



## The effect of two fixation methods (TAF and DESS) on morphometric parameters of *Aphelenchoides ritzemabosi*

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Identification of nematode species by using conventional methods requires fixation of the isolated material and a suitable preparation for further analyses. Tentative identification using microscopic methods should also be performed prior to initiating molecular studies. In the literature, various methods are described for the preparation of nematodes from the genus *Aphelenchoides* for identification and microscopic studies. The most commonly used fixatives are formalin (Timm 1969; Szczygieł & Cid del Prado Vera 1981, Crozzoli *et al.* 2008, Khan *et al.* 2008), FAA (Wasilewska 1969; Vovlas *et al.* 2005, Khan *et al.* 2007) and TAF (Hooper 1958, Chizhov *et al.* 2006, Jagdale & Grewal 2006).

Few studies have focused on the effects of preparation on nematodes. Comparison of techniques suggests that both the method of killing and of fixing nematodes affects the quality of mounted specimens, as well as their morphometric parameters (Lamberti & Sher 1969, Grewal *et al.* 1990). According to Brown & Topham (1984), the same method of preparation should be used for comparing morphometry between closely related species. However, the variety of techniques applied by different authors often makes it impossible to compare new observations with literature data. According to research conducted by Grewal *et al.* (1990), morphology of nematodes after TAF fixation was changed the least, implying that measurements are more reliable than those obtained using other processing techniques, including the commonly applied FAA method. However, for a simultaneous morphological and molecular analysis, where only one specimen is examined, the most appropriate technique is to use DESS (Yoder *et al.* 2006).

For molecular analyses, nematodes are commonly rinsed in water without any fixative (Jiang *et al.* 2007, Khan *et al.* 2012). However, subsequent procedures must be carried out immediately after nematode isolation. DESS (a solution of dimethyl sulphoxide, disodium EDTA and NaCl) allows one to perform DNA isolation even after several months of storage at room temperature (Yoder *et al.* 2006). We thought it important to investigate whether material fixed in this solution could be also used for classical identification methods. The aim of the present study was to compare the morphometric characteristics of female specimens of *Aphelenchoides ritzemabosi* (Schwartz, 1911) Steiner & Buhner, 1932 taken from the same population, by using two methods of fixation: in TAF and in DESS solution.

Female *A. ritzemabosi* were isolated from the leaves of *Weigela florida* DC. variety ‘Bristol Ruby’, from Poland. Nematodes were isolated using a modified Baermann tray extraction method (Luc *et al.* 2005), and then killed with hot water at a temperature of 60–70 °C. For fixation, TAF (Courtney *et al.* 1955) and DESS (Yoder *et al.* 2006) were used. After a one year storage of preserved nematodes at room temperature, glycerin mounts were prepared using the Seinhorst method (Seinhorst 1962, 1966). A Nikon Eclipse 80i light microscope was employed for nematode species determination, using differential interference contrast (Nomarski DIC). Morphometric measurements were performed at 100X magnification. After identification of specimens preserved in DESS genomic DNA from three specimens was extracted using a Nucleo Spin Tissue XS (Macherey-Nagel, Germany) following the manufacturer’s instructions. For each sample, DNA was eluted in 30 µl H<sub>2</sub>O. 3 µl of extracted DNA was used as template for polymerase chain reactions (PCR). The 28S rDNA sequence fragment was amplified with primers Aph\_F (5’-AGAGAGTGCAAGAGAACGTGA-3’) and Aph\_R (5’-ACCATCTTTCGGGTCTCAAA-3’). PCR primers were designed using the online PRIMER 3 software (<http://primer3.ut.ee>) after aligning sequences of closely related in GenBank. Primers were purchased from Sigma-Aldrich (Sigma-Aldrich, Milwaukee, WI, USA). Amplification of 28S rDNA and sequencing was performed as described previously (Dobosz *et al.* 2013) with modified polymerase (12.5µl Taq PCR Master Mix Gdansk, Poland) and

primer concentrations (1µl of 10 µM each primer). Cycling was performed by a Veriti 96-Well Thermal Cycler as follows: an initial denaturation step of 94°C for 3 min, followed by 40 cycles of 94°C for 30 s, 58°C for 30 s and 72°C for 1 min and a final extension step of 72°C for 5 min. 28S rDNA sequences were aligned in MEGA ver. 6.06 program software using default parameters (Tamura *et al.* 2011). The sequences reported in this study have been deposited in Gen Bank with accession numbers KR261601, KR261602, KR261603. The length of sequences for *A. ritzemabosi* ranged from 453 bp to 528 bp. Sequences were compared to each other and deletion of one nucleotide in one of the samples was observed.

**TABLE 1.** Morphometric measurements of females of *A. ritzemabosi*. Measurements are in µm and in the form: range, mean±SD.

	♀♀-DESS n=30, mean±SD	♀♀-TAF n=30, mean±SD
L (µm)	683.7–830.3 767.7±33.2	693.7–824.0 774.1±30.0
head-vulva length	471.0–562.0 526.9±23.5	483.7–566.0 535.3±21.4
vulva-anus length (v-a)	174.2–224.9 199.5±11.7	166.5–217.8 198.9±12.7
maximum body diameter	15.6–19.4* 17.9±0.9	14.2–18.1 16.3±1.0
tail length	36.5–47.0 41.3±2.9	35.2–45.9 40.0±2.9
anal body diameter	9.2–12.* 10.8±0.8	8.2–11.1 9.9±0.8
postvulval uterine sac (PUS)	73.6–136.6 109.2±14.3	96.8–134.1 111.5±11.2
PUS - % v-a	42.3–66.5 54.7±6.6	46.4–62.7 56.1±4.3
PUS - xV	4.0–7.5* 6.1±0.8	6.0–8.2 6.9±0.7
V%	66.9–70.8 68.6±1.0	67.1–70.8 69.2±0.9
a	36.9–47.2* 43.0±2.3	42.2–54.8 47.6±3.3
c	17.1–20.4 18.7±1.1	16.0–22.5 19.5±1.6
c'	3.2–4.5 3.9±0.3	3.2–5.5 4.1±0.5
b	7.4–8.6 8.1±0.4	7.3–9.2 8.1±0.6
b'	4.4–5.2* 4.7±0.2	4.6–5.5 5.0±0.3
pharynx length	89.7–99.8 95.0±3.9	85.7–103.5 95.6±6.4
pharynx length (with glands)	148.1–179.7* 164.4±8.0	140.1–168.3 154.1±7.9
stylet length	10.8–13.4 12.5±0.8	11.6–13.2 12.5±0.5
excretory pore from head end	82.7–96.5* 88.8±4.7	76.3–103.9 97.2±6.1
lateral field width	2.6–4.4 3.5±0.5	2.7–3.8 3.2±0.3

\* values marked with a star in columns are significantly different according to Tukey test ( $p < 0.05$ )

Analysis of variance (ANOVA) was applied to the morphometric results. For comparison averages, a Tukey test was used, assuming a  $p = 0.05$  level of significance. The analysis was performed in STATISTICA v. 10 (Stat Soft Inc., 2011). Statistical analysis revealed significant differences between methods of fixation in 7 out of 20 tested parameters. Females preserved in DESS were characterized by a greater body width, both in the widest part of the body and at the level of the anus, as well as a longer pharynx with glands compared to females fixed in TAF. An inverse dependency was observed in the position of excretory pore from the head end (Table 1).

FAA is one of the more commonly used fixatives. Grewal *et al.* (1990), comparing different fixation methods, showed that FAA causes many changes in morphometric parameters. According to these authors, TAF causes shrinking of the specimens, although to a lesser extent than other fixatives. In the present study, there were 7 disagreements in parameters between TAF and DESS, which indicate that DESS had less effect on in the morphometric changes than TAF. Furthermore, as shown by Yoder *et al.* (2006), morphological characteristics of nematodes preserved in DESS are comparable to those fixed in formalin, which is employed frequently in the methodology of nematode fixation. The utility of DESS in both morphological examination and molecular analyses was also shown for animal-parasitic nematodes (Naem *et al.* 2010) and other groups of invertebrates (Strona *et al.* 2009). It should be noted, however, that a disadvantage of this fixation method is reduced sharpness of specimen morphological characteristics as relates to nematodes with gently outlined diagnostic features. This does not change the fact that fixation in DESS after killing nematodes with hot water allows one to perform both regular morphometric measurements of nematodes and molecular analyses.

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