

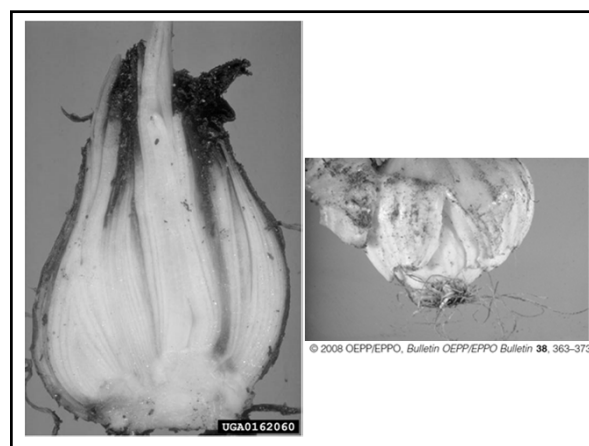
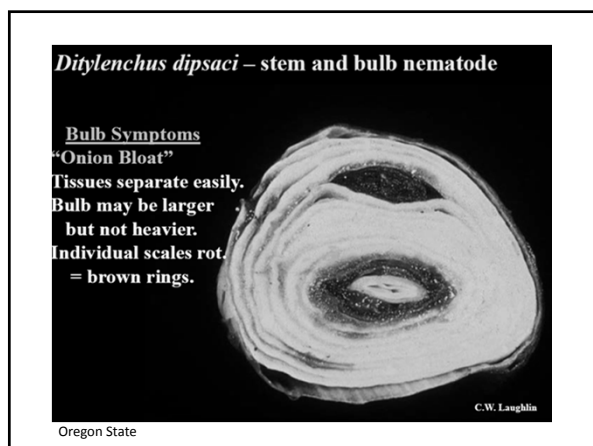
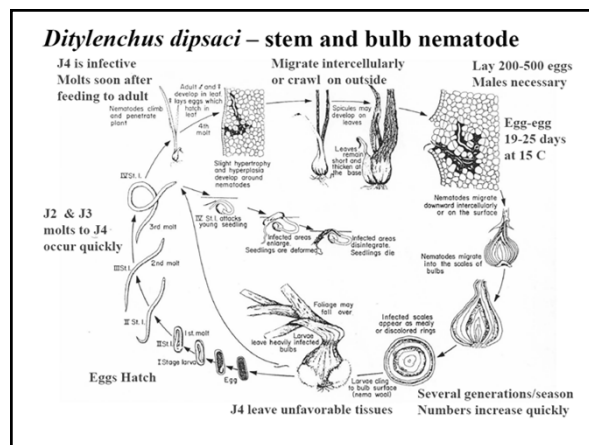
- more than 80 species
- difficult to distinguish morphologically
- many host races
- host range of many "races" overlap

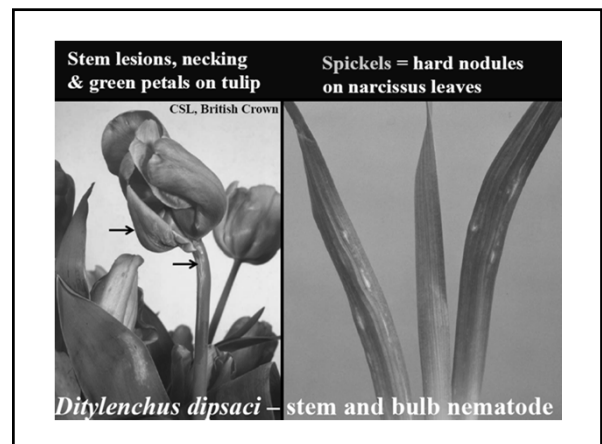
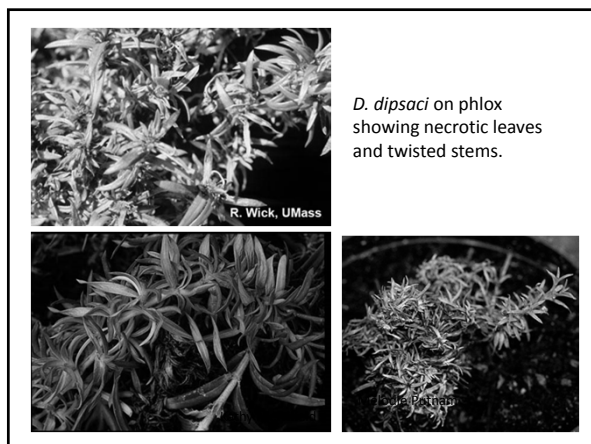
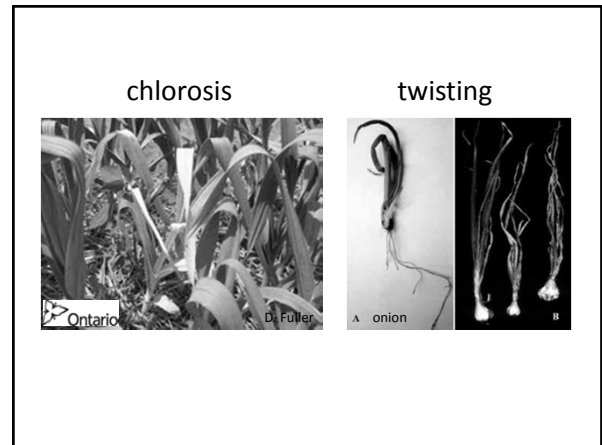
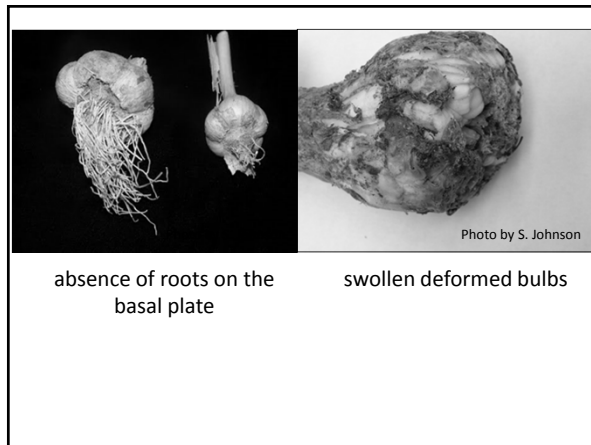
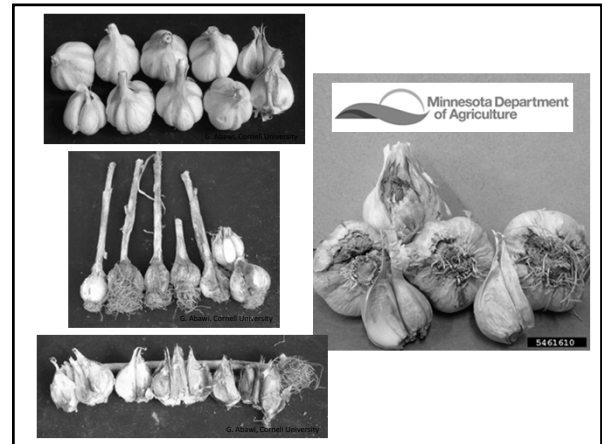
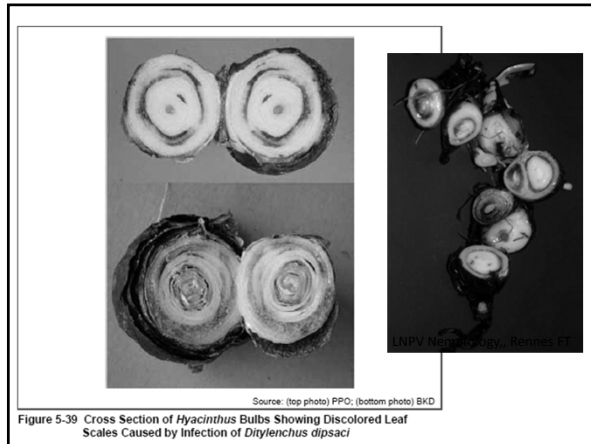
Table 1. Host range of the two tested *Ditylenchus dipsaci* populations

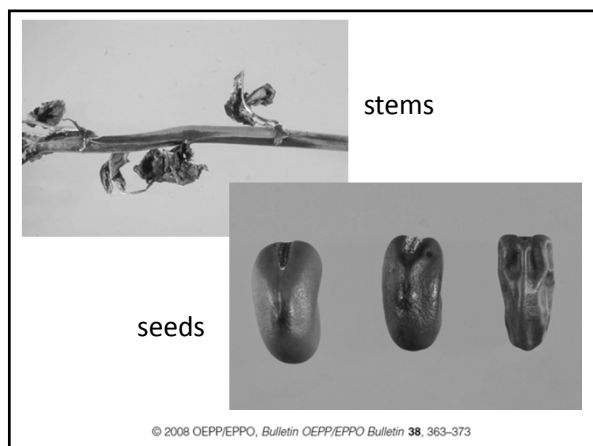
| | Onion | Garlic | Leek | Spinach | Chicory |
|-------------------------|-------|--------|------|---------|---------|
| Population from chicory | 0 | 1 | 0 | 1 | 2 |
| Population from garlic | 2 | 2 | 0 | 0 | 1 |

0 - no nematodes survived; 1 - only single adult individuals survived; 2 - life cycle of the parasite completed

DOUDA O. (2005): Host range and growth of Stem and Bulb Nematode (*Ditylenchus dipsaci*) populations isolated from garlic and chicory. *Plant Protect. Sci.*, 41: 104-108.







Cultural practices:

- plant clean, uninfested cloves
- grow nonhost crops
- do not bring infested soil into fields

Chemical control:

- pre-plant fumigation
- oxymyl

Recovery and Identification

Incubation Basics

Cover plant tissues with water.
To obtain a clean prep, place
plant tissues on Kleenex -
nematodes will crawl through
the tissue and collect in the
water reservoir.

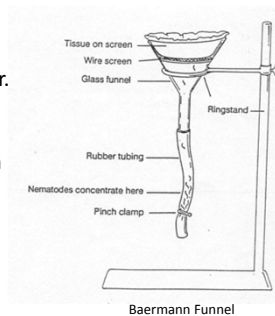


Table 1 Key to distinguish *Ditylenchus* spp. from other tylenchid genera (modified from Brzeski, 1998)

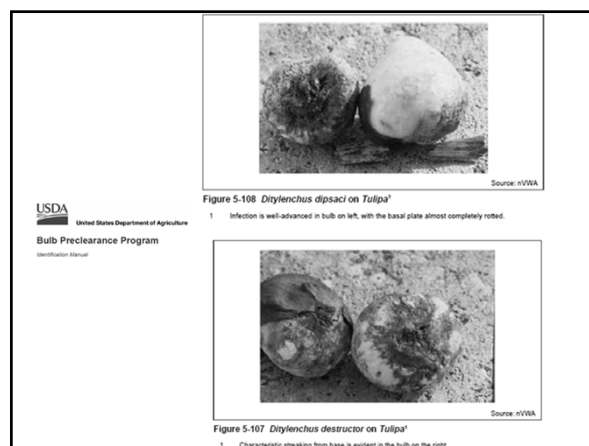
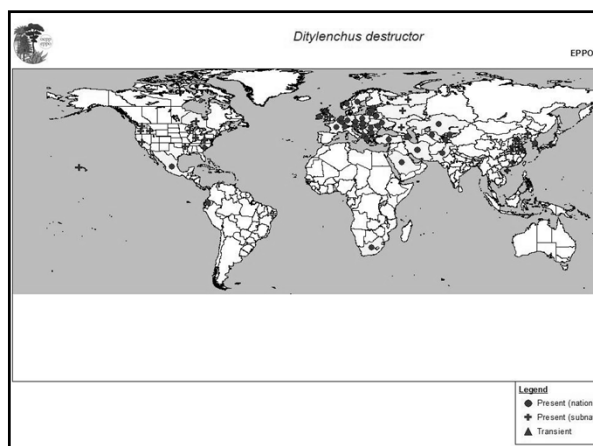
| | | |
|---|---|--------------------|
| 1 | Females mobile | 2 |
| | Females swollen, globose or lemon-shaped | Other genera |
| 2 | Female gonads prodigic and outstretched | 3 |
| | Female gonads paired or reflex when prodigic | Other genera |
| | Pharyngeal glands offset from intestine or slightly overlapping it | 4 |
| | Pharyngeal glands distinctly overlapping intestine | Other genera |
| 4 | Metacarpus offset from procorpus, metacarpal plates short or absent | 5 |
| | Procorpus gradually expands into large metacarpus, metacarpal plates long | Other genera |
| 5 | Sperm large, head usually low, spermatheca not off-set | <i>Ditylenchus</i> |
| | Sperm small, head usually high, spermatheca off-set in most genera | Other genera |

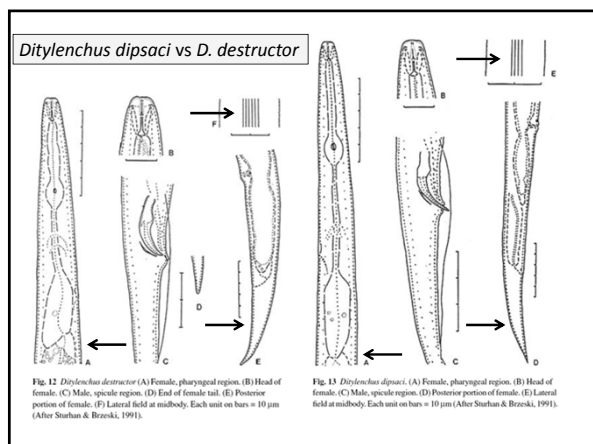
Table 2 Discriminating morphological characteristics of *D. destructor*, *D. dipsaci*, *D. comallariae* and *D. myceliophagus* (after Decker, 1969)

| | <i>D. destructor</i> | <i>D. dipsaci</i> | <i>D. comallariae</i> | <i>D. myceliophagus</i> |
|-----------------------------------|---|----------------------------|--------------------------------|-----------------------------|
| A ratio 2 | 32 (18-41) | 37 (36-40) | 42 (32-54) | 30 (23-44) |
| Body length (mm) | 1.0 (0.8-1.4) | 1.1 (0.9-1.3)* | 1.1 (0.9-1.3) | 0.9 (0.6-1.0) |
| Stylet length (µm) | 10-12 | 11-13 | 11-13 | 7-10 |
| Posterior bulb | short, dorsally overlapping | not overlapping | not overlapping | short, dorsally overlapping |
| Number of lateral lines | 6 | 4 | 6 | 6 |
| Valva position (%) | 80 (78-83) | 82 (79-82) | 77 (74-79) | 82.5 (74-90) |
| Post-valva sac length | 2/3-3/4 of valva-anus distance | 1/2 of valva-anus distance | 1/4-1/2 of valva-anus distance | 2-2 1/4 valva-anus distance |
| Valva-anus length | 1 3/4-2 1/3 tail length | 1 3/4-2 1/4 tail length | 2-2 1/4 tail length | 2-2 1/4 tail length |
| Tail shape | conoid, usually slightly bent to ventral side in posterior part | conoid | conoid | broadly conoid |
| Tail tip | finely rounded | sharply pointed | sharply pointed | finely rounded |
| Spiculum length (µm) | 9-12 | 10-12 | 8-11 | 9 |
| Length of conchotal stylet length | About 50% | About 50% | << 50% | << 50% |

*The 'giant' race of *D. dipsaci* in *Faba* bean can be up to 2.0 mm long.

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Appendix 1 ITS-RFLP analysis according to Wendt *et al.* (1993)

Appendix 2 Specific SCAR-PCR according to Esquibet *et al.* (2003)

Appendix 3 Specific PCR according to Subbotin *et al.* (2005)

Appendix 4 Specific PCR according to Marek *et al.* (2005)

Appendix 5 Specific PCR according to Kerkoud *et al.* (2007)

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Madani *et al.*, 2015
Can. J. Plant Pathol., 2015
Vol. 37, 212 - 220

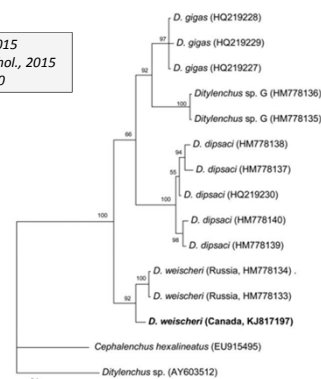
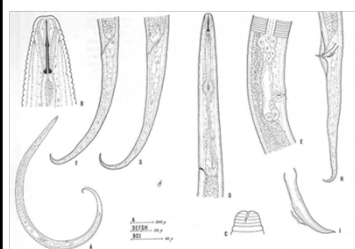
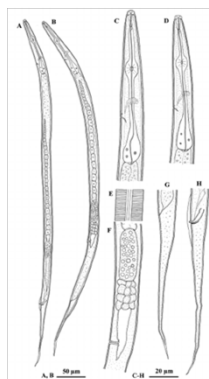


Fig. 2 Phylogenetic relationships of *D. weischeri* with other species of the genus *Ditylenchus* as inferred from the Bayesian analysis of the *hsp90* gene sequences under the GTR+I+G model. Posterior probabilities are given on appropriate clades.

Don't confuse these genera
with *Ditylenchus*.....



Tylenchus



Filenchus

