the long road to evaluate the role of human foodborne contamination by *Echinococcus* spp.**
- **Realization of only washing step by each participants**
 - ✓ **Illustrated SOP** (lettuces and berries)
 - ✓ **increases nb of participants** (no new method to perform)
 - ✓ **Facilitate process of fresh samples**
- **Filtration and molecular detection in one lab**
 - ✓ **Easily comparable results:** same method
 - ✓ **Great confidence and robustness** in the results: LOD, reproducibility



Discussion

High proportions of *Em* and *Eg* in lettuces and berries in Europe

- ✓ Even higher in Tunisia and Pakistan
- ✓ In accordance with known high endemic areas
- ✓ Quite similar proportions between European countries
- ✓ Berries significantly more contaminated than lettuces for *E. multilocularis*

Detection of DNA but:

- No observation of eggs
 - ✓ But low sensitivity VS molecular biology
 - ✓ DNA supposed to be from eggs regarding the method
- Estimation of the number of *Echinococcus* eggs (dPCR)
 - ✓ According LOD: **generally 1 to 3 eggs in positive food samples**

No proof of viability of the eggs



Acknowledgments

All the EJPOH MEmE consortium and external partners

