

MPsizeBase: a database for particle size distributed and size aligned environmental microplastic data

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Abstract

Understanding the sources, dispersion, and health impacts of microplastic (MP) pollution requires quantitative observations in terms of MP number and/or mass concentrations. Measurements of environmental MP fragments or fibers typically target variable size spans within the formal 1 to 5000 μm range, due to different sampling and detection techniques, and are therefore not directly comparable. In addition, the majority of measurements by microscopy techniques are number based, whereas life cycle analysis, numerical models for MP dispersion and policy evaluation, and plastic additive exposure estimates are mass based. Data analysis frameworks, based on measured MP particle number size distribution (PSD), have been developed to align and extrapolate measurements to a common MP size range (e.g. 1-5000 or 1-300 μm), which in turn allows direct data-data and data-model inter-comparison. In this study we present the MPsizeBase, a database of published MP number concentrations and PSD in the environment, including land, ocean and atmosphere. MPsizeBase uses the power law to fit observed cumulative MP number PSDs, and extrapolate and align observations over a common size range of choice. Assuming MP shape and volume approximations, MPsizeBase also estimates size-aligned MP mass concentrations. We illustrate how the MPsizeBase can be used for inter-comparison studies, and end with recommendations on MP size, shape, and PSD reporting, including length, width and surface area. We invite scientists to contribute old and new data to MPsizeBase for additional environmental compartments, including soils, sediment and biota.

Introduction

Recent unsustainable growth in plastic production, use and mismanaged waste is confronted with the ecological, social and health impacts it makes [1]. The petroleum and polymer industries are responsible for nearly 4% of modern global greenhouse gas emissions, a figure that may rise to 15% by 2050 [2] under the anticipated redirection of petroleum resources from the transport to the polymer sector. The miracle properties of synthetic polymers such as plasticity and durability do not last, and degradation of plastic produces microplastic (MP) particles [3]. MP, from one micrometer to five millimeters in size, is now found in all Earth surface environments, from the highest mountains [4] to the deepest ocean trenches [5]. MP enters our food, and the air we breathe, reaching our internal tissues, from the brain to our reproductive organs [6,7]. In the environment and inside our body, plastics release toxic chemicals such as additives and monomers that disrupt endocrine function and increase risk for neurodevelopmental disorders, reproductive birth defects, infertility, cardiovascular disease, and cancers [8]. In short, the unsustainable growth in plastic production and mismanaged waste has culminated in a plastic Ponzi scheme because economic activities are distanced,

spatially, temporally and socially from their environmental impacts [9]. A globally-binding policy instrument to curb plastic pollution is currently under negotiation at the United Nations.

Over the past ten years numerous sampling and analysis protocols have been developed for MP identification and classification. Many of these are microscopy based, including manual microscopy, fluorescence microscopy after Red-Nile staining [10], Fourier-transform infra-red (FTIR) and Raman micro-spectroscopy [11]. Microscopy techniques typically involve contrast-based digital image or spectral image analysis, resulting in statistical data on particle length and width, and a measure of MP ‘number’ concentrations expressed as MP counts per volume or per mass of sample matrix. In addition, FTIR and Raman micro-spectroscopy provide chemical identification of the polymer type, such as polyethylene, polystyrene etc. MP number concentrations can be converted to MP mass concentrations if the particle density and volume are known. Alternative techniques, such as pyrolysis-gas chromatography – mass spectrometry (pyGCMS) provide a direct measurement of polymer mass concentration, but do not provide information on particle size [12]. Studies that do provide MP particle size statistics generally do so for different size ranges, making the reported MP number concentrations or fluxes difficult to compare. Standardized data intercomparison and inter-conversion methods, from number to mass concentrations and vice versa, are therefore needed to address MP pollution and exposure [13,14]. MP mass concentrations, stocks and mass fluxes are for example needed for life cycle analysis of plastic and MP (OECD), for environmental plastic and MP dispersal budgets [15], for plastic additive exposure estimates [16].

In this contribution we present the CNRS MPsizeBase, a database of peer-reviewed MP number concentrations and particle number size distributions (PSD) in the environment. By extending established particle size alignment approaches [17,18], we illustrate how the measured PSD of individual studies can be used to extrapolate and intercompare MP concentration data. MPsizeBase currently contains 58 studies with 109 PSDs, representing 203 MP concentration datapoints for common microplastic shapes (fragments, fibers, films) in aerosols, atmospheric deposition, surface ocean, and subsurface ocean. The MPsizeBase also contains key metadata on sample type, sampling methods, geographical location, and many additional parameters, providing a long-needed catalogue of knowledge gathered by and for the microplastics community. We briefly discuss data uncertainty, limitations of PSD-based MP concentration extrapolation, and end with recommendations on PSD data reporting. Ideally the database will be expanded to include all environmental media, including sediments, soils and biota amongst others. Scientists have access to MPsizeBase via the web portal (<https://www.get.omp.eu/mpsizebase/>) and via Zenodo (10.5281/zenodo.17380284) [19], and can contribute new and published data via the web portal.

Particle size distribution and the power law

Intercomparing microscope-based MP number concentrations is challenging, because studies generally target different MP size ranges within the formal 1 to 5000 μm range. The observed range is generally dictated by the methods’ limit of detection (LOD) on the lower end, and by the sample volume and occurrence of less abundant larger MP on the higher end. In addition, studies tend to bin MP number observations using widely variable bin size ranges. Figure 1a,b illustrates an example where an imaginary (simulated) Raman ‘study A’ reports 888 $\text{MP}_{10-90\mu\text{m}} \text{m}^{-3}$ of air in the 10-90 μm range (10 μm bin size range), and an FTIR ‘study B’ reports 34 $\text{MP}_{250-1750\mu\text{m}} \text{m}^{-3}$ for the same sampling site in the larger 250-1750 μm size range (250 μm bin size range). Concluding that study A found higher MP concentrations than study B, at the same site, is potentially incorrect and symptomatic of comparing proverbial apples and oranges. Intercomparison of studies A and B must be done over identical size ranges.

An important starting point to intercompare MP studies is to examine the measured MP particle number size distribution (PSD), illustrated in Figure 1a,b for studies A and B as histograms on linear scales, and in Figure 1c on log scales. Observed particle number size distributions in nature have been commonly represented by a power law, $y = bx^a$ [20], with $a < -1$ which was first illustrated for marine MP [21] and observed by many studies since then. In this formulation, y is the number of particles having length in the infinitesimal range x to $x + dx$. An important property of the power law is that its exponential behavior (Figure 1a,b) is linearized on log scales ($\log(y) = a \cdot \log(x) + \log(b)$, Figure 1c.), so that intercept, b , and slope, a , can be obtained by linear regression. The slope reflects the relative abundance of small vs. large particles, with steeper negative slopes corresponding to more small particles. The intercept reflects the total number of particles, with higher values corresponding to more particles.

Data analysis and alignment frameworks, based on power law fits of measured MP PSD, have been developed to extrapolate MP concentration measurements to identical MP size range (e.g. 1 to 5000 or 1-300 μm), which in turn allows direct data inter-comparison [14,17]. Figure 1c illustrates that the individual PSD of studies A and B are both part of the same broader PSD once we normalize MP concentrations for (i.e. divide by) reported MP bin width (10 μm in A, 250 μm in B). In other words, both datasets can be fitted with the same power law equation $y = bx^a = 10,000x^{-2}$, where x is the representative MP length of a bin (estimated as the geometric mean of the bin boundaries in μm), and y is the bin-normalized MP number concentration (MP $\text{m}^{-3} \mu\text{m}^{-1}$). Now that we have fitted the bin-normalized BN-PSDs, we can extrapolate the observations of both studies A and B to the formal 1-5000 μm MP size range by integrating the surface area under the power law BN-PSDs in Figure 1c:

$$MP_{1-5000\mu\text{m}}^{\#} = \int_{1\mu\text{m}}^{5000\mu\text{m}} bx^a dx = \frac{b}{1+a} (5000^{1+a} - 1^{1+a}) \quad (\text{Eq.1})$$

where $b = 10,000$, and $a = -2$. Doing so will return identical $MP_{1-5000\mu\text{m}}$ concentrations of 9998 MP/m^3 for both studies A and B. Reported PSDs are therefore critical in normalizing and extrapolating MP concentration data so that we can intercompare studies over any MP size range. In the case of our example, studies A and B sampled the same air, but observed different parts of the MP PSD, and are therefore complementary, delivering identical MP concentrations after alignment. Although beyond the scope of this contribution, understanding variability in PSDs, in particular the exponent a that typically varies between -1 and -3, informs us on plastic fragmentation mechanisms, and on natural particle sorting processes such as sedimentation, emission and deposition [22–24].

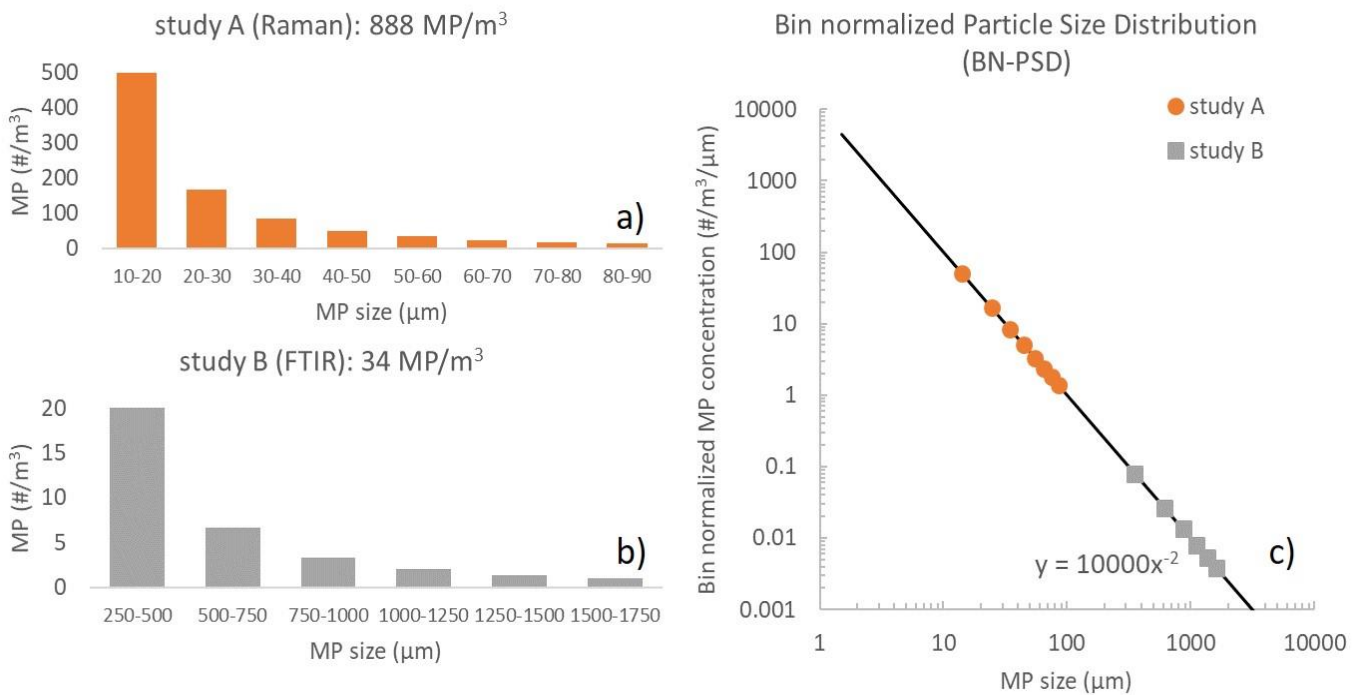


Figure 1a-c. Two hypothetical studies that observe MP number concentrations in air at the same site, but using different micro-spectroscopy techniques. a) Study A, using Raman, observes high a number concentration of small $MP_{10-90\mu\text{m}}$ (888 MP m^{-3}). b) Study B, using FTIR, observes a lower number concentration of larger $MP_{250-1750\mu\text{m}}$ (34 MP m^{-3}). By normalizing the observed MP particle size distributions (PSD) in a and b to their respective bin size (10 and 250 μm), we can see in c) that both studies define an identical bin-normalized BN-PSD, fitted with a power law as $y = 10,000x^{-2}$. Consequently, extrapolating the limited observed MP range, to the full, formal 1-5000 μm range (using Eq.1) aligns the observations and returns identical $MP_{1-5000\mu\text{m}}$ concentrations of 9998 MP m^{-3} for both studies A and B .

Binning bias in power law fitting

Real world, published MP PSD histograms do not behave like the ideal PSDs in the above hypothetical example (Figure 1), and it has been observed that power law slope b , and intercept a depend directly on the

143 chosen PSD bin size [18,24]. That is problematic because incorrect power law coefficients a and b lead to
144 erroneous MP concentration extrapolation using Eq.1, undermining the entire data alignment method. Binning
145 bias can technically be avoided by fitting raw MP number data [23], yet very few studies have published such
146 raw data and instead report binned PSD histograms. In MPsizeBase we apply a practical solution to the binning
147 bias problem by transforming the regular binned PSD to a complementary cumulative binned PSD [24,25].
148 The cumulative PSD (C-PSD hereafter) gives the number of particles having length $\geq x$. It can be computed
149 from a binned PSD by successively adding the bin's non-normalized MP number concentrations, starting with
150 the bin containing the largest particles and proceeding to the bin containing the smallest particles. This
151 results in a new log-log distribution that is linear for most of the PSD and is far less sensitive to bin width and
152 binning bias [24]. C-PSDs are shown in Figure 2 for three published datasets compiled in MPsizeBase,
153 alongside their BN-PSDs. Log-log C-PSDs have three important properties [24]: 1) the bin size, x is no longer
154 represented by the geometric mean, but the lower bin limit, 2) the C-PSD slope is shifted by +1 (less steep)
155 compared to the original non-cumulative PSD or BN-PSD, and 3) the right tail of the C-PSD distribution,
156 representing low counts for large MP, does not follow the power law, and therefore has to be omitted before
157 fitting. A drop in the right tail also occurs for regular PSDs and BN-PSDs, and will be discussed below in the
158 context of an upper limit of MP detection.

159 As mentioned, in log-log space, the C-PSD power law has a slope $a' = a+1$ (meaning the original BN-
160 PSD slope is shifted by +1) and an intercept $b' = -b/(a+1)$ [24]. The parameters a and b are therefore computed
161 from the slope a' and log-intercept, $\log 10^{(b')}$, obtained from linear regression of the C-PSD: $a = a'-1$ and $b =$
162 $-b'a'$. The extrapolated MP number concentration within the size range 1 to 5000 μm is again given by Eq.1
163 above. One can adjust the formal 1-5000 μm size range in Eq.1 to any size range of choice, such as 1-10, 10-
164 100, or 1-300 μm .

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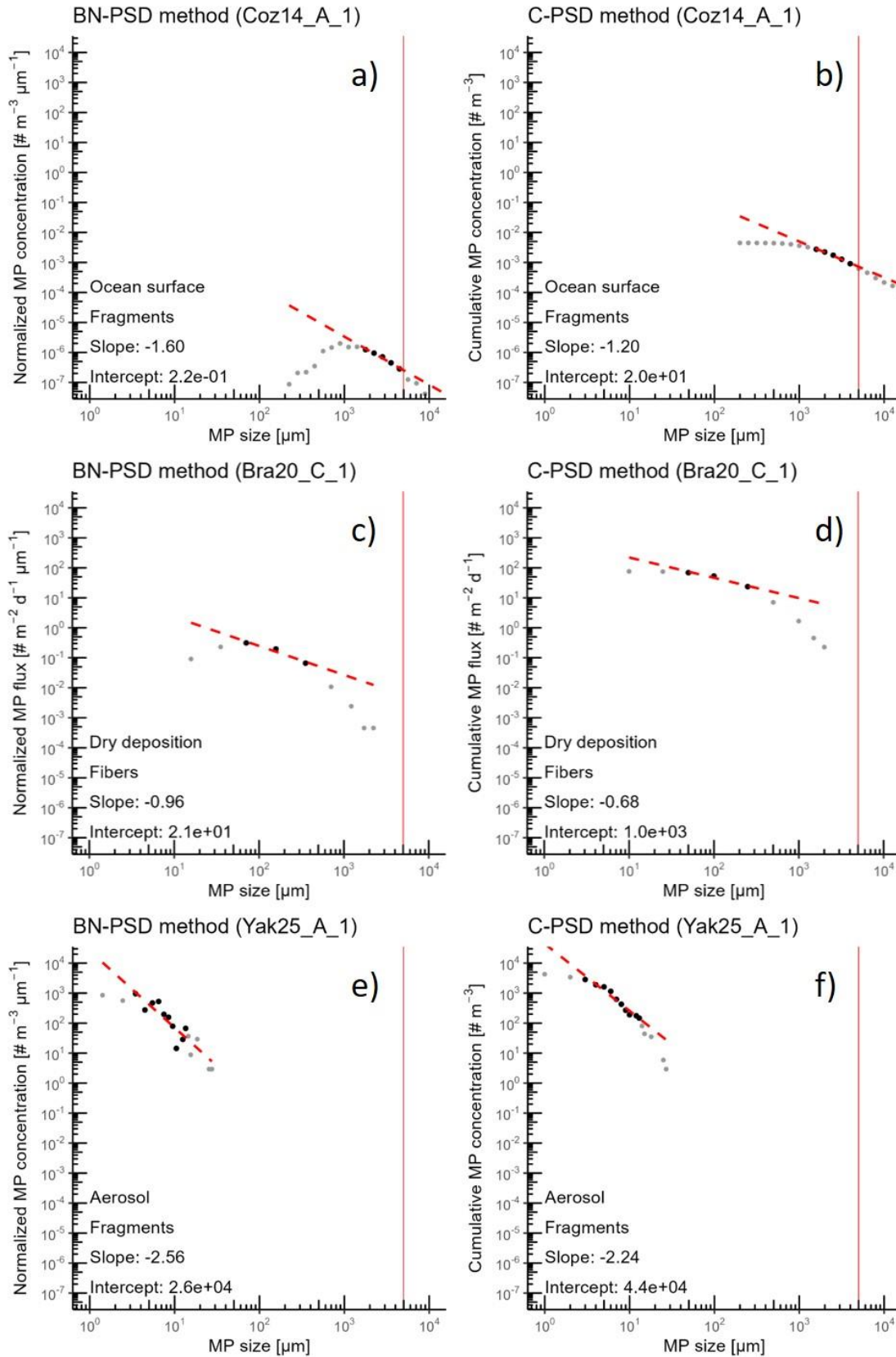


Figure 2a-f. Three published MP fragment particle size distributions (PSD) illustrating the bin-normalized particle size distribution (BN-PSD, left panels) and the complementary cumulative C-PSD (right panels). The three data sets illustrate common behavior for three different MP size ranges observed with three different techniques: a) and b) Coz_A_1 from 225-550,000 μm by digital microscopy [21], c) and d) Bra_C_1 from 18-2250 μm by FTIR microscopy [26], and e) and f) Yak_A_1 from 2-28 μm by Raman microscopy [27]. Fitted data points are shown in black, and delimit the lower and upper LOD, with data excluded from fitting in grey, according to criteria in Table 1. The vertical red lines highlight the formal MP range from 1 to 5000 μm . BN-PSD power law slopes are generally lower (steeper) due to binning bias, which is avoided in the C-PSD fits.

MPsizeBase data QAQC

The studies currently integrated into the CNRS MPsizeBase are all peer-reviewed studies from the recent scientific literature, covering sampling years from 2010 to 2023. Typically 2 to 3 reviewers, in addition to editors, have verified quality assurance and quality control (QA/QC) criteria. The majority of studies publish a single ‘generic PSD’ histogram for fibers and/or fragments that includes the MP counts of all measured samples in the study. We combine this generic PSD with the mean or median MP number concentration at a given location to estimate the absolute MP number concentration for each observed particle size bin. MPsizeBase metadata contains a ‘comment’ column, where we cite the original figures, tables, or text that provided PSD and MP concentrations. MPsizeBase includes all originally reported PSD data, except for ‘0’ MP counts that were removed. When studies report MP number concentrations at different geographical locations, or for different time periods at the same location (seasonality for ex.), or as a function of ocean depth, then we try to retain this spatiotemporal variability but repeatedly apply the same, single reported PSD to generate particle size distributed MP number concentrations. Reported MP number concentrations at multiple sites were often found to be non-normally distributed, and we therefore converted reported mean values to medians when necessary.

We included QAQC indicators, which are the number of MP counts included in the published PSD, and the approximate number of MP counts measured for each individual sample (filter). The former is often explicitly reported in Methods sections or with the PSD, while the latter was calculated indirectly by multiplying the reported mean/median MP number concentration, the volume of sample taken and the % of filter analyzed. Statistical considerations indicate that at least 96 particles need to be measured in order to obtain a 10% uncertainty on a single parameter of study, i.e. particle size [28]. In practice, true MP counts per sample are often lower than 96, and therefore studies generally sum up all MP observations and report a single size distribution. We currently do not exclude studies with low MP counts and consider that database users can use the QAQC indicators and additional statistics to decide if bias occurs, and if outlier studies should be removed from further data analysis. Database users are welcome to explore different LOD definitions and QAQC indicators.

Lower and upper LOD considerations during C-PSD fitting

As mentioned above, the measured PSD range is generally dictated by a methods’ limit of detection on the lower MP size end (lower LOD), and by the sample volume and occurrence of less abundant larger MP on the higher MP size end (upper LOD). Authors generally do not formally define their LODs and report all MP counts made in their PSD. This often generates lower than expected counts for the smaller PSD bins (lower LOD bias), and zero or low MP counts for upper size bins (upper LOD bias). We illustrate this in Figure 2 for three published PSDs acquired with three types of microscopy, stereo, FTIR and Raman, and over three corresponding size ranges. Both lower and upper LOD are important in fitting a C-PSD, and will be discussed in the following.

Lower LOD bias, originally observed for nearly all surface ocean water net tow samples, has historically been explained by a loss mechanism for small MP, such as sinking, biological uptake, or selective transport [21,29]. Recently, MP fragmentation models suggested that the smallest fragments are fewer because the “energy” required to produce small fragments is higher [30]. In parallel, analytical studies increasingly recognize that the various microscopy techniques, including manual stereo or digital microscopy, and FTIR and Raman micro-spectroscopy, have LODs related to the physical principles of the technique, operator bias, and sample matrix complexity [11,24,31]. Optical microscopy, and FTIR and Raman spectroscopy have approximate lower size LODs (i.e. the limit of accurately identifying all MP of a given size in a sample) of 50-500 μm , 10-50 μm , and 1-10 μm respectively. Figure 2b,d,f illustrates that lower LOD bias, i.e. the lower than expected MP counts in smaller bins, occurs across the entire MP size range and for all techniques. In particular, FTIR and Raman studies show that power law behavior of MP occurs across the full MP size range, sometimes down to the 1 μm lower limit. Experimental plastic fragmentation similarly shows that optical and electron microscopy and nanoparticle tracking analysis all show successive lower LOD bias, while at the same time documenting a MP and NP power law distribution across the full 0.1 – 5000 μm range [31].

Upper LOD bias, which is the deviation of large MP counts from power law behavior, manifests itself as noise, and a drop in counts in the righthand tail of a BN-PSD, and similarly as a drop in cumulative counts in the corresponding C-PSD (Figure 2a-f). To a large extent the upper LOD of the BN-PSD and C-PSD is

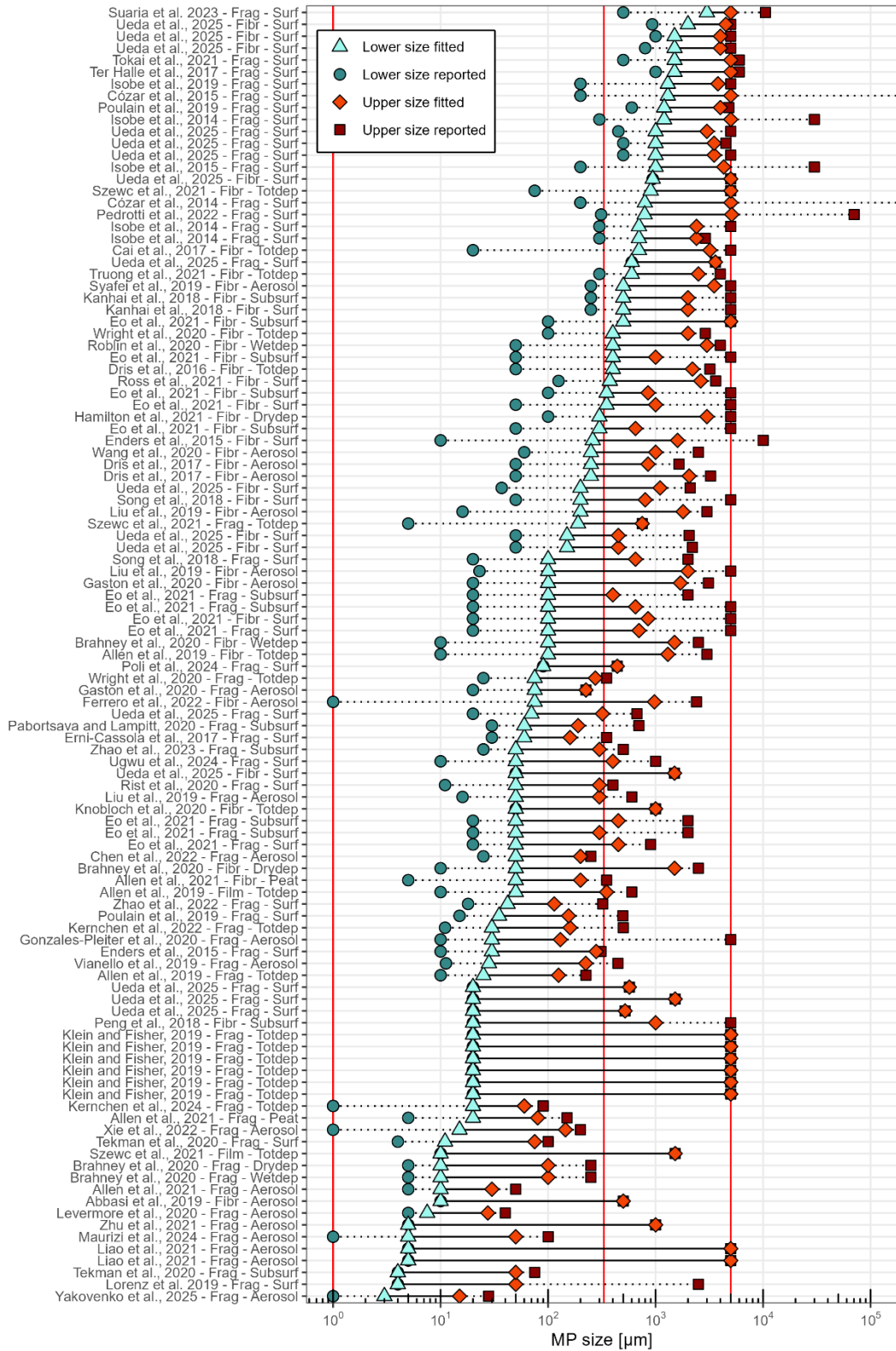
related to two factors: 1) The sampling volume and the % of the sample (i.e. filter area) analyzed. These two factors influence the probability of detecting 1 or more MP of large size. For example, if surface ocean MP particles of 4 to 5 mm are present at a concentration of 1 MP/m³, then a low volume 0.1 or 0.5 m³ pumped sample will not recover sufficient MP, thereby underestimating the true concentration. Such a lower than expected concentration causes the PSD tails to slump downward. Conversely, if a 1 m³ pumped or 100 m³ plankton net tow sample were analyzed, then 1 and 100 MP would be recovered respectively, potentially producing an accurate MP count and concentration. If, however, the 1 MP from the 1 m³ sample ends up on a FTIR filter, but only 10% of the filter is analyzed, then again chances are high that the single MP will not be found, biasing the 4-5mm MP counts low. 2) The second factor that causes a slump in the right-hand tail of the C-PSD is related to the sum of all large particles that have not been counted.

For fitting purposes, we use the algorithm of Segur et al. (2025), to identify the most linear part of the power law C-PSD in log-log space, and to assign the lower and upper LOD. The algorithm identifies log-normal outlier data < lower LOD and > upper LOD by minimizing residuals between C-PSD data points and an ordinary least squares regression fit. Finally, biased small MP data are identified with respect to the maximum MP count in the BN-PSD and removed from fitting, similar to previous studies [14]. In Figure 2a,c,e the MP counts above the upper LOD clearly define a slumping tail in the C-PSD. For datasets to be fitted and included in MPsizeBase, at least 3 bins must remain after LOD assignment. Table 1 summarizes the PSD data analysis and fitting pipe line for published datasets included in the MPsizeBase. Figure 3 gives an overview of the reported MP size range for all 109 PSDs and the assigned LODs.

Although there is today growing consensus that the full 1-5000 μm size range of MP obeys the power law [23,24,31], there are numerous physical processes that induce environmental MP particle sorting (sedimentation, emission, deposition), which potentially affects the PSD shape and slope [22]. Log-normal (i.e. curved) PSD shapes are in fact common phenomena in geology [32], while atmospheric aerosols other than MP generally follow a multimodal size distribution represented as the combination of multiple log-normal distributions [33]. We alert MP scientists that our attribution of a lower LOD to the data bin at or above the maximum MP count in the BN-PSD is in fact an argued assumption. This assumption, that environmental MP PSDs follow a power law, facilitates data alignment, yet we should, as a community, remain receptive to changes in this assumption and paradigm in the future.

Table 1. PSD data analysis and fitting pipe line for published datasets included in the CNRS MPsizeBase

Step	Action
1	Recover published MP number PSD from Figures (digitize) or Tables; recover MP number concentration, and metadata. Have the extracted data validated by a 2nd person.
2	Convert the original, reported PSD to both a bin-normalized BN-PSD, and to a cumulative C-PSD. Visually examine the BN-PSD on a log-log scale, to examine log-linear (straight) and log-normal (curved) features of the PSD in relation to lower and upper LODs.
3	Use the automated C-PSD algorithm from Segur et al. (2025) to assign lower and upper LODs, and fit the C-PSD on a log-log scale with the linearized power law. Convert the C-PSD fitted slope a' and intercept b' to regular PSD slope a and intercept b as follows: $a = a' - 1$ and $b = -b'a'$.
4	Use the PSD fit parameters a and b to extrapolate, via Eq.1, the observed MP number concentration to any size range of interest, such as the formal MP _{1-5000μm} window, or other 1-10, 10-100 μm windows.
5	Use a and b in Eqs. 5-7 to derive MP mass concentrations for the size range of interest.



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Figure 3. Size range reported by all 109 PSDs currently compiled in MPsizeBase. The initial lower size marker (dark blue circle) is the minimum MP size reported by a study. The filtered lower size (light blue triangle) is the lower MP size used for power law fitting and represents the lower limit of detection (LOD). The upper MP size reported is shown as brown squares, and the upper LOD fitted as red diamonds. Operational size ranges for small microplastics (SMP, 1 to 300 μm), large microplastics (LMP, 300 to 5000 μm) and meso and macroplastics (P, > 5000 μm) are delimited by vertical red lines. LMP correspond approximately to the plankton/neuston net mesh for surface water MP sampling and the upper size range of MP fragments that can be emitted by oceans via the bubble bursting mechanism. The sample types are Surf (surface ocean waters), Subsurf (subsurface ocean waters) Aerosol (atmospheric MP) for studies reporting concentrations, and Totdep (Total deposition), Wetdep (Wet deposition), Drydep (Dry deposition) and Peat (peat archive total deposition) for studies reporting atmospheric MP deposition.

MP number to mass conversion

Numerous studies in the literature convert MP number to MP mass concentrations by assuming an average MP weight, which is typically based on the measured average MP size. Because environmental MPs display a continuum of size and shape properties, MP volume and weight can vary by 11 orders of magnitude, from ~0.3 pg for the smallest 1 μm MP fragments, to ~10 mg for the largest 5000 μm MP fragments. Assuming an average MP weight, without taking into account the full PSD, therefore leads to very large uncertainties in MP mass concentration estimates, because both small and large MP are assigned the same average weight.

It is therefore more accurate to do a number to mass conversion by integrating over and extrapolating beyond the observed PSD using the power law. To convert a MP number concentration to a MP mass concentration, the particle density and volume need to be known for the measured MPs. Polymer density, ρ , varies relatively little (0.9 to $1.4 \times 10^{-6} \mu\text{g}/\mu\text{m}^3$) and is set here to $1.14 \times 10^{-6} \mu\text{g}/\mu\text{m}^3$ based on [17], but volume varies substantially according to particle shape and size. MPsizeBase includes observations of three common microplastic shapes: fragments, fibers, and films. Fragment MP volume depends on MP length, width (L , W , measured) and height (H , generally not measured). The majority of MP studies included in MPsizeBase unfortunately only reports L distributions (and not W). Dedicated MP morphological studies have observed that the median fragment W/L ratio is 0.67 ± 0.03 , indicating an ellipsoid MP shape [13,23,34–36]. It is often further assumed that ellipsoid $H/W = W/L$ [14,36], with a correction for surface irregularities based on the measured MP perimeter [13]. Recently, an ellipsoid volume model was calibrated for small MP and the special case when only L is known, and displayed $H/W = 0.40 \pm 0.08$ [35]. Therefore, in MPsizeBase, when MP fragment L is explicitly reported as the major MP axis, or major Feret diameter, we approximate V based on the ellipsoid volume:

$$V_{\text{ellipsoid}} = \pi/6 \times L \times W \times H = 0.097 \times L^3 \quad (\text{Eq.2})$$

, where $W = 0.68 \times L$ and $H = 0.40 \times W$ [37]. When fibers are reported, studies typically provide the PSD for fiber length, but rarely report the fiber diameter (D). If fiber D is not reported, we assume it to be 15 μm [38], which is a typical mean D measured in select studies [39]. For fibers with L shorter than 45 μm, we assume $D = L/3$, in order to respect the common definition of fiber aspect ratio, $L/W > 3$. Similar to previous studies we also assume environmental MP fibers to have a 40% void fraction [13,36]. MPsizeBase users are welcome to explore alternative volume and mass estimates in their studies. For MP film data we assume film thickness (H , height) to be 17 μm, which is the standard thickness of grocery bags, the most common littered film item [40]. Fiber and film volume expressions in MPsizeBase are then defined as:

$$V_{\text{fiber}} = \pi \cdot (D/2)^2 \cdot L \cdot 0.6 \quad \text{Eq.(3)}$$

$$V_{\text{film}} = \frac{\pi}{4} L \cdot W \cdot H \quad \text{Eq.(4)}$$

, where for reported film data, $W=L$, and 0.6 refers to the non-void fraction of fibers. New studies reporting MP number PSDs are suggested to provide both L and W size distributions, or even better: single particle raw data in the Supporting Information, including L , W , area, and perimeter (and H , if at all possible). This will greatly aid in making more accurate MP count to mass conversions, which currently have a substantial uncertainty.

In MPsizeBase we estimate aligned, extrapolated MP mass concentrations by extending Eq.1 to include density and volume approximations (Eq.2-4), slope and intercept from the C-PSD fitting strategy, and then integrate (see [24] for details):

$$\text{fragments: } MP_{1-5000\mu\text{m}}^{\text{mass}} = \rho \cdot 0.097 \cdot \frac{b}{4+a} (5000^{4+a} - 1^{4+a}) \quad (\text{Eq.5})$$

$$\text{fibers: } MP_{1-5000\mu\text{m}}^{\text{mass}} = \rho \cdot \pi \cdot 0.6 \cdot (D/2)^2 \cdot \frac{b}{2+a} (5000^{2+a} - 1^{2+a}) \quad (\text{Eq.6})$$

$$\text{films: } MP_{1-5000\mu\text{m}}^{\text{mass}} = \rho \cdot \frac{\pi}{4} H \cdot \frac{b}{3+a} (5000^{3+a} - 1^{3+a}) \quad (\text{Eq.7})$$

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326 where $MP_{1-5000\mu m}^{mass} [\mu g m^{-3}]$ is the MP mass concentration within the size range 1 to 5000 μm , ρ is the MP
327 density of $1.14 \times 10^{-6} \mu g/\mu m^3$, D is fiber diameter, H is film height, and a and b are the slope and intercept
328 derived from C-PSD fitting. Note that we assume fibers have length $\geq 3xD$ and films have length $\geq 10xH$.
329 Again, one can adjust the formal 1-5000 μm size range in Eqs.5-7 to any size range of choice.

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331 **Data alignment uncertainty**

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333 It is widely acknowledged by the MP community that concentration measurements are difficult to make, due
334 to the variability and complexity of particles in the natural environment. Generally, MP are minority particles
335 that need to be separated and pre-concentrated before identification is possible by microscopy. The many field
336 and laboratory manipulations involved in this pre-concentration increase both the risk of contamination and
337 MP loss. While most studies quantify and correct for contamination, procedural MP loss is rarely monitored,
338 and standard operating procedures and certified reference materials remain under development [41,42].
339 Moreover, identification of MPs in natural samples often relies on comparison with spectral libraries (FTIR,
340 Raman, PyGCMS) which are incomplete and contain select polymers or additives only [11,43]. Multiple
341 reference libraries exist and thus, two studies reporting MP counts or mass and polymer type may not have
342 used the same libraries. The published MP data compiled in MPsizeBase, covering the period 2014-2023,
343 therefore reflects the ensemble of sampling and measurement uncertainties, even before C-PSD data
344 alignment.

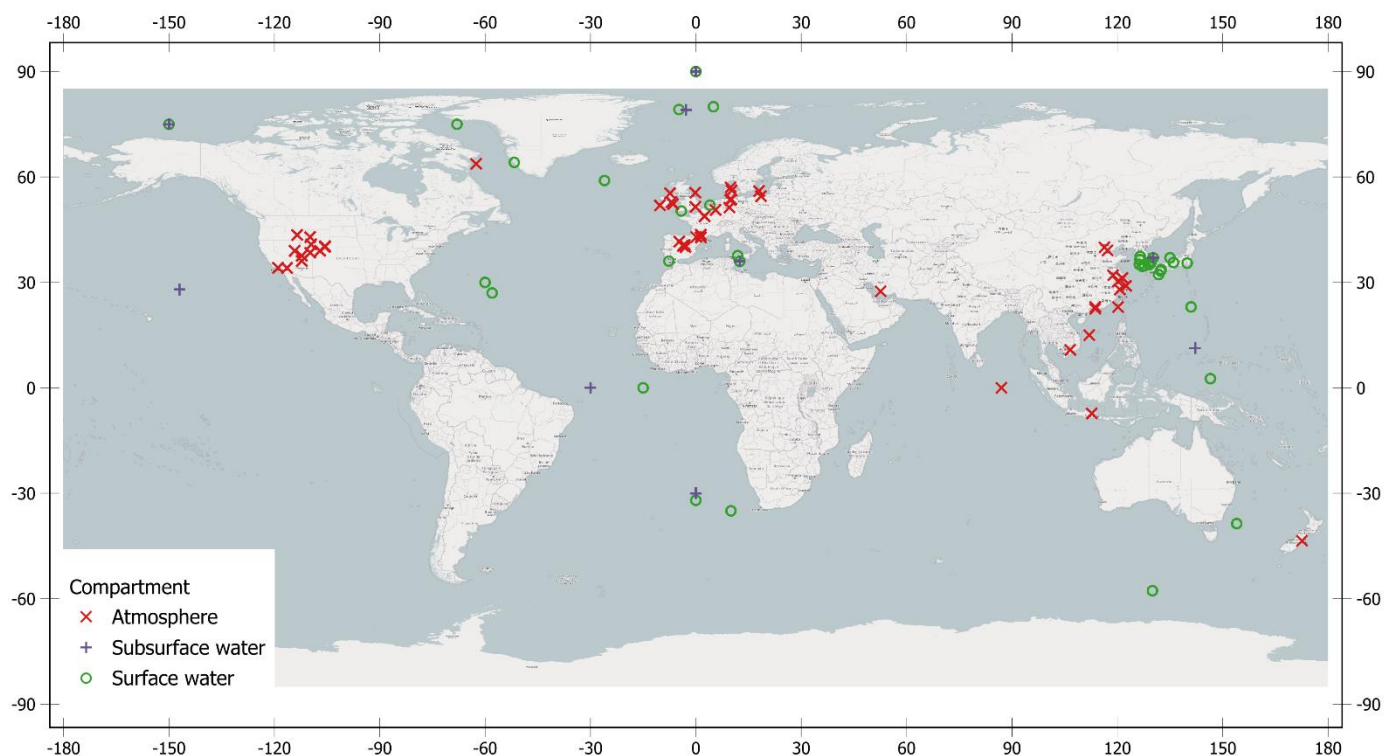
345 In MPsizeBase we currently estimate the standard error (SE) of C-PSD slope and intercept, and
346 propagate those SEs during extrapolation from observed to expanded $MP_{1-5000\mu m}$ number and mass
347 concentrations. Across all data, this leads to an average uncertainty of around 100% on $MP_{1-5000\mu m}$ number
348 concentrations, and around 55% on $MP_{1-5000\mu m}$ mass concentrations. The reason for the difference is that MP
349 numbers are dominated by the mostly unobserved 1-10 μm size range, requiring extrapolation and thus higher
350 uncertainty, while MP mass is dominated by the typically observed 500-5000 size range, requiring less
351 extrapolation. Alternative, more complex Monte Carlo based uncertainties can be evaluated by including or
352 excluding additional data points in the C-PSD power law fit. We consider however that presently the largest
353 contribution to the uncertainty of an aligned $MP_{1-5000\mu m}$ concentration is not the C-PSD power law
354 extrapolation, but the inherent methodological variability of the original measurements. By compiling as many
355 datasets as possible, MPsizeBase users can identify outlier data with standardized statistical tests among the
356 aligned datasets.

357

358 **Metadata**

359

360 The MPsizeBase includes different classes of metadata: Full reference and doi; General sample type (ocean,
361 atmosphere, sediment, soil, biota,...), various subsample types (surface, subsurface, indoor, outdoor), and
362 operational descriptors for fragments, fibers, films or pellets; Method details, including MP sampling,
363 extraction and detection details, and sample ID; Geographical information (Figure 4), including latitude,
364 longitude, altitude, continent, ocean basin, country, location, sampling dates; Ancillary parameters, such as
365 population density etc. Scientists contributing new datasets can suggest columns with additional parameters.
366



367
368 *Figure 4. Map showing the geographical locations of ocean and atmosphere samples included in MPsizeBase*
369 *v1.0. Ocean surface water includes the 0-50m surface mixed layer; ocean subsurface water refers to all*
370 *samples below 50m depth. Because most studies pool all sample MP counts into a single PSD, much*
371 *geographical MP concentration variability (e.g. an oceanographic cruise track) is assigned a single location,*
372 *or grouped into 2-4 geographical zones.*

373 374 375 **Results and Discussion**

376
377 Supporting Information 1 (SI-1) provides a compact summary of all 58 MPsizeBase studies, 109 individual
378 PSDs, metadata, and extrapolated $MP_{1-5000\mu m}$ number and mass concentrations. MPsizeBase users can adjust
379 the standard 1-5000 μm extrapolation size range in SI-1 to integrate over a different size range. Alternatively,
380 users can adjust the size range in the most up to date online version of MPsizeBase. Table 2 summarizes MP
381 PSD properties, C-PSD fitting results and extrapolated $MP_{1-5000\mu m}$ number and mass concentrations and
382 atmospheric deposition fluxes for the major and sub-environmental MP categories in MPsizeBase (e.g. fibers,
383 fragments, coastal, offshore, etc). Several key features are: 1) The systematically higher extrapolated MP_{1-}
384 $5000\mu m$ number and mass concentrations compared to the original reported observations for ocean and
385 atmosphere. In general, the larger in size the observed MP number data are, the higher the extrapolation factor
386 is, due to accounting for non-observed small MP. 2) The significantly higher fragment $MP_{1-5000\mu m}$ number (8.8
387 $10^4 MP m^{-3}$) and mass ($1131 \mu g m^{-3}$) concentrations in coastal surface waters compared to offshore surface
388 waters ($787 MP m^{-3}$, $10 \mu g m^{-3}$). 3) The significantly different PSD slope between fibers (-1.86 ± 0.36) and
389 fragments (-2.66 ± 0.68) in all major data groups, which partly reflects the dimensionality of MP fragmentation
390 [24].

391 Figure 5 visualizes measured atmospheric MP deposition fluxes in relation to extrapolated size-aligned
392 $MP_{1-5000\mu m}$ fluxes for the formal 1-5000 μm MP range. It illustrates how aligned $MP_{1-5000\mu m}$ data are generally
393 higher than the original MP measurements. Similar visual comparisons between measured and extrapolated
394 $MP_{1-5000\mu m}$ MPsizeBase data for the surface and subsurface ocean, and aerosols can be found in Segur et al.
395 [24]. Aligning different published MP measurements to the same MP range, whether it is 1-10, 1-300, 1-5000
396 μm or other ranges, makes it possible to intercompare data sets, and compare observed MP concentrations to
397 modeled MP concentrations. For the atmosphere, the upper size limit of extrapolation can be adapted
398 according to different environments: Table 2 reports that both fiber and fragment deposition, especially in
399 urban environments, includes large MP up to 3100 and 1650 μm on average. Aerosol MP fragment

observations indicate a lower average size limit around 200 μm , which would justify a lower extrapolation limit of perhaps 500 μm as observed also in experimental marine MP emissions studies [44].

The average PSD slopes for fibers and fragments reported in Table 2 can also be used to extrapolate and align MP number and mass concentrations for datasets without PSDs, following published methods [17]. A straightforward way to use such a ‘generic’ slope (Table 2) to size-align MPsizeBase or other datasets, is to apply the generic slope a , observed MP number concentration and size range in Eq.1, and calculate the intercept b . Next, slope a , and intercept b can be used in Eq.1 and Eq’s. 5-7 to estimate the size-aligned MP number and mass concentrations for a size range of choice. Technically, a small MP bias correction must be made for reported MP number concentrations, because it is generally underestimated due low counting bias (Figure 2). In addition, the generic PSD slopes (Table 2) can also be used to align MP mass concentrations by thermo-analytical mass concentration measurements (pyGCMS or other) provided that the analyzed MP size range, by pyGCMS, is known from field and laboratory filtration procedures, such as lower and upper filtration cutoff.

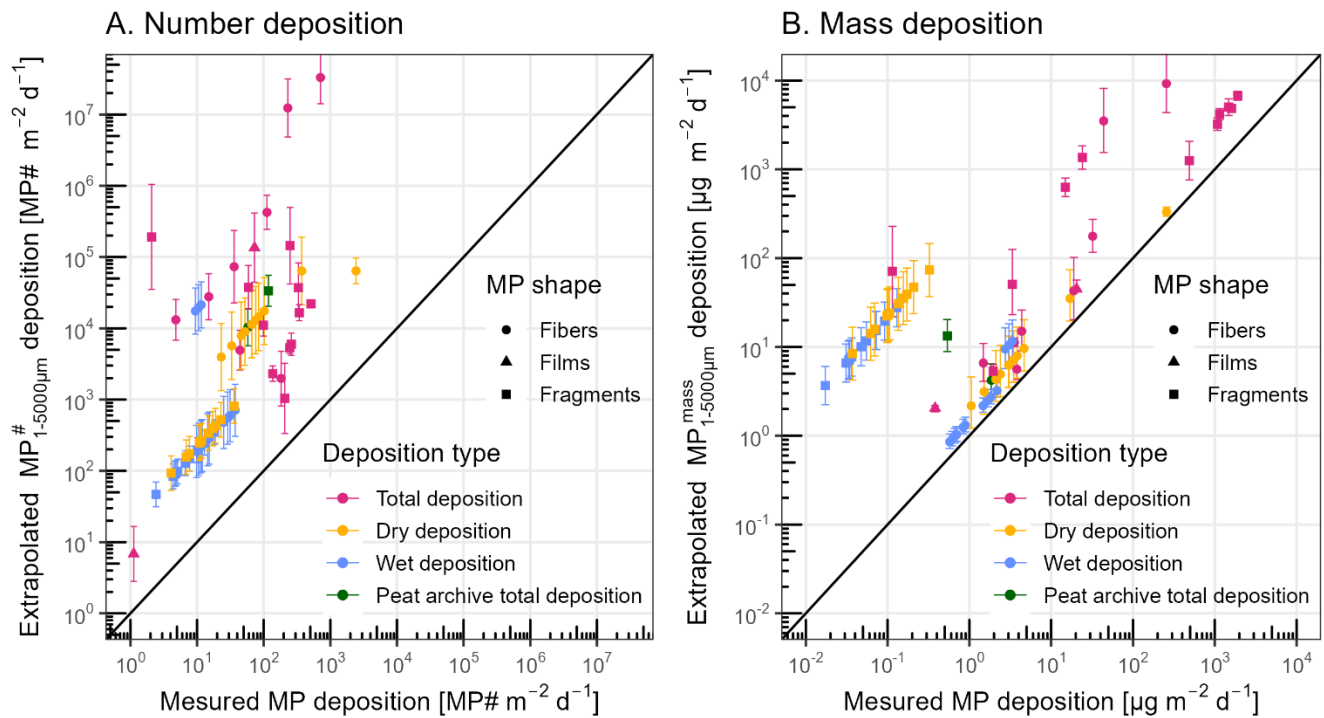


Figure 5. Comparison of reported, measured atmospheric MP deposition flux to extrapolated $\text{MP}_{1-5000\mu\text{m}}$ deposition flux in A. number ($\# \text{ m}^{-2} \text{ y}^{-1}$) and B. mass ($\mu\text{g} \text{ m}^{-2} \text{ d}^{-1}$). Because only a fraction of the MP size spectrum between 1 and 5000 μm is measured, the extrapolated MP number and mass fluxes are generally higher.

Table 2. Summary of C-PSD data alignment for MP number and mass concentrations in surface ocean (coastal and offshore), subsurface ocean and atmosphere (aerosols, and total deposition, i.e. sum of wet and dry deposition). ‘Coastal’ corresponds to locations over the continental shelf (<140 m depth), n data points = number of samples or sites included, n PSD = number of particle size distributions across all data, rep = reported, est = estimated, # = number, IQR = interquartile range.

		n data points	n PSD	PSD min	PSD max	#MP/PSD	#MP/sample	PSD slope	PSD intercept		MP#	$\text{MP}_{1-5000\mu\text{m}}$		$\text{MP}^{\text{mass}}_{1-5000\mu\text{m}}$		
				rep	rep	rep	est	est	estimated	reported	estimated	estimated				
				median	median	median	median	mean	SD	median	IQR	median	median	IQR	median	IQR
		#	#	μm	μm	#	#					$\#/\text{m}^3$	$\#/\text{m}^3$			$\mu\text{g}/\text{m}^3$
Surface Ocean																

Fiber	Coastal	20	12	50	3 500	118	22	-2.13	0.42	2.3 10 ⁴	3.9 10 ³	5.7 10 ⁴	40	1.9 10 ⁴	4.6 10 ³	5.0 10 ⁴	20	4	42
Fiber	Offshore	4	3	125	3 500	117	4	-1.91	0.38	2.1 10 ⁴	1.0 10 ⁴	4.6 10 ⁴	51	1.9 10 ⁴	9.6 10 ³	4.0 10 ⁴	15	8	27
Fragment	Coastal	28	21	30	2 300	432	198	-2.73	0.58	1.5 10 ⁵	2.4 10 ⁴	9.1 10 ⁶	27	8.8 10 ⁴	1.8 10 ⁴	3.7 10 ⁶	1131	493	3719
Fragment	Offshore	15	11	200	4 950	543	4	-2.44	0.58	1.1 10 ³	306	1.1 10 ⁴	0.06	787	204	5.7 10 ³	10	5	81
Subsurface Ocean																			
Fiber		14	7	100	3 500	82	4	-1.92	0.39	2.2 10 ⁴	1.2 10 ⁴	3.7 10 ⁴	42	1.9 10 ⁴	1.7 10 ⁴	3.1 10 ⁴	17	14	26
Fragment		18	8	20	1 000	412	18	-2.81	0.62	3.3 10 ⁵	3.8 10 ⁴	2.7 10 ⁶	116	1.9 10 ⁵	2.5 10 ⁴	1.2 10 ⁶	1281	808	2390
Atmosphere - Aerosol																			
Fiber		11	9	23	2900	114	16	-1.91	0.41	73	5	335	0.9	58	9	488	0.13	0.03	1.8
Fragment		22	14	5	206	436	55	-2.70	0.88	4.2 10 ³	710	3.3 10 ⁴	37	2.9 10 ³	416	1.8 10 ⁴	56	6	942
Atmosphere - Deposition																			
Fiber		122	9	50	3100	556	34	-2.23	0.38	3.4 10 ⁴	7.2 10 ³	5.9 10 ⁵	91	2.8 10 ⁴	1.2 10 ⁴	4.2 10 ⁵	15	8	177
Fragment		87	12	20	1653	437	40	-2.47	0.46	1.9 10 ⁴	4.6 10 ³	5.7 10 ⁴	192	1.4 10 ⁴	4.6 10 ³	3.7 10 ⁴	1305	66	4354

Reporting MP data in the literature

Based on our experience in compiling MP observations, and developing the C-PSD data alignment framework, the following recommendations can be made for publication of MP number and PSD data:

- Provide raw MP particle size data in numerical format (.xls, .csv or other) as part of the supporting information or in an online data repository. Raw data, unbinned, provides the best estimate of MP number PSD slope and intercept [23,24].
- Provide separate PSD's for MP fragments, fibers, films and other shapes (pellets, foams).
- If possible, report particle size data for each sampling location (provided the sample size is large enough)
- When observing fiber PSD, also report the mean/median fiber diameter, or better, the diameter size distribution. For MP films, the thickness would be of interest to measure and report.
- Provide, if possible, MP fragment length, width, area, perimeter and circularity measurements and PSDs (as raw data, and as binned histograms), which allows better estimates of height and volume, thereby limiting uncertainty of the number to mass concentration conversion [13,34,35].
- Provide estimates of the measurement technique lower and upper LODs.
- Provide metadata in practical format, including latitude and longitude in decimals (not degrees)
- Scientists are welcome to use the online MPsizeBase data template directly as Supporting Information with MP papers. This will facilitate rapid transfer to MPsizeBase and data use by the community.

In theory, a different PSD can be reported for every single sample, if sufficient MP counts are made to make that PSD robust. We recall that at least 96 particles need to be measured in order to obtain a 10% uncertainty on a single parameter of study, i.e. particle size [28]. We recommend that if sufficient MP counts are made, it could be of great scientific interest to report raw data PSDs as a function of time, location, water depth, soil depth etc, in order to understand how physico-chemical processes sort and fractionate MP size, and where and how MP fragmentation takes place. An example of detailed raw data PSD analysis and slope variability in the aquatic environment can be found in Kooi et al. (2021). We emphasize that binning raw data leads to information loss on PSD and on MP size sorting (i.e. changes in PSD) to the point that we do not detect significant differences in PSD slope between aerosols, atmospheric deposition, surface ocean, and subsurface ocean. Understanding size sorting of MP in the environment is key because it controls both plastic mass transport of large MP, and biota exposure and uptake of small MP. Ideally, over the next decade the MP community consistently publishes raw PSD data, so that we can collectively tap into the full information contained within PSDs, and test data alignment frameworks that go beyond the power law.

Most spectroscopy or pyGCMS studies report MP polymer identity, which is a parameter that is under development in MPsizeBase, and therefore not discussed in detail here. In principle, if spectral polymer identification is accurate, there should be little intercomparison issues to be expected between FTIR, Raman or pyGC-MS. In reality, spectral identification by comparison to a spectral library is fraught with uncertainty because biological polymers and additives have similar spectral features as plastic polymers. Incorrect assignment of measured particles as MP polymers, or not, directly leads to over- or underestimation of MP

number and mass concentrations. Expert spectroscopists are welcome to help add polymer identity to MPsizeBase, explore the associated uncertainty aspects, and explore potential relationships between polymer identity, PSDs and metadata.

Database use and development

Users will have open access to the latest validated version of the MPsizeBase here: <https://www.get.omp.eu/mpsizebase/> and Zenodo (<https://doi.org/10.5281/zenodo.17380284>) [19]. Users are asked to cite this publication and the doi associated with the version of the database; this in turn helps us secure the funding necessary to expand and maintain the database. Users can submit previously published and new peer-reviewed binned PSDs and raw PSDs on the MPsizeBase website. Users can also become active members of the MPsizeBase development group, and propose new approaches, concepts or parameters to include and explore.

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