

## INTRODUCTION

The stem and bulb nematode, *Ditylenchus dipsaci*, causes swelling and deformation of aerial parts and necrosis or rot of stem base. A race, on 30 biological race described, distinguished from *D. dipsaci*, owing to a greater body size was named “giant race” and described as a new species named *D. gigas* (Volvas *et al*, 2011). The detection and identification of *D. dipsaci* and *D. gigas* in seed lots is an obligatory part of the sanitary control and regulation in Europe (import, export and sale of seed lots to farmers) on alfalfa seeds. Few studies have reported the difference on morphological characters between the two pathogens. However molecular methods have been recently developed to confirm the *Ditylenchus* subsp. (Esquibet *et al*, 2003; Kerkoud *et al*, 2007; Volvas *et al*, 2011). The aim of the TESTA project was to harmonize and validate at international level a detection method of *D. dipsaci* and *D. gigas* .



## GOAL

- To compare performance of the biological and molecular protocols currently used in Europe.
- To validate a method which enables the detection of *D. dipsaci* and *D.gigas* and propose it as an official ISTA and EPPO protocol.

## COMPARISON OF PROTOCOLS

### Filtration protocol vs Decantation protocol

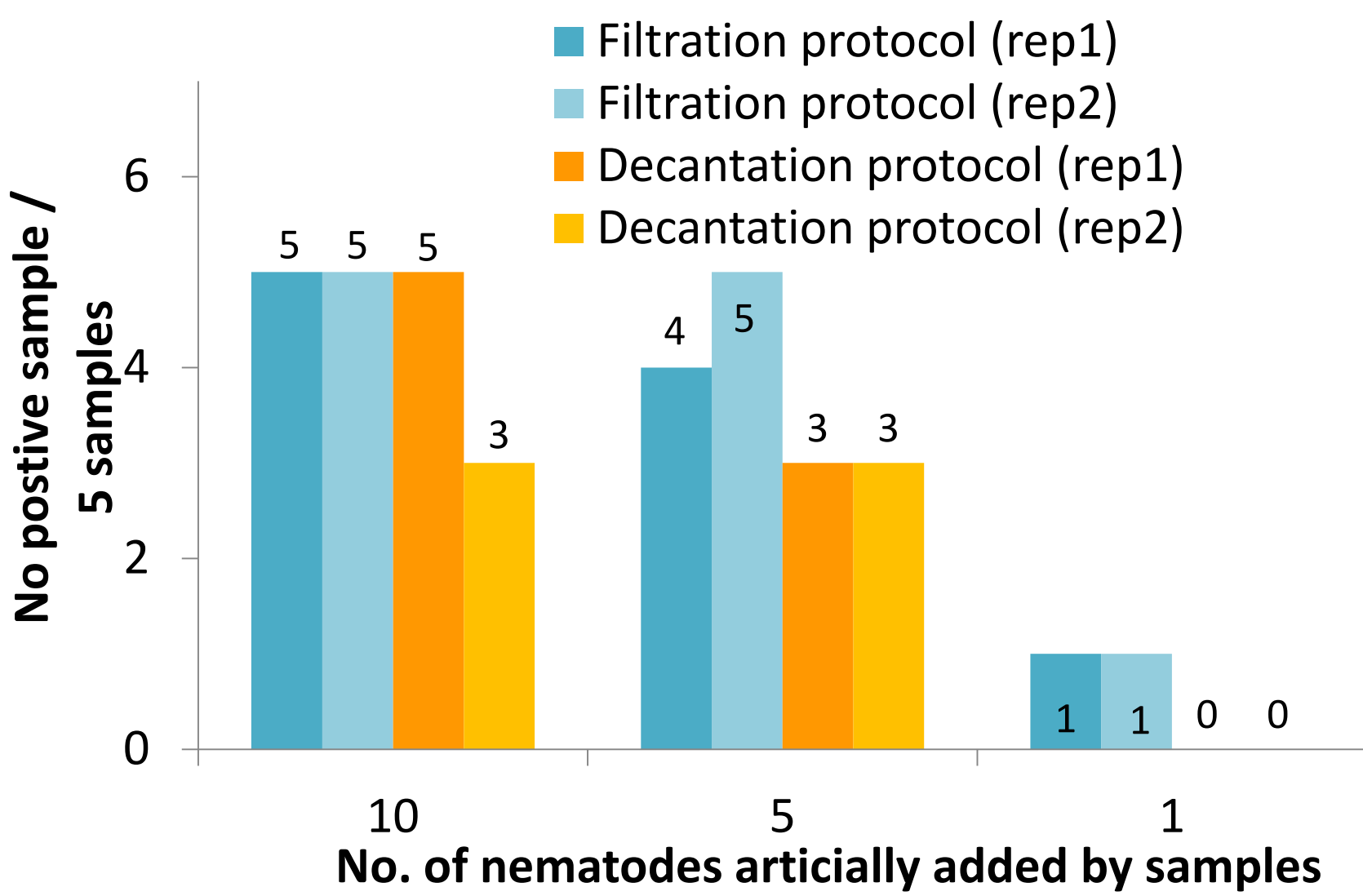
#### Contamination rate in seed lots

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

**Filtration protocol = concentrate population of nematode**  
 High capacity of detection in low infected seed lots

**Decantation protocol = discarded supernatent**  
 Detection of *Ditylenchus* sp. in supernatant  
 → False negative sample

#### Detection limit

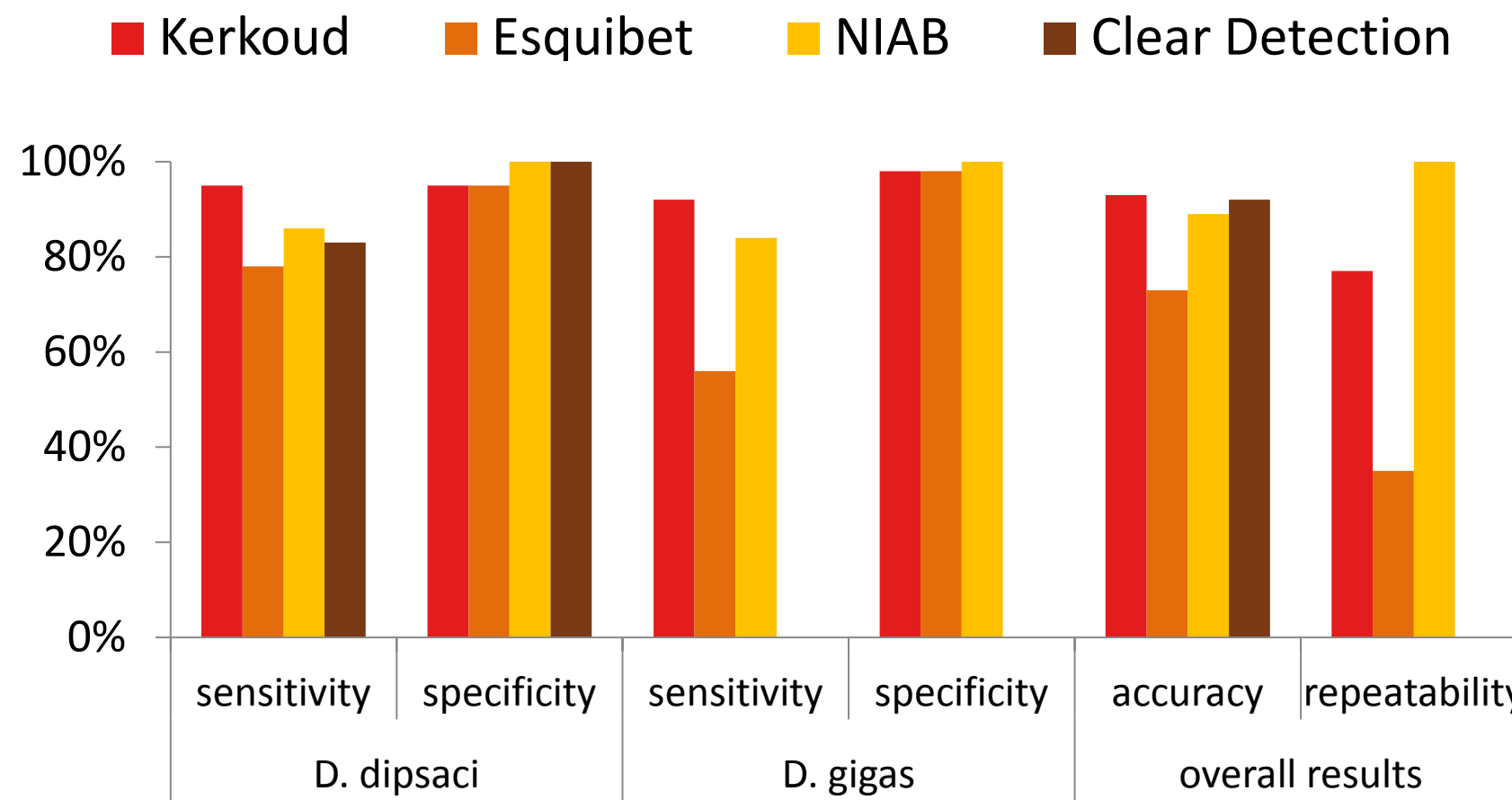


**Filtration protocol**  
limit of detection at 10 nematodes (higher for decantation protocol)  
→ **Method chosen**

### Confirmation by PCR

#### Primers available

Accuracy and repeatability calculated for the four evaluated methods



#### Kerkoud primers

Best results  
89 % accuracy

#### Wood primers

Identification of both species in mixed samples.

## VALIDATION TESTS

Three seed lots (healthy, medium and high infected) of alfalfa and Faba bean were tested for homogeneity and confirmed by qualitative analysis. 8 participants from France, Germany, United Kingdom, Czech Republic and Slovenia tested the filtration method and the Kerkoud and Wood primers. **Performance criteria were studied : Diagnostic specificity, Diagnostic sensitivity, Accuracy, Accordance (repeatability), Concordance (reproducibility)**

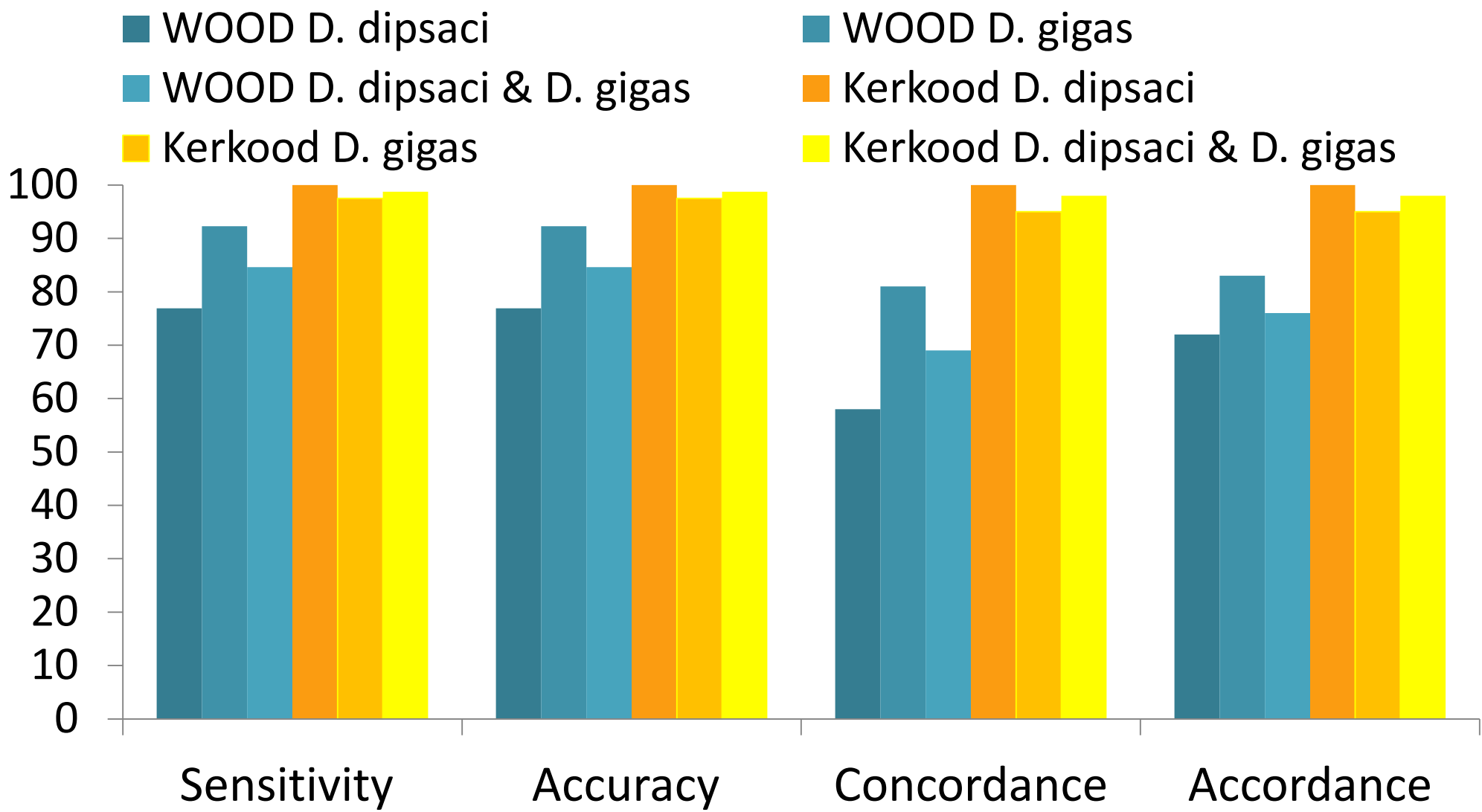
### Homogeneity & Stability test

10 samples were tested per seed lot

		Homogeneity test		Stability test	
Alfalfa	Lot A	0+/10		0+/10	
	Lot B	10+/10		10+/10	
	Lot C	10+/10		10+/10	
Faba bean	Lot D	0+/10		0/10	
	Lot E	9+/10		9+/10	
	Lot F	8+/10		10+/10	

**Conform to expected results. For heterogenous seed lot E and F**  
No. of positive subsample conformed to expected positive subsamples out of the 10 tested at a 95% confidence

### PCR

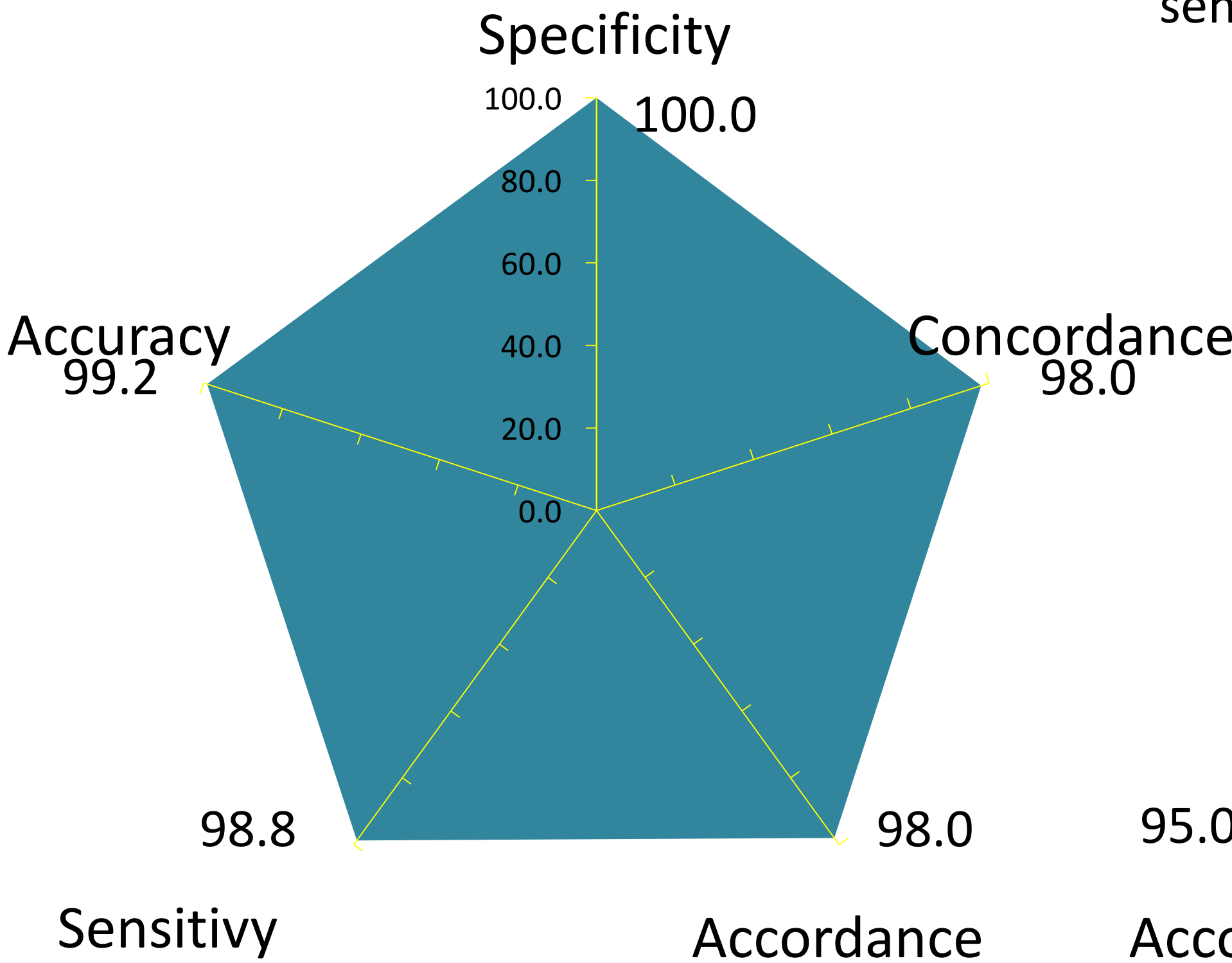


**Kerkoud primers → Better results**  
97,5 % Accuracy / 98,8% Sensitivity  
98,0% Concordance / 98,0 % Accordance

### Results of the validation test

**Results obtained by participants conform to expected results**

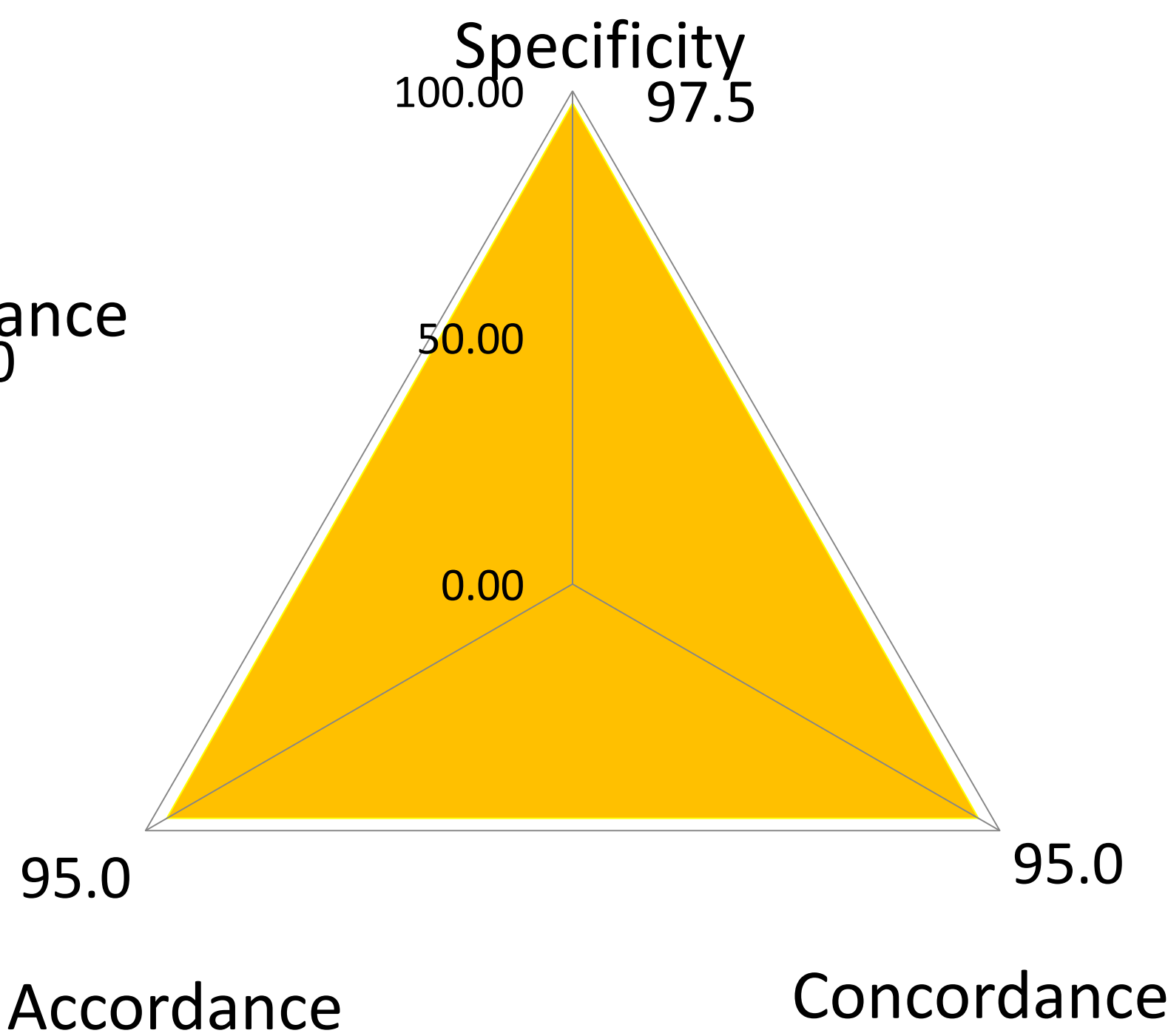
#### Detection protocol on alfalfa



**Very good results higher than 95 %**

#### Detection protocol on Faba bean

Due to heterogeneity of seed lots, diagnostic sensitivity and accuracy were not calculated



**Very good results higher than 95 %**

Combination of all these results proved that a detection method using filtration in alfalfa and faba bean seed lot was validated.

## CONCLUSION

A detection method using sieving at 20µm 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*.