

TESTA WP5

Task 5.2:

Comparison of detection methods for *Ditylenchus* in alfalafa and faba bean seed lots and method validation



GEVES

Expertise & Performance



Background

● Stem and bulb nematode

- Caused by *Ditylenchus dipsaci*
- Hosts: wide range of crops (including *Alfalfa* and faba bean)
- Worldwide distribution
- Symptoms on plant:

swelling and distortion of aerial parts,
necrosis or decaying at the base of the stems



Background

- **Sanitary control.**
 - Transmitted by seeds, seed dust, soil and weeds
 - Quarantine pathogen on alfalfa seeds in Europe **Regulation**

Detection threshold at 0 *D. dipsaci*

Alfalafa

Nematodes on seeds and dust

D. dipsaci

Many different others nematodes

Mainly stage J4 present

Faba bean

Nematodes under the seed coat

D. Dipsaci & *D.gigas*

Less others nematodes

All stages present (juveniles, adults)

Background

- Stem and bulb nematode
- 30 biological races of *D. dipsaci* described
- A giant race, distinguished from *D. dipsaci*
 - ↳ New species named *D. gigas* (Volvás *et al.* 2011)
- Molecular methods have been recently developed to confirm the *Ditylenchus* species.
 - Esquibet *et al.*, 2003; Kerkoud *et al.*, 2007; Volvás *et al.*, 2011.



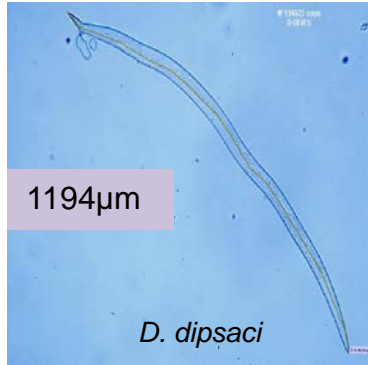
Identification

Morphological criteria difference

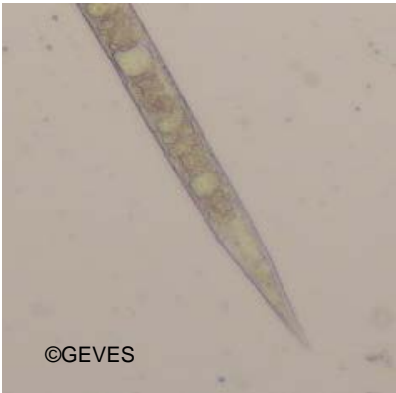
| Criteria | <i>D. Dipsaci</i> (OEPP bulletin 38, 2008) | <i>D. gigas</i> (Volvas et al., 2011) |
|-------------------------|---|--|
| Stylet length (µm) | 10-12µm | 10-12µm |
| Number of lateral lines | 4 | 4 |
| Body length (µm) | 1000-1300µm | 1380-1950µm |
| Body width | 36-41µm | 34—63µm |



Body size



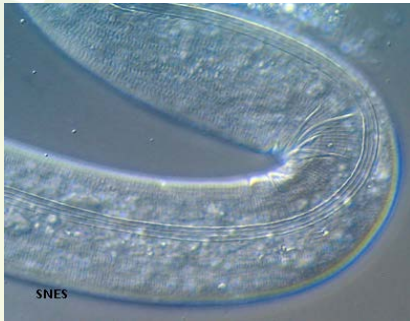
Shape of tail



Shape of lip



Number of line



Aim & Objectives

● Aim

- To harmonize and validate at international level a detection method of *D. dipsaci* and *D. gigas*.

● Objectives:

- To compare performance of the existing different biological and molecular protocols
- To validate a method that enable the detection of the *D. dipsaci* and *D.gigas* : **performance characteristics and interlaboratory test**
- Propose it as an official ISTA and EPPO protocol.



Comparison of methods

| | Decantation method | Filtration method |
|---------------------------|---|--|
| Reference | NIAB : 013 STNEM beans v5 | ANSES : MOA13 part A |
| Sample size (g) faba bean | 500 | 200 |
| Sample size (g) alfalfa | Not tested | 70 |
| Mousseline | Yes dirty samples | Using a paper in the deep of the sieve |
| Soaking | Overnight (17 hours min) | Overnight (24 hours min) |
| Sieving | No, leave 4h water standing, pour off the top liquid, | 250µm and 20µm sieve, |
| Examination | keep 100 ml in Petri dish, x25 | on the remaining liquid x60 |
| Identification | Morphological. In case of doubt → PCR on a pool of nematodes to confirm <i>D. gigas</i> (Wood primers) | Morphological. In case of doubt → PCR on individual nematodes no difference <i>D. dipsaci</i> , <i>D.gigas</i> .(Kerkoud primers) |
| Counting | semi quantitative: light (1-15), medium (15-50), heavy (50-500), very heavy contamination (>500) | No, presence/absence only |



Comparison of methods

● Pretest on alfalfa and faba bean

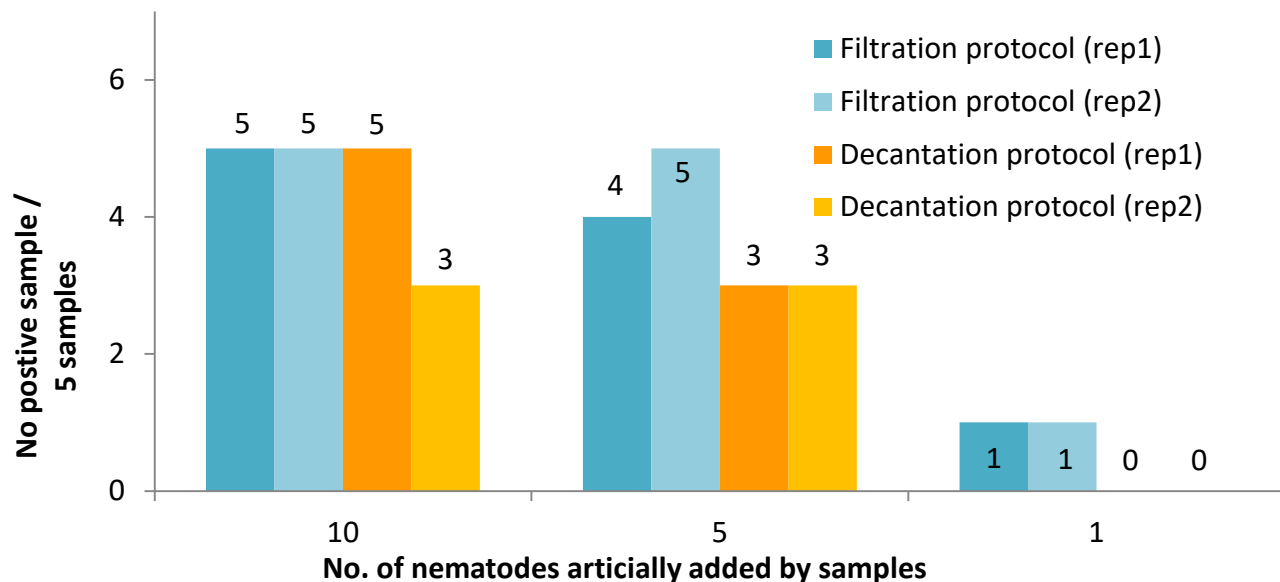
| Seed lots | Filtration protocol (Sieving) | | Decantation protocol (bottom) | | Decantation protocol (Supernatant) | |
|-------------|-------------------------------|---------|-------------------------------|------------|------------------------------------|------------|
| | Nb positive/ nb samples | Couting | Nb positive/ nb samples | Estimation | Nb positive/ nb samples | Estimation |
| Alfaalfa 1 | 5/5 | 843 | 5/5 | 50 to 500 | 5/5 | 1 à 15 |
| Alfaalfa 2 | 5/5 | 1204 | 5/5 | > 500 | 5/5 | 50 à 500 |
| Alfaalfa 3 | 5/5 | 1404 | 5/5 | >500 | 5/5 | 50 à 500 |
| Faba bean 1 | 5/5 | 10190 | 5/5 | > 500 | 5/5 | > 500 |
| Faba bean 2 | 4/5 | 454 | 4/5 | 50 to 500 | 4/5 | 15 to 50 |
| Faba bean 3 | 5/5 | 124 | 4/5 | 15 to 50 | 5/5 | 15 to 50 |

False negative

● Limit of detection

Filtration protocol
High capacity of detection
in low infected seed lots

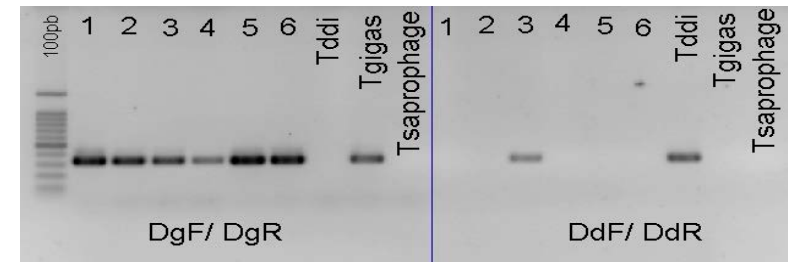
Decantation protocol
limit of detection
higher than 10
nematodes



Comparison of methods

● Primers availables

| Primers | <i>Ditylenchus dipsaci</i> | | <i>Ditylenchus gigas</i> | |
|-----------------------------------|----------------------------|-----------|--------------------------|-----------|
| | | | | |
| Kerkoud <i>et al</i> 2007 | DdpS1 (F) | rDNA2 (R) | DdpS2 (F) | rDNA2 (R) |
| Esquibet <i>et al</i> 2003 | H05 (F) | H06 (R) | D09 (F) | D10 (R) |
| Wood <i>et al</i> 2013 | Dd (F) | Dd (R) | Dg (F) | Dg (R) |

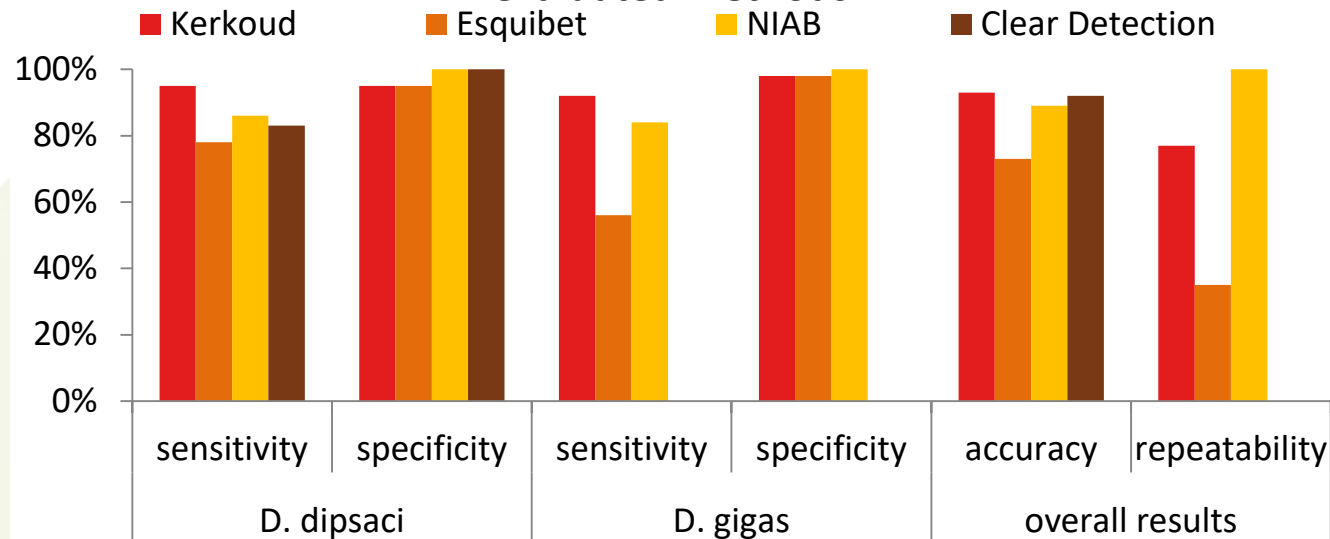


● Obtained results

Kerkoud primers
best results
89 % accuracy

Wood primers
allowing identification
of both species in
mixed samples.

Accuracy and repeatability calculated for the four evaluated methods



Interlaboratory test

- 8 laboratories, filtration method + PCR
- 3 alfalfa and 3 faba bean seed lots (healthy, medium, high)
- Homogeneity test
 - Done on 3 alfalfa and 3 faba bean seed lots provided by the seed industry
 - 10 samples were analyzed per seed lots.

Qualitative analysis of results obtained after the homogeneity test

| | Seed lots | Nb positive samples/total |
|-----------|-----------|---------------------------|
| Alfalfa | Lot A | 0/10 |
| | Lot B | 10/10 |
| | Lot C | 10/10 |
| Faba bean | Lot D | 0/10 |
| | Lot E | 9/10 |
| | Lot F | 8/10 |

Result proved the homogeneity for seed lot A, B, C, and F

Seed lot E and F were not homogenous

The seed lots were used for the validation test

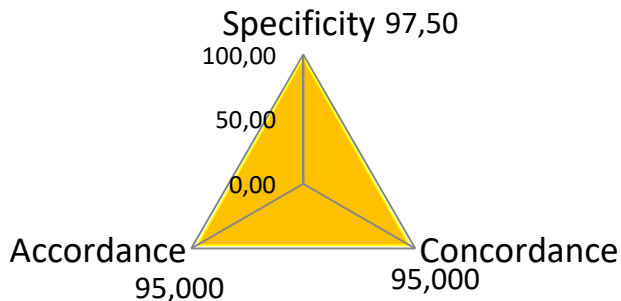
Validation test

● Results

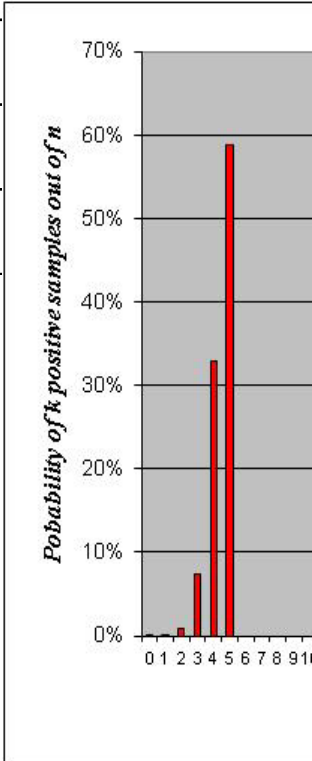
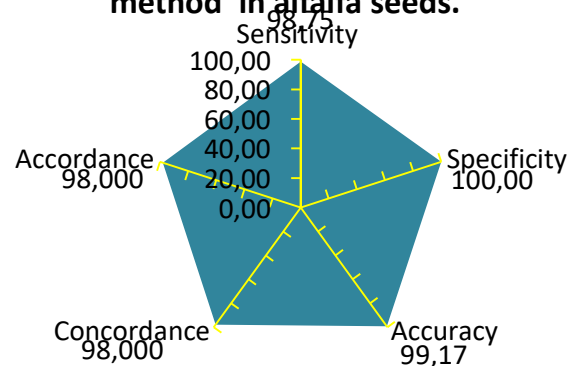
Conform to expected results

| | | | | No. of obtained positive seed subsamples | | | | | | | |
|-----------|---------------------|--|---|--|---|---|---|---|---|---|---|
| | | | | Laboratories | | | | | | | |
| Seed Lot | Contamination Level | No. of expected positive seed subsamples | | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 |
| Alfalfa | Lot A | Healthy | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Alfalfa | Lot B | Low | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 4 |
| Alfalfa | Lot C | Medium | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 |
| Faba bean | Lot D | Healthy | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 1 | 0 |
| Faba bean | Lot E | Low | 5 | 5 | 4 | 5 | 5 | 5 | 5 | 5 | 4 |
| Faba bean | Lot F | Medium | 5 | 5 | 5 | 5 | 5 | 5 | 4 | 5 | 5 |

Detection of Ditylenchus spp. by sieving method in faba bean seeds.



Detection of Ditylenchus spp. by sieving method in alfalfa seeds.



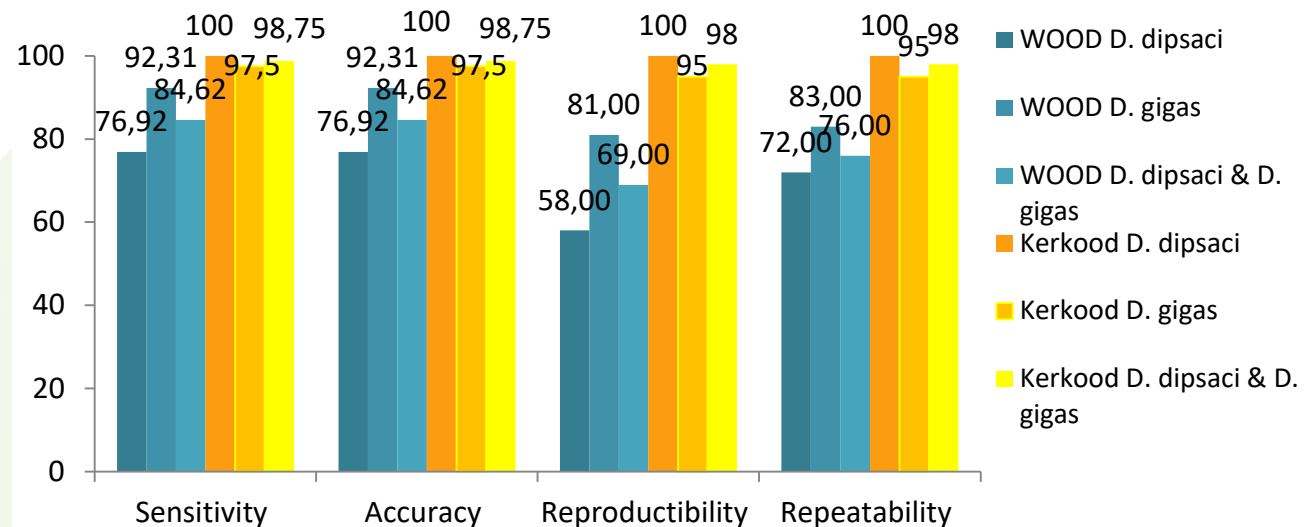
Validation test

Results PCR confirmation

| Primers | Nematodes tested | No. of expected positive sample in PCR | No. of obtained positive seed subsamples | | | |
|---------|-------------------|--|--|----|----|----|
| | | | Laboratories | | | |
| | | | 1 | 2 | 3 | 4 |
| Wood | <i>D. dipsaci</i> | 10 | 5 | 10 | 10 | 5 |
| Wood | <i>D. gigas</i> | 10 | 9 | 10 | 10 | 7 |
| Kerkoud | <i>D. dipsaci</i> | 10 | 10 | 10 | 10 | 10 |
| Kerkoud | <i>D. gigas</i> | 10 | 10 | 10 | 10 | 9 |
| Wood | Both | 20 | 14 | 20 | 20 | 12 |
| Kerkoud | Both | 20 | 20 | 20 | 20 | 19 |

Kerkoud primers best results

97,5 % Accuracy
98,8% Sensitivity
98,0% Concordance
98,0 % Accordance



Summary

- Filtration method tested by 8 laboratories on 30 samples
- Results showed a very good sensitivity, specificity, accuracy, concordance and accordance
 - On alfalfa seed lots (healthy, medium and high infected)
 - On faba bean seed lots (healthy, medium and high infected)

| | Sensitivity | Specificity | Accuracy | Concordance | Accordance |
|--|-------------|-------------|----------|-------------|------------|
| Detection of <i>Ditylenchus</i> spp. by filtration method | >95% | >95% | >95% | >95% | >95% |

- Primers from Kerkoud allow a better distinction between *D. dipsaci* and *D. gigas* only on individual nematode.

Conclusion

- A detection method using filtration (sieve at 20 μ m) in order to concentrate population of nematodes present in a alfalfa and faba bean seed lot is validated.
- A PCR method using Kerkoud primers is validated in order to confirm the species of *Ditylenchus* between *D. dipsaci* and *D. gigas*



GEVES

Groupe d'Étude et de contrôle
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Thanks for your attention

