# **CHAPTER 2 ANNEX I**

# DISTRIBUTION OF THE INVASIVE CTENOPHORE *MNEMIOPSIS LEIDYI* IN THE

# BELGIAN PART OF THE NORTH SEA

#### Adapted from:

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#### **Abstract**

The invasive ctenophore *Mnemiopsis leidyi* was recorded for the first time in northern Europe in summer 2005, while the first records in the North Sea date back to the summer of 2006. The first sightings in the Belgian part of the North Sea were made in August 2007 in the port of Zeebrugge, but most probably M. leidyi had already been present for a longer period in this area. The high densities in the port of Zeebrugge suggest that M. leidyi entered the Belgian marine waters via ballast water transport, comparable with the invasion in the Black and Caspian Seas and the Dutch part of the North Sea. In the period 2009–2011, M. leidyi was found in all ports and all along the Belgian coastline, up to 27 km offshore. Further offshore, no M. leidyi were found in zooplankton samples and small meshed otter trawl samples. Sightings of adult individuals in the coldest winter months imply that the species can survive Belgian winters. Highest densities (17 ind.m<sup>-3</sup>) were found in the Sluice dock in the port of Oostende. Along the coastline, average densities of 0.4 ind.m<sup>-3</sup> were recorded. As M. leidyi might previously have been misidentified on the basis of morphological features alone, we also identified the species with genetic identification tools. Taking into account the notorious impact of this species in its native and in other invaded waters, it is recommended to continue the monitoring of M. leidyi populations in the Belgian part of the North Sea.

Keywords: *Mnemiopsis leidyi*, Belgian ports and coast, gelatinous zooplankton, non-native species, ballast water, DNA sequencing, molecular probe

#### 1. Introduction

The lobate ctenophore *Mnemiopsis leidyi* naturally occurs along the Atlantic coast of North and South America (Mayer 1912, Gesamp 1997). This ctenophore species can survive in water temperatures ranging from 0 to 32 °C and in salinity of 2–39 PSU (Kremer 1994). *Mnemiopsis leidyi* matures after 13 days and can produce *ca.* 8000 eggs in 23 days (Baker and Reeve 1974). Additionally, the species shows hermaphroditism, which implies that a single comb jelly in principle can produce a complete new population. As the densities of *M. leidyi* can increase very fast, they might form a serious threat for the zooplankton, fish eggs and fish larvae, which are considered as their main prey (Vinogradov and Shushkina 1992, Sullivan *et al.* 2001).

The introduction of *M. leidyi* into the Black Sea is considered a textbook example of the deleterious consequences of marine bioinvasions. In addition to overfishing and eutrophication, the introduction of M. leidyi in the early 1980's (probably through ballast water) lead to shifts in the pelagic food web and severe economic losses for the anchovy Engraulis encrasicolus (Linnaeus, 1758) fishery in the Black Sea (Kideys 2002, Knowler 2005). In 1989, the biomass of *M. leidyi* peaked at 1.5–2 kg.m<sup>-2</sup> in the Black Sea (Vinogradov et al. 1989). Because of this notorious history, the introduction of M. leidyi in other areas is of major concern. The first records of M. leidyi in the North Sea date back to 2005, where it was found in the Skagerrak area (Oliveira 2007). In August 2006, blooms of M. leidyi were observed in the Dutch Wadden Sea and the Scheldt estuary (Faasse and Bayha 2006). Prior to these sightings, autumn blooms of lobate ctenophores -identified as Bolinopsis infundibulum- were reported in the Netherlands (Holsteijn 2002). However, B. infundibulum is a cold water species considered rare along the Dutch coasts. As morphological identification is indeed difficult for related ctenophores, it remains uncertain whether these earlier sightings of B. infundibulum could have been misidentified M. leidyi. It is recommended that 'older' ctenophore observations in the North Sea are considered with some precaution. This illustrates the need for an unambiguous identification with genetic tools. Based on different nuclear and mitochondrial genetic biomarkers, these tools can be used for species identification and even the delineation of different populations (Gorokhova and Lehtiniemi 2010, Reusch et al. 2010).

In autumn 2006, *M. leidyi* was also found in the German part of the North Sea near Helgoland (Boersma *et al.* 2007) and the Baltic Sea near Kiel (Javidpour *et al.* 2006). *Mnemiopsis leidyi* was observed for the first time in the Belgian part of the North Sea (BPNS) in August 2007, where it was found at high numbers in the port of Zeebrugge (Dumoulin 2007).

In this Annex, we present the observation of *M. leidyi* in several docks within the ports of Nieuwpoort, Oostende and Zeebrugge, and at ten mesozooplankton monitoring stations in the BPNS (Fig. 1). Ctenophores were caught using plankton nets, fish trawls and dip nets. The individual species were identified both morphologically and genetically.

#### 2. Materials and Methods

#### 2.1 Ctenophore sampling

Ctenophores were sampled in the ports, coastal and offshore areas of the Belgian part of the North Sea with a variety of methods (Addendum 2). Between August 2008 and March 2011, qualitative data were monthly gathered by the Belgian Marine Life Field Study Group in the Ports of Nieuwpoort, Oostende and Zeebrugge (Fig. 1). These data mainly consist of non-standardized visual inspections from the shore, mostly combined with hand gathering using dip nets. In the Sluice dock (1.5 m deep, situated in the port of Oostende), an additional quantitative sampling was performed in October 2010 at four sampling locations, with a rigid inflatable boat towing a WP2 plankton net (57 cm diameter, 1 mm mesh size, fitted with a flow meter) at 2 knots just below the water surface. Biovolumes of *M. leidyi* in the Sluice dock were measured with a graduated cylinder.

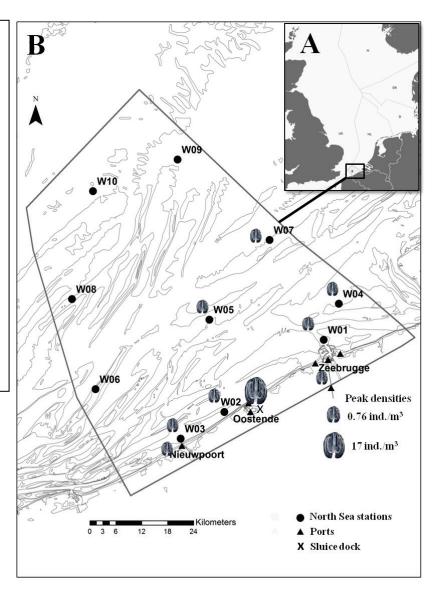
From January 2010 to December 2010, monthly mesozooplankton samples were taken with a WP2 net (57 cm, 200 µm mesh size, fitted with a flow meter) towed in an oblique haul from bottom to surface at ten monitoring stations in the Belgian part of the North Sea (Fig. 1). Although similar samples were gathered in 2009, these could not be used for this study, as the whole zooplankton samples were fixed in 4 % formaline solution, in which *Mnemiopsis* species dissolve. In 2010, the sampling protocol was adapted, with all ctenophores extracted and quantified prior to sample fixation.

Between January 2009 and December 2010, an otter trawl was used at the same ten sampling locations on the BPNS to investigate whether there were dense aggregations of

Mnemiopsis close to the seabed. The trawl has a net opening of 3\*1 m, net mesh size 10 mm, and was dragged over the seafloor for 30 minutes at 3 knots. Ctenophores were directly identified onboard. A binocular stereomicroscope was used for small or damaged specimens. From January 2011 to March 2011, quantitative data on *M. leidyi* were gathered with a WP3 net (1 m diameter, 500 μm mesh size, fitted with a flow meter) at stations W01-04-07-09 in the BPNS (Fig. 1). The WP3 net was towed along a curvilinear trajectory (net going down and up the water column (undulating) twice, at a net speed of 3 knots) to sample a representative cross section of the water column. The WP3 net is more suited to look for *Mnemiopsis* occurring in low densities than a WP2 net, as it filters much bigger water volumes.

Figure 1: The port, nearshore and midshore peak densities of *Mnemiopsis leidyi* found in this study, with highest densities in the Sluice dock of Oostende.

- A) North sea Exclusive Economic Zones;
- B) Belgian part of the North with ten stations (nearshore W01-04, midshore W05-07, offshore W08-10) sampled monthly zooplankton (black dots), and ports specifically sampled for Mnemiopsis leidyi (triangles). The Sluice dock is part of the Oostende port (cross). Locations where Mnemiopsis was found are indicated with a ctenophore miniature symbol.



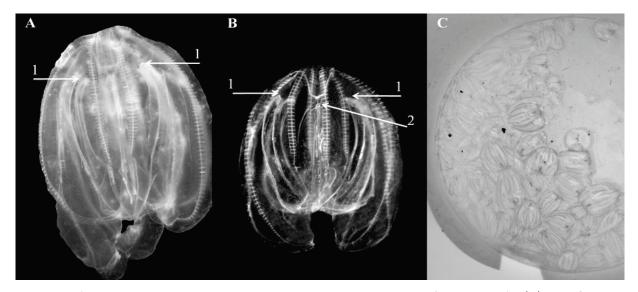


Figure 2: A) *Mnemiopsis leidyi* photographed at the marina in the port of Zeebrugge (23/9/2009, © Hans Deblauwe) B) *Mnemiopsis leidyi* photographed in the German Bight (29/1/2010, © Karl Van Ginderdeuren). The most important morphological feature characterizing *Mnemiopsis leidyi* is the extent of the oral lobes (1). In this species the oral lobes extend over almost the entire body and reach the statocyst (2) C) *Mnemiopsis leidyi* photographed at the Sluice dock in Oostende (22/10/2010, © Karen Soenen).

# 2.2 Morphological species identification

All ctenophores were identified alive onboard. Intact ctenophores from plankton trawls and dip nets were measured (total length in mm, including lobes) to determine the size range. The most important morphological feature characterizing *M. leidyi* is the extent of the oral lobes (Fig. 2). The oral lobes of *M. leidyi* extend over almost the entire body and reach the statocyst (the apical sense organ), whilst in the similar species *Bolinopsis infundibulum*, the oral lobes terminate between the mouth and the statocyst. The extent of the oral lobes was verified for each individual lobed ctenophore.

## 2.3 Molecular analysis

Ctenophores from locations W01-02-03-05 sampled in September 2010 were selected for molecular analysis (Addendum 2). Each individual was preserved separately in 70–100 % ethanol and stored at 4 °C till genetic analysis. Genomic DNA was extracted from the gelatinous lobe tissue using the Genomic DNA extraction Wizard® (Promega). Molecular species identification was done by amplifying and sequencing the internal transcribed spacer (ITS) DNA sequence as a genetic biomarker with primers KN8-9-11 (Fuentes *et al.* 2010, Ghabooli *et al.* 2011).

Base calling and sequence assembly for *ca.* 420bp fragments was done with the Bionumerics software 6.5™ (Applied Maths). Positions showing two peaks were coded as degenerated

and given a DNA IUB ambiguity code. Sequence identity was evaluated performing a NCBI-BLASTN search against the sequences present in the GenBank.

#### 3. Results

Between August 2008 and March 2011, *Mnemiopsis leidyi* were seen in the ports of Nieuwpoort, Oostende and Zeebrugge, the nearshore stations W01-02-03-04 and the midshore stations W05 and W07 (Fig. 1, Addendum 2). No individuals of *M. leidyi* were caught in midshore station W06 and the offshore stations W08-10 (> 30 km from the coast). Highest densities were noted in the Sluice dock in Oostende. In all cases the oral lobes of the ctenophores reached the position of the statocyst, which positively identified them as *M. leidyi* (Fig. 2). The individuals that were measured ranged between 1 and 6 cm in total length. *Bolinopsis infundibulum* was not found among the ctenophores.

The specimens that were genetically characterized, showed a similarity score of 99–100 % with the ITS sequences from *M. leidyi* clones present in the Genbank. Query coverages ranged from 80–100 %. Other hits showing lower similarity scores included *Pleurobrachia* sp. (96 % similarity, 43 % query coverage), *Bolinopsis* sp. (95 % similarity, 100 % query coverage), and *Beroe* sp. (92 % similarity, 72 % query coverage). In agreement with the morphological identification, the molecular results also identified the ctenophores as *M. leidyi*.

*Mnemiopsis leidyi* was not found between November 2009 and June 2010. For the rest of the studied period, the species was found in all seasons. Even in the coldest winter months, adult *Mnemiopsis* were present in the ports and near/midshore, at water temperatures as cold as 2 °C. Peak densities were observed in October 2009 and 2010, with *M. leidyi* found at all sampled locations except offshore stations (Addendum 2).

Based solely on the WP2 zooplankton samples where *M. leidyi* occurred in 2010, a yearly average density of 0.4 (± SD 0.2) ind.m<sup>-3</sup> was calculated. The WP3 samples from January and February 2011 showed very low densities of *M. leidyi*, on average 0.01 ind.m<sup>-3</sup>. The highest density at sea was 0.8 ind.m<sup>-3</sup> at the nearshore station W01 (Zeebrugge) and midshore station W05 (Gootebank) in October 2010.

*M. leidyi* was recorded in the otter trawl samples only in October 2009 and 2010. In October 2010, densities were similar to those in the zooplankton samples: on average 0.37 and 0.29

ind.m<sup>-3</sup>, respectively. In the Sluice dock (port of Oostende), *M. leidyi* appeared to be common in Autumn 2010, with an average density of  $9.7 \pm 6.5$  ind.m<sup>-3</sup> (peak density 16.8 ind.m<sup>-3</sup>) on 22nd October, and an average biovolume of  $24.8 \pm 17.6$  ml.m<sup>-3</sup>.

## 4. Discussion

Marine invasions are considered a major threat for the world's oceans. Ballast water conveys marine species on a worldwide scale. It is estimated that the annual amount of 3.4 billion tons of ballast water moves some 7000 species around the world at any given time (Carlton 1985, Clarke *et al.* 2003, Globallast 2007). Transferred species can establish reproductive populations in the host environment, outcompeting native species and multiplying into pest proportions. Currently, 71 marine non-indigenous species (e.g. algae, crustaceans, cnidarians) have established persistent populations in the Belgian part of the North Sea (Kerckhof *et al.* 2007, Vandepitte *et al.* 2012).

It is possible that *Mnemiopsis leidyi* has been introduced into the North Sea through a secondary invasion from the Baltic Sea (Reusch *et al.* 2010). However, the fact that *M. leidyi* was first seen within the port of Zeebrugge strongly supports the hypothesis that the species was brought in directly via ballast water transport in cargo ships, comparable with the invasions of *M. leidyi* in the Black and Caspian Seas (Vinogradov *et al.* 1989, Ivanov *et al.* 2000), and as indicated for the Dutch part of the North Sea (Faasse and Bayha 2006). Given the high densities upon discovery in the port of Zeebrugge, it is likely that the species was already present for several months before August 2007.

Molecular analysis confirmed the morphological identification of *M. leidyi* in the Belgian part of the North Sea. However, efforts must be undertaken to increase the genetic data on ctenophores, as the number of corresponding ITS sequences in GenBank for different ctenophores such as *Beroe* sp., *Bolinopsis* sp. and *Pleurobrachia* sp. remains scarce. Also, the development of real time PCR probes might contribute to an easy and fast detection of ctenophore species -including hard-to-identify eggs and larvae of ctenophores- in bulk samples.

It is known that jellyfish and ctenophores can thrive in areas with high anthropogenic impacts such as overfishing, eutrophication and habitat modification (Mills 2001, Purcell *et al.* 2007, Richardson 2008, Richardson *et al.* 2009). Moreover, temperate jellyfish species

such as *M. leidyi* can benefit from the effects of global warming (Purcell 2005). *Mnemiopsis leidyi* exhibits typical characteristics of a pest species, e.g. extensive temperature and salinity ranges, high reproduction rates, survival in very eutrophic and polluted waters, and the ability to rapidly colonize foreign and neighboring areas (Kremer 1994, Faasse and Ligthart 2007). Our results indicate that *M. leidyi* might have established a viable population along the Belgian coast. The 2009-2010 winter was the coldest in fifteen years (KMI 2010) with an average estimated SST of 4.1 °C on the BPNS (OSTIA, Stark *et al.* 2007). Yet, *M. leidyi* was present all along the coastline and in all ports during summer and autumn of 2010. This suggests that, in terms of temperature regimes, *M. leidyi* is likely to remain present in the BPNS. However, as data on eggs and larvae of *M. leidyi* are not available yet, it is not possible to state if the species really reproduces in the BPNS or if it is introduced yearly from source populations in adjacent regions (e.g. the Westerschelde estuary or the Dutch coast). Riisgård *et al.* (2012) describe how in years with a pronounced water exchange between the North Sea and the Baltic Sea, *Mnemiopsis* ctenophores are imported to the Kattegat to establish temporary populations.

Extensive molecular analyses, *cf*. the study on the central Baltic Sea by Schaber *et al.* (2011), and the development of molecular identification probes could help to unravel this.

In Narragansett Bay (US), *M. leidyi* biovolumes of up to 100 ml.m<sup>-3</sup> were noted (Kremer 1994 in Purcell *et al.* 2001). Baker and Reeve (1974) described *M. leidyi* densities up to 50 ind.m<sup>-3</sup> in native areas. In Limfjorden (Denmark) densities reached 800 ind.m<sup>-3</sup>, with biovolumes ranging around 100–300 ml.m<sup>-3</sup> (Riisgård *et al.* 2007). With few predators present, *M. leidyi* can reach biovolumes of up to 600 ml.m<sup>-3</sup> (Purcell *et al.* 2001). It should be noted that it is not always clear if the above mentioned values are solely based on adult densities. The highest densities observed in our study (animals > 1 cm) were found in the Sluice dock (port of Oostende), with a maximal biovolume of 25 ml.m<sup>-3</sup>. Since the 1960's, the Sluice dock has been of high economic importance for oyster farming (Curé *et al.* 2002). *Mnemiopsis leidyi* is classified as a zooplanktivorous predator (Vinogradov and Shuskina 1992, Sullivan *et al.* 2001). The high densities of this species in the Sluice dock potentially could have a profound effect on the oyster farm, as *M. leidyi* probably also feeds on oyster larvae and may compete with adult oysters for planktonic food sources.

The presence of *M. leidyi* can pose a serious threat on the pelagic ecosystem and the anthropogenic activities in the Belgian part of the North Sea. The most important target fish species of the Belgian fishing fleet are plaice *Pleuronectes platessa* and sole *Solea solea*. *Mnemiopsis leidyi* might have a profound negative effect on the pelagic larvae and eggs of these coastally spawning fish through predation and competition (Reeve *et al.* 1978, Kremer 1979, Hamer *et al.* 2011). Especially for sole, which spawns from April to September (Munk and Nielsen 2005), as there is an overlap between the presence of *M. leidyi* adults and sole larvae in the water column. Competition for zooplankton food sources could form a major problem for a successful recruitment of plaice and sole juveniles. In 2009, the Belgian sole and plaice fisheries resulted in a catch value of 16 million Euro (Comm. Department of agriculture and fisheries – Sea fisheries service). This income could potentially be severely reduced due to the presence of *M. leidyi*.

Through natural predation, the negative impact of the *M. leidyi* invasion might be kept to a minimum. Another ctenophore *Beroe gracilis* is known to predate on *Pleurobrachia pileus* (Greve 1970, Greve and Reiners 1988). Preliminary lab experiments point out that *Beroe gracilis* from the Belgian part of the North Sea is also capable of eating *M. leidyi* (Vansteenbrugge, Pers. Obs.). This corresponds with lab experiments performed by Hosia *et al.* (2011). Not only gelatinous predators prey upon *M. leidyi*, also several fish species are known to feed on ctenophores. Mianzan *et al.* (1996) found that 20 of 69 investigated fish species in Argentina had ctenophores (including *M. leidyi*) in their gut, while Schaber *et al.* (2011) reported *M. leidyi* to be present in stomachs of cod *Gadus morhua* in the Baltic Sea. Further studies are needed to investigate which jellyfish and fish species prey upon *M. leidyi* in the BPNS.

Since the discovery in the port of Zeebrugge less than six years ago (Dumoulin 2007), *M. leidyi* is now known to occur along the entire Belgian coast. The observed peak densities and biovolumes are still lower than in Denmark, the Black Sea or the natural habitat in the US (Decker *et al.* 2004, Riisgård *et al.* 2007). The southern North Sea is a very different water body compared to the Black Sea, with other food web interactions, differences in water quality and other fish communities and fisheries. For the time being, the deleterious

ecosystem scenario similar to that in the Black Sea might be less likely in the Belgian part of the North Sea and the adjacent water bodies, but cannot be ruled out.

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