**Abstract:** The Marine Strategy Framework Directive (MSFD) aims to reduce the marine debris burden in the marine environment by 2020. This requires an assessment of the actual situation, which includes the occurrence as well as the caused impacts. Information on both is scarce when it comes to top predators like marine mammals and the burden of microplastic. This is hampered by the limited access to free ranging marine mammals for collecting samples, as well as sample handling. The present study investigated gastrointestinal tracts and faecal samples of harbour seals (*Phoca vitulina*) and grey seals (*Halichoerus grypus*) regularly occurring in the German North Sea and Baltic Sea with the aim of gaining information on the occurrence of microplastics. In total, 255 particles ≥100 µm (70 fibres, 185 fragments) were found in exemplary ten intestine and nine faecal samples. The findings ranged from zero fibres and six fragments, up to 35 fibres and 55 fragments per sample. Additionally, this study established a protocol for sample handling, microplastic isolation (≥100 µm) and quantification of gastrointestinal tracts and faecal samples of marine mammals with a low share of contamination. This approach helps to quantify the presence of microplastics in free-ranging marine mammals and is therefore applicable to assess the real burden of microplastic presence in the marine environment.

**Keywords:** microplastic; plastic isolation; plastic ingestion; gastrointestinal tract; marine mammals

1. Introduction

The challenging nature of marine debris pollution is well known [1]. Due to its widespread use, its specific characteristics and discard virtue, synthetic polymers, so-called “plastics” make up a high share of marine debris in our oceans [2–4]. Microplastic (MP) includes synthetic polymers in the form of particles smaller than 5 mm [5]. These particles originate either from large plastic items cracking down into smaller fragments due to various forces (secondary MP); or are intentionally produced in those small sizes (primary MP) [6–9]. The awareness of microplastics started back in the 1970s [10,11] but has only recently been brought more into focus of environmental research [12,13].

MPs were revealed to be ingested by organisms of lower trophic levels, and even a trophic transfer was detected [14–16]. A variety of international studies confirmed the presence of different sized plastic particles in the gastrointestinal tract (GIT) of various marine mammalian species [17–21].

Dealing with faunal samples implies the initial elimination of organic matter enclosing the particles; their detection otherwise would be obscured [22,23]. Therefore, the biogenic compounds have to be removed prior to further investigations in order to obtain convincing and reliable results [22,23].
Furthermore, it needs to be ensured that the risk of secondary contamination during sampling, storing and sample treatment is kept as low as possible. Contamination of samples might occur during their handling and origin, e.g., from storage containers, laboratory cloths, gloves and the working environment in general [24–28].

Previous studies already presented various digestion protocols of biota samples in recent years [17,20,29]. Enzymes were predominantly used to remove the organic matter [23,30], in some cases acids or alkaline solutions were applied to digest organic remains [26,29,31]. Subsequently, MP identification techniques such as the staining and fluorescence microscopic approach [32,33] and spectroscopic analyses like Fourier transform infrared (FT-IR) [34–36] or Raman spectroscopy are commonly used [33,37,38]. A combination of both methods is considered to be a reliable option regarding the identification of polymer compositions [39].

However, information on the presence of MP and its impacts on marine top predator species is scarce. Therefore, the present study focusses on establishing a protocol to isolate particles, evaluate potential contamination and propose a method to discriminate between MP and other particles in faecal and intestinal samples of seals in order to investigate the presence of MP in marine mammals in further research. Samples were taken from harbour seals (Phoca vitulina) and grey seals (Halichoerus grypus), which regularly occur in the North and Baltic Seas. This study was realised within the framework of the project “Assessment and implementation for long-term monitoring of pollution of diverse marine compartments and biota with marine litter” (Federal Environment Agency Germany), the aims of which were to optimize sample handling and to reduce the risk of secondary contamination for detecting microplastic particles in intestinal samples of marine mammals.

2. Materials and Methods

2.1. Sample Collection

The Institute for Terrestrial and Aquatic Wildlife Research (ITAW) at the University of Veterinary Medicine Hannover (Foundation Germany) regularly conducts necropsies of stranded, bycaught or euthanised marine mammals found along the coastline of the Federal State of Schleswig-Holstein, Germany [19,40,41]. These are predominantly the three species regularly occurring in German waters: harbour porpoise (Phocoena phocoena), harbour seal and grey seal.

Necropsy Sampling

Concerning microplastic analyses in the GIT of marine mammals, the rectum of harbour seals and grey seals has been sampled since 2014. Out of these, ten pinniped samples were used for this study. The caudal part of the rectum was tied off with a drawstring (Figure 1A). Subsequently, in a cranial direction, an 8–10 cm section was tied off. Both ends were cut off behind the drawstrings to prevent the loss of faeces (Figure 1B). With the help of metallic tweezers, the samples were transferred to cleaned and disinfected glass jars. Afterwards, the glass jars were stored at −20 °C until further processing (Figure 1C,D).

Sandbank Sampling

In the course of a regularly conducted health monitoring of live animals at the Lorenzenplate [42], a tide-dependent submerged sandbank in the North Sea, further faecal samples of free-ranging harbour seals and grey seals have been collected since 2012. Out of these, nine faecal samples were analysed. No genetic analyses have been performed to distinguish between harbour seal and grey seal faeces. Hence, this study consecutively refers to them as seal faecal samples.

The following steps were conducted in a closed acrylic box (see Supplementary Materials) and are applicable for all further investigations on intestinal and faecal samples. The box was equipped with two holes, which formed the only openings within the box, these being necessary for processing the samples. During the whole procedure, the samples were solely processed with glass or metallic instruments. Furthermore, cotton gloves were worn over nitrile ones during all processing steps.
Figure 1. Sampling of intestinal samples: (A) The caudal part of the rectum is tied off; (B) An 8–10 cm section of the rectum is measured in cranial direction and tied off a second time; (C) The intestinal sample is cut with a scalpel and placed with metal tweezers in a glass jar (D).

2.1.1. Preparation

After defrosting the samples in the glass jars, the closed intestinal parts were rinsed with MilliQ water (Millipore), measured and opened inside the washing sachets on a glass cutting board to prevent the loss of faeces and potential MP.

For separating the MP from the biogenic matter, the intestine and faeces samples were washed in the above mentioned self-sewn double layer washing sachets in a commercial washing machine (OK., OWM 15012 A1). Each sample was placed in an inner bag (mesh size 300 µm) which was then placed in the outer bag (mesh size 100 µm). The sachets were made of nylon cloth and were sewn together in the acrylic box with a conventional sewing machine (Singer, Tradition TM 2282) using black cotton yarn. Furthermore, each nylon sachet was used only once to prevent a cross-contamination.

2.1.2. Washing Procedure

The washing procedure of the samples was based on the protocol established by Bravo Rebolledo et al. [17]. To remove biogenic organic matter, enzyme-based washing powder (Biotex®, stain removing powder, biological detergent, bleach free; 35 g) was added in the prewash cycle. Subsequently, conventional detergent (35 g, Gut & Guenstig Classic, Edeka Zentrale AG & Co. KG, Hamburg, Germany) was used in the main wash cycle, assisting the cleaning procedure. The samples were washed in a delicate wash cycle without spinning at 60 °C. A delicate wash cycle without spinning was chosen in order to prevent particles from escaping from the sachets and thus not being available for further analysis. For a valid evaluation, all samples were weighed before and after the washing procedure, to quantify the loss of biogenic compounds (e.g., blood, faeces, soft tissue).
Prior to each wash cycle, an empty wash cycle was run using disinfectant (Impresan, Brauns-Heitmann GmbH & Co.KG, Warburg, Germany) at 90 °C. This step cleaned the washing machine and additionally reduced the likelihood of cross-contamination.

2.1.3. Isolation

After the washing procedure, the washing sachets containing the sample residues were covered in aluminium foil and stored in a half-closed box under a fume hood for drying overnight. The outsides of the sachets were cleaned with MilliQ water prior to opening. The residue was rinsed with a filtered, saturated sodium chloride (NaCl) solution (350 g of table salt dissolved in 1 l of MilliQ water) in glass beakers. The solution was left overnight for density separation.

Approximately 20 to 30 mL of the surface of the supernatant was pipetted onto a cellulose filter (Rotilabo®, Typ11A, Ø 55 mm, retention 12–15 µm) using a Buechner funnel (Rotilabo®, porcelain, volume: 70 mL; Ø 55 mm) attached to a vacuum pump. The remaining solution and the sediment were decanted onto a second filter. Both surface and bottom sample suspensions were investigated separately in order to avoid an MP underestimation. The filtration steps resulted in a total of four filters for each sample (inner bag: surface water and water from the beaker bottom; outer bag: surface water and water from the beaker bottom).

The filters were stored in glass petri dishes (STERIPLAN®, Ø 60 mm) and dried in a heating cabinet for three hours at 50 °C. Until further processing, the closed petri dishes including the sample filter were stored in a closed and dry environment.

2.1.4. Pre-Trials to Verify the Procedure

The methodical efficiency was evaluated, focusing on potential losses of particles and secondary contaminations. The use of washing sachets was assessed for potential loss of particles and loss of nylon fibres and thus the recovery rate. To determine the loss of potential MP, polyethylene microbeads (fragment size: 150–800 µm) were placed in the nylon sachets and processed in the same manner as the intestine and scat samples.

Prior to sample processing, five sachets were solely washed without any detergent and sample (at 60 °C, delicate cycle, no spinning cycle as mentioned above). A Cora ball®, which traps loose fibres, was added and showed no attached fibres after the washing procedure.

2.1.5. Polymer Identification

For identifying possible MP, the filters were stained with Nile Red diluted in chloroform (1 mg/mL in chloroform) after Tamminga et al. [32]. The stained filters remained covered for drying for at least 24 h, before being analysed under a plan fluorescence microscope (Kern OBN- 148 and Zeiss AX10) using a TRITC HC filter set (F36-503, AHF Analyseotechnik, Tuebingen, Germany) in accordance with Tamminga et al. [32].

Potential MP particles were photographed (Kern ODC 832 and Canon EOS 80D) under the fluorescence microscope. Subsequently, the photos were visually evaluated, and the suspected MP were measured and counted by using Adobe Photoshop (Version 21.0.3). After visual identification, single particles were isolated with tweezers and pins, and placed on microscopic slides for further identification by µRaman spectroscopy (ThermoFisher Scientific Inc., Waltham, MA, USA; DXR2xi Raman Imaging Microscope).

In general, direct comparisons among different studies are hampered due to the varying sample treatment protocols and mesh sizes being used. To address this problem, the present study followed the sample processing conducted by Bravo Rebolledo et al. [17] and focussed on particle sizes ≥ 100 µm. In this study, the term MP included synthetic fragments and fibres. The classification of fibres was used, if a fragment has an elongated cylindrical shape and a thickness of up to 23 µm.

All particles larger than 100 µm were taken into account for valid results due to the used mesh size (100 µm) of the outer washing sachet to avoid an overestimation. Two researchers independently
investigated the photos taken of particles to identify the potential MP. The following parameters were used for discrimination: fluorescence intensity and appearance of the particle (surface, isolation, completeness, etc.). For additional information, see the Supplementary Materials.

Based on the aforementioned discrimination parameters, we used recursive partitioning to explore decision rules for predicting whether the particle was plastic or not. The decision tree was created in RStudio (Version 1.1.447) with the package “rpart” R package rpart.plot [43] with 1185 characterised particles (biogenic or synthetic). The number of particles resulted from available MP samples including pinnipeds and cetaceans from the North and Baltic Seas, where this study group was involved in. It builds the basis for the used parameters in the decision tree. The original dataset (n = 1185) was randomly split into training (80%) and test (20%) subsets. The training subset was used for model tuning and training and the performance of the final model was compared using the Test subset.

2.2. Evaluation of the Methodical Efficiency

In the present study, the entire working space was located in an acrylic box situated under a fume hood to decrease the potential secondary contamination of samples and used items. All equipment and instruments were rinsed several times with MilliQ water prior to usage. The box was washed with MilliQ water and wiped between each step with cotton cloths, which were exchanged regularly.

For monitoring secondary contamination, procedural blanks of the applied chemicals and detergents were used in accordance with Hengstmann et al. [44] and Nuelle et al. [45]. Procedural blanks helped to estimate the number of particles being introduced during the processing; particles on the procedural blank filters were separately quantified, measured and qualified. Contamination of cloths or instruments was deducted after the quantitative analysis as performed in former studies.

To prove the possible contamination of plastic bags, five faecal samples which had been frozen in low-density polyethylene (LDPE) bags since 2012 were used. Nine faecal samples were additionally included in this study. These samples were divided: half of each sample was stored in LDPE-bags, the other half in glass jars for three, six and 12 months. The plastic bags in which the samples are stored were not disinfected or cleaned prior usage. The comparison of the found number of particles in glass jars and plastic samples was performed using a Wilcoxon signed-rank test, due to the relation between the samples and the assumption of a non-normal distribution. To test whether the findings of suspected MP increased over time, the plastic samples taken from different periods were analysed by a non-parametric ANOVA using a Kruskal-Wallis test.

3. Results

3.1. Evaluation of the Methodical Efficiency

3.1.1. Procedural Efficiency of Washing Sachets

The pre-trials showed that a loss of 0.358 g (2.99%) was determined for those sachets, which were closed with a hook-and-loop fastener. Due to this small percentage loss, the usage of nylon cloth was maintained, but the hook-and-loop fastener was discarded and replaced by a seam. For further details, refer to the Supplementary Materials.

By comparing the dry weights of the samples (before and after the washing procedure), the average reduction in biogenic matter in intestinal samples differed between 9.97 and 0.22 g (mean = 1.57 g). In the faecal samples, the average reduction varied between 5.07 and 48.55 g (mean = 13.33 g).

3.1.2. Efficiency of Contamination Control Measures

Procedural blanks analysed for the background contamination of the working space showed a share ranging from zero to two particles per sample (M ± SD = 0.9 ± 0.99). Only one fiber larger than 100 μm was found on the working environment controls (n = 12). One to two fragments were found within the NaCl blanks (n = 5), which also included a potential contamination by the used
plastic laboratory bottles (M ± SD = 1.4 ± 0.99). Furthermore, five control washing sachets showed a contamination of two to at most eight particles (M ± SD = 4.6 ± 2.7). Therefore, a mean number of seven particles ≥100 µm (six fragments and one fiber) per sample was considered as contamination. Figure 2 shows the share of suspected MP particles found in the procedural blanks.

Figure 2. Suspected microplastics (MPs) from procedural blank controls (≥100 µm). Working environment: 12 control filter; Washing sachets: Five blank samples; NaCl Solution: Five control samples.

3.1.3. Comparison of Suitable Storage Methods for Avoiding Contamination

Each sample archived in a glass jar showed a lower amount of suspected MP compared with those stored in plastic bags (Figure 3). The Wilcoxon test revealed significant differences, showing that plastic stored samples included eight particles (≥100 µm) more compared with those stored in glass jars (particles ≥100 µm: p = 0.012; particles ≤100 µm: p = 0.0104).

Figure 3. Differential values in the quantity of suspected MPs (≥100 µm) in samples stored in plastic bags, compared to ones stored in glass jars from different storage times.
The direct comparison of samples stored for different periods in plastic bags (three, six and 12 months) revealed no significant increase in microplastic particles over time (Figure 3). This was due to a high variability in each period group. However, a comparison with samples, which had been stored in plastic bags for eight years, showed no significant differences to the present samples (Figure 4).

3.2. Protocol Validation

The results of the used decision tree (Figure 5) revealed the following parameters to be decisive: (1) If a particle could be associated with a biogenic shape or structure (e.g., crustacean limbs or algae). If yes, it was classified as no MP (No-Plastic). If not, the fluorescent colour of the particle (orange including red and yellow or white) defined the next classification (2). If the fluorescent colour was white, the particle was classified as suspected MP. If the colour was yellow or orange, the next category was the appearance: (3) blurred, melted or defined appearance. If it was defined, the particle was classified as plastic; if not it was identified as No-Plastic. If the fluorescent colour the second category was white, the particle was categorised as plastic. A catalogue of examples and categories is presented in the Supplementary Materials. Potential microplastic particles identified by using this decision tree are then analysed using µRaman spectroscopy.

The calculated sensitivity (true positive rate, TPR) of 84%, that measures the proportion of particles that were identified as possible MP and the specificity (true negative rate, TNR) of 69%, which measures the proportion of particles that were identified as being non-plastic, revealed a good performance of the decision tree.

3.3. Isolation of Potential MP in Intestinal and Faecal Samples

In total, 653 potential plastic particles were determined in ten intestinal samples of free-ranging harbour and grey seals, and nine seal faecal samples where the described procedure (sample collection, isolation and identification) was used. Due to the used mesh sizes of the washing sachets (100 µm and 300 µm), all particles smaller than 100 µm were excluded from the study, resulting in 255 identified suspected MP particles (70 fibres and 185 fragments) in 18 out of a total of 19 samples. The share of fibres ranged from zero to 35 per sample (M ± SD = 6 ± 7.4) when taking all specimens stored in glass jars into consideration (≥100 µm). The number of fragments (≥100 µm) in each sample stored in glass jars varied from five to 55 parts (M ± SD = 13.3 ± 11.3).
Some MP findings are shown in Figure 6. For a total of 31 suspected plastic particles, the polymer composition was verified via µRaman spectroscopy. 28 particles (90%) of these 31 particles were identified as plastic polymer. The following polymers were determined: Polyethylene (PE, \( n = 14 \)), polyethylene terephthalate (PET, \( n = 5 \)), ethylene-vinyl acetate (EVA, \( n = 6 \)), polyamide (PA, \( n = 1 \)) and polypropylene (PP, \( n = 1 \)). Those particles were collected randomly from all 19 named samples, thus 11% of the whole quantity of suspected MPs were analysed.

Figure 5. Example of a decision tree of the test set-up of 510 particles showing the decisive parameters.

Figure 6. Identified microplastic particles from intestinal samples stained with Nile red photographed under a fluorescence microscope using a TRITC filter: (A) A nylon fibre of the used nylon cloth; (B) a polyethylene terephthalate (PET)-fibre found in a grey seal; (C) an ethylene-vinyl acetate (EVA) particle found in a harbour seal; (D) a polypropylene (PP) particle found in a grey seal; (E) a polyethylene (PE) particle found in a harbour seal.
Figure 7 shows the higher quantity of particles found in intestinal samples of grey and harbour seals compared with the collected faecal samples where the species is unknown. It is significant that the quantity of suspected MP exceeded the maximal assumed secondary contamination of seven particles per sample (see dotted line in Figure 7).

**Figure 7.** Suspected microplastics in intestinal samples of harbour \( (n = 5) \) and grey seals \( (n = 5) \), in comparison to faecal samples from seals collected on the sandbank (Lorenzenplate, North Sea) \( (n = 9 \), no determined species per sample: faeces originated from either a grey or a harbour seal). All samples were stored in glass jars. The dotted line shows the maximal presumption of secondary contamination of seven particles per sample.

### 4. Discussion

An initial and essential step for studying the effects of MP particles in GITs of marine mammals is the detection and evaluation of their current presence and distribution. Apart from assessing the presence of microplastics in marine mammals, the main focus of this study was to improve the avoidance of secondary contamination during sample handling. This problematic nature of secondary contamination, which was also shown in previous studies \[17,26,29,44,46\], needs to be focused on to achieve genuine results. The presented protocol combines established protocols for isolating MP from biota samples \[17,29\] with new approaches for specifically avoiding secondary contamination.

The first step for avoiding contamination was to store the samples in glass jars, since they are prone to contamination when stored in conventional plastic bags. The present study presented evidence of this aforementioned fact. The results of the direct comparison between archived samples in plastic bags and in glass jars showed that a contamination by plastic bags is likely. The number of plastic particles did not significantly increase over time, but the basis of contamination from external sources seems to be higher when archiving samples in plastic bags instead of storing them in glass jars. It has to be taken into account that the used samples already show a basic microplastic contamination. This differs from sample to sample, resulting in the fact that the plastic burden of the samples also varies between the time periods. A conceivable parameter besides the polymer structure of plastic bags is the cleaning aspect. Glass jars were cleaned and disinfected prior to usage; plastic bags were used in their original state. Based on the results of this study, it is recommended to use glass jars instead of plastic bags or containers.

A closed working environment was created to improve the measures undertaken in previous studies to prevent secondary contamination. The procedural blanks analysed for the background contamination of the working space showed a very low number of particles (0–2 per sample).
Furthermore, the share of fibres was significantly lower in this study compared to similar approaches in marine mammals [20,21,47,48] or other marine species [33,49,50]. Additionally, it has been previously highlighted that fibres in particular are an indicator of secondary contamination, originating from the laboratory clothing and working environment [25,29,44–46,51]. Therefore, the acrylic box safeguarded the samples from direct contact with the laboratory environment and personnel.

The presented study revealed that closed sealing, like the washing sachet, reduces the loss of particles. To minimise the risk of losing particles as well as reducing the contamination of the samples, it is advisable to place the samples into a nylon cloth, which is then sewn together. Pre-trials revealed the application of hook-and-loop fasteners to be unsatisfactory (see Supplementary Materials); similar results are noted for rubber bands [52].

The usage of nylon cloths was inevitable due to the favourable mesh sizes and the fact that no solid material can be used in the washing machine, although polyamide (nylon) is one of the most abundant polymers [53,54]. Indeed, fibres of the nylon washing sachets can be easily identified showing a distinct fibre pattern and were therefore not included in our study. Thus, there was no need to exclude polyamides from this study if these polymers were found. To validate this assumption, a found fibre showing a distinct structure and fluorescence intensity compared with the fibres of the used nylon cloths was identified as polyamide (nylon) by μRaman spectroscopy. Therefore, the polymer identification via μRaman spectroscopy was used to manifest the presence of microplastics and is an already established and reliable application in microplastic analysis [37,55–57]. Thus, the benefits of this applied cleaning procedure (1. time and cost effective, 2. valuable results) outweigh the disadvantages of the used nylon cloths. The usage of a conventional washing machine and detergent was demonstrated as a successful application to speed up the purification procedure resulting in clean residues. Removing biogenic substances with the help of the washing procedure preserves the original particles, and it is less time- (approx. 90 min) and budget-consuming compared with the application of enzymatic or chemical solutions for the digestion of biogenic material. Thus, the reduction of biogenic compounds (e.g., faeces, blood, and soft remains of prey species) was effective to decrease the sample to the remaining hard parts (e.g., sand grains, fish bones and lenses, beaks, and microplastics), as also seen by the compared dry weights. These results are also shown by the comparison of dry weights before and after the washing procedure. In the intestinal samples, the intestinal tissue and the hard parts remained. A mean difference of 1.75 g was determined. In comparison to the faecal samples where a mean difference of 13.33 g was ascertained. The weights were mostly dependent on the quantity of hard parts (e.g., sand grains, fish otoliths, and lenses, bones or squid beaks) within the respective sample. Furthermore, the weight of pure faeces was higher in the sandbank faecal samples (mean = 19.52 g) compared to faeces included in the intestinal tissue (mean = 15.06 g). Firstly, these differences can be explained by the intestinal physiology of phocid species, with a short length of the large intestine where the chymus is dehydrated and stored as faeces until excretion [58,59]. Secondly, passage times of marine mammals have to be taken into account if an assessment of the microplastic burden in marine mammals is required [60]. The passage times of harbour seals and grey seals were estimated to last for 6.5 to 30 h [61,62]. Thus, the sandbank samples may include the faeces of a full egestion of the rectum. In contrast, the intestinal samples represent only a part of the actual excretion, which is here further analysed. Furthermore, a reduction in size and loss of microplastics is conceivable as is confirmed for otoliths [63]. Combining collected faecal and intestinal samples is advisable for further investigations as this will display the burden of microplastics in an extended way.

Besides the method, there are more side parameters which have to be considered while collecting samples from wild ranging animals. This study only investigated in a part of the intestinal system instead of examining the whole GIT or the stomach as performed in recent studies [20,47,64]. This step was chosen to enable a potential contamination-free sample although a complete examination of the residual GIT and the carcass can be conducted. The reason for this is the usage of one organ for only one assessment is applicable in the fewest of cases during necropsies. Another benefit of this strategy
is the unhindered necropsy of the whole individual, which enables an entire overview of the health status and possibly provides more background information.

The number of positive samples (suspected MP \( \geq 100 \, \mu m \), in glass jars) is higher than in similar studies of faecal samples of seal species from the northern and southern Pacific Ocean [21,65] despite it is being conformable with studies investigating marine mammals of the north-eastern North Atlantic region [20,47,66]. As a consequence of the results of the used statistical applications (precision of 94% and sensitivity of 84%), in combination with the 90% chance of a correct classification as MP, this study is assumed to be a valuable approach in MP monitoring in marine mammals.

The establishment of a decision tree helps to categorize different variables, which are helpful for untrained researchers to gain results that are comparable between research groups. This is essential to compare the burden in different marine environments and biota. Furthermore, the decision tree helps to narrow down suspected microplastic particles out of a bunch of a variety of particles in a sample, which are then used for further analysis. This is cost- and time-effective and a useful tool for encouraging decision makers to support necessary monitoring programmes.

In addition, the applied methods comprise a combination of interdisciplinary analyses: identifying microplastics and their consumed prey (diet analysis of hard parts). This is based on the conciliatory washing procedure; the retained hard parts can be used for prey species identification [52,67]. Since it is proven that microplastics are transported across the food web [14,16,68], it is assumed that in particular the contained contaminants accumulate due to absorption and leaching [69] in top predator species as it is already evidenced, for example, for organochlorides [70,71]. This allows new insights into the correlation between microplastic occurrence in marine top predator species and their prey, which serve as a vector.

5. Conclusions

The sample protocol applied within this study is applicable for isolating and identifying microplastics from samples of gastrointestinal tracts of marine mammals and is comparable with other approaches like diet analysis. Low numbers of detected fibres led to the conclusion that secondary contamination is low. This seems most likely to be an effect of using a closed environment, the acrylic box and the usage of glass jars as storage container. Furthermore, the used isolation procedure is less cost-intensive, more gentle and faster in comparison to an enzyme or chemical digestion. However, every working environment and samples of different species and different sizes pose different challenges for microplastic analysis.

It can be assumed that the current approach provides reliable results, reflecting the actual microplastic burden in the analysed species. Thus, this study presents a possible application for microplastic monitoring in marine mammal samples, which has to be extended to a higher number of samples to achieve convincing findings in different species and water bodies. Nevertheless, this method is also transferable to other mammalian species.

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