Evaluation of three preparation methods for living diatoms at a sandy river–mouth tidal flat: conventional acid–cleaning, nuclear staining, and sieving

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Abstract: Three preparation methods for the identification of diatom flora at a river mouth tidal flat were evaluated in terms of species richness, diversity, and composition using a conventional acid–cleaning method, a nuclear staining method, and a newly introduced sieving method. Twenty–five diatom samples were collected from Fujimae Tidal Flat sediments, Nagoya, Japan, in July 2014. Three methods respectively showed different aspects of the diatom flora. The acid–cleaning method indicated the highest species diversity, but it extracted dead diatoms with living ones indiscriminately. In comparison with the acid–cleaning method, the staining method effectively extracted diatoms with fragile and small–sized cells. The sieving method scraped the sessile diatoms selectively and supplemented the staining method in which sessile diatoms were mostly neglected. A combination of these three methods enables to describe precise diatom flora of a tidal flat.

Key words: diatom flora, Hematoxylin, nuclear staining method, river mouth tidal flat, sieving method

INTRODUCTION

Benthic diatoms often become dominant at the sediment–water interface where biogeochemical cycles are highly active. They play an important role in the food web as primary producers, especially in sandy and muddy sediments without vegetation (Smith & Underwood 1998; Underwood & Kromkamp 1999). Tidal flats are one such habitat where benthic diatoms are usually dominant. Evaluating diatom species composition is therefore very important for understanding the tidal flat ecosystem.

Our understanding of diatom flora at river mouth tidal flats is complicated by allochthonous diatoms. These diatoms are transported by river or tidal flows and are usually found as dead frustules or valves without cytoplasm (Kosugi 1986a, b; Sawai 2001; Chiba et al. 2011). We cannot separate these dead diatoms from living ones with conventional preparation methods, which include the chemical cleaning of frustules or valves (Gotoh 1988; Kosugi 1988; Sawai 2001). Valve cleaning is necessary for precise identification of species based on their fine structures; therefore, alternative observation techniques must be combined as described below.

Nuclear staining methods are useful in distinguishing living cells from dead ones. Among these methods, DNA–specific fluorochrome 4',6–diamidino–2–phenylindole (DAPI) staining is well known in cell biology. However, it is not suitable for the evaluation of living diatoms because the simultaneous observation of nuclear fluorescence and the fine structures of valves is impossible. Other staining methods such as Acetocarmine staining (Mayama & Kobayasi 1982), Hematoxylin and Eosin staining (Kosugi 1985), and Hematoxylin staining (Gotoh 1988) enable the observation of fine valve structures and therefore are suitable for both living and dead diatoms. Of these methods, the simplest and most practical is the Hematoxylin staining method developed by Gotoh (1988).

Unfortunately, these staining methods cannot detect most epipsammic diatoms firmly attached to sand grains, because these methods do not decompose the mucilage by which diatoms attach to the substrata. This may lead to a profound underestimation of diatom diversity in tidal flats. Round (1996) extracted epipsammic diatom assemblages from a riverbed by mixing, settling, and decanting specimens repeatedly. However, he reported that loss of epipsammic diatoms and contamination by diatoms from other habitats were unavoidable. In the present study, we used a sieve rather than mixing, settling, and decanting to extract the attached epipsammic diatoms selectively. The dead frustules would theoretically be washed out while sieving, because they are loosely attached or merely trapped among the sand grains, whilst living ones adhere firmly to sand grains by their mucilage.
Thus, we attempted to extract living epipsammic diatoms through a sieving preparation method.

We aimed to achieve a clear picture of the living diatom flora in a sandy tidal flat. To achieve this, we combined multiple methods for evaluating diatom species composition with different characteristics. We applied three different preparation methods: conventional preparation of acid cleaned materials, the nuclear staining method using Hematoxylin solution (Gotoh 1988), and a newly introduced sieving method for the same samples. The different species compositions obtained via these three methods were evaluated in terms of species richness, diversity, and composition based on statistical analyses and the ecological characteristics of the observed species.

Finally, we will propose a synthesis of the three methods to identify the living diatom flora in a sandy tidal flat.

**Materials and Methods**

**Study site.** Samples were collected from the Fujimae Tidal Flat at Nagoya Port in Aichi Prefecture, central Japan (36° 50‘15”E, 35° 04’49”N) (Fig. 1). The Fujimae Tidal Flat is located at the mouth of three rivers: the Shonai River, Shin River, and Nikko River. The tidal flat has a total area of 3.23 km². The average monthly tidal range (based on tide observations for over 12 months) is approximately 2.6 m with sediments consisting of 20% silt and 80% fine sand (Shonai River Office 2006). Diurnal variation of salinity approximately ranges from 4 to 16 psu (Yagi et al. 1999). The Shonai River and Shin River feature some salt marshes consisting of patchy colonies of the reed, Phragmites australis. Because the Nikko River is a high-bedded river, it has a lock gate to prevent the backflow of seawater with stagnant backwater.

**Sampling procedure.** Sediment samples were collected during the exposure period at low tide in July 2014. Surface samples down to a 1.5 cm depth were collected using a tip cut syringe with a 2.7 cm diameter. We set a sampling area of 14,400 m² and collected 25 samples from the area. Each sample was fixed with a 2% formaldehyde solution and preserved in a refrigerator.

**Conventional acid–cleaning method.** We made conventional permanent slides of cleaned diatoms, following the method of Akiba & Suto (2013). Cells collected via this method should contain all diatoms, living or dead, in the sample. Fixed sample material was boiled in a 10% hydrogen peroxide (H₂O₂) solution to remove organic matter, and then 1 N hydrochloric acid (HCl) was added to remove calcareous materials. After washing with distilled water, a sodium pyrophosphate solution (Na₃P₂O₇·10H₂O, 0.01 N) was added to remove clay particles that could hinder the preparation of permanent slides. The sample was again washed with distilled water to obtain cleaned frustules. The solution containing cleaned frustules was deposited onto the cover glass, dried on a hot plate at 60 °C, and mounted using Pleurax (Mountmedia, Wako Pure Chemical Industries, Tokyo, Japan) at approximately 200 °C to make a permanent slide.

**Nuclear staining method to distinguish living diatoms from dead ones.** This method allows the observation of the fine structure of the diatom frustules and the evaluation of the state of the cells by staining the nuclei and decolorizing the plastids (Gotoh 1988). The fixed sample was stirred, and then the suspension was repeatedly centrifuged for five minutes at 1600 rpm and rinsed to remove the formaldehyde. The diatom nuclei were then stained with filtrated Mayer’s Hematoxylin solution (Wako Pure Chemical Industries, Tokyo, Japan) for 15 minutes at room temperature. After this treatment, the samples were washed three times with distilled water and centrifuged (1600 rpm for five minutes), and the liquid layer was discarded. The samples were then decolorized in ethanol for five minutes. After centrifuging (1600 rpm for five minutes), the solution was decanted twice, and the liquid was replaced with 96% ethanol. Each treated sample was placed onto a cover glass, then dried at the room temperature, and mounted using Pleurax at approximately 80 °C to make a permanent slide.

**Sieving method for the extraction of epipsammic diatoms.** This method selectively detects epipsammic diatoms attached to sand grains. The limit for particle size separating sand and silt is defined as 63 μm in the Wentworth scale. Wet sediment sample was washed with distilled water using a sieve with a mesh aperture of 63 μm. Epipsammic diatoms firmly attached to sand grains remained in the sieve, whilst loosely attached or entangled dead diatoms were washed out along with silt and clay. The residue was dried in an oven at 60 °C. It was then boiled with 10% H₂O₂ solution for 20 minutes to decompose organic matter, and then 1 N HCl was added to decompose calcareous matter. The sample was then washed with distilled water by settling and decanting three times, to remove remaining HCl and calcium ions. The permanent slides were made in the same way as in the conventional cleaning method.

![Fig. 1. Location of the Fujimae Tidal Flats. Dotted areas in Map C indicate intertidal areas. Sampling site is indicated by a symbol of star.](Modified from the map in Shonai River Office 2006)
Species identification and counting. Living diatoms always retain intact frustules (i.e., a set of upper and lower valves, usually with girdle bands) in the stained samples while those that are dead or acid–cleaned ones are often separated into unilateral valves. Therefore, we counted 150 cells from each sample for the staining method and 300 valves for both acid–cleaning and sieving methods, respectively. They are equivalent in the number of valves. We observed the slides with a light microscope (BX50, Olympus, Tokyo, Japan), with an objective lens of ×100 (N. A. = 1.3, oil immersion) and equipped with a digital camera (DP70, Olympus, Tokyo, Japan). As the micrographs of nucleus–stained specimens were sometimes not clear enough for precise identification, they were compared with the subjected acid cleaned valves to identify. Some of the cleaned diatoms were identified using a scanning electron microscope (SU6600, Hitachi, Tokyo, Japan).

Evaluation of the three methods through statistical analyses. We expected that the acid–cleaning method would extract all diatoms alive or dead, while the staining method would detect only living diatoms (excluding epipsammic diatoms), and that the sieving method would selectively detect the epipsammic diatoms. Therefore, the assessed species richness, diversity, and composition of a sample might different depending on the method used.

We tested the three methods for differences in their assessment of species richness and diversity. The Shannon–Wiener index \( H' \) was applied for diversity, which is calculated by following equations

\[
H' = -P\sum_{i=1}^{s} \frac{N_i}{N} \log_2 \frac{N_i}{N},
\]

where \( s \) is the number of species, \( N_i \) is the detected number of species \( i \), and \( N \) is the total counted number (Ohgaki 2008). We performed two statistical analyses to identify the differences; the Friedman test and the Wilcoxon signed rank test. Since the three methods were applied to the same sample and the number of samples was small (25 samples for each method) irrespective of the non–normality of the data, we conducted the Friedman test, which is applicable to non–parametric, paired multiple comparison tests (Kurihara 2011). When significant differences were detected, post hoc test for Friedman tests were applied for comparison between two groups.

We compared relative frequencies of each abundant species between the three preparation methods using the Wilcoxon signed rank test (Shimizu 2004). In order to reduce the risk of type I error (null hypothesis is rejected even though it is actually true and should not be rejected), we adjusted the significance level by Bonferroni correction (Shaffer 1995), thus the significance of 5% was divided by the number of groups (3 groups) and a significance of 1.7% was applied. We targeted taxa which were detected in more than 10 samples among the 75 samples observed. All statistical analyses in the present study were performed using R. 3.4.0 (R–project) and RStudio ver. 1.1.453.

Habitat salinity and life form of diatom species. We focused on criteria concerning habitat salinity: fresh, brackish (including fresh/brackish, brackish/marine, and fresh/brackish/marine), and marine, and diatom life form; planktonic, motile, and sessile. In tidal flats, motile (sometimes called “unattached” or “free–living”) diatoms are usually epipelic. Sessile diatoms attach firmly to the substrata, and are also referred as prostrate, haptobenthic, epipsammic, epiphytic, and epilithic. Habitat salinity and life form information were found from the references (e.g. Carpelan 1978; Vos & De Wolf 1993; Van Dam et al. 1994; Chiba & Sawai 2014; Guiry, M.D. & Guiry, G.M 2018; Miranda & Guiry 2018). When detailed descriptions for a species was lacking in the literature, we consulted information on organism’s genus instead (Round et al. 1990).

Results

Number of species detected by three methods. We detected 316 taxa using the three methods. The number of taxa detected by each of the methods are shown in Fig. 2. The staining method highlighted the nuclei in the diatom cells (Figs 3, 5, 7, 9, 13, 15). Among the three methods, the conventional acid cleaning method detected the largest number of taxa: 249 taxa belonging to 70 genera, followed by the sieving method: 236 taxa belonging to 64 genera, and the staining method: 54 taxa belonging to 24 genera. Although each method used the same material, there are multiple taxa detected solely by one of the preparation procedures: 63 taxa for the cleaning method, 16 taxa for the staining method, and 43 taxa for the sieving method.

The prediction states that the acid–cleaned samples should include all the taxa that are present in either stained or sieved ones, and there should be no taxa that are present in both stained and sieved ones. However, from the acid–cleaned samples we could not detect 24 taxa those were detected from stained ones, and 51 taxa from sieved ones. In addition, 37 taxa were common in both stained and sieved samples. The mean relative frequency of each species using the three methods, their habitat salinity, life form, and references are exhibited in Supplementary table (Table S1).

The dominant species identified by the three methods were as follows: for the acid–cleaning method,
Staurophora dubitabilis (17.5%, Figs 3, 4), Cyclotella meneghiniana (10.1%, Figs 5, 6), and Aulacoseira granulata (5.5%, Figs 7, 8); for the sieving method, Hippodonta sp. 1 (5.9%, Figs 9, 10), Staurophora dubitabilis (5.8%), and Planothidium frequentissimum (4.5%, Figs 11, 12); and for the staining method, Pseudostaurosira trainorii (36.8%, Figs 13, 14), Skeletonema costatum (19.1%, Fig. 15), and Staurophora dubitabilis (10.8%).

Species diversity at 25 sites identified by each method. The differences in species richness and the Shannon–Wiener index of the 25 samples obtained using the three methods were compared as shown in Figs. 16 and 17. The Friedman test detected a significant difference in both species richness and Shannon–Wiener index between the three methods (p < 0.001). The post hoc tests for species richness showed no significant difference between cleaned and sieved samples (p = 0.086), but the staining method was significantly smaller than in both the sieved and acid–cleaned samples (p < 0.001). The Shannon–Wiener index indicated significant differences between each combination of the three groups (cleaned < sieved, p = 0.003; sieved > stained, p < 0.001; cleaned > stained, p < 0.001).

The frequency of appearance of each species for the three methods. Using 75 subsamples (25 samples×3 methods), 127 taxa were detected in 10 or more subsamples. In total, a Wilcoxon signed rank test determined that significant differences existed in the detection of 90 species between at least one combination of the three preparation methods with a p–value of less than 1.7%. In terms of relative frequencies, 50 taxa exhibited the most abundant in the sieved samples, 42 in the acid–cleaned samples, and 5 in the stained samples.

Fig. 16. Boxplot of the number of diatom species identified at 25 sites through three methods.

Fig. 17. Boxplot of the Shannon–Wiener index determined for 25 sites through three methods.


**DISCUSSION**

**The differences in identified species diversity between the methods**

The Friedman tests clearly showed differences between the three methods in terms of species richness and Shannon–Wiener’s diversity (H’), although the difference between acid–cleaned and sieved samples was not distinct. It was expected that species diversity in acid–cleaned samples would be higher than in the stained ones because most of the dead and/or epipsammic diatoms would be excluded in the latter. In addition, the acid–cleaning procedures separate the valves from the frustules and increase the probability that rare species were counted as single valves. This partly explains the higher species richness and diversity in the acid–cleaned and the sieved methods compared to the stained one, although it cannot be the sole reason for the marked differences. The small difference between acid–cleaned samples and stained ones is notable, because the latter also should have excluded most of the dead and/or epipelic diatoms. This issue will be discussed below through a detailed comparison between the species compositions.

**Diatom species extracted by the acid–cleaning method**

The freshwater diatoms which were present in the acid–cleaned samples but were much rarer or absent in both the stained and sieved samples, were presumed to be allochthonous. Among the 63 taxa which were found only in the acid cleaned samples, 34 were freshwater species belonging to genera such as *Eunotia*, *Gomphonema*, and *Luticola* (Fig. 2, Table S1). They were presumed to have been dead by the time of sampling, unless some of them were still living but not detected because of their rarity. Among the 42 species which were significantly more abundant in terms of relative frequency in the acid cleaned samples than in the others, 29 taxa were freshwater diatoms (e.g. *Amphora* *copulata*, *Diadesmis* *confervacea*, *Encyonema* *simile*, *E. ventricosum*, *Eunotia* *metamonodon*, *E. sedina*, *Gomphonema* *contrattriss*, *G. inaequilongum*, *G. sp.5*, *Pinnularia* *oriunda*, *P. parvulissima*, *P. schoenefelderi*, and *P. sp.2*; Table S1). Because most of them were absent or very rare in the stained and sieved samples, they were also considered to be from the river.

Some previous studies have reported that allochthonous frustules brought by tidal currents and river inflow becomes significant contaminants in coastal wetlands (e.g. *Kosugi* 1986b; *Vos & de Wolf* 1993; *Sawai* 2001). *Gotoh* (1988) also stated that “conventional cleaning method leads to misunderstanding of the diatom assemblage when the sample contains many dead or allochthonous cells.”

However, many brackish motile diatoms, which were the most abundant in the acid–cleaned sample, were also rare in the stained samples. They correspond to *Bacillaria* *paxillifera* var. *tumidula*, *B. urve–millerae*, *Entomoneis japonica*, *Hantzschia amphioxxys*, *Navicula degitoradiata*, *Parlibellus cruciculoides*, *Petroneneis marina* and *Tryblionella apiculata*. Except *Staurophora dubitabilis* (10.8%), these brackish motile diatoms were rare in the stained samples in terms of relative frequency (0%–0.7%), irrespective of their nature as a potential component of tidal flat diatoms. We hypothesize that while they used to be live in the tidal flat, they were already dead by the time of sampling.

**Diatom species extracted by the sieving method**

Diatoms identified through the sieving method were typically of the sessile type (Table S1). They included many monoraphid species belonging to *Achnanthidium*, *Cocconeis*, *Planothidium*, and other *Achnanthes* sensu lato, which are monoraphid and strongly attach to substrata through secreted mucilage on their raphe valves *(ROUND* *et al.* 1990). Many species were not found or were very rare in the stained samples. For instance, *Achnanthidium exiguum*, *A. subhudsonis*, *Cocconeis lineata*, *Planothidium frequentissimum* and *P. rostratum* were all found to have 0.0% mean relative frequencies in the stained samples, while they were all abundant in the sieved samples (>3%). Small and prostrate *Sellaphora nigri* were also showed a similar tendency (0.0% in stained samples vs. 4.4% in sieved ones). These results suggest that the sieving method can scrape the epipsammic diatoms selectively and supplements the staining method, in which epipsammic diatoms are mostly neglected.

Many freshwater diatoms were abundant in the sieved samples irrespective of the brackish nature of the Fujimae Tidal Flat (4–16 psu; *Yagi* *et al.* 1999). For instance, *Achnanthidium exiguum* (0.8%), *A. subhudsonis* (3.1%), *Karayevia clevei* (0.2%), *Nitzschia fonticola* (2.6%), and *Planothidium frequentissimum* var. *magnum* (0.8%) are usually referred to as freshwater taxa in the past literature. They may be resistant to brackish environments, and travel via rivers attached to sand grains. These contaminations of freshwater species might partly explain the relatively higher species richness in the sieved samples.

Unexpectedly, some planktonic diatoms such as *Cyclotella* *atoms*, *C. cryptica* and *C. mediana* were significantly more abundant in the sieved samples than in the acid–cleaned ones. These diatoms can be trapped within the biofilm on the sand grains and sometimes attach to the substrate through their own mucilage, which is likely why they occurred in the sieved samples. Their relative abundances, however, should be higher in the acid–cleaned samples than in the sieved ones. A possible explanation of why the reverse occurred is that the procedure to remove clay particles in the acid–cleaning method washed away these small diatoms, while the sieving method did not employ this process.

**Diatom species extracted by the staining method**

The staining method showed an advantage in that it could not only selectively detect living diatoms but also fragile and/or small ones. Some diatom taxa with fragile
frustules (e.g. Chaetoceros minimus, Cylindrotheca graci-
lis, Entomoneis aequabilis, E. paludosa, Skeletonema
potamos) and/or small–sized cells (e.g. Cyclotella ato-
mus, C. cryptica, Discostella wolterski) were detected
only by the staining method. Previous studies point out
that the acid solution and the decanting procedure of
the acid–cleaning method sometimes break the cells
(e.g. BEYENS & DENYS 1982; HIROSE et al. 2013; CHIBA
2014). We also suspect that small–sized cells remained
suspended in the supernatant water during the cleaning
due to the use of a dispersing agent and were therefore
washed out. Our results show that the staining method
can detect diatoms with fragile frustules and small–sized
cells more effectively than the conventional acid–clean-
ning method.

Five taxa were significantly more abundant in
samples evaluated using the staining method than in
either of the other two methods. These species were
autochthonous with high abundance (e.g., 36.8% for
Pseudostaurosira trainorii, 19.1% for Skeletonema
costatum, and 5.7% for Navicula arenaria var. rostellata).
Among them, P. trainorii is considered to be a freshwater
species. However, this might be a misidentification of
Staurosira sopotensis (Witkowski 1994), and it is also
possible that they belong to the same species.

**Evaluation of the three methods**

The conventional acid–cleaning method alone should
not be used for floristic studies of the tidal flat diatoms,
although it is necessary to identify these diatoms pre-
cisely. In this study, the diatom flora identified through
the acid–cleaning method contained many species to be
allochthonous, belonging to freshwater genera such as
Eunotia, Gomphonema, and Luticola. These subsamples
also contained species which were possibly autochtho-
nous, but apparently have died at the sampling time,
such as Hantzschia amphioxys, Bacillaria paxillifera
var. tuidula, and Navicula digitatoradiata. In addition,
this method cannot detect many diatoms with fragile
and/or small cells (see above).

The staining method selectively extracts living cells,
especially fragile and small species such as Chaetoceros
minimus, Cylindrotheca gracilis, Entomoneis aequabilis,
Cyclotella atomus, C. cryptica, and Discostella wolterski,
which are frequently overlooked by the acid–cleaning
method. Conversely, this method overlooked most of the
sessile or epiphasmic diatoms which were attached to
the sand grains, and thus caused an underestimated
of the species diversity.

The diatoms detected by the sieving method were
characterized by the sessile type, such as Achnanthes
dispar, Cocconeis, Planonothidium, and Sellaphora, which the
staining method could hardly detect. This method also
has an advantage over the conventional acid–cleaning
method in that the procedure to remove clay particles
can be omitted, and small–sized diatoms can thus be
detected efficiently.

The combination of the three methods employed
in this study can be used to more precisely identify the
living diatom flora of tidal flats than the approaches used
in previous studies.

**Three advantages of the combined method**

In this section, we discuss three advantages that our
combined method to evaluate diatom flora has over
methods used in the past studies.

The first advantage of our combined method is
the elimination of allochthonous diatoms. As discussed
above, the acid–cleaning method cannot eliminate
allochthonous diatoms. Previous studies on tidal–flat diatom
flora have often documented many freshwater diatoms
irrespective of the high salinity. For instance, salinity
ranges from 8–35 psu in the Yaquina Estuary (RIZNYK
1973) and approximately 25–30 psu in the Ariake Sea
(OHTSUKEYA 2005; PARK et al. 2012). As both areas feature
much higher salinity than that of the Fujimae Tidal Flat
(4–16 psu), most of the freshwater diatoms identified
there are presumed to be allochthonous. Other methods
such as the cover–glass technique (OPPENHEIM 1991) and
lens–tissue one (COLIN & DUKEMA 1981) can select liv-
ing diatoms, but they may also exclude many planktonic
or sessile diatoms with low motility.

The second advantage is the detection of fragile
and/or small diatoms. The staining method in the present
study revealed a novel and unexpected aspect of tidal–flat
diatom flora: many diatoms with fragile frustules lived
there. We found fragile diatoms such as Chaetoceros
minimus, Cylindrotheca gracilis, Entomoneis aequabilis,
E. paludosa, and Skeletonema potamos from the surface
of the sand. Such diatoms have been scarcely detected in
most previous studies (RIZNYK 1973; COLIN & DUKEMA
1981; OPPENHEIM 1991; OHTSUKEYA 2005; PARK et al.
2012), with the exception of SABBRE (1993), which de-
tected the fragile diatom Nitzschia closterium as a major
component. It is probable that these fragile diatoms are
destroyed during the cleaning process. Small diatoms
could be washed out if a suspending agent is used to
remove clay particles, as the present study demonstrated.
In principal, the cover–glass technique (OPPENHEIM
1991) or the lens–tissue technique (COLIN & DUKEMA
1981) can identify such diatoms, but in reality, such
diatoms are rarely detected because of their non–motile
nature or weak adhesive power.

The third advantage of the combined method
is the effective detection of sessile diatoms that attach
to sand grains. Because sessile diatoms usually move
more slowly than motile ones (HARPER 1977), both the
cover–glass technique (OPPENHEIM 1991) and the lens–
tissue technique (COLIN & DUKEMA 1981) are assumed
to collect them less effectively than our sieving method.
This assumption, however, might not apply to diatoms
such as Planonothidium and Halanaphora, because these
diatoms were also reported by COLIN & DUKEMA (1981)
and OPPENHEIM (1991). Although the conventional acid–
cleaning method can also detect these sessile diatoms,
it cannot separate the allochthonous ones from those
living on the sand grains in the tidal flats.

We are convinced that we can assess the living diatom community on a tidal flat more thoroughly than previous diatom studies through the combination of these three preparation methods. Such combined methods are necessary to precisely describe diatom flora, especially those of depositional environments where dead diatom frustules are abundant and multiple microhabitats are present even within a sample.

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Supplementary material

The following supplementary material is available for this article:

Supplementary Table S1.

This material is available as part of the online article (http://fottea.czechphycology.cz/contents)