

Morphology, 18S rRNA gene sequence and life history of a new *Polydora* species (Polychaeta: Spionidae) from northeastern Japan

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ABSTRACT: A new species of spionid polychaete, *Polydora onagawaensis*, is described from mollusk shells in Pacific waters of northeastern Japan. Its nuclear 18S rRNA gene sequence as well as its morphology, reproductive features, life history and infestation characteristics are reported. *Polydora onagawaensis* sp. nov. belongs to the *Polydora ciliata/websteri* group and has a moderate size and variable black pigmentation on the palps and body. Up to 115 worms were found boring in a single scallop shell from suspended cultures in Onagawa Bay, with significantly higher numbers in the right than in the left valve. Females repeatedly deposited a string of egg capsules from around October to June (seawater temperature was below 15°C). The larvae developed inside the egg capsules for 2 wk (10°C, laboratory conditions), until the 3-chaetiger stage, before being released as planktonic larvae. The main spawning occurred in December, recruitment onto the shells increased after January, and most large worms disappeared between July and October. Thus, the estimated life span is around 1.5 yr after settlement. Details on biology and gene information not only contribute to distinguishing the species from other polydorids similar in morphology, but also allow control of polydorid infestation in mollusk aquaculture.

KEY WORDS: *Polydora onagawaensis* sp. nov. · Spionidae · Polychaeta · Reproduction · Morphology · Life history · Mollusk aquaculture · 18S rRNA gene

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INTRODUCTION

The species of *Polydora* and related genera (Polychaeta: Spionidae), also known as polydorids, have a characteristically modified fifth chaetiger (Blake 1996). Many of them are widely known as borers, as they bore holes into mollusk shells (Blake & Evans 1972). Polydorid infestation may cause depreciation in commercial value, reduction of growth rate and meat yield, and heavy mortality of commercially important mollusks. Commercially important mollusk species affected by polydorid infestation include scallops (e.g. Sato-Okoshi 1994, 1999, Radashevsky & Olivares 2005), oysters (e.g. Handley & Bergquist 1997, Sato-Okoshi 1999, Sato-Okoshi & Takatsuka 2001), pearl oysters (Mohammad 1972, Sato-Okoshi 1999), abalone (Sato-Okoshi 1999, Radashevsky &

Olivares 2005, Sato-Okoshi et al. 2008), mussels (Kent 1979, 1981, Ambariyanto & Seed 1991) and clams (Caceres-Martinez et al. 1999). Measures to prevent or control such infestation of cultured mollusks have been suggested (Handley 1993, 1997, 2002, Diggles et al. 2002, Simon et al. 2010), but the problem often remains unsolved. In particular, heavy polydorid infestations require the identification of both the species and its infestation process before attempts can be made to reduce the impact on mollusk aquaculture (Leonart et al. 2003, Dunphy et al. 2005, Simon et al. 2010).

Among polydorids, there are distinct species with similar morphological characteristics, while others show high intraspecific morphological variation. Morphological, biological and ecological approaches are mainly used to identify polydorid species, along with

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differences in methyl green staining (Read 2010) and information from allozyme analyses (Manchenko & Radashevsky 1993, 1998, 2002, Mustaquim 1988, Radashevsky & Pankova 2006). A molecular biological approach has been suggested in order to distinguish the species with closely similar morphology, but there have been few such studies in the Spionidae, particularly in polydorids (Bastrop et al. 1998, Rice et al. 2008).

The recent increase in mollusk aquaculture has resulted in the worldwide distribution of certain commercially important mollusks (Cohen & Carlton 1998). Consequently, some associated organisms, such as sessile and boring species, have also been distributed artificially and unintentionally outside their native regions (Elton 1958, Carlton 1975, Bailey-Brock 2000, Sato-Okoshi et al. 2008). Oysters, for instance, have been introduced globally to 73 countries and are vectors for many non-native species, including disease-causing organisms (Ruesink et al. 2005). Ecological disturbances caused by invasive organisms (including polychaetes) associated with these mollusks are currently seen as a major cause of biodiversity losses worldwide (Mack et al. 2000, Miura 2007).

In the present study, specimens of *Polydora* sp. were found for the first time in shells of the scallop *Patinopecten yessoensis* Jay, 1857 cultured to reach a commercial size in Onagawa Bay (northeastern Japan), but originally produced and cultured in Obira (Sea of Japan, coast of Hokkaido, Japan) as seed, a system typically seen in Japanese mollusk aquaculture. Here we describe this species of *Polydora*, which corresponds in part to *Polydora* sp. previously reported from Japan (Sato-Okoshi 1999), as a new species, with a focus on its morphological variability, molecular data, biology and ecology.

MATERIALS AND METHODS

Specimen collection

Polydora sp. were collected from suspended shells of the cultured scallop *Patinopecten yessoensis*, the wild oyster *Crassostrea gigas* (Thunberg, 1793) and the wild turban snail *Omphalius rusticus* (Gmelin, 1791) at 3 sites (Onagawa Bay, Gobu-ura, and Sasuhama, in Miyagi Prefecture, northeastern Japan; Fig. 1). The morphological characteristics were observed under a Leica MZ 9.5 stereo-microscope, both of live specimens and after fixation in 10% neutral formalin solution. The type specimens are deposited in the National Museum of Nature and Science, Tokyo (NSMT).

DNA extraction, PCR amplification, sequencing and phylogenetic analysis

Six individuals were chosen for molecular sequence analysis (Table 1, Fig. 2). Genomic DNA was directly extracted from live specimens. All animal tissues were washed several times in filtered (pore size 0.2 μm) seawater and distilled water to remove as much extraneous matter as possible. PCR tubes (0.2 ml), each containing 50 μl of 10% Chelex suspension (Bio-Rad Laboratories) and animal tissues were heated at 95°C for 20 min to extract genomic DNA according to Richlen & Barber (2005). Extracted DNAs were used as templates to amplify the target regions. All PCRs were performed on a thermal cycler in a reaction mixture (25.0 μl) containing 1.0 μl of template DNA, 0.2 mM of each deoxynucleotide triphosphate (dNTP), 1 \times PCR buffer, 2.0 mM MgSO_4 , 0.4 U of KOD-Plus-ver. 2 DNA polymerase (Toyobo; with intensive 3'→5' exonuclease activity) and 0.2 μM of each primer. Three primer pairs (18S-1F1/18S-1R632, 18S-2F576/18S-2R1209 and 18S-3F1129/18S-R1772) were used to amplify the nuclear 18S rRNA gene of specimens of the new species described here (Table 2). The PCR cycling conditions were as follows: initial denaturation at 94°C for 2 min; 38 cycles at 94°C for 15 s, 54°C for 30 s, and 68°C for 45 s. Results of PCR amplification were checked on 1.5% agarose gels, with ethidium bromide staining. To remove unincorporated PCR primers and dNTPs, 1.5 μl of the PCR products were then treated with 0.6 μl exonuclease I and shrimp alkaline phosphatase (Exo SAP-IT; USB) at 37°C for 15 min, followed by incubation at 80°C for 15 min to inactivate the enzymes. The PCR products were sequenced directly with an automated DNA sequencer (ABI 3100;

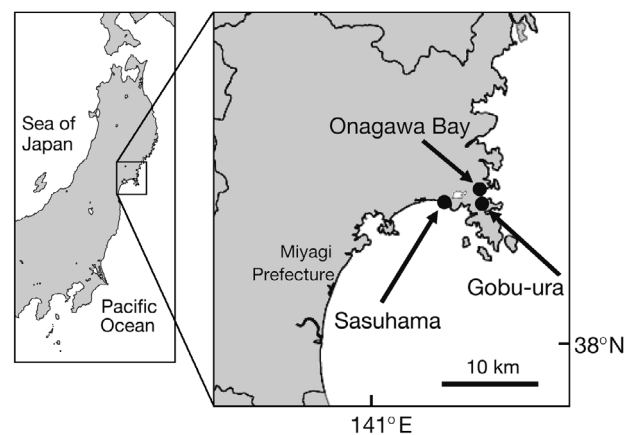


Fig. 1. Sampling sites in Onagawa Bay, Sasuhama, Gobu-ura and Obira, the locality at which the seed scallops were originally cultivated

Table 1. Pigmentation characteristics of individuals of *Polydora onagawaensis* sp. nov. applied for molecular analysis of 18S rRNA gene. The superscripted 'c' and 'w' indicate cultured and wild specimens, respectively

Individual	Host shell	Locality	Palps	Pigmentation Body	Pygidium	Figures
PO-1	<i>Patinopecten yessoensis</i> ^c	Onagawa Bay	Transparent	Transparent	Semi-transparent	Fig. 2A
PO-2	<i>Crassostrea gigas</i> ^w	Onagawa Bay	Transparent	Partially black	Transparent	Fig. 2B–D
PO-3	<i>Crassostrea gigas</i> ^w	Gobu-ura	Partially black	Black all over	Black all over	Fig. 2E–G
PO-4	<i>Crassostrea gigas</i> ^w	Gobu-ura	Transparent	Transparent	Partially black	Fig. 2H
PO-5	<i>Omphalius rusticus</i> ^w	Sasuhama	Black along edge	Black all over	Black all over	
PO-6	<i>Omphalius rusticus</i> ^w	Sasuhama	Transparent	Transparent	Transparent	



Fig. 2. *Polydora onagawaensis* sp. nov. (A) Living worm with no black pigmentation. (B–D) Living worm with black pigmented anterior chaetigers: (B) whole body; (C) anterior body, dorsal view; (D) posterior end and pygidium, lateral view. (E–G) Living worm with entire body black, strongly pigmented: (E) whole body; (F) anterior body, dorso-lateral view; (G) posterior end and pygidium, dorso-lateral view. (H) Living worm with black pigmented pygidium. (I) Burrows excavated by these individuals seen from inner surface of shell. Scale bars: A, I = 1 mm; B, E, H = 600 µm; C, F = 300 µm; D, G = 100 µm

Applied Biosystems). The forward and reverse complementary sequences were combined and 3 nucleotide sequences obtained with the 3 primer pairs were aligned using GENETYX software (Genetyx). Sequences of the nuclear 18S rRNA gene were aligned

with the sequences of other related species obtained from GenBank using the Clustal W algorithm (Thompson et al. 1994).

Maximum-likelihood analyses were performed with PhyML3.0 software (Guindon & Gascuel 2003, Guin-

Table 2. Details of the primers used in this study. Target region is nuclear 18S rDNA. All primers refer to Nishitani et al. (2012)

Primer name	Sequence (5'–3')	Annealing site (nt)
18S-1F1 ^a	AACCTGGTTGATYCTGCCAG	1–20
18S-1R632 ^a	ACTACGAGCTTTTAAACYGCARC	610–632
18S-2F576 ^a	GGTAATTCCAGCTCYAATRG	576–595
18S-2R1209 ^a	AAGTTTTYCCCGTGTTGARTC	1190–1209
18S-3F1129 ^a	GCTGAAACTTAAAGRAATTGACGGA	1129–1153
18S-1R1772 ^b	TCACCTACGGAAACCTTGTTACG	1749–1772
^a <i>Dinophysis acuta</i> (AJ506973); ^b <i>Chlamydomonas reinhardtii</i> (M32703)		

don et al. 2010) using an input tree generated by BIONJ with a general time-reversal model (Rodríguez et al. 1990) that incorporated invariable sites and a discrete gamma distribution (8 categories) (GTR+I+Γ). PhyML bootstrap trees (1000 replicates) were constructed using the same parameters as the individual maximum-likelihood trees. Posterior probabilities of Bayesian trees were also estimated using MrBayes 3.1.1 (Huelsenbeck et al. 2001, Ronquist & Huelsenbeck 2003) under GTR+I+Γ. One cold and 3 heated Markov chain Monte Carlo (MCMC) chains were run for 500 000 generations to sample log-likelihoods and trees at 100-generation intervals (5000 trees were saved during MCMC with a burn-in of 1250 trees).

Biological and ecological analysis

Seven to sixteen 3- to 4-year-old cultured scallops originating from Obira (Sea of Japan coast of Hokkaido, north Japan), where they were produced and cultured for 18 to 22 mo from seeds, were collected monthly from July 2008 to December 2010 from a depth of 5 to 10 m in Onagawa Bay (Fig. 1). The vertical profile of temperature was determined using a salinity temperature depth recorder at the same time. Neither polydorid worms nor burrows were observed in the scallop shells before they were hung in Onagawa Bay or prior to their use in this study.

After removing the scallop flesh, the shell was broken into small pieces using cutting pliers and the spionids were removed from their burrows with a helve needle and insect pin under a stereo-microscope. The presence of spermatocytes and oocytes was examined *in vivo*. The length of the smallest worm with gametocytes was considered the minimum size at maturity. Therefore, from this length up, worms were considered adults (including those with gametocytes absent, i.e. non-reproducing adults), whereas smaller worms

were considered juveniles. The relationships between width of chaetiger 5 and body length and total number of chaetigers were estimated on the basis of selected complete specimens (N = 162). Width was further measured under a stereo-microscope to estimate body size.

Egg capsules were carefully removed from the burrows with an insect pin and placed in culture plates containing GF/F filter-sterilized seawater at 10°C. Seawater was changed every 2 to 3 d.

RESULTS

Taxonomic account

Family Spionidae Grube, 1850

Genus *Polydora* Bosc, 1802

Polydora onagawaensis sp. nov. (Figs. 2 & 3)

Polydora sp.: Sato-Okoshi (1999), p. 836

Material

Holotype (NSMT-Pol H-572) from the shell of a living cultured scallop *Patinopecten yessoensis*, Onagawa Bay, Miyagi Prefecture (38° 44' N, 141° 46' E), W. Teramoto coll., 22 January 2010; 10 paratypes (NSMT-Pol H-573), same host species, location, collector and date as holotype; 10 paratypes (NSMT-Pol H-574) from shells of the wild oyster *Crassostrea gigas*, Gobu-ura, Miyagi Prefecture (38° 40' N, 141° 47' E), W. Teramoto coll., 6 June 2012; 10 paratypes (NSMT-Pol H-575) from shells of the wild turban snail *Omphalius rusticus*, Sasuhama, Miyagi Prefecture (38° 41' N, 141° 37' E), W. Teramoto coll., 6 June 2012.

Diagnosis

A mid-sized *Polydora* with modified chaetiger 5 possessing thickened falcate spines with lateral tooth, accompanying pennoned chaetae; with dorsal and ventral tufts of winged capillary chaetae. High variation in pigmentation: absent or black pigment on palps, prostomium, peristomium, body and pygidium; no black-bar pigment on palps. Prostomium weakly incised; caruncle extended back to chaetiger 4. Peristomium small and short. Branchiae from chaetiger 7 and continue to middle or almost end of body. Posterior chaetigers without modified noto-chaetae. Pygidium disc-like, with dorsal gap.

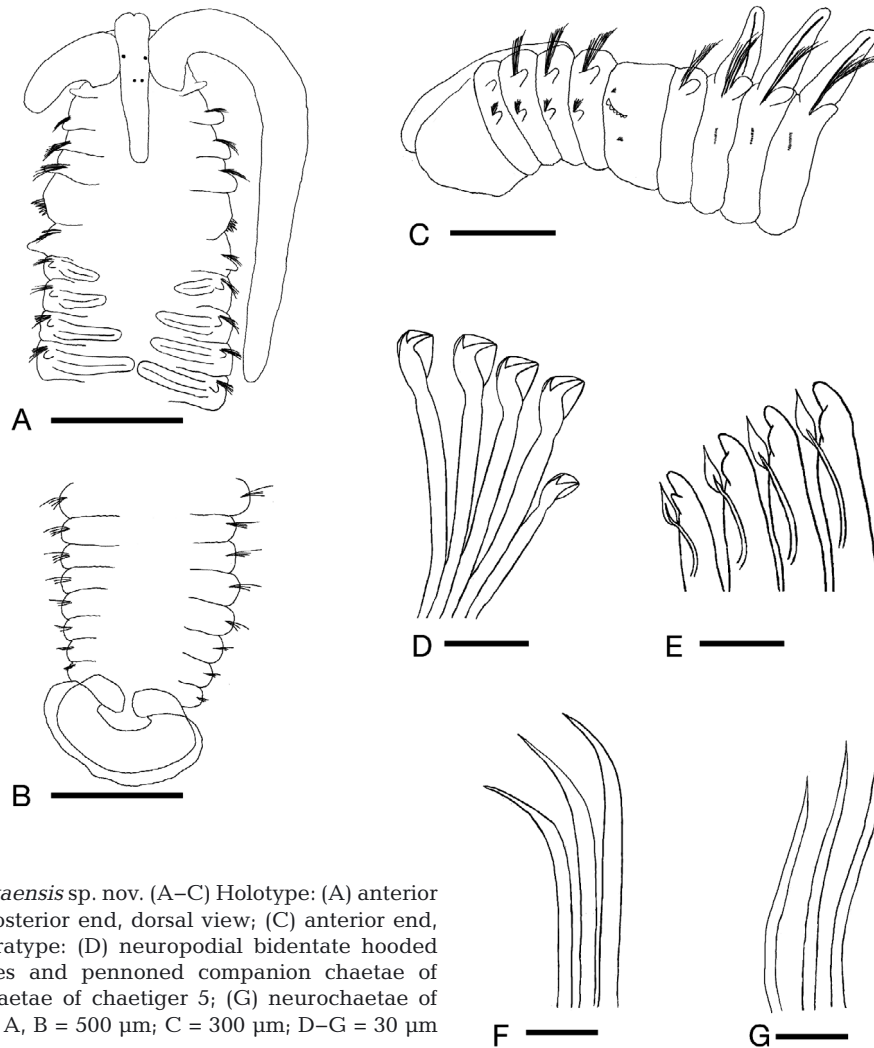


Fig. 3. *Polydora onagawaensis* sp. nov. (A–C) Holotype: (A) anterior end, dorsal view; (B) posterior end, dorsal view; (C) anterior end, lateral view. (D–G) Paratype: (D) neuropodial bidentate hooded hooks; (E) major spines and pennoned companion chaetae of chaetiger 5; (F) notochaetae of chaetiger 5; (G) neurochaetae of chaetiger 5. Scale bars: A, B = 500 µm; C = 300 µm; D–G = 30 µm

Holotype

Length 14.5 mm, 0.7 mm wide at chaetiger 5, with 86 chaetigers. Color light tan without conspicuous pigmentation. Prostomium weakly incised anteriorly and continued as caruncle to end of chaetiger 3 (Fig. 3A). No occipital tentacle. Palps transparent. Four eyes present, arranged trapezoidally.

Chaetiger 1 with short notopodial lobe, with neurochaetae only (Fig. 3C). Winged capillary notochaetae arranged in 2 rows from chaetigers 2–4, 6 and succeeding ones (Fig. 3F). No special notochaetae in posterior chaetigers. Chaetigers 2–4 and 6 with winged capillary neurochaetae arranged in 2 rows; from chaetiger 7 (Fig. 3G), replaced by bidentate hooded hooks without accompanying capillaries. Up to 8 hooded hooks with constriction on shaft, main fang at right angle to shaft and at acute angle with apical tooth (Fig. 3D).

Chaetiger 5 modified, with 5 major spines, falcate, with lateral tooth or flange, alternating with pen-

noned companion chaetae (Fig. 3E). Dorsal and ventral winged capillary chaetae present.

Branchiae from chaetiger 7, attaining full size at chaetiger 10, gradually diminishing in posterior chaetigers, absent from last 10 chaetigers (Fig. 3B).

Pygidium disc-like with dorsal gap (Fig. 3B).

Variability

Mid-sized worms measuring up to 16.2 mm long and 1.1 mm wide at chaetiger 5, with up to 111 chaetigers. Color in life light tan, with variable black pigmentation on whole body, or partially on palps (Fig. 2). Palps transparent or brown to black on and along margin of palp edge. Dark to un-pigmented on prostomium, peristomium, along both sides of prostomium, and on both dorsal and ventral sides of chaetigers through anterior to posterior. Pygidium with or without black pigmentation. Occurrence of

worms with slight to strong black pigmentation fluctuated throughout the year; less than half of the total number of individuals examined (9533) during all years were pigmented.

Caruncle extension size-dependent, reaching posterior end of chaetiger 2 to 4. Eyes absent or present; when present up to 4, trapezoidal in arrangement. Branchiae from chaetiger 7, reaching full size from chaetiger 10 or posteriorly, then decreasing in size and continuing to half of body to near the posterior end.

Modified spines of chaetiger 5 falcate, with lateral tooth or sheath, numbering 5 to 8 per row plus approximately 2 spines buried inside, alternating with slender, pennoned accompanying chaetae. Winged capillary noto- and neurochaetae present on chaetiger 5, ca. 2 to 4 antero-dorsal and ca. 5 to 7 postero-ventral, respectively. Bidentate hooded hooks numbering from 5 to 10 per row up to the mid-body, then decreasing in number to only 1 in posterior chaetigers.

Pygidium disc-like with dorsal gap, with or without black pigmentation; if present, partially black along margin or all over.

Remarks

Polydora sp. (Sato-Okoshi 1999) shows a highly variable black pigmentation both on palps and body. Some of the specimens did not possess distinct black bars on palps, but had various degrees of black pigmentation on part or all of the body, or lacked pigmentation. These specimens are here considered as belonging to *P. onagawaensis* sp. nov. *Polydora* sp. measured up to 16 mm long, had eggs of 80–100 µm in diameter, and the early 3-chaetiger larva reached up to 200–225 µm long, characteristics that also agree with *P. onagawaensis* sp. nov. (see 'Reproduction and development' below). Moreover, the geographical location for *Polydora* sp. includes that for *P. onagawaensis* sp. nov.

Etymology

The specific name *onagawaensis* refers to Onagawa Bay, the type locality of the new species.

Molecular biological analysis

Morphological variation, particularly presence or absence of pigmentation, and degree of pigmentation, was observed both within and between sampling

sites and/or host shells (Table 1, Fig. 2). However, the nuclear 18S rRNA gene sequences (1771 bp) were identical in all analyzed individuals of *P. onagawaensis* sp. nov. (GenBank accession number AB691768).

Species belonging to the genus *Polydora*, excluding *Polydora ciliata*, formed a generic group (Fig. 4). *Polydora onagawaensis* sp. nov. was arranged closely to *Polydora calcarea*.

Habitat and ecology

Polydora onagawaensis sp. nov. was found in *Patinopecten yessoensis*, *Crassostrea gigas* and *Omphalius rusticus* in Onagawa Bay. *Polydora onagawaensis* sp. nov. forms U-shaped burrows that never branch (Fig. 2I). The presence of up to 115 individuals in a single shell often caused the host scallop to secrete abnormal brown or black shell on its inner surface. The polychaetes were most often found to bore weakly in a concentric pattern, around annual rings, and to aggregate in the ear of the scallop shell (Fig. 5). The mean number of individuals per shell tended to increase from December to March, and then decrease towards November (Fig. 6). Moreover, the infestation was significantly higher in the right than in the left valve during the 2 yr of the study ($t_{220} = 10.7$, $p < 0.01$; Fig. 6).

Reproduction and development

The spermatozoa of *Polydora onagawaensis* sp. nov. observed in June 2011 had an elongated head, a head and middle region of ca. 10 µm long, and a tail of ca. 50 µm long.

Females deposited up to 49 egg capsules in a single string, attached to the burrow wall, each of them usually containing 60 to 70 eggs, with major and minor axes ca. 110 and 100 µm, respectively. Early eggs were light yellow/orange in color and developed at 10°C to the 1-chaetiger stage after 2–8 d (mean 4.5 d), the 1- to 2-chaetiger stage after an additional 3–5 d (mean 3.5 d), and the 2- to 3-chaetiger stage (ready to hatch) after a further 2–8 d (mean 4.2 d). All eggs developed simultaneously; no unfertilized or nurse eggs were found. From the 1- to the 3-chaetiger stage, the larvae contained yolk, which appeared partially depleted, in the center of the body. The larvae had 3 sets of eyespots, with the 2 lateral pairs gradually joining during the ontogeny.

The 3-chaetiger larvae measured 200 to 250 µm long, had only the lateral and median eyespots, con-

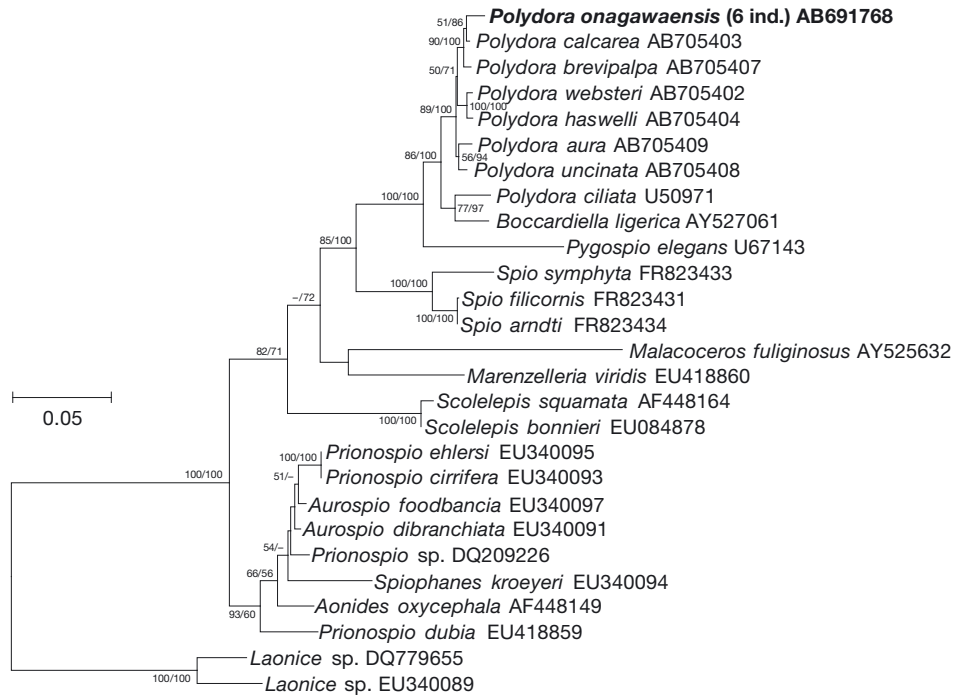


Fig. 4. Maximum likelihood tree inferred from the nuclear 18S rRNA gene sequences of spionid polychaetes. The sequence of *Polydora onagawaensis* sp. nov. is in bold. Left: bootstrap support; right: posterior probability; -: values < 50. Scale bar: number of substitutions per site

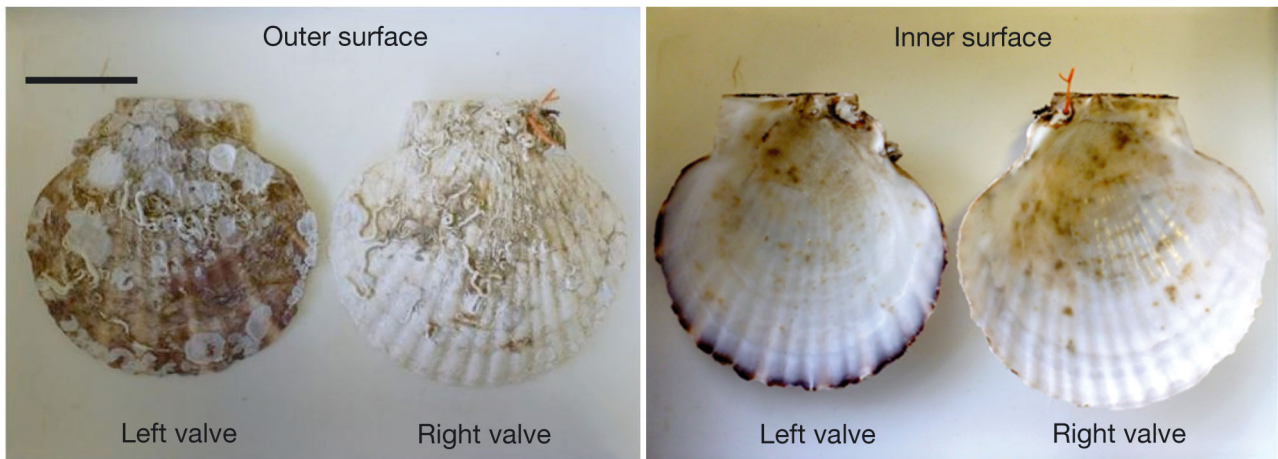


Fig. 5. Outer (A) and inner (B) surface of cultured scallop shells infested by *Polydora onagawaensis* sp. nov. Scale bar: 50 mm

tained little or no yolk and pigmentation was absent. They were able to feed and swim immediately after hatching. Attempts to culture larvae beyond the 3-chaetiger stage were unsuccessful.

Brooding worms were present from around October to June, with a peak in December each year and 2 minor increases in March and May. Spawning was rarely observed during summer (July to September; Fig. 7). The smallest worm with gametocytes was 400 μ m wide.

Population dynamics

The width of chaetiger 5 is positively correlated with both body length and total number of chaetigers (Fig. 8). Males, females, juveniles and non-reproducing adults have the same size relationship. The analysis of the size–frequency distributions of *Polydora onagawaensis* sp. nov. (Fig. 9) shows that the number of worms larger than 600 μ m wide tended to decrease from July to October during the first and

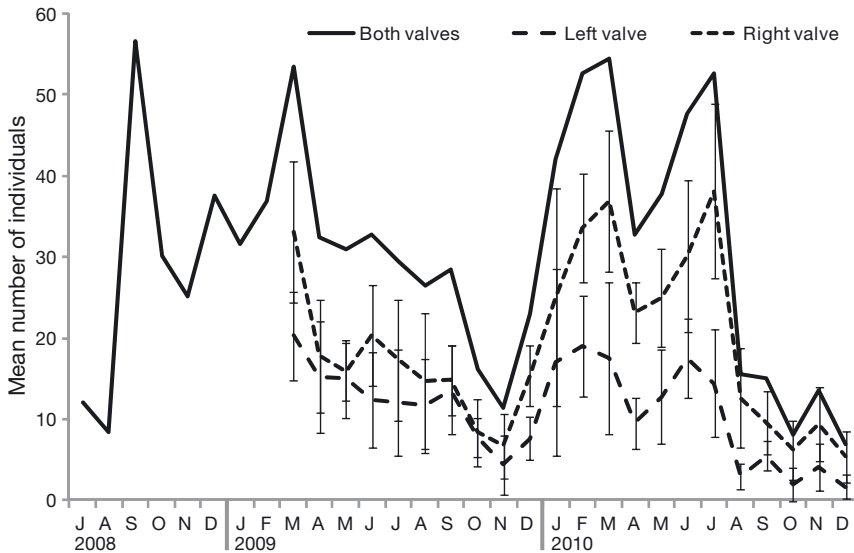


Fig. 6. *Polydora onagawaensis* sp. nov. Mean (\pm SD) number of worms extracted per shell from July 2008 to December 2010

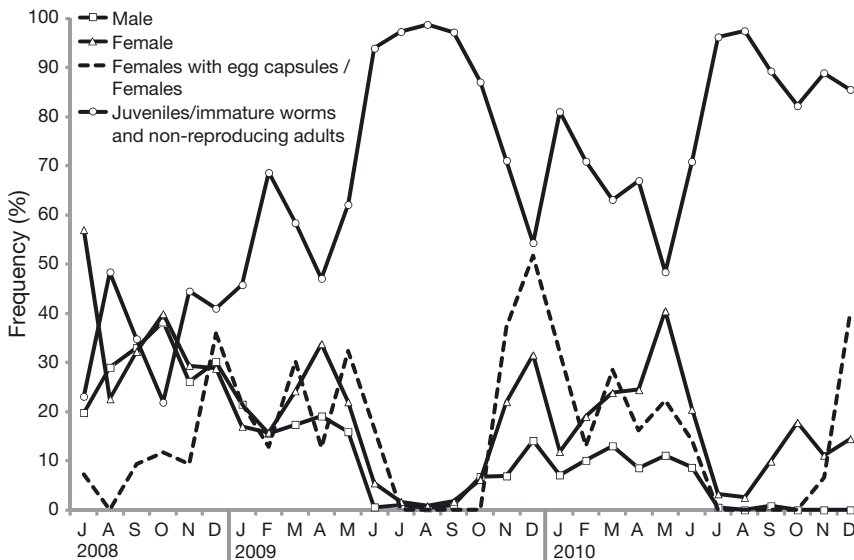


Fig. 7. *Polydora onagawaensis* sp. nov. Frequencies of sexually mature worms, females with egg capsules, and juveniles/immature worms and non-reproducing adults from July 2008 to December 2010

second years only, while the number of mature worms 450–600 μ m wide tended to increase from October to December each year. From January to March (and particularly in February), the number of juveniles smaller than 400 μ m wide tended to increase and the number of mature worms tended to decrease.

DISCUSSION

Taxonomy

Walker (2011) included 147 species within the 9 currently recognized genera of the *Polydora* complex worldwide: *Polydora* Bosc, 1802 (44 species); *Dipolydora* Verrill, 1879 (36); *Pseudopolydora* Czerniavsky, 1881 (18); *Boccardia* Carazzi, 1895 (21); *Polydorella* Augener, 1914 (5); *Tripolydora* Woodwick, 1964 (1); *Boccardiella* Blake & Kudenov, 1978 (7); *Carazziella* Blake & Kudenov, 1978 (13); and *Amphipolydora* Paterson & Gibson, 2003 (2). In Japan, 26 polydorid species have been described, including 12 belonging to *Polydora*: *P. brevipalpa*, *P. ciliata*, *P. curiosa*, *P. flava orientalis*, *P. websteri*, *P. aura*, *P. uncinata*, *P. glycymerica*, *P. cornuta*, *P. haswelli*, *P. calcarea* and *Polydora* sp. (Sato-Okoshi 1999, 2000; Sato-Okoshi & Abe 2012, in press). However, based on our surveys to date, there may be several different species included within this *Polydora* sp. Only some individuals of *Polydora* sp. that lack black bars on the palps correspond to the new species here described, presently only known from Miyagi (Pacific coast of Japan).

Polydora onagawaensis sp. nov. resembles *P. haswelli* Blake & Kudenov and *P. neoacaeca* Williams & Radashkevsky in many major morphological characters. Both possess black pigmentation on the palps, dorso-laterally on the prostomium, on the dorsal and ventral peristomium, and on both the dorsal and ventral sides of the anterior chaetigers; more than 80% of the specimens of *P. neoacaeca* have black pigmentation on the anterior body (Williams & Radashkevsky 1999). Radashkevsky et al. (2006) described *P. cf.*

haswelli from Brazil with variable pigmentation on the palps and anterior chaetigers. Sato-Okoshi & Abe (in press) demonstrated a variable degree of pigmentation on the palps, peristomium, prostomium and anterior chaetigers of *P. haswelli* from Japan. However, the 2 previous species, *P. haswelli* and *P. neoacaeca*, have distinct black bars on the palps, more

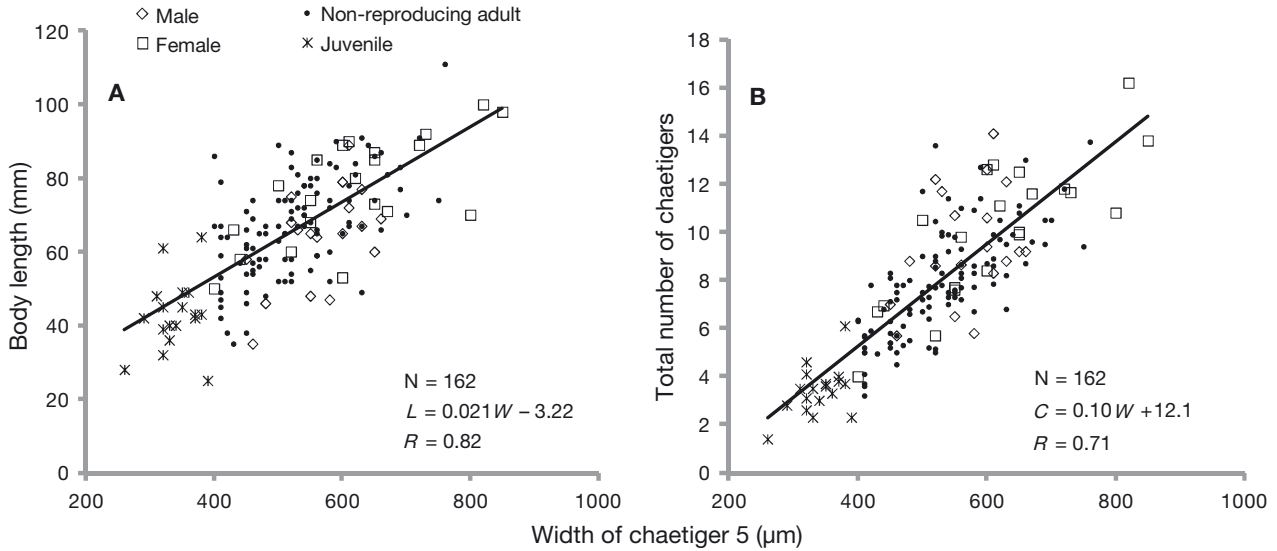


Fig. 8. *Polydora onagawaensis* sp. nov. Relationship between width of chaetiger 5 (W) and (A) body length (L) and (B) total number of chaetigers (C)

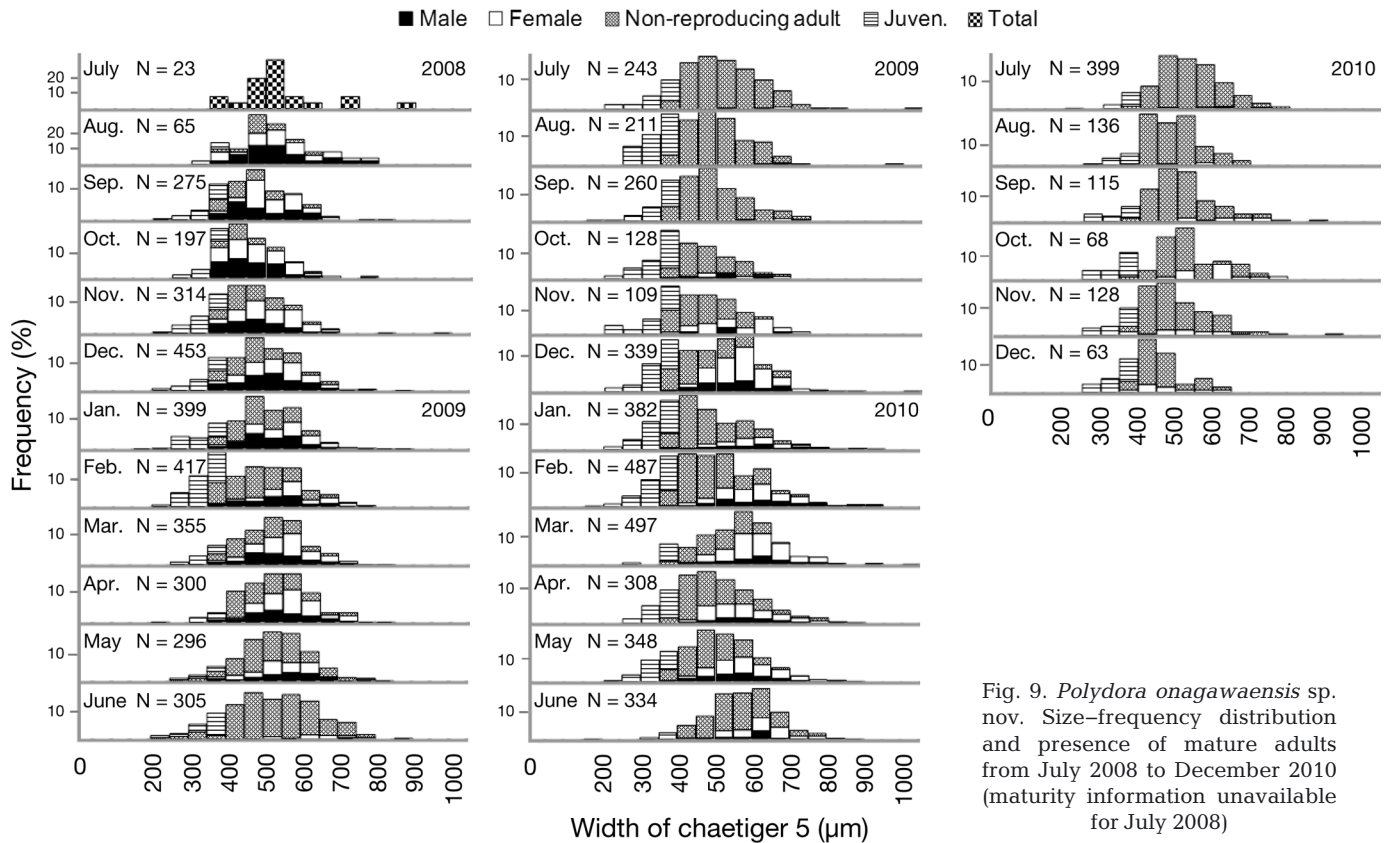


Fig. 9. *Polydora onagawaensis* sp. nov. Size-frequency distribution and presence of mature adults from July 2008 to December 2010 (maturity information unavailable for July 2008)

apparent black pigmentation with restricted distribution in part of the body (e.g. dorsal anterior chaetigers 1–4), and more slender bodies. In contrast, a low frequency of *P. onagawaensis* sp. nov. specimens have pigmentation over the whole body, including the palps (less than 50%). If pigmented on the palps,

some specimens have black pigmentation longitudinally along the edge of the palps, while others have diffuse black color all over the palps, but none have black bars. Moreover, *P. onagawaensis* sp. nov. shows variation in black pigmentation over the whole body, including the posterior region. *Polydora neocaeca* and

P. haswelli lack black pigmentation of posterior chaetigers and pygidium (Blake & Kudenov 1978, Williams & Radashevsky 1999, Sato-Okoshi & Abe in press), while the Brazilian specimens of *P. cf. haswelli* have white pygidia and yellow patches in the posterior region (Radashevsky et al. 2006). The largest specimen of *P. neocaeca* (originally from the east coast of North America) measured 29.5 mm long and 0.7 mm wide (at chaetiger 7) and had 133 chaetigers (Williams & Radashevsky 1999), whereas in *P. onagawaensis* sp. nov. the largest specimen measured 16.2 mm long and 1.1 mm wide at chaetiger 5 and had 111 chaetigers. *Polydora haswelli* has been reported as similar in size to *P. neocaeca* from New Zealand (Read 2010) and Japan (Sato-Okoshi & Abe in press), and as larger in size from South Korea (Sato-Okoshi et al. 2012). *P. onagawaensis* sp. nov. is smaller and wider in body shape.

Polydora calcarea (Templeton, 1836) most closely resembles the new species in pigmentation pattern and it is difficult to distinguish the two from their morphology. However, *P. calcarea* can be distinguished by having fewer branchial chaetigers, continuing until one-half of the body in specimens from Japan and Australia (Sato-Okoshi & Abe in press) and one-half to two-thirds of the body in specimens from Russia (cf. Radashevsky & Pankova 2006). In addition, the caruncle extends to the end of chaetiger 2 in *P. calcarea*, while it extends to the end of chaetiger 4 in the new species, despite the larger size in *P. calcarea*. *Polydora websteri* Hartman, in Loosanoff & Eagle, 1943 also typically shows black pigmentation along the palps, but no other black pigmentation (see Read 2010 for a review of variation in its pigmentation; Sato-Okoshi & Abe in press), a longer peristomium and fairly long branchiae continuing until almost the end of the body (Sato-Okoshi & Abe in press). Careful morphological observation is essential for accurate species discrimination.

As for egg capsules, *Polydora onagawaensis* sp. nov. has up to 49 capsules per egg string, with 10–83 eggs per capsule, with major and minor axes of 110 and 100 μm mean diameter, respectively, and the 3-chaetiger larvae are 200–250 μm long. In contrast, *Polydora neocaeca* has 13–24 per egg string, with 8–47 eggs per capsule, an average egg diameter of $116.2 \pm 10.0 \mu\text{m}$, and the early and late 3-chaetiger larvae are 240 and 340 μm long, respectively. The egg diameter of *P. cf. haswelli* from Brazil measured ca. 100 μm (Radashevsky et al. 2006). These differences are considered robust enough to support the description of the Japanese species as new.

As stated above, the well-known morphological variability of polydorid species often leads to difficulties in their morphological description and classification. To date, molecular sequences for several genes have been published for only 13 of the 147 polydorid species in GenBank: *Polydora brevipalpa*, *P. uncinata*, *P. aura*, *P. websteri*, *P. calcarea*, *P. haswelli*, *P. triglanda*, *P. ciliata*, *P. cornuta*, *P. giardi*, *Dipolydora carunculata*, *Boccardia proboscidea* and *Boccardiella ligerica* (Walker 2011, Sato-Okoshi & Abe 2012, in press), with the data for the 18S rRNA gene sequences only being available for *P. brevipalpa*, *P. uncinata* and *P. aura* (Sato-Okoshi & Abe 2012) and *P. websteri*, *P. calcarea* and *P. haswelli* (Sato-Okoshi & Abe in press) from Japan and Australia, *P. triglanda* from Taiwan (V. V. Pankova & V. I. Radashevsky unpubl.), *P. ciliata* (source locality unstated) (S. Nadot & A. Grant unpubl.), *P. giardi* from France (Rousset et al. 2004), *Dipolydora carunculata* from Russia (V. V. Pankova & V. I. Radashevsky unpubl.) and *Boccardiella ligerica* from Germany (Struck & Purschke 2005). Owing to the short lengths of their partial sequence data, *P. giardi*, *P. triglanda* and *D. carunculata* were not included in the phylogenetic tree (Fig. 4).

Although *Polydora onagawaensis* sp. nov. closely resembles *P. calcarea* and *P. haswelli* in terms of morphology, the 3 species can be sharply distinguished by the 18S rRNA gene analysis. Molecular biological approaches can provide a key tool for the accurate discrimination of polydorid species, and for elucidating genealogical relationships among the taxa.

Excavating pattern

Polydora onagawaensis sp. nov. preferably infested right valves of the suspended cultured scallop shells in Onagawa Bay (Fig. 6). In general, however, sown cultured scallop shells examined were rarely or never infested in the right valve (Sato-Okoshi & Nomura 1990). Obviously, right valves of scallops sown on the sea bottom might seldom have the chance of being colonized by settling polydorid larvae, as these valves are usually the lower ones and are buried in the bottom. By contrast, both valves are exposed to colonization in suspended scallops, which, in Onagawa Bay, are arranged with the left valves facing out and the right ones facing in (Fig. 10). Thus, left valves might be more exposed to currents, while the right ones may offer a more sheltered settling environment to planktonic larvae. Further experiments must be addressed to test this

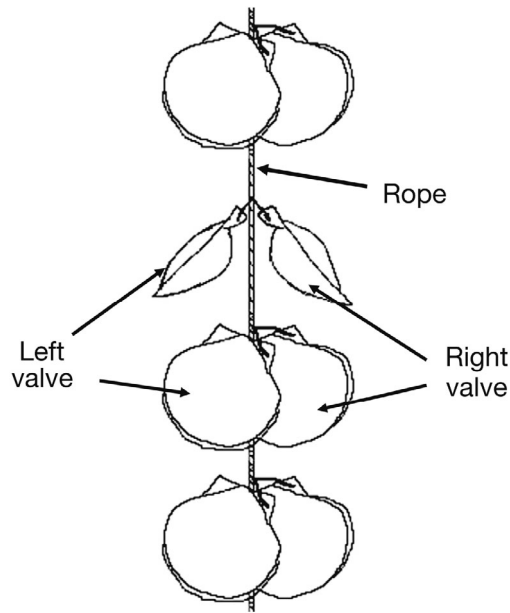


Fig. 10. Illustration of a suspended scallop culture in Onagawa Bay. Scallops are attached to the rope with ears, with the left and right valves outwards and inwards, respectively

hypothesis, which may have relevant implications for larval settlement behavior.

Dead sown and suspended cultured scallops also differ in their infestation pattern. Living *Polydora* specimens have seldom been extracted from dead shells of wild and sown scallops (Sato-Okoshi 1994), while in suspended scallops, dead shells contain approximately the same number of *Polydora* as living shells. It is likely that dead shells of wild and sown scallops may be either transported to unsuitable areas or buried in mud, thus negatively affecting the suspension-feeding activity of *Polydora*. We strongly suggest that the worms are capable of surviving in dead suspended cultured scallop shells since, in our study, the environmental conditions did not change in comparison with those of the living shells (i.e. the shells were not buried in the mud).

The mean number of worms extracted per shell tended to increase towards March with increasing numbers of recruits, and then decreased towards October as the number of large worms decreased. Exceptionally, the number of worms decreased in July to August 2010. In August 2010, many barnacles were attached to the scallop shells because of the continuous heat wave. The settlement by barnacles may have inhibited polydorid feeding, leading to their eventual death. Data in July and August 2008 may be biased as only a few individual worms were examined.

Reproduction, development and life history

Many papers describe the reproduction of *Polydora* species (reviewed by Blake 2006). In general, mature females deposit egg capsules in their burrows, where the eggs develop until they are released as planktonic larvae. The larvae further develop, settle, metamorphose and may bore in mollusk shells or other calcareous substrates (Dorsett 1961, Daro & Polk 1973, Gudmundsson 1985, Zajac 1992, Sato-Okoshi 1994, Williams 2001). Direct development is also known for some species (Blake 1996, Sato-Okoshi et al. 2008). Also, *Polydora* species are most likely polytelic (i.e. reproducing more than once in a season) and show a high fecundity (Gudmundsson 1985, Sato-Okoshi et al. 1990, Blake & Arnofsky 1999, Williams 2001, Radashevsky et al. 2006). For example, *Polydora robi* is a rapid breeder, producing successive broods within less than 7 d; the eggs develop (~6 d) until they are released at the 3-chaetiger stage, when the females spawn again (Williams 2001). In *Polydora onagawaensis* sp. nov., brooding worms were present from around October (autumn) to June (early summer), with a peak in December (Fig. 7). Brooding females already contained the next generation of oocytes, proving the species to be polytelic.

Temperature affects polychaete breeding patterns (Schroeder & Hermans 1975; Olive & Clark 1978); for instance, the reproductive activity of *Streblospio benedicti* Webster, 1879 was shown to decrease during the fall, when water temperatures dropped below 12.5°C (Levin & Creed 1986), while *Marenzelleria viridis* (as *Scolecopides viridis*) Verrill, 1873 first spawned after the water temperature dropped below 15°C (Bochert & Bick 1995). During the study period, mean water temperature ranged from 6.3 to 22.7°C at 5–10 m depth in Onagawa Bay, with the lowest temperatures in April and the highest in August or September (Fig. 11). The seawater temperature was below ca. 15°C during the entire spawning period (Fig. 11), so this may be considered the upper temperature limit supporting spawning for *Polydora onagawaensis* sp. nov.

Many spionids tend to reproduce during periods of high temperature (Blake 1969, Levin 1984, Levin & Creed 1986, Sato-Okoshi et al. 1990). In Japan, there are various reports that *Polydora brevipalpa* mainly spawns from August to October (seawater temperature ca. 16–17°C), *P. websteri* from May to June and October to November (ca. 6 and 7°C, respectively), *P. curiosa* from April to June, *P. uncinata* in October, *P. aura* in September and *Polydora* sp. from April to June Sato-Okoshi 1999. In contrast, the spawning

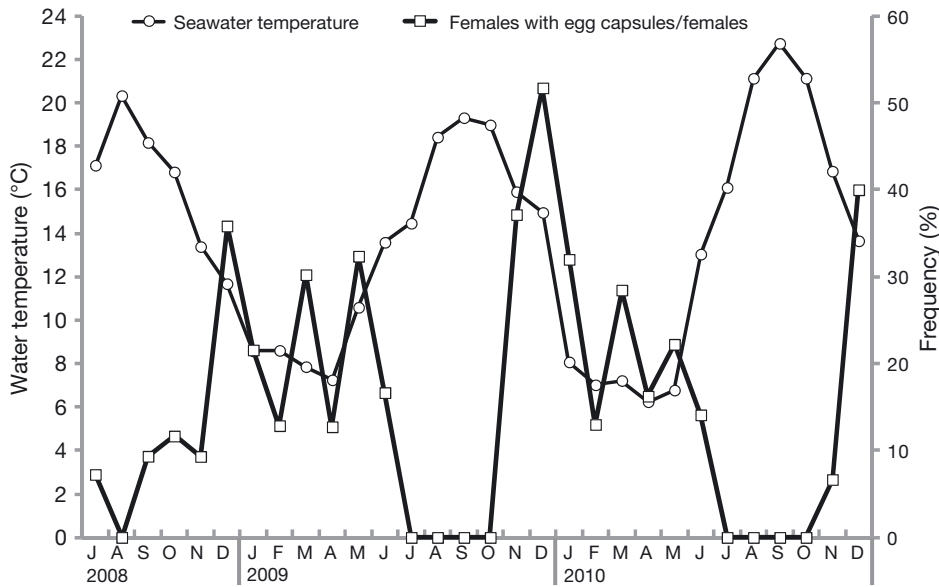


Fig. 11. *Polydora onagawaensis* sp. nov. Frequency of females with egg capsules and mean seawater temperature at 5–10 m depth in Onagawa Bay throughout the study period

peak of *Polydora onagawaensis* sp. nov. occurred in December. It is interesting to note that a large number of planktonic larvae of the genus *Polydora* were observed from December to February (winter, water temperature from 14 to 7°C) in Onagawa Bay, which seemed coincident with the forthcoming spring phytoplankton bloom (January to March/April). Conversely, most planktonic polychaete larvae, excluding *Polydora*, tended to synchronize with summer phytoplankton increases (July or August) and autumn blooms (September to October) in the near-surface water in Onagawa Bay (Abe et al. 2011). Therefore, we postulate that this segregation may consequently allow the timing of *Polydora* larvae release to coincide with the spring bloom, in contrast to other larvae.

Although planktonic larvae of spionid polychaetes abounded in Onagawa Bay, adults in the muddy bottom were not correspondingly numerous (Abe 2010). Therefore, the relationships between these spionid larvae and other *Polydora* species present in the bay, such as the new species described here, require further clarification.

Polydora onagawaensis sp. nov. larvae spent ca. 2 wk developing inside the capsule until hatching at the 3-chaetiger stage under laboratory conditions. However, the planktonic larval period in nature is still undefined. In other *Polydora* species with planktonic larvae, the larval incubation period is 1–2 mo (Woodwick 1960, Dorsett 1961, Hatfield 1965, Day & Blake 1979, Sato-Okoshi 1994). In *Polydora brevipalpa* (as *P. variegata* in Sato-Okoshi 1994), larvae

were planktonic until they reached the 17-chaetiger stage, a process requiring more than 2 mo at 15°C. This long duration may indicate the unsuitable conditions for growing and settling. The attempts to culture larvae of *P. onagawaensis* sp. nov. until settling and metamorphosing were unsuccessful. However, the smallest recruit found 'in situ' was a 25-chaetiger stage. Therefore, like other polydorids, the new species seemed to settle and metamorphose around the 20-chaetiger stage. Further assays with different culturing conditions will be required to determine the conditions adequate to induce larvae of *P. onagawaensis* sp. nov. to settle and metamorphose.

A peak of abundance of recently settled juveniles on scallop shells was not recorded during the study period (Fig. 9). However, we assume that the main recruitment period is from January to March, due to the increase in the mean number of worms extracted per shell and the relatively high presence of juveniles during this period (Figs. 6 & 9). If so, the larvae could start settling on shells approx. 1 mo after spawning in December. However, spawning in some females, and the consequent presence of recruits, occurred continuously throughout the year.

Many species of *Polydora* have a 1-yr life span and one or several spawning events (Söderström 1920, Lambeck & Valentijn 1987). Some species, however, may survive for more than 1 yr in natural conditions, so that they may contribute to the next spawning event (Gudmundsson 1985). Moreover, some *Polydora* species may survive over 2 (Radashevsky et al. 2006) or 2.5 yr in suitable conditions (Sato-Okoshi 1994). The life span of *P. onagawaensis* sp. nov. appears to be approximately 1.5 yr, as settlement occurred in January, continuous spawning occurred from around October to June (with a peak in December), and death is common in August (Figs. 7 & 9).

Origin of *Polydora onagawaensis* sp. nov. and association with its scallop host

The borings of *Polydora onagawaensis* sp. nov. were abundant in the scallop shells in Onagawa

Bay during the study period. However, the previous report on this species (as *Polydora* sp.) did not mention such an infestation (Sato-Okoshi 1999), as it reported infestations in other bivalves (*Crassostrea gigas*, *Ostrea circumpicta*, *Saccostrea kegaki*, *Mytilus edulis*, *Chlamys nobilis*, *Pinctada fucata* and *Anomia chinensis*) and gastropods (*Haliotis discus hannai*, *Batillu cornutus*, *Lunella coronate coreensis* and *Chlorostoma argyrostoma lischkei*). However, different *Polydora* species have been previously reported as borers into the shells of suspended cultured scallops (Sato-Okoshi & Nomura 1990).

During our survey, *Polydora onagawaensis* sp. nov. infested the scallop shells along with the large-sized *P. brevipalpa*, which is common in Hokkaido as well as in the Aomori Prefecture, making this southern population the first record of *P. brevipalpa* for the region. We recorded 25 individuals (1 juvenile and 24 mature adults, including 1 brooding female) found between 2008 and 2010. Despite its rarity, the presence of deposited egg capsules was observed. *Polydora brevipalpa* has been reported as a specific borer of scallop shells in Japan (Sato-Okoshi 1999). Chemical cues driving larval settlement of gregarious marine invertebrates are often related to the presence of conspecifics (Pawlik 1986, Toonen & Pawlik 1996, Bryan et al. 1997), and this may also be the case for *P. brevipalpa*. Considering the small number of individuals, the specific borer *P. brevipalpa* may still be flourishing and might well increase in abundance in the future. Further investigations are needed in order to follow its possible success in the area.

Polydora onagawaensis sp. nov., as well as *P. brevipalpa*, could be non-indigenous, introduced species in Onagawa Bay as they could have come from Obira, in the Sea of Japan off the coast of western Hokkaido, together with the young scallops purchased for culture in the bay over the last several decades. *Polydora onagawaensis* sp. nov. seems better adapted to low temperatures as compared with other native polychaetes from Onagawa Bay, since its reproductive period coincided with seawater temperatures below 15°C (Fig. 11).

There is no information on infestation by *Polydora onagawaensis* sp. nov. (neither for Onagawa Bay nor for Obira) prior to the start of the present study in 2008. Therefore, further studies on possible environmental changes in these regions and on the ecology and habitats of the species are required to assess possible threats to the regional aquaculture industry, as well as to confirm the invasive behavior attributed to a presumably non-indigenous species.

Considering the species as a pest may be an anthropogenic interpretation, as it frequently occurs with other boring species where only the cosmetic commercial value of the host mollusks may be affected (Martin & Britayev 1998). Currently, no direct or obvious damage to the fitness of scallops has been demonstrated, so the species cannot be formally considered as a parasite. Further studies on the physiological implications of the presence of the worms are required to clarify the characteristics of the association.

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