

***Balanion masanensis* n. sp. (Ciliophora: Prostomatea) from the Coastal Waters of Korea: Morphology and Small Subunit Ribosomal RNA Gene Sequence**

JAE SEONG KIM,^a HAE JIN JEONG,^a DENIS H. LYNN,^b JAE YEON PARK,^a YOUNG WOON LIM^c and WOONGGHI SHIN^d

^aSchool of Earth and Environmental Science, College of Natural Sciences, Seoul National University, Seoul 151-747, Korea, and

^bDepartment of Integrative Biology, University of Guelph, Guelph, Ontario, Canada N1G 2W1, and

^cNational Institute of Biological Resources, Incheon 404-708, Korea, and

^dDepartment of Biology, Chungnam National University, 220 Gung-Dong, Yuseong-Gu, Daejeon 305-763, Korea

ABSTRACT. The planktonic ciliate *Balanion masanensis* n. sp. is described from living cells, from cells prepared by quantitative protargol staining (QPS), scanning electron microscopy (SEM), and transmitted electron microscopy (TEM) preparations, and the sequence of its nuclear small subunit rDNA (SSU rDNA) is reported. This species is almost ovoid with a flattened anterior oral region when the cells are alive and stained. The flattened anterior region of a living cell often forms a dome with the perimeter receded in a groove, and this region is easily inflated or depressed. In SEM photos, a brosse of six to nine monokinetids (or possibly three to five dikinetids) was observed inside the circumoral dikinetids. In TEM photos, circumoral microtubular ribbons were observed below the oral cilia, which along with the oral flaps were 8–16 µm in length. The cytostome is a slight funnel-like central depression on the flattened anterior end. The morphological characteristics of this ciliate are identical to those of the genus *Balanion* (Order Prorodontida). The ranges (and mean ± standard deviation) of cell length, cell width, and oral diameter of living cells (n = 23–26) were 27–43 µm (35.2 ± 4.6), 25–32 µm (28.6 ± 2.3), and 25–30 µm (27.6 ± 1.3), respectively, while those of the QPS-stained specimens (n = 70) were 23–37 µm (30.6 ± 3.5), 26–35 µm (30.7 ± 2.2), and 26–33 µm (29.5 ± 1.5), respectively. Forty-six to 55 somatic kineties (SKs) were equally spaced around the cell body and extended from the oral to near the posterior regions with 24–50 monokinetids per kinety. Each kinetid bore a cilium 2.8–7.2 µm long. A caudal cilium (ca 14 µm long) arose on the posterior end. The single ellipsoid macronucleus is 6.8–13.4 × 6.8–10.5 µm, accompanied by a single micronucleus (2.0–2.8 × 1.5–2.5 µm) visible only in QPS specimens. Because, the cell size, the number of SKs, and the number of kinetosomes per SK of this ciliate were much greater than those of *Balanion comatum* and *Balanion planctonicum*, the only two *Balanion* species so far reported, we have established *B. masanensis* n. sp. When properly aligned, the sequence of the SSU rDNA of *B. masanensis* n. sp. (GenBank Accession No. AM412525) was approximately 9% different from that of *Coleps hirtus* (Colepidae, Prorodontida) and 12% different from that of *Prorodon teres* (Prorodontidae, Prorodontida).

Key Words. Brosse, ciliate, DNA, plankton, prostome ciliate, protozoa.

PROSTOME ciliates are often dominant heterotrophic protists in freshwater (Müller 1989; Sommaruga and Psenner 1993; Sonntag et al. 2002; Weisse and Müller 1998) and marine environments (Smetacek 1989; Stoecker, Davis, and Provan 1983). However, the taxonomy and ecology of these ciliates is still poorly known, even though they are sometimes abundant. There are two orders established in the Class Prostomatea—Order Prostomatida and Order Prorodontida—distinguished by whether a ciliate has “brosse” kinetids around the oral region or not (Lynn and Small 2002). This oral structure, which is often difficult to find, can be identified by careful examination with specialized methods, such as silver impregnation technique or scanning electron microscopy (SEM) after deciliation (Bardele 1999).

DNA sequences of only a few prostome species have been reported, and there are no sequences for the Order Prostomatida, and only three sequences for the Order Prorodontida for the genera *Coleps*, *Prorodon*, and *Cryptocaryon* (Hirt et al. unpubl. observ.; Leipe et al. 1994; Stechmann, Schlegel, and Lynn 1998; Wright and Colorni 2002).

The species belonging to the genus *Balanion* (Order Prorodontida) are sometimes abundant in both freshwater and marine environments (Jakobsen and Hansen 1997; Müller and Geller 1993; Stoecker and Evans 1985; Weisse et al. 2001). They are known to be effective grazers on nanoflagellates and red-tide dinoflagellates (Jakobsen and Hansen 1997; Kenter, Zimmermann, and Müller 1996; Müller and Schlegel 1999; Stoecker et al. 1983; Stoecker and Evans 1985). So far, only two species have been reported in this genus—*Balanion comatum* and *Balanion planctonicum* (Bardele 1999; Foissner, Berger, and Kohmann 1994; Foissner, Oleksiv, and Müller 1990; Jakobsen and Montagnes 1999; Small and Lynn 1985). Wulff (1919) provides an early morphological study

of *B. comatum*. In their redescription of *B. comatum*, Jakobsen and Montagnes (1999) provided the key characters for the genus *Balanion* as (1) the presence of the circumoral microtubular ribbons (COMT), (2) the prominent oral dikinetid–flap complex, and (3) the brosse lying internal to the circumoral dikinetids.

During red tides dominated by the mixotrophic dinoflagellate *Heterocapsa triquetra* in Masan Bay, Korea, we found a new prostome ciliate that appeared to be a *Balanion* sp. In this study, we describe this is a new species of *Balanion*, *Balanion masanensis* n. sp. and establish its phylogenetic position within the Class Prostomatea based on its small subunit rDNA (SSU rDNA) sequence.

MATERIALS AND METHODS

Preparation of samples. Plankton samples collected with water samples were taken from a pier in Masan Bay, Korea, from December 2005 to February 2006 when the water temperature and salinity were 3.1 °C–14.1 °C and 23.3–32.7 psu, respectively. The samples were transported to the laboratory within 30 min, gently screened through a 202-µm nitex mesh, and divided into 270-ml polycarbonate (PC) bottles and 500-ml polyethylene (PE) bottles. Some of a dense culture of *H. triquetra* (concentration = ca 15,000 cells/ml) was added to the 270-ml PC bottles as food. The bottles were placed on plankton wheels rotating at 0.9 rpm and incubated at 5°–10 °C under an illumination of 20 µE/m²/s of cool white fluorescent light on a 12:12 h light–dark cycle. Two days later, aliquots of the enriched water were transferred to six-well tissue culture plates and a clonal culture of *B. masanensis* n. sp. was established by two serial single-cell isolations. Once dense cultures of this ciliate were obtained, they were transferred every 2 days to 270-ml PC bottles of fresh *H. triquetra* as prey. For the observation of living cells and the analysis of DNA sequence, the cells maintained by this method were used.

For the quantitative protargol stain (QPS) method (Montagnes and Lynn 1987), samples in the PE bottles were fixed with a

Corresponding Author: H. J. Jeong, School of Earth and Environmental Science, College of Natural Sciences, Seoul National University, Seoul 151-747, Korea—Telephone number: +82 2 880 6746; FAX number: +82 2 874 9695; e-mail: hjeong@snu.ac.kr

Table 1. The list of the nucleotide sequences used in this study.

Species	Accession number	References
<i>Amphisiella magnigranulosa</i>	AM412774	Schmidt et al. (2007)
<i>Amylovorax dehorityi</i>	AF298817	Cameron, Adlard and O'Donoghue (2001)
<i>Anophryoides haemophila</i>	U51554	Ragan et al. (1996)
<i>Anoplophrya marylandensis</i>	AY547546	Affa'a et al. (2004)
<i>Astylozoon enriquesi</i>	AY049000	
<i>Blepharisma americanum</i>	M97909	Greenwood et al. (1991)
<i>Bresslaia vorax</i>	AF060453	Lynn et al. (1999)
<i>Bryometopus sphagi</i>	AF060455	Lynn et al. (1999)
<i>Bursaria truncatell</i>	U82204	Stechmann, Schlegel, and Lynn (1998)
<i>Caenomorpha uniserialis</i>	U97108	
<i>Cardiostomatella vermiforme</i>	AY861632	Li et al. (2006)
<i>Chattonidium setense</i>	AM295495	Modeo et al. (2006)
<i>Chilodonella uncinata</i>	AF300281	
<i>Chlamydonodon excocellatus</i>	AY331790	Snoeyenbos-West et al. (2004)
<i>Climacostomum virens</i>	X65152	Hammerschmidt et al. (1996)
<i>Coleps hirtus</i>	U97109	
<i>Colpes</i> sp.	DQ487194	
<i>Colpes</i> sp.	X76646	Stechmann et al. (1998)
<i>Colpoda inflata</i>	M97908	Greenwood et al. (1991)
<i>Colpoda steinii</i>	DQ388599	
<i>Cryptocaryon irritans</i>	AF351579	Wright and Coloni (2002)
<i>Dasytricha ruminantium</i>	U57769	Wright and Lynn (1997)
<i>Dextiostrichides pangii</i>	AY212805	Song, Ma, and AL-Rasheid (2003)
<i>Didinium nasutum</i>	U57771	Wright and Lynn (1997)
<i>Discophrya collini</i>	L26446	Leipe et al. (1994)
<i>Dysteria derouxi</i>	AY378112	
<i>Ephelota gemmipara</i>	DQ834370	
<i>Eufolliculina uhligi</i>	U47620	Hammerschmidt et al. (1996)
<i>Furgasonia blochmanni</i>	X65150	Bernhard et al. (1995)
<i>Gonostomum strenuum</i>	AJ310493	Bernhard et al. (2001)
<i>Gruberia</i> sp.	L31517	Hirt et al. (1995)
<i>Halteria grandinella</i>	AF194410	Shin et al. (2000)
<i>Heliophrya erhardi</i>	AY007446	
<i>Hemiophrys macrostoma</i>	AY1102173	
<i>Homalozoon vermiculare</i>	L26447	Leipe et al. (1994)
<i>Isochona</i> sp.	AY242116	Snoeyenbos-West et al. (2004)
<i>Loxodes striatus</i>	U24248	Hammerschmidt et al. (1996)
<i>Macropodinium ennuensis</i>	AF298820	
<i>Maristentor dinoferus</i>	AY630405	Miao et al. (2005)
<i>Metopus contortus</i>	Z29516	Hirt et al. (1995)
<i>Miamiensis avidus</i>	AY550080	Jung et al. (2005)
<i>Nyctotheroides deslierresae</i>	AF145353	
<i>Obertrumia georgiana</i>	X65149	Bernhard et al. (1995)
<i>Ophryoglena catenula</i>	U17355	Wright and Lynn (1995)
<i>Paramecium tetraurelia</i>	X03772	Sogin and Elwood (1986)
<i>Pelagostrobilidium neptuni</i>	AY541683	Agatha et al. (2005)
<i>Peritroums kahli</i>	AJ537427	
<i>Phacodinium metchnikoffi</i>	AJ277877	Shin et al. (2000)
<i>Philasterides dicentrarchi</i>	AY642280	
<i>Plagiostoma lumbrici</i>	AY547545	Affa'a et al. (2004)
<i>Platyophrya vorax</i>	AF060454	Lynn et al. (1999)
<i>Polycosta roundi</i>	AF298819	
<i>Prorocentrum micans</i>	AY585526	Zhang, Bhattacharya, and Lin (2005)
<i>Prorodon teres</i>	X71140	Leipe et al. (1994)
<i>Prorodon viridis</i>	U97111	
<i>Pseudoplatyophrya nana</i>	AF060452	Lynn et al. (1999)

Table 1. (Continued).

Species	Accession number	References
<i>Rimostrombidium lacustris</i>	DQ986131	Agatha and Strueder-Kypke (2007)
<i>Schizocaryum dogieli</i>	AF527756	Lynn and Strueder-Kypke (2002)
<i>Sirloxophyllum utriculariae</i> (submitted as <i>Loxophyllum</i>)	L26448	Leipe et al. (1994)
<i>Sorogena stoianovitchae</i>	AF300287	Lasek-Nesselquist and Katz (2001)
<i>Strombidinopsis jeokjo</i>	AJ628250	Jeong et al. (2004)
<i>Strombidium inclinatum</i>	AJ488911	Modeo et al. (2003)
<i>Symbiodinium pilosum</i>	X62650	Medlin et al. (1988)
<i>Tetrahymena canadensis</i>	M98022	Sogin et al. (1986)
<i>Tintinnopsis dadayi</i>	AY143562	Strueder-Kypke and Lynn (2003)
<i>Tracheloraphis</i> sp.	L31520	Hirt et al. (1995)
<i>Trithymostoma steini</i>	X71134	Leipe et al. (1994)
<i>Urocentrum turbo</i>	AF255357	Strueder-Kypke et al. (2000)

All are available from the GenBank/EMBL databases and have the accession numbers.

modified concentrated Bouin's solution (final concentration 10%, v/v) in which most of the cells were well preserved.

For SEM, a 20 ml aliquot of a dense culture of *B. masanensis* n. sp. was fixed with osmic acid (final concentration 2%, w/v) in seawater for 1.5 h and then the fixed cells were collected on a PC membrane filter (pore size = 5 µm) without additional pressure. Fixed cells were rinsed three times with distilled waters to remove the salt, dehydrated through an ethanol series, and finally dried using critical point dryer (Bal-Tec, CPD 300, Balzers, Germany). The dried filters were mounted on a stub and coated with gold-palladium. Cells were viewed with a JSM-840A SEM (JEOL Ltd., Tokyo, Japan) and photographed using a digital camera connected by a computer. To observe the brosse, an important key to the genus *Balanion*, deciliated cells were observed when fixed for SEM with 2% (w/v) osmium tetroxide.

For transmission electron microscopy (TEM), 500 ml monocultured cells were fixed for 1.5 h in 4% (v/v) glutaraldehyde in culture medium. Cells were centrifuged, and the pellet was embedded in agar. After several rinses with medium, the cells were postfixed in 1% (w/v) osmium tetroxide in deionized water. Dehydration was accomplished using a graded ethanol series (50%, 60%, 70%, 80%, 90%, and 100% ethanol, followed by two 100% ethanol steps). The material was embedded in Spurr's low-viscosity resin (Spurr 1969). Sections were obtained with a RMC MT-XL ultramicrotome (Boeckeler Instruments Inc., Tucson, AZ) and post-stained with 3% (w/v) aqueous uranyl acetate followed by lead citrate. Stained sections were viewed with a JEOL-1010 electron microscope (JEOL Ltd.).

Measurement of specimens. We measured cell length (the maximum longitudinal linear distance excluding cilia), cell width (the maximum diameter), and oral diameter of living cells, using an image analysis system mounted on a compound microscope (Image-Pro Plus 4.5, Media Cybernetics, Silver Spring, MD). We measured morphometric characteristics of cells stained using QPS as follows: more than 1 wk after being fixed with Bouin's solution, the samples were stained using QPS, and examined with a compound microscope at a magnification of 400–1,000X. Measurement of the cells followed the recommendations of Lynn and Small (2002), Foissner (1984), Foissner et al. (1994), and Jakobsen and Montagnes (1999). We measured number and length of

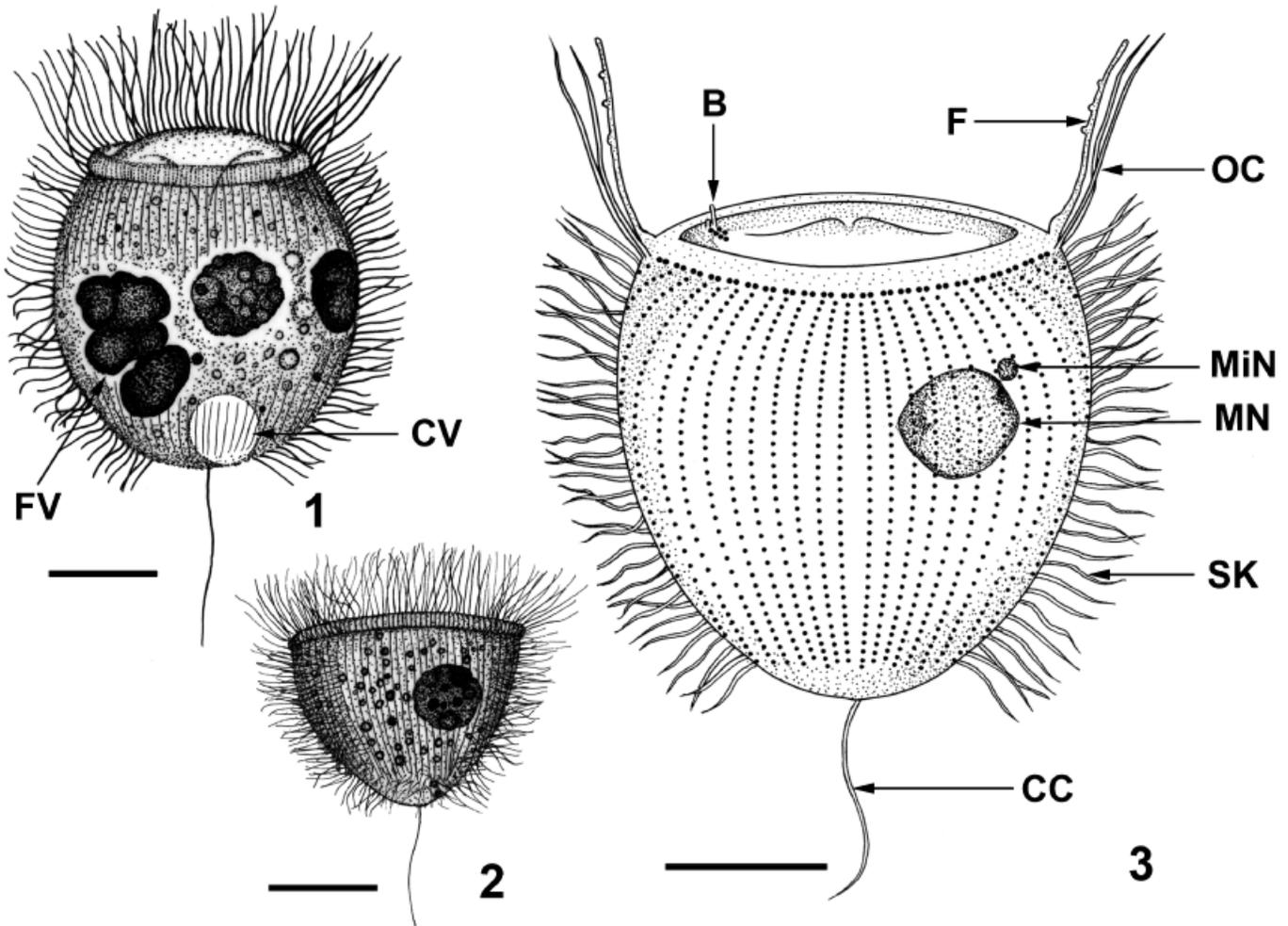


Fig. 1–3. Schematic diagrams of *Balanion masanensis* n. sp. 1, 2. Lateral view based on living cells in a natural population. 1. General cell shape having eaten prey. 2. Starved cell. 3. Lateral view based on live, protargol-stained, and scanning electron microscopic preparations indicating the position and shape of the oral cilia (OC), oral flaps (F), brosse (B), somatic kineties (SKs), caudal cilium (CC), macronucleus (MN), and micronucleus (MiN). CV, contractile vacuole; FV, food vacuoles. Scale bars = 10 μ m.

oral cilia and length of somatic cilia using the samples fixed by formalin solution (final concentration 5%, v/v) and SEM specimens.

DNA extraction, PCR amplification, sequencing and data analysis. Genomic DNA was extracted from ca 100 cells from a clonal culture with a DNeasy Tissue kit (Qiagen, Stanford, CA) following the manufacturer's instructions. The extracted DNA was divided into two tubes and used to conduct independent PCR reactions for the SSU rDNA, which were performed under the following conditions: one cycle of 3 min at 94 °C; 15 cycles of 30 s at 94 °C, 40 s at 56 °C, and 3 min at 72 °C; 25 cycles of 30 s at 94 °C, 40 s at 52 °C, and 3 min at 72 °C; and a final cycle of 5 min at 72 °C. For the full-length fragment, the universal eukaryotic forward primer EukA (5'-CTG GTT GAT CCT GCC AG-3') and the reverse primer EukB (5'-TGA TCC TTC YGC AGG TTC-3') were used (Petroni et al. 2002). For the internal 600 bp fragment (approximately 600–1,200 bp downstream from the 5'-end of the gene), forward primer Gas+600 (5'-CGG TAA TKC CAG CTC CAA TAG CG-3') and reverse primer Gas-1220 (5'-CCT GGT GGT GCC CTT CCG TC-3') were used. The PCR fragments were inserted into a pGEM-T vector (Promega, Madison, WI) and at

least three colonies for each PCR fragment were sequenced in both directions with EukA, EukB, Gas+600, Gas-1220 primers, and two additional primers, Gas+1390 (5'-CTG GTT AAT TCC GAT AAC G-3') and Gas-1540 (5'-GGG CAT CAC AGA CCT GT-3') using an ABI PRISM[®] 3700 DNA Analyzer (Applied Biosystems, Foster City, CA).

Sequence availability and phylogenetic analysis. The nucleotide sequences are available from the GenBank/EMBL databases (Table 1). The sequences were aligned by eye, with the alignment based on a secondary structure using the SSU rDNA of *Tetrahymena canadensis* (Wuyts, Perriere, and Van de Peer 2004) in the Genetic Data Environment (GDE 2.2) program (Smith et al. 1994). A total of 1,291 nucleotides were included in the phylogenetic analyses under maximum parsimony (MP), neighbour-joining (NJ), maximum likelihood (ML) criteria, and Bayesian analyses.

For ML, NJ, and Bayesian analyses, we performed a likelihood ratio test using MODELTEST 3.7 (Posada and Crandall 1998) to determine the best available model for the data. The selected model was a Tamura and Nei model (TrN, Tamura and Nei 1993) with a gamma correction for among-site rate variation

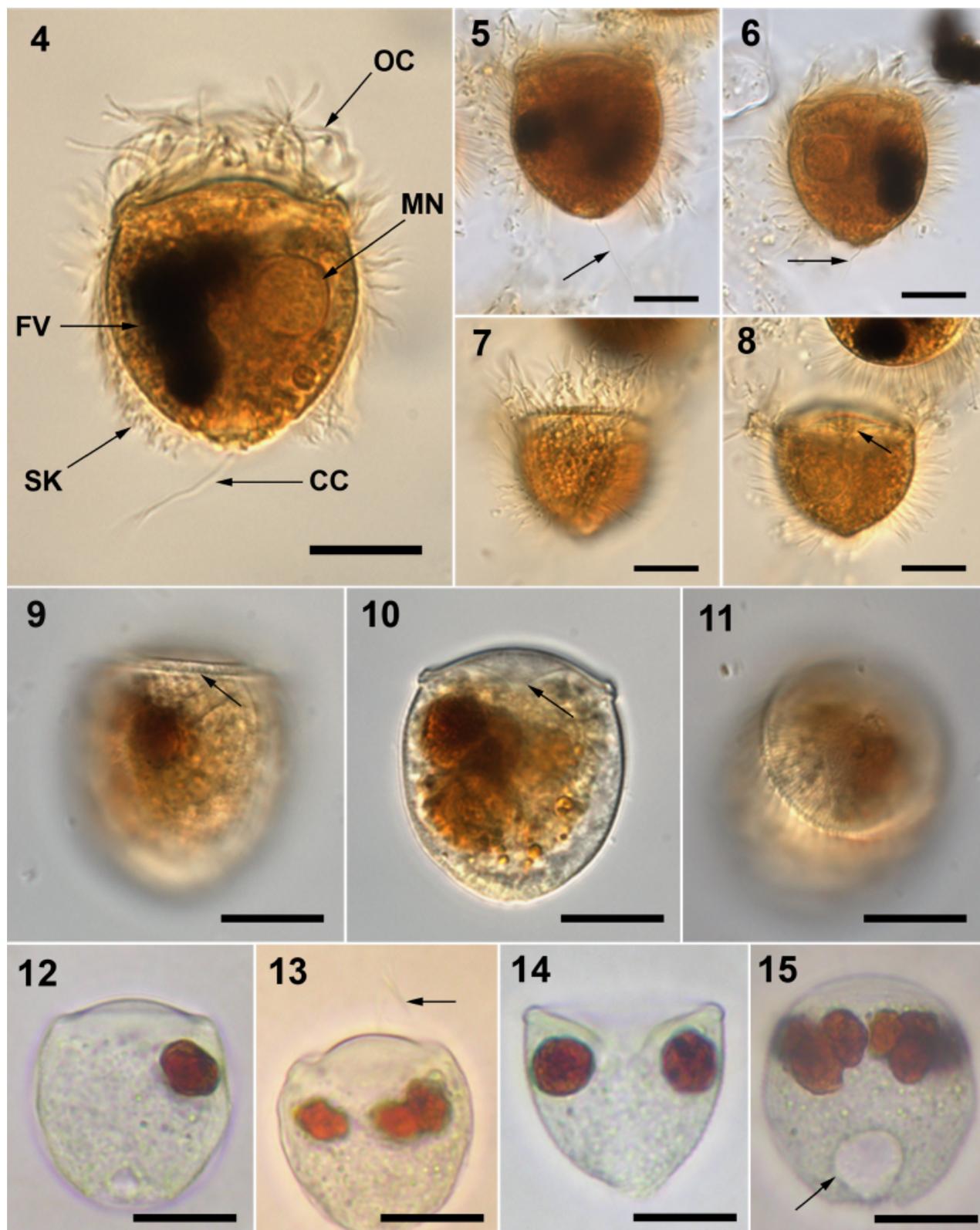


Fig. 4–15. Micrographs of cells fixed by Lugol’s solution (Fig. 4–11) and living cells (Fig. 12–15) of *Balanion masanensis* n. sp. All particles in the cells colored black or dark brown were prey (*Heterocapsa triquetra*). 4–8. Lateral view showing general cell shapes (Fig 4–6: cells satiated with prey, Fig. 7–8: starved cells), oral cilia (OC), macronucleus (MN), somatic kinetids (SKs), caudal cilium (CC), and food vacuoles (FV). Arrows in Fig. 5, 6 indicate the CC and the arrow in Fig. 8 indicates the oral depression. 9, 10. Lateral view of deciliated cell (Fig. 9: “dorsal” focal plane, Fig. 10: middle focal plane) showing the distinct “lip” (arrow in Fig. 9) surrounded by dense oral dikinetids and surrounding a central oral depression (arrow in Fig. 10). 11. Semi-anterior view showing the flattened oral region. 12. Lateral view showing general shape. 13. Dorsal–lateral view indicating the position and shape of the OC (arrow). 14. Lateral view showing the oral depression. 15. Dorsal–posterior view showing the position and shape of contractile vacuole (arrow). Scale bars = 10 μ m.

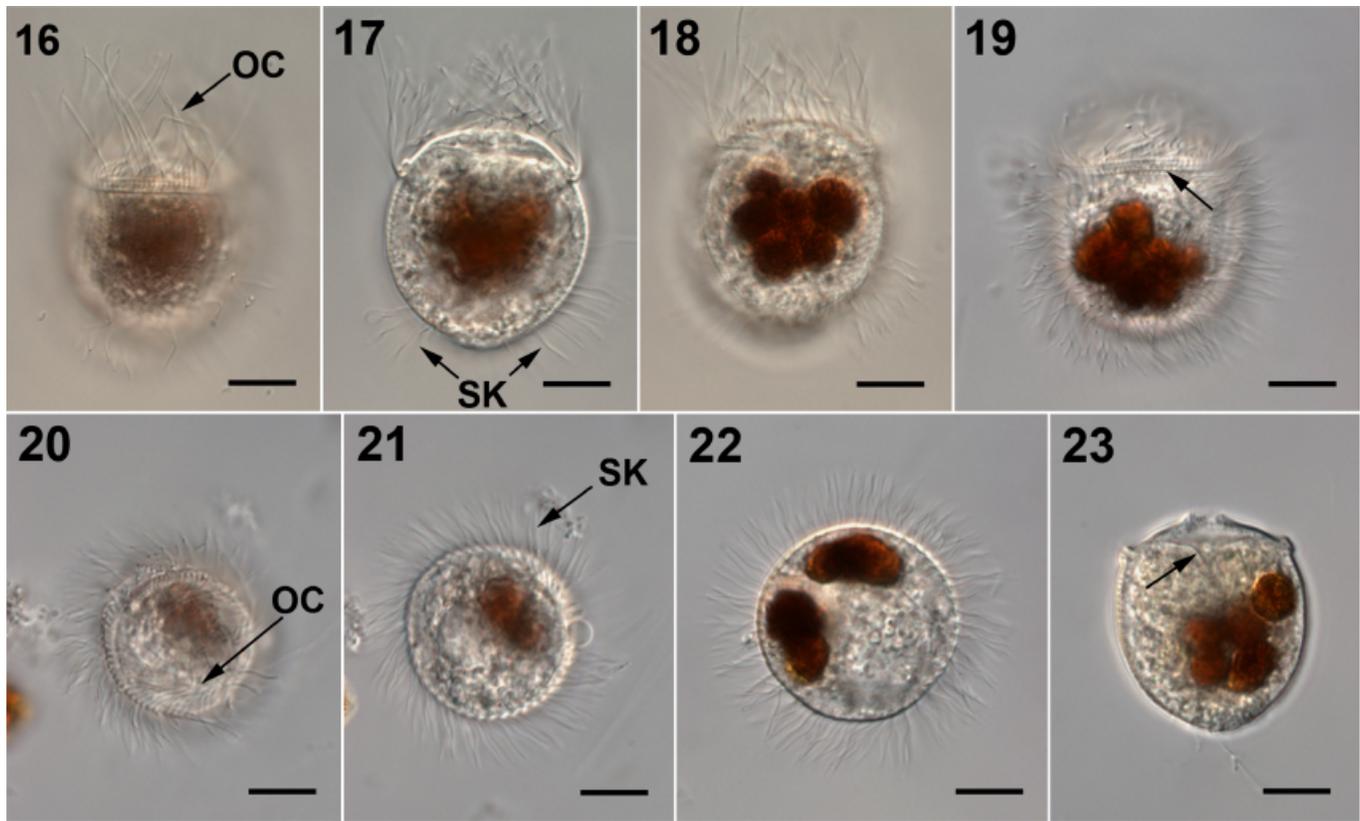


Fig. 16–23. Micrographs of *Balanion masanensis* n. sp. fixed by formalin solution. 16–19. Lateral views showing the oral cilia (OC), cilia of the somatic kinetids (SKs), and distinct ‘lip’ (arrow in Fig. 19). 20–22. Anterior views demonstrating the OC and cilia of the SKs. 23. Lateral view showing the oral depression (= cytostome) of a deciliated cell. Scale bars = 10 μ m.

($\alpha = 0.5528$) and invariant sites ($I = 0.3165$). The phylogenetic analyses were performed by ML, MP, and NJ implemented with PAUP 4.0b10 (Swofford 2002) for MacPro OS X. The ML, NJ, and Bayesian analysis used a single model (TrN+I+ Γ) for the data set: (BaseFreq = (0.3139, 0.1853, 0.2372), RateMatrix = (1.0000, 2.5048, 1.0000, 1.0000 3.8650), $\Gamma = 0.5528$, Invariant = 0.3165). Maximum likelihood and MP analyses used heuristic searches with a branch-swapping algorithm (tree bisection–reconnection). Distance matrices for NJ analyses were calculated via ML. The robustness of branches was tested by bootstrap analyses (Felsenstein 1985) using 1,000 (NJ and MP) and 250 (ML) replications. Ten heuristic searches in MP and NJ and two in ML (starting trees via random stepwise addition) were used for each bootstrap replicate. Maximum likelihood bootstrapping was performed on 10 processors. Bayesian analyses were performed with a TrN model with rate set to invgamma and nucleotide frequencies set to dirichlet (A = 0.3139, C = 0.1853, G = 0.2372, T = 0.2637). Four Markov chains (2,000,000 generations per chain) were run and trees were saved every 1,000 generations with the first 800 trees discarded. A majority-rule consensus tree was created from the remaining 1,201 trees to examine the posterior probabilities of each clade.

RESULTS

Balanion masanensis Kim, Jeong, & Lynn, n. sp.

Description. *Balanion masanensis* n. sp. is oval with a flattened anterior oral region, which is encircled by oral dikinetids bearing two cilia, and a rounded or bluntly pointed posterior end

when the cells are both alive and stained (Fig. 1–15, 24, 25). Living cells are transparent except for food vacuoles (Fig. 12–15). Their flattened anterior region often forms a dome with the perimeter receded in a groove, and this part easily changes shape, becoming either inflated or depressed (Fig. 13, 14). The cytostome, a slight funnel-like central depression on the flattened anterior end, is observed well in SEM specimens (Fig. 32). In lateral views, the cytostome was often observed in live cells and those fixed by formalin and Lugol’s solution (Fig. 8, 10, 14, 23). The flattened oral region was covered by radial ridges in Lugol’s, SEM, and TEM specimens (Fig. 11, 32, 35–40, 41). At the edge of the flattened oral region of deciliated SEM specimens, the ‘brush,’ called a ‘brosse’ by Jakobsen and Montagnes (1999), was observed (Fig. 32, 35–40). The brosse consisted of a relatively long kinety (1.5–2 μ m; three to five kinetids) and relatively short kinety (ca 0.5 μ m; three to six kinetids). With our preparations, we were unable to determine whether these were organized as dikinetids.

The number of oral dikinetids was 51–63 (Table 2). Oral flaps were observed inside the oral dikinetids, encircling the oral region. The range of the length of oral dikinetid cilia was 8.4–15.5 μ m ($n = 24$), measured in the SEM specimens. This oral cilia complex was easily lost in QPS specimens (Fig. 24–30), but was well preserved in specimens fixed with formalin (final concentration 5%, v/v) or in Lugol’s (1%–2%) solutions. The flaps in some SEM specimens were swollen (Fig. 34).

The ranges of cell length, cell width, and oral diameter of living cells were 27.1–43.3, 24.8–31.7, and 24.8–29.6 μ m, respectively, while those of the QPS-stained specimens were 22.8–37.3,

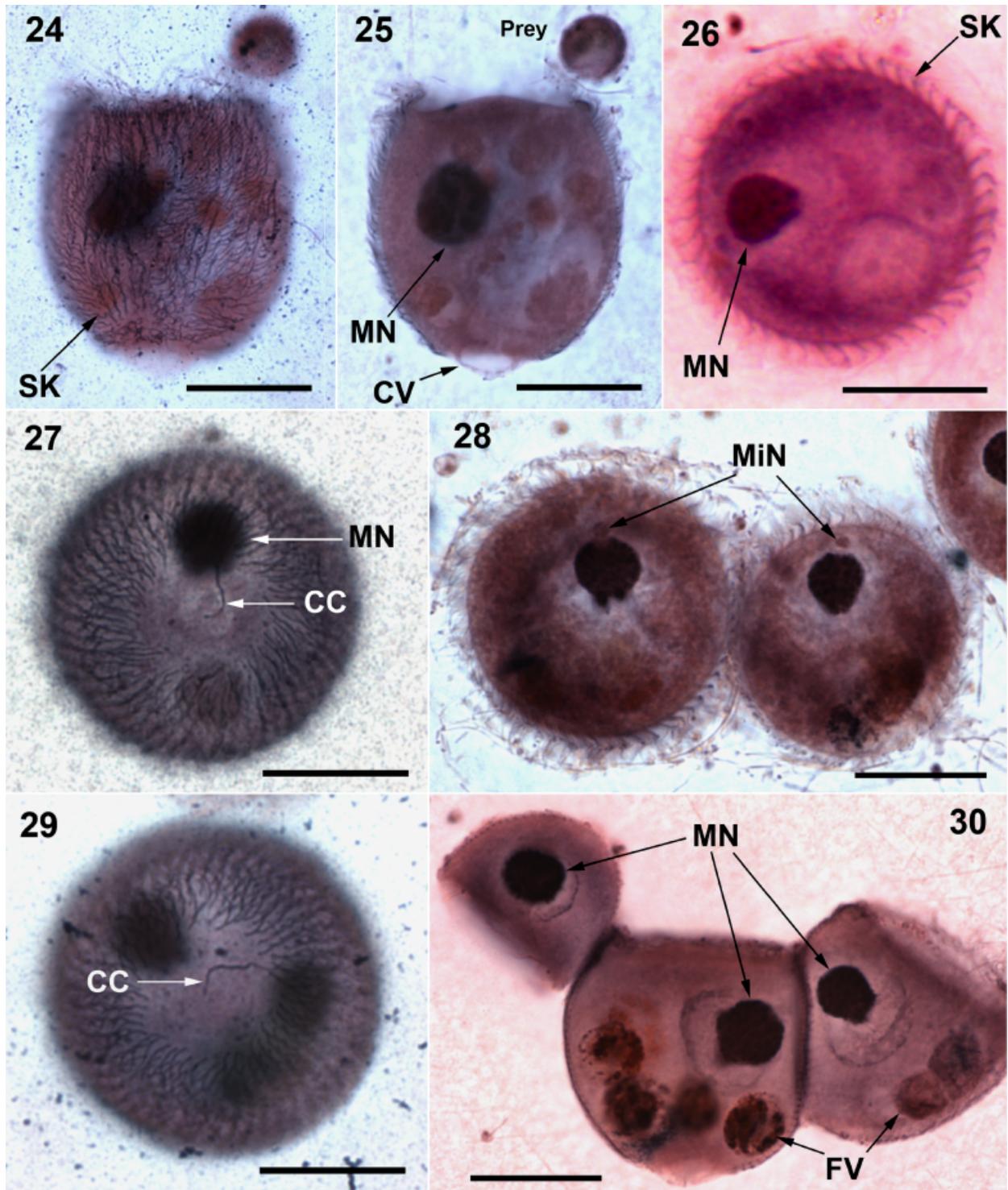


Fig. 24–30. Micrographs of protargol-stained specimens of *Balanion masanensis* n. sp. 24, 25. Lateral views (Fig. 24: “dorsal” focal plane, Fig. 25: middle focal plane) of a cell demonstrating the somatic kineties (SKs), macronucleus (MN), and contractile vacuole (CV). 26. Anterior view taken in the middle of the cell showing the position and shape of MN and SKs. 27, 29. Posterior views showing caudal cilium (CC) and the bare area of the end of the cells. 28. Anterior view taken in the middle of the cell showing the position and shape of MN and micronucleus (MiN). 30. Lateral views showing variations in cell size. Note MN and food vacuoles (FV). Scale bars = 10 μ m.



Fig. 31–34. Scanning electron micrographs of *Balanion masanensis* n. sp. 31, 33. Lateral views showing oral cilia (OC) and somatic kineties (SKs). Arrow in Fig. 33 indicates the oral anlage. 32. Anterior views showing the shape and position of the cytostome and “brosse” (B). 34. Oral dikinetids (black arrow heads) are forming the “complex” with oral flaps (white arrows). Scale bars = 10 μm for Fig. 31–33, and 1 μm for Fig. 34.

26–5.4, and 26–33 μm, respectively (Table 2). Forty-six to 55 longitudinally oriented somatic kineties (SKs) were equally spaced around the cell body, and extended from the oral end to near the posterior end. Typically, the posterior one-fifth of the cell was bare except for a caudal cilium (ca 14 μm long) (Fig. 4–6, 27, 28, Table 2). Each SK consisted of 24–50 monokinetids (Fig. 24,

25, 31, 33), and each kinetid bore cilia of 10–15 μm long. The numbers of oral dikinetids, SKs, and monokinetids in a SK of cells starved for 1 day were not different from those of cells satiated with *H. triquestra*.

This ciliate had one ellipsoid macronucleus of 6.8–13.4 × 6.8–10.5 μm and one micronucleus of 2.0–2.8 × 1.5–2.5 μm, visible

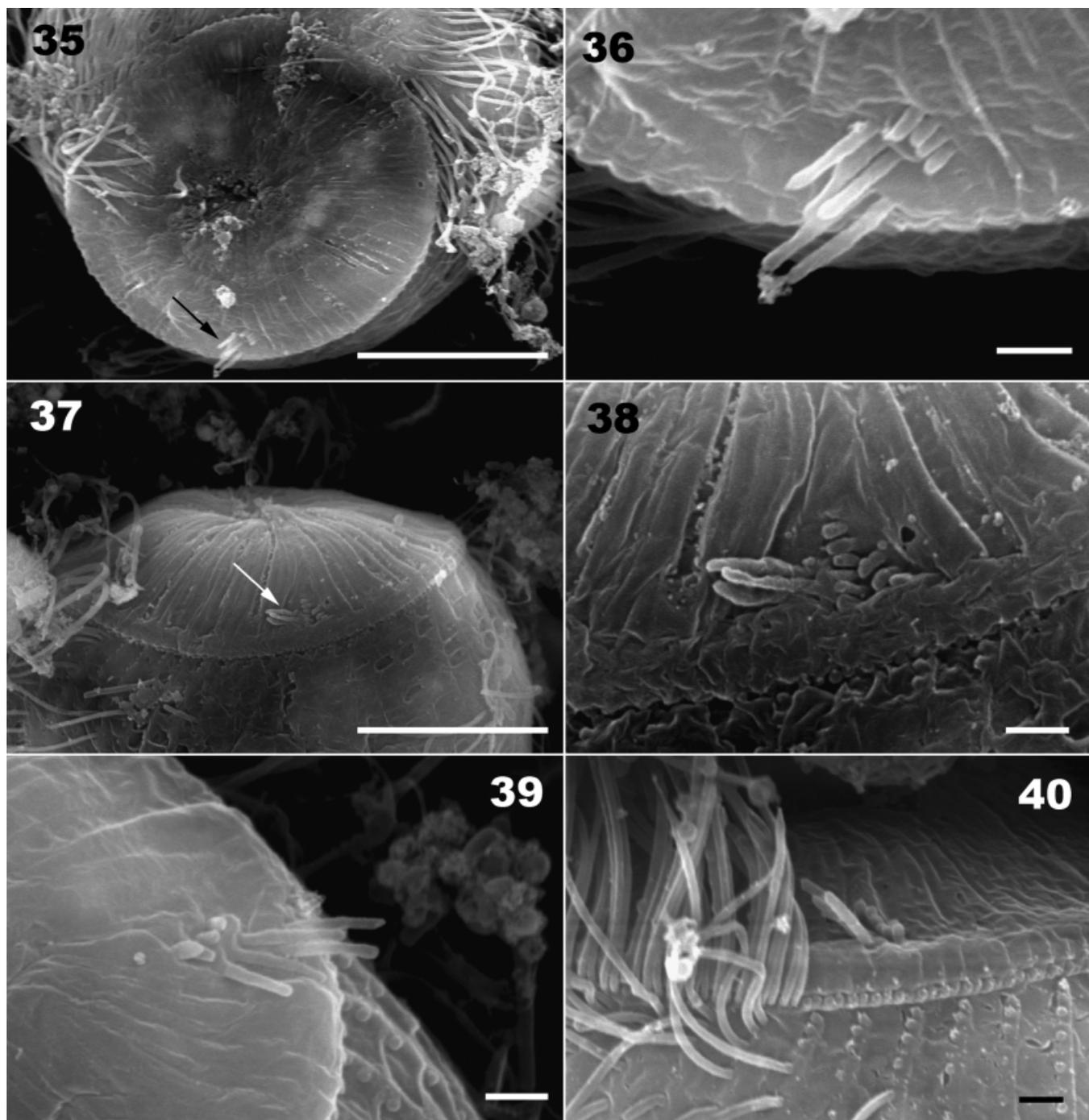


Fig. 35–40. Electron micrographs of *Balanion masanensis* n. sp. showing the brosse located inside the circumoral dikinetids. Scale bars = 10 μ m for Fig. 40, 42, and 44, and 1 μ m for Fig. 41, 43, and 45. We are unable to determine whether the brosse is organized as monokinetids or dikinetids.

only in QPS-stained specimens (Fig. 28). The macronucleus was located near the center of the cell and the micronucleus located on its outer side (Fig. 28).

In TEM photos, a circumoral microtubular ribbon (COMT) was observed adjacent to the oral kinetids (Fig. 42, 45, 46). The number of microtubules inside oral flaps was approximately 7–9 (Fig. 43, 44).

Gene sequence. The SSU rRNA gene sequence (Genbank Accession No. AM412525) of this new isolate has a length of 1,752

nucleotides and a GC content of 43.6%. Its sequence shows <90.9% similarity with closely related species of the Class Prostomatea, but shows the highest SSU rRNA gene sequence similarity to *Cryptocaryon irritans* (Cryptocaryonidae, Prorodontida; 90.9%), followed by *Prorodon teres* (Prorodontidae, Prorodontida; 90.5%) and *Prorodon viridis* (Prorodontidae, Prorodontida; 89.7%). The phylogenetic analyses using Bayesian, ML, MP, and NJ grouped the new isolate with members of the Class Prostomatea, but its phylogenetic placement was ambiguous. In Bayesian

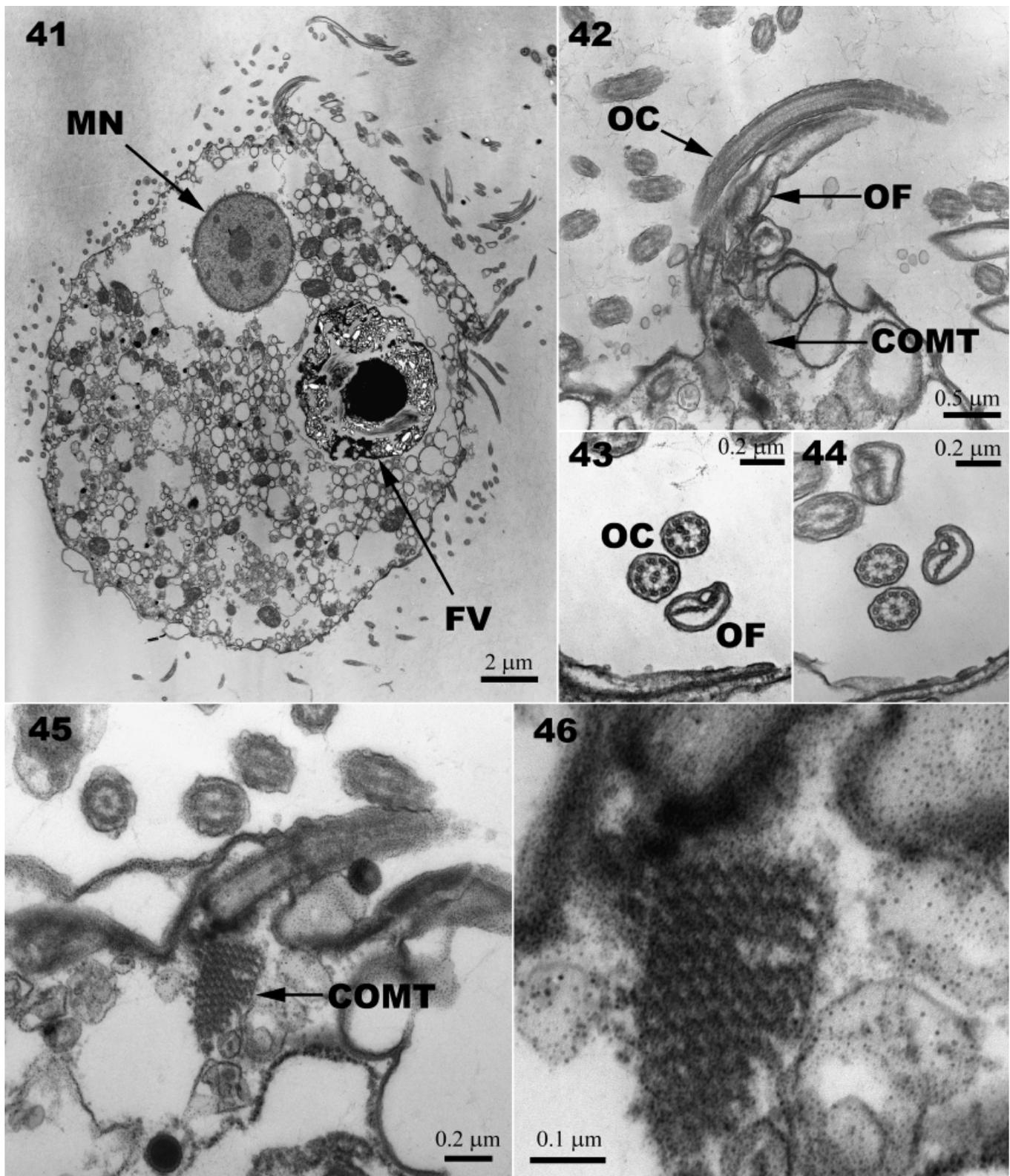


Fig. 41–46. Transmission electron micrographs of *Balanion masanensis* n. sp. 41, 42, 45, 46. Anterior longitudinal section showing macronucleus (MN), oral cilia (OC), oral flap (OF), and circumoral microtubular ribbon (COMT). 43, 44. Cross-sections of the OC and OF.

Table 2. Morphometric characterization of *Balanion masanensis* n. sp. based on protargol-stained specimens (and living cells in parenthesis) and specimens observed by scanning electron microscopy (*).

Character	Range	Average	SD	n
Cell length (μm)	22.8–37.3 (27.1–43.3)	30.6 (35.2)	3.5 (4.6)	70 (23)
Cell width (μm)	26.0–35.4 (24.8–31.7)	30.7 (28.6)	2.2 (2.3)	70 (26)
Oral diameter (μm)	26–33 (24.8–29.6)	29.5 (27.7)	1.5 (1.3)	70 (25)
No. of oral dikinetids	63–51	57.8	3.9	10
Length of oral cilia* (μm)	8.4–15.5	12.5	2.0	24
No. of kinetids in long brosse kinety	3–5	4.2	0.8	6
No. of kinetids in short brosse kinety	3–6	4.5	1.0	6
Length of long brosse kinety (μm)	1.5–2	1.7	0.2	15
Length of short brosse kinety (μm)	0.4–0.6	0.5	0.1	14
No. of somatic kinety	46–55	51	2.5	22
Length of somatic kinety (μm)	19.8–27.4	24.1	7.4	13
No. of cilia per somatic kinety	24–50	38.6	7.4	15
Length of somatic cilia* (μm)	2.8–7.2	5.1	1.0	20
Somatic kinety as % of total cell length	73.5–81.8	78.1	2.8	13
No. of macronucleus	1	1	0	68
Length of macronucleus (μm)	6.8–13.4	9.3	1.0	68
Width of macronucleus (μm)	6.8–10.5	8.1	0.7	68
Length of micronucleus (μm)	2.0–2.8	2.4	0.3	10
Width of micronucleus (μm)	0.5–2.5	1.9	0.3	10

and ML analyses, the new species clustered together with *C. irritans* to form a monophyletic clade with weak support (0.9 posterior probability, 50% BP) (Fig. 47).

Time and locality of isolation. Our new isolate was found in Masan Bay, which is located in southern coastal Korea ($35^{\circ}12'N$, $128^{\circ}34'E$). It was often observed from November to early March, when the ranges of the water temperatures and salinities were 3.1°C – 14.1°C and 23.3–32.7 psu, respectively.

Remarks on culturing and behavior. The new isolate has been cultured for > 1 yr by providing the red-tide dinoflagellate *H. triquetra* as prey. The maximum growth rate of this ciliate on *H. triquetra*

was $\sim 2.5\text{ day}^{-1}$ (our unpubl. data). When starved, this ciliate swims in straight paths, but when satiated, it swims in a large circle.

DISCUSSION

Morphological and genetical characteristics. Our isolate from Masan Bay is definitely a prostome ciliate, based on the sequence of its SSU rDNA and on its general morphology. It can be assigned unambiguously to the genus *Balanion* because it has (1) a brosse inside the circumoral ring of dikinetids, (2) shows the circumoral microtubular ribbon (COMT), and (3) has elongated oral flaps that are supported by a ribbon of microtubules. However, its body size and the numbers of oral cilia, SKs, kinetosome per SK, and microtubules inside oral flaps of this ciliate were considerably different from those of *B. comatum* and *B. planctonicum*, the only other *Balanion* species, so far reported (Table 3). The mean cell length, cell width, and oral diameter of the QPS-stained specimens (30.6, 30.7, and 29.5 μm , respectively) were considerably larger than those of *B. comatum* and *B. planctonicum* (13.1–20.0, 10.0–13.5, and 9–13.3 μm , respectively). The number of oral dikinetids of Masan Bay *Balanion* (range = 51–63) was greater than that of *B. comatum* (35–46) and *B. planctonicum* (29–39). The six to nine brosse monokinetids of our new isolate were greater than the two kinetids of *B. comatum*, but similar to the eight kinetids of *B. planctonicum*. We were unable to determine whether the brosse kinetids of *B. masanensis* n. sp. are organized as monokinetids or dikinetids. In addition, the brosse appears to consist of two different-sized kineties, which has not been previously described. The 51–63 SKs of the Masan Bay isolate was greater than for both *B. comatum* (20–29) and *B. planctonicum* (21–29). The 24–50 monokinetids per SK of the Masan Bay isolate were also greater than for both *B. comatum* (12–26) and *B. planctonicum* (7–12). For these reasons, we establish the new species *B. masanensis* n. sp.

Class Prostomatea Schewiakoff, 1896

Order Prorotontida Corliss, 1974

Family Balanionidae Small and Lynn, 1985

Balanion masanensis Kim, Jeong, & Lynn n. sp.

Table 3. Comparison of morphological characteristics of *Balanion masanensis* n. sp. with other *Balanion* spp. based on specimens observed by the protargol-staining, scanning electron microscopy (*), and transmission electron microscopy (***) specimens.

Character	<i>B. planctonicum</i>		<i>B. comatum</i>	<i>B. masanensis</i>	
Cell length (μm)	13.1 (10–18)	20	14 (12–26)	30.6 (23–37)	
Width (μm)	13.5 (10–17)	15	12.1	30.7 (25–32)	
Oral end diameter (μm)	10.1 (7–13)	13.3	9	29.5 (26–33)	
Oral dikinetid number*	33.6 (29–38)	34 (29–39)	NA	57.8 (51–63)	
Oral cilia length* (μm)	10.3 (8–13)	8	NA	12.5 (8.4–15.5)	
SK number	24.7 (21–28)	26 (23–29)	NA	51 (46–55)	
SK length (μm)	6.8 (4–10)	8	9	24.1 (16.8–27.4)	
SK as % of cell length	54	60	63	78.1 (73.5–81.8)	
Kinety per SK	11.3 (7–12)	(8–11)	NA	38.6 (24–50)	
Somatic cilia length (μm)	6.4	NA	6–7	7.7 (6–10)	
Brosse kinetids*	0	4	NA	At least 2 (n = 1) 6–9	
Caudal cilia	1	1	1	1	
MN number	1	1	1	1	
MN length (μm)	4.5 (3–6)	(8–11)	5	9.3 (7–13)	
MN width (μm)	3.6 (3–5)	(7–8)	4.6	8.1 (7–11)	
Microtubules in oral flap**	NA	4	NA	(9–10) NA	
Presence of oral depression	NA	Food induced	Present	Rare	Present
Habitat	Freshwater	Freshwater	Marine/brackish	Marine/brackish	Marine/estuary
Reference	Foissner et al. (1990)	Bardele (1999)	Wulff (1919)	Jakobsen and Montagnes (1999)	This study

NA, not available; MN, macronucleus; SK, somatic kinety.

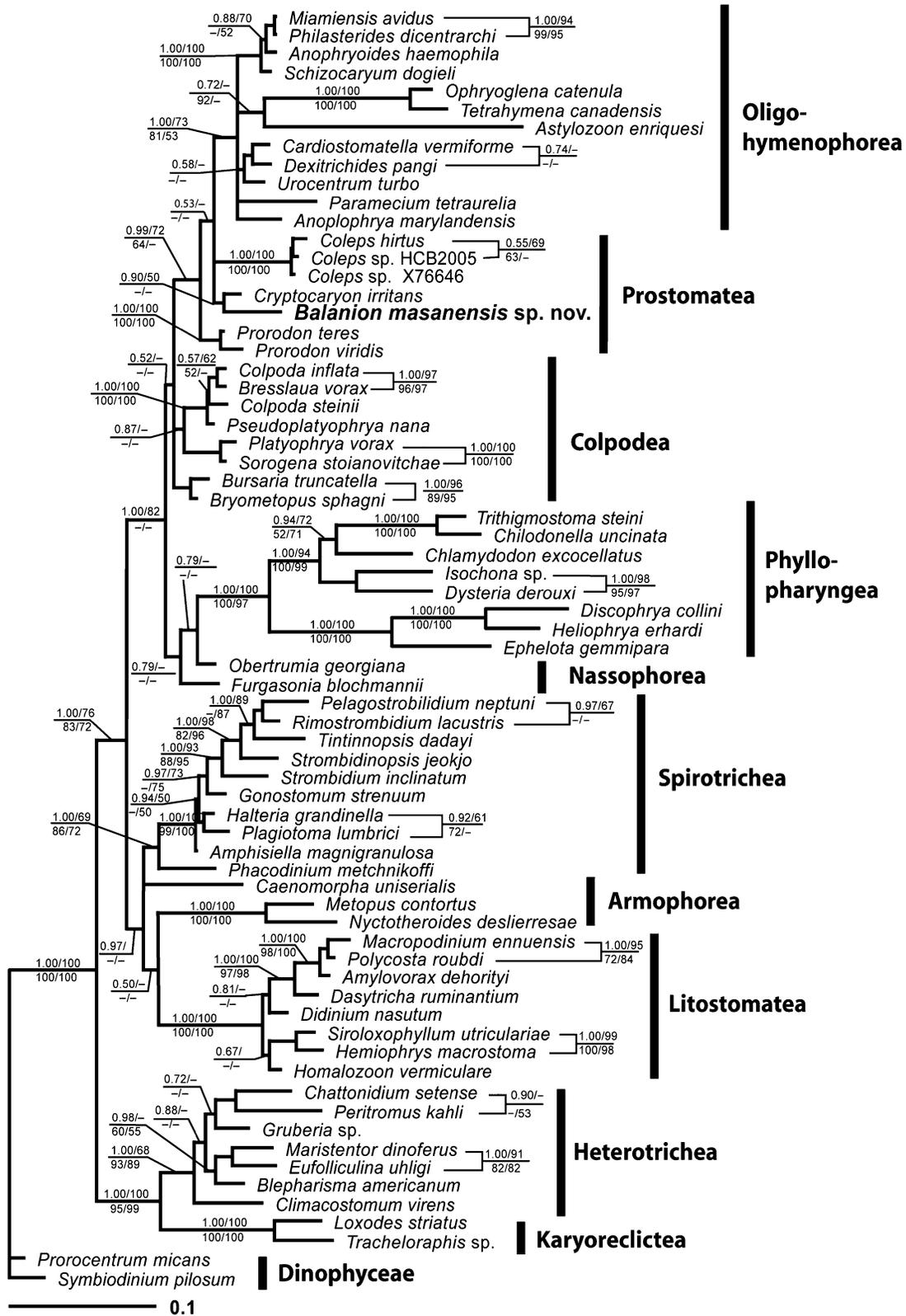


Fig. 47. Consensus Bayesian tree (–ln L = 13,579.625) based on 1,291 aligned positions with *Prorocentrum micans* and *Symbiodinium pilosum* as outgroup taxa. Numbers above branches indicate the posterior probability (left) from Bayesian analysis and bootstrap values (right) from maximum likelihood (ML) analysis. Numbers below branches indicate bootstrap values for neighbor-joining (NJ) (left) and maximum parsimony (MP) (right) analyses. Bootstrap values $\geq 50\%$ and posterior probabilities ≥ 0.5 are shown.

Diagnosis. Ovoid with a flattened anterior oral region and a rounded or bluntly pointed posterior end when the cells are both alive and stained. Brosse inside the circumoral dikinetids, which are supported by circumoral microtubules and accompanied by elongate flaps. Cell length, cell width, and oral diameter of living cells are 27.1–43.3, 24.8–31.7, and 24.8–29.6 μm , respectively, while those of the QPS-stained specimens are 22.8–37.3, 26–5.4, and 26–33 μm , respectively. The numbers of oral dikinetids are 51–63. The six to nine brosse monokinetids are arranged into two different-sized kineties. The 51–63 SKs have 24–50 monokinetid per kinety.

Etymology. The specific epithet *masanensis* refers to the type locality—Masan Bay, Korea.

Deposition of type material. A hapantotype slide as USNM slide 1107776 of protargol-stained cells has been deposited in the Ciliate Type Specimen Slide Collection, US Natural History Museum, Smithsonian Institution, Washington, DC.

Gene sequence. The SSU rRNA gene sequence—GenBank Accession No. AM412525.

ACKNOWLEDGMENTS

We are grateful to Jong Heok Kim and Nam Seon Kang for technical assistance. This paper was funded by grants from a National Research Laboratory grant from MOST & KOSEF (M1-0302-00-0068) and Sooteuk program from MOMAF & KMIST awarded to H. J. Jeong, KOSEF (R01-2006-000-10207-0) awarded to W. Shin, and a NSERC Canada Discovery Grant awarded to D. H. L.

LITERATURE CITED

- Affa'a, F.-M., Hickey, D. A., Strüder-Kypke, M. & Lynn, D. H. 2004. Phylogenetic position of species in the genera *Anoplophrya*, *Plagiotoma*, and *Nyctotheroides* (Phylum Ciliophora), endosymbiotic ciliates of annelids and anurans. *J. Eukaryot. Microbiol.*, **51**:301–306.
- Agatha, S. & Strüder-Kypke, M. C. 2007. Phylogeny of the order Choreotrichida (Ciliophora, Spirotricha, Oligotricha) as inferred from morphology, ultrastructure, ontogenesis, and SSrRNA gene sequences. *Eur. J. Protistol.*, **43**:37–63.
- Agatha, S., Strüder-Kypke, M. C., Beran, A. & Lynn, D. H. 2005. *Pelagostrobilidium neptuni* (Montagnes and Taylor, 1994) and *Strombidium biarmatum* nov. spec. (Ciliophora, Oligotricha): phylogenetic position inferred from morphology, ontogenesis, and gene sequence data. *J. Protistol.*, **41**:65–83.
- Bardele, C. E. 1999. Ultrastructure and morphogenesis of the prostome ciliate *Balanion planktonicum*. *Eur. J. Protistol.*, **35**:17–33.
- Bernhard, D., Leipe, D. D., Sogin, M. L. & Schlegel, M. 1995. Phylogenetic relationships of the Nassulida within the phylum Ciliophora inferred from the complete small subunit rRNA gene sequences of *Furgasonia blochmanni*, *Obertrumia georgiana*, and *Pseudomicrothoras dubius*. *J. Eukaryot. Microbiol.*, **42**:126–131.
- Bernhard, D., Stechmann, A., Foissner, W., Ammermann, D., Hehn, M. & Schlegel, M. 2001. Phylogenetic relationships within the class Spirotrichea (Ciliophora) inferred from small subunit rRNA gene sequences. *Mol. Phylogenet. Evol.*, **21**:86–92.
- Cameron, S. L., Adlard, R. D. & O'Donoghue, P. J. 2001. Evidence for an independent radiation of endosymbiotic litostome ciliates within Australian marsupial herbivores. *Mol. Phylogenet. Evol.*, **20**:302–310.
- Felsenstein, J. 1985. Confidence limits on phylogenies: an approach using the bootstrap. *Evolution*, **39**:783–791.
- Foissner, W. 1984. Morphologie und Infraciliatur einiger limnischer Ciliaten (Protozoa: Ciliophora). *Schweiz. Z. Hydrol.*, **46**:210–223.
- Foissner, W., Berger, H. & Kohmann, F. 1994. Taxonomische und ökologische Revision der Ciliaten des Saprobiensystem's— Band III: Hymenostomata, Prostomatida, Nassolida. *Informationsberichte des Bayer. Landesamtes für Wasserwirtschaft*, Heft 1/94:548.
- Foissner, W., Oleksiv, I. & Müller, H. 1990. Morphologie und Infraciliatur einiger Ciliaten (Protozoa: Ciliophora) aus stagnierenden Gewässern. *Arch. Protistenkd.*, **138**:191–206.
- Greenwood, S. J., Schlegel, M., Sogin, M. L. & Lynn, D. H. 1991. Phylogenetic relationships of *Blepharisma americanum* and *Colpoda inflata* within the phylum Ciliophora inferred from complete small subunit rRNA gene sequences. *J. Protozool.*, **38**:1–6.
- Hammerschmidt, B., Schlegel, M., Lynn, D. H., Leipe, D. D., Sogin, M. L. & Raikov, I. B. 1996. Insights into the evolution of nuclear dualism in the ciliates revealed by phylogenetic analysis of rRNA sequences. *J. Eukaryot. Microbiol.*, **43**:225–230.
- Hirt, R. P., Dyal, P. L., Wilkinson, M., Finlay, B. J., Roberts, D. M. & Embley, T. M. 1995. Phylogenetic relationships among karyorelictids and heterotrichs inferred from small subunit rRNA sequences: resolution at the base of the ciliate tree. *Mol. Phylogenet. Evol.*, **4**:77–87.
- Jakobsen, H. H. & Hansen, P. J. 1997. Prey size selection, grazing and growth response of the small heterotrophic dinoflagellates *Gymnodinium* sp. and the ciliate *Balanion comatum*—a comparative study. *Mar. Ecol. Prog. Ser.*, **158**:75–86.
- Jakobsen, H. H. & Montagnes, D. J. S. 1999. A redescription of *Balanion comatum* Wuff, 1919 (Prorodontida, Ciliophora), with notes on its cultivation and behavior. *J. Eukaryot. Microbiol.*, **46**:198–205.
- Jeong, H. J., Kim, J. S., Kim, S. H., Song, J. Y., Lee, I. H. & Lee, G. H. 2004. *Strombidinopsis jeokjo* n. sp. (Ciliophora: Choreotrichida) from the coastal waters off Western Korea: morphology and small subunit ribosomal DNA gene sequence. *J. Eukaryot. Microbiol.*, **51**:451–455.
- Jung, S. J., Kitamura, S., Song, J. Y., Joung, I. Y. & Oh, M. J. 2005. Complete small subunit rRNA gene sequence of the scuticociliate *Miamiensis avidus* pathogenic to olive flounder *Paralichthys olivaceus*. *Dis. Aquat. Org.*, **64**:159–162.
- Kenter, U., Zimmermann, U. & Müller, H. 1996. Grazing rates of the fresh-water ciliate *Balanion planktonicum* determined by flow cytometry. *J. Plank. Res.*, **18**:1047–1053.
- Lasek-Nesselquist, E. & Katz, L. A. 2001. Phylogenetic position of *Sorogena stoianovitchae* and relationships within the class Colpodea (Ciliophora) based on SSU rDNA sequences. *J. Eukaryot. Microbiol.*, **48**:604–607.
- Leipe, D. D., Bernhard, D., Schlegel, M. & Sogin, M. L. 1994. Evolution of 16S-like ribosomal RNA genes in the ciliophoran taxa Litostomatea and Phyllopharyngea. *Eur. J. Protistol.*, **30**:354–361.
- Li, L., Song, W., Warren, A., Wang, Y., Ma, H., Hu, X. & Chen, Z. 2006. Phylogenetic position of the marine ciliate, *Cardiostomatella vermiforme* (Kahl, 1928) Corliss, 1960 inferred from the complete SSrRNA gene sequence, with establishment of a new order Loxocephalida n. ord. (Ciliophora, Oligohymenophorea). *Eur. J. Protistol.*, **42**:107–114.
- Lynn, D. H. & Small, E. B. 2002. Phylum Ciliophora. In: Lee, J. J., Leedale, G. F. & Bradbury, P. (ed.), *The Illustrated Guide to the Protozoa*. 2nd ed. Society of Protozoologists, Lawrence, KS. p. 371–655.
- Lynn, D. H. & Strüder-Kypke, M. 2002. Phylogenetic position of *Licinophora*, *Lechriopyla*, and *Schizocaryum*, three unusual ciliates (phylum Ciliophora) endosymbiotic in echinoderms (phylum Echinodermata). *J. Eukaryot. Microbiol.*, **49**:460–468.
- Lynn, D. H., Wright, A. D., Schlegel, M. & Foissner, W. 1999. Phylogenetic relationships of orders within the class Colpodea (Phylum Ciliophora) inferred from small subunit rRNA gene sequences. *J. Mol. Evol.*, **48**:605–614.
- Medlin, L., Elwood, H. J., Stickel, S. & Sogin, M. L. 1988. The characterization of enzymatically amplified eukaryotic 16S-like rRNA-coding regions. *Gene*, **71**:491–499.
- Miao, W., Simpson, A. G., Fu, C. & Lobban, C. S. 2005. The giant zooxanthellae-bearing ciliate *Maristentor diniferus* (Heterotricha) is closely related to Folliculinidae. *J. Eukaryot. Microbiol.*, **52**:11–16.
- Modeo, L., Petroni, G., Rosati, G. & Montagnes, D. J. 2003. A multidisciplinary approach to describe protists: redescription of *Novistrombidium testaceum* Anigstein 1914 and *Strombidium inclinatum* Montagnes, Taylor, and Lynn 1990 (Ciliophora, Oligotrichia). *J. Eukaryot. Microbiol.*, **50**:175–189.
- Modeo, L., Rosati, G., Andreoli, I., Gabrielli, S., Verni, F. & Petroni, G. 2006. Molecular systematics and ultrastructural characterization of a forgotten species: *Chattonidium setense* (Ciliophora, Heterotricha). *Proc. Jpn. Acad. B Phys. Biol. Sci.*, **82**:359–374.
- Montagnes, D. J. S. & Lynn, D. H. 1987. A quantitative protargol stain (QPS) for ciliates: method description and test of its quantitative nature. *Mar. Microb. Food Webs*, **2**:83–93.

- Müller, H. 1989. The relative importance of different ciliate taxa in the pelagic food web of Lake Constance. *Microb. Ecol.*, **18**:261–273.
- Müller, H. & Geller, W. 1993. Maximum growth rates of aquatic ciliated protozoa: the dependence on body size and temperature reconsidered. *Arch. Hydrobiol.*, **126**:315–327.
- Müller, H. & Schlegel, A. 1999. Responses of three fresh-water planktonic ciliates with different feeding modes to cryptophytes and diatom prey. *Aquat. Microb. Ecol.*, **17**:49–60.
- Petroni, G., Dini, F., Verni, F. & Rosati, G. 2002. A Molecular approach to the tangled intrageneric relationships underlying phylogeny in *Euplotes* (Ciliophora, Spirotrichea). *Mol. Phylogenet. Evol.*, **22**:118–130.
- Posada, D. & Crandall, K. A. 1998. MODELTEST: testing the model of DNA substitution. *Bioinformatics*, **14**:817–818.
- Ragan, M. A., Cawthorn, R. J., Despres, B., Murphy, C. A., Singh, R. K., Loughlin, M. B. & Bayer, R. C. 1996. The lobster parasite *Anophryoides haemophila* (Scuticociliatida: Orchitophryidae): nuclear 18S rDNA sequence, phylogeny and detection using oligonucleotide primers. *J. Eukaryot. Microbiol.*, **43**:341–346.
- Schmidt, S. L., Bernhard, D., Schlegel, M. & Foissner, W. 2007. Phylogeny of the Stichotrichia (Ciliophora; Spirotrichea) reconstructed with nuclear small subunit rRNA gene sequences: discrepancies and accordances with morphological data. *J. Eukaryot. Microbiol.*, **54**:201–209.
- Shin, M. K., Hwang, U. W., Kim, W., Wright, A.-D. G., Krawczyk, C. M. & Lynn, D. H. 2000. Phylogenetic position of the ciliates *Phacodinium* (Order Phacodiniida) and *Protocruzia* (Subclass Protocruziidia) and systematics of the spirotrich ciliates examined by small subunit ribosomal RNA gene sequences. *Eur. J. Protistol.*, **36**:293–302.
- Small, E. B. & Lynn, D. H. 1985. Phylum Ciliophora Doflein, 1901. In: Lee, J. J., Hutner, S. H. & Bovee, E. C. (ed.), *An Illustrated Guide to the Protozoa*. Society of Protozoologists, Lawrence, KS. p. 393–575.
- Smetacek, V. 1989. The annual cycle of protozooplankton in the Kiel Bright. *Mar. Biol.*, **63**:1–11.
- Smith, S. W., Overbeek, R., Woese, C. R., Gilbert, W. & Gillevet, P. M. 1994. The genetic data environment an expandable GUI for multiple sequence analysis. *Comp. Appl. Biosci.*, **10**:671–675.
- Snoeyenbos-West, O. L. O., Cole, J., Campbell, A., Coats, D. W. & Katz, L. A. 2004. Phylogenetic relationships amongst phyllopharyngean ciliates inferred from SSU r-DNA gene genealogies. *J. Eukaryot. Microbiol.*, **51**:441–450.
- Sogin, M. L. & Elwood, H. J. 1986. Primary structure of the *Paramecium tetraurelia* small-subunit rRNA coding region: phylogenetic relationships within the Ciliophora. *J. Mol. Evol.*, **23**:53–60.
- Sogin, M. L., Ingold, A., Karlok, M., Nielsen, H. & Engberg, J. 1986. Phylogenetic evidence for the acquisition of ribosomal RNA introns subsequent to the divergence of some of the major *Tetrahymena* groups. *EMBO J.*, **5**:3625–3630.
- Sommaruga, R. & Psenner, R. 1993. Nanociliates of the order Prostomatida: their relevance in the microbial food web of a mesotrophic lake. *Aquat. Sci.*, **55**:179–187.
- Song, W., Ma, H. & AL-Rasheid, K. 2003. *Dextrotrichides pangi* n. sp. (Protozoa, Ciliophora, Scuticociliatida), a new marine ciliate from the north China sea. *J. Eukaryot. Microbiol.*, **50**:114–122.
- Sonntag, B., Posch, T., Klammer, S., Griebler, C. & Psenner, R. 2002. Protozooplankton in the deep oligotrophic Traunsee (Austria) influenced by discharges of soda and salt industries. *Water Air Soil Pollut. Focus*, **2**:211–226.
- Spurr, A. R. 1969. A low viscosity epoxy resin embedding medium for electron microscopy. *J. Ultrastruct. Res.*, **26**:31–42.
- Stechmann, A., Schlegel, M. & Lynn, D. H. 1998. Phylogenetic relationships between prostome and colpodean ciliates tested by small subunit rRNA sequences. *Mol. Phylogenet. Evol.*, **9**:48–54.
- Stoecker, D. K. & Evans, G. T. 1985. Effects of protozoan herbivory and carnivory in a microplankton food web. *Mar. Ecol. Prog. Ser.*, **25**:159–167.
- Stoecker, D. K., Davis, L. H. & Provan, A. 1983. Growth of *Favella* sp. (Ciliata: Tintinnia) and other microzooplankters in cages incubated in situ and comparison to growth in vivo. *Mar. Biol.*, **75**:293–302.
- Strüder-Kypke, M. C. & Lynn, D. H. 2003. Sequence analyses of the small subunit rRNA gene confirm the paraphyly of oligotrich ciliates sensu lato and support the monophyly of the subclasses Oligotrichia and Choreotrichia (Ciliophora, Spirotrichea). *J. Zool. (London)*, **260**:87–97.
- Strüder-Kypke, M. C., Wright, A. D., Fokin, S. I. & Lynn, D. H. 2000. Phylogenetic relationships of the Subclass Peniculia (Oligohymenophorea, Ciliophora) inferred from small subunit rRNA gene sequences. *J. Eukaryot. Microbiol.*, **47**:419–429.
- Swofford, D. L. 2002. PAUP*. Phylogenetic Analysis Using Parsimony (*and Other Methods). Version 4.0b10. Sinauer Associates, Sunderland, MA.
- Tamura, K. & Nei, M. 1993. Estimation of the number of nucleotide substitutions in the control region of mitochondrial DNA in humans and chimpanzees. *Mol. Biol. Evol.*, **10**:512–526.
- Weisse, T. & Müller, H. 1998. Planktonic protozoa and the microbial food web in the Lake Constance. *Arch. Hydrobiol. Spec. Issues Adv. Limnol.*, **53**:223–254.
- Weisse, T., Karstens, N., Meyer, V. C. L., Janke, L., Lettner, S. & Teichgräber, K. 2001. Niche separation in common prostome fresh-water ciliates: the effect of food and temperature. *Aquat. Microb. Ecol.*, **26**:167–179.
- Wright, A.-D. G. & Colorni, A. 2002. Marine fish parasite *Cryptocaryon irritans* finds taxonomic home. *Eur. J. Protistol.*, **37**:375–378.
- Wright, A. -D. G. & Lynn, D. H. 1995. Phylogeny of the fish parasite *Ichthyophthirius* and its relatives *Ophryoglena* and *Tetrahymena* (Ciliophora, Hymenostomatia) inferred from 18S ribosomal RNA sequences. *Mol. Biol. Evol.*, **12**:285–290.
- Wright, A. -D. G. & Lynn, D. H. 1997. Monophyly of the trichostome ciliates (Phylum Ciliophora: Class Litostomatea) tested using new 18S rRNA sequences from the vestibuliferids, *Isotricha intestinalis* and *Dasytricha ruminantium*, and the haptorian, *Didinium nasutum*. *Eur. J. Protistol.*, **33**:305–315.
- Wulff, A. 1919. Über das Kleinplankton der Barentssee. *Wiss. Meeresunters. Helgoland (N.F.)*, **13**:97–125.
- Wuyts, J., Perriere, G. & Van de Peer, Y. 2004. The European ribosomal RNA database. *Nucleic Acids Res.*, **32**:D101–D103.
- Zhang, H., Bhattacharya, D. & Lin, S. 2005. Phylogeny of dinoflagellates based on mitochondrial cytochrome *b* and nuclear small subunit rDNA sequence comparisons. *J. Phycol.*, **41**:411–420.

Received: 06/11/07, 09/11/07; accepted: 09/12/07