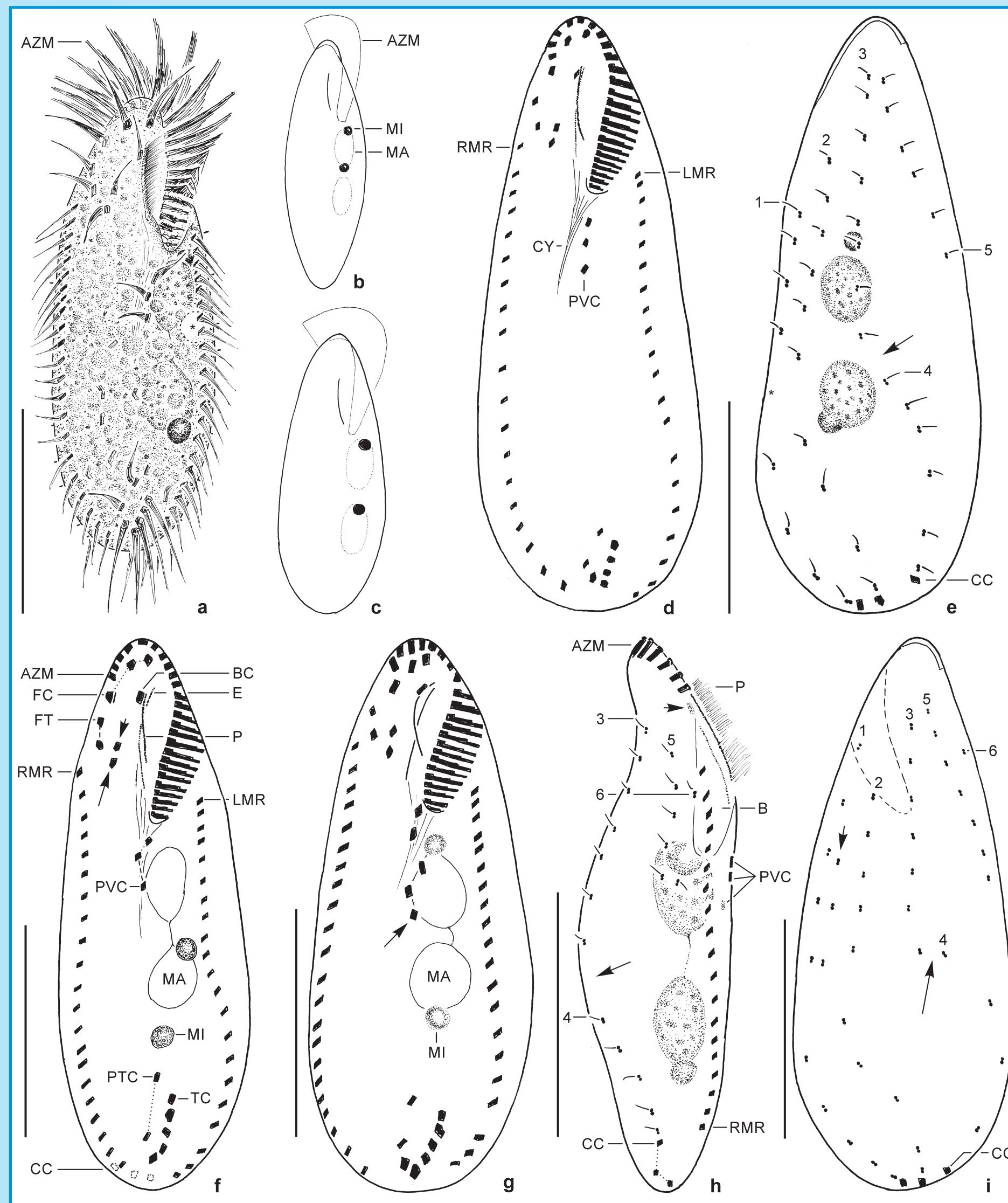


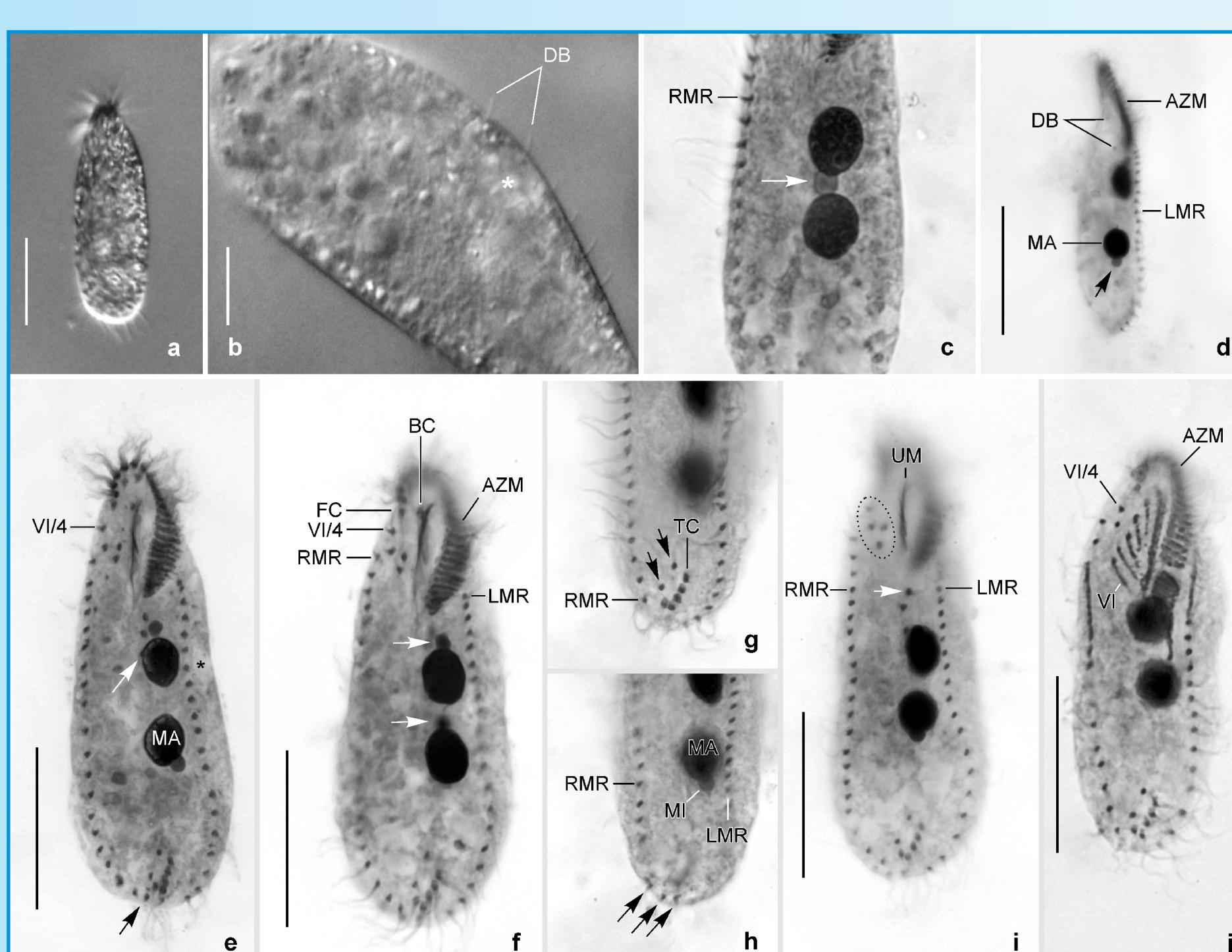
# Morphology, phylogenetic relationships, and pH response of two *Oxytricha*-like species (Ciliophora, Hypotricha)

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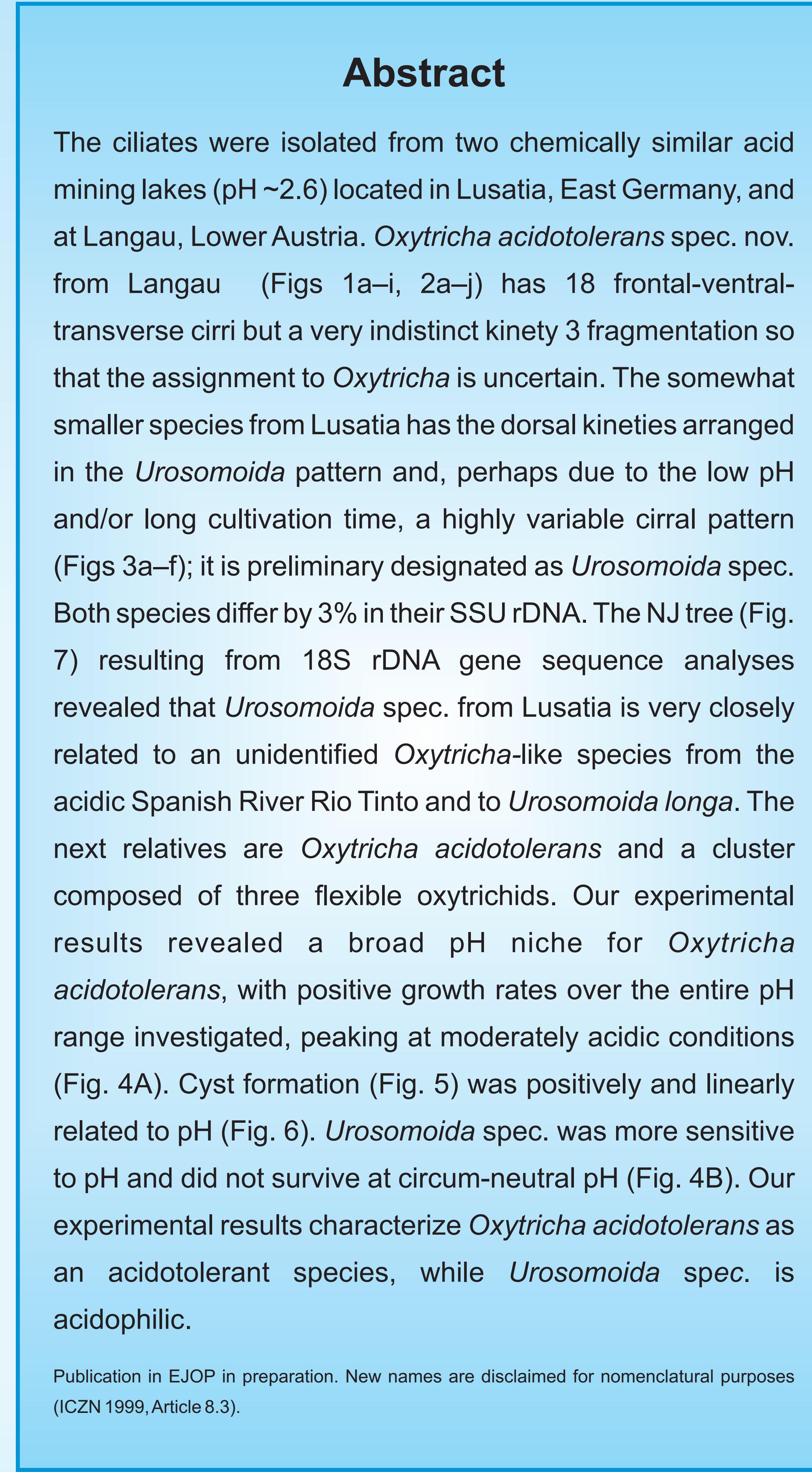
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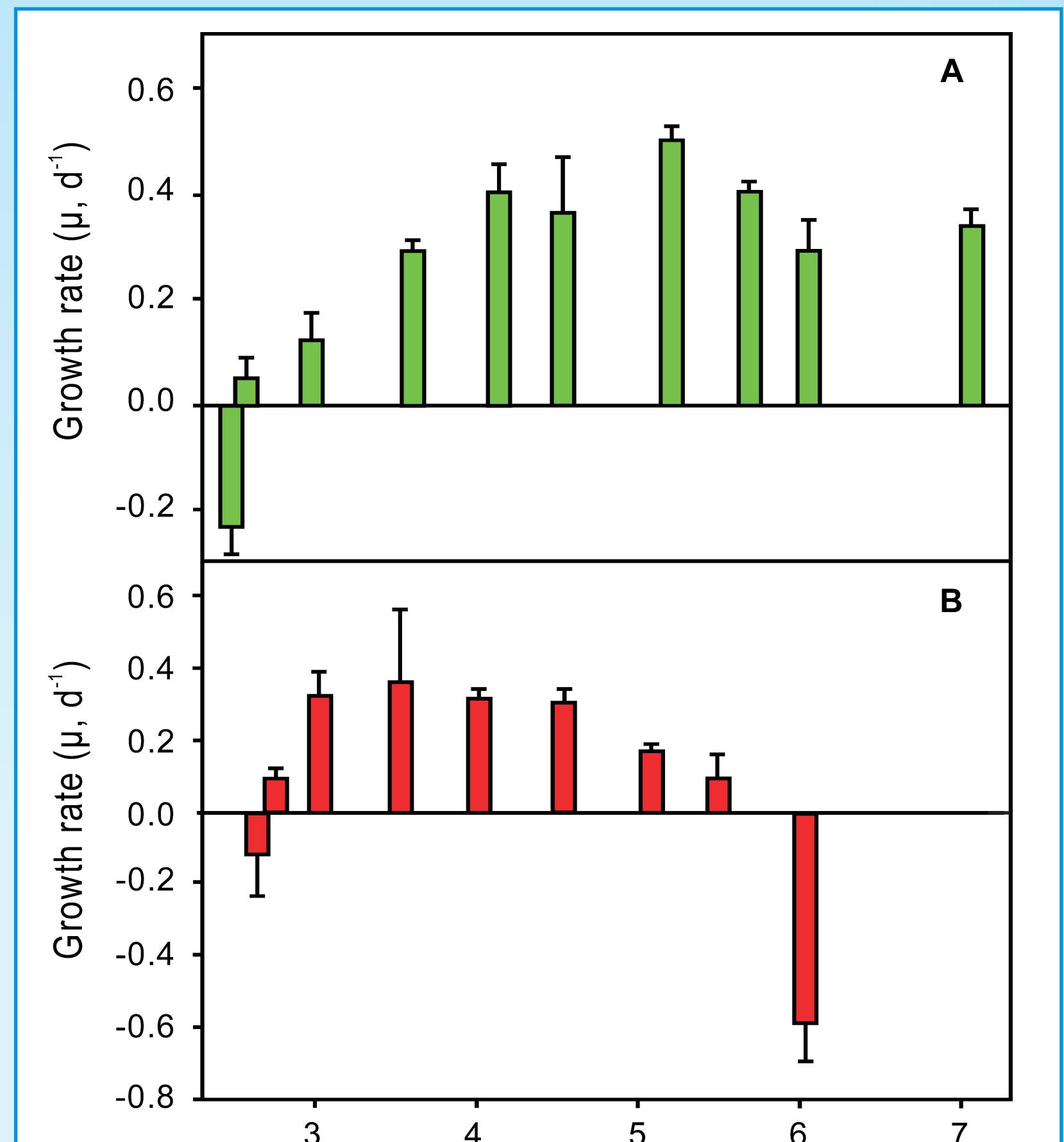
**Figs 1a–i.** *Oxytricha acidotolerans* spec. nov. in vivo (a–c) and after protargol impregnation (d–i). a: Ventral view of representative specimen. Asterisk marks contractile vacuole. b, c: Spindle-shaped and slightly oviform shape variants with two rare arrangements of micronuclei. d, e: Infraciliature of ventral and dorsal side and nuclear apparatus of holotype specimen. Asterisk in (e) marks break in dorsal kinety 1, arrow denotes indistinct "fragmentation" of kinety 3. f: Specimen with almost linearly arranged transverse cirri. Short arrow marks frontoventral cirrus III/3, long arrow denotes cirrus IV/3. g: Specimen with five postoral ventral cirri (rearmost marked with arrow) and six transverse cirri. h: Right lateral view. Short arrow marks slightly subadjacent buccal cirrus, long arrow marks "fragmentation" of kinety 3. i: Infraciliature of dorsal side of a specimen with an additional (remnant of previous generation?) short row of dorsal bristles (short arrow); kinety 6 is composed of one (rarely two) basal body pair only, if present at all. Long arrow marks "fragmentation" of kinety 3. AZM, adoral zone of membranelles; B, buccal cavity; BC, buccal cirrus; CC, caudal cirri; CY, pharyngeal fibres; E, endoral; FC, frontal cirri; FT, frontoterminal cirri (= cirri VI/3, VI/4); LMR, left marginal row; MA, macronuclear nodule; MI, micronucleus; P, paroral; PVC, postoral ventral cirri; PTC, pretransverse ventral cirri; RMR, right marginal row; TC, transverse cirri; 1–6, dorsal kinetics. Scale bars: 30 µm.



**Figs 2a–j.** *Oxytricha acidotolerans* spec. nov. in vivo (a, b; differential interference contrast; video frames) and after protargol impregnation (c–j). a: Dorsal view of a freely motile specimen. b: Optical section showing, inter alia, dorsal bristles (about 3 µm) and contractile vacuole (asterisk). c: Rare specimen with a single micronucleus (arrow) in between the two macronuclear nodules. d: Right lateral view showing that the nuclear apparatus is rather close to the ventral side (arrow marks micronucleus). e: Holotype specimen (composite of eight micrographs). White arrow marks rearmost postoral ventral cirrus (V/3), black arrow denotes a caudal cirrus. Asterisk indicates area of contractile vacuole. f: Specimen with micronuclei (arrows) at anterior end of macronuclear nodules. g, h: Posterior end showing, inter alia, pretransverse ventral cirri (arrows in g) and caudal cirri (arrows in h). i: Specimen with anterior postoral ventral cirrus (arrow) slightly shifted leftwards. Frontoventral cirri circled. j: Ventral view of reorganizer. AZM, adoral zone of membranelles; BC, buccal cirrus; DB, dorsal bristles; FC, right frontal cirrus; IV/4, anterior frontoterminal cirrus; LMR, left marginal row; MA, macronuclear nodule; MI, micronucleus; PVC, postoral ventral cirri; PTC+TC, pretransverse ventral cirri and transverse cirri; RMR, right marginal row; TC, transverse cirri; UM, undulating membranes; VI, frontal-ventral-transverse cirri anlage VI. Scale bars: 30 µm (a, d–f, i, j), 10 µm (b).



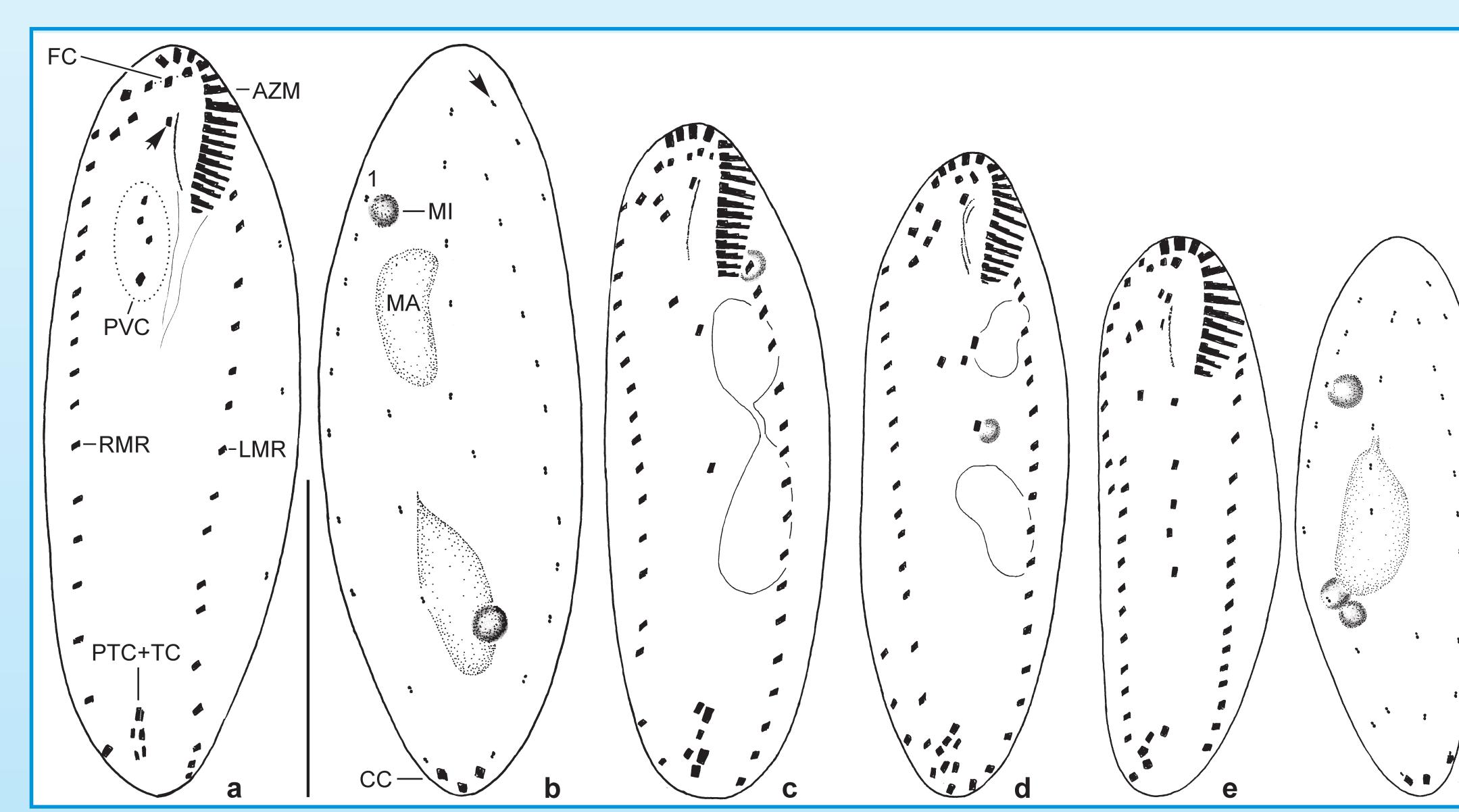
Publication in EJOP in preparation. New names are disclaimed for nomenclatural purposes (ICZN 1999, Article 8.3).



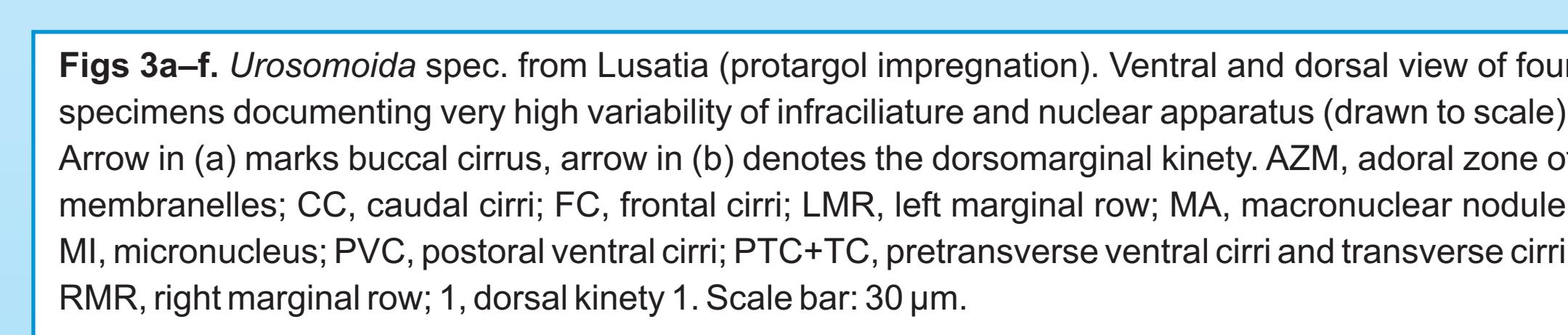
**Figs 4A, B.** Population growth rate of *Oxytricha acidotolerans* (A) and *Urosomoida* spec. (B) in relation to pH. Bars represent means of triplicates, the error bars denote 1 SD.



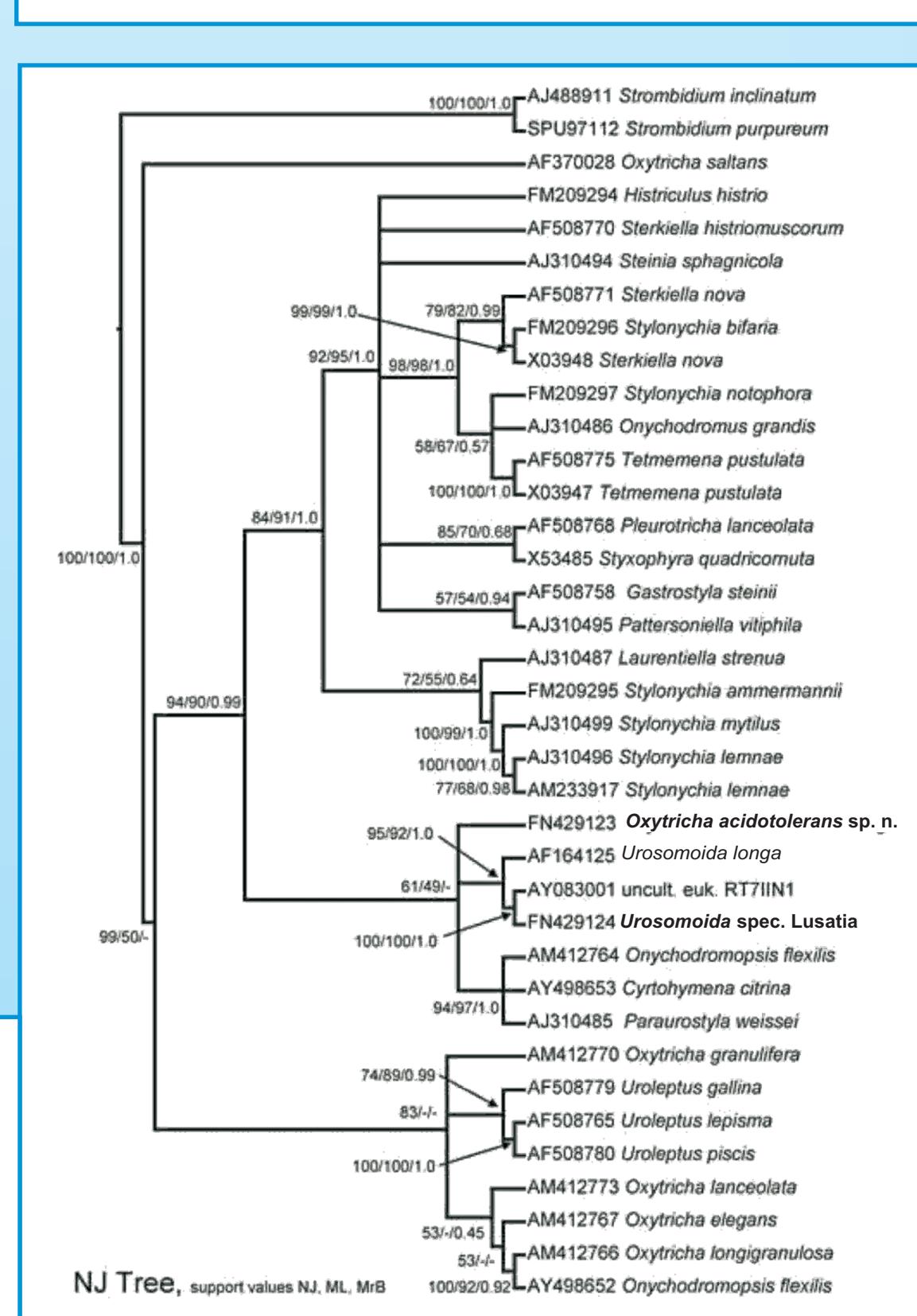
**Fig. 5.** A motile cell and cysts of *Oxytricha acidotolerans* after Lugol's fixation. Scale bar = 50 µm.



**Fig. 6.** The percentage of cysts in the population of *Oxytricha acidotolerans* in relation to pH. The line represents the linear regression of cysts (%) vs. pH.



**Fig. 7.** Phylogenetic NJ based consensus tree of the SSU rDNA of hypotrich ciliates. The species used in this study are shown in bold face. Support values at the nodes represent bootstrap percentages of the Neighbor-joining (first number) and Maximum-likelihood (second number) analyses and posterior probabilities of the Bayesian analysis (third number). Arrows denote support values to corresponding nodes.



## Acknowledgements

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