

Intercomparison of dissolved trace elements at the Bermuda Atlantic Time Series station

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1	Intercomparison of Dissolved Trace Elements at the Bermuda Atlantic Time Series
2	Station
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- 30
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37 Abstract

38 The international GEOTRACES program is emphasizing the use of the GEOTRACES 39 reference samples program and the importance of cross-over or baseline stations where all 40 aspects of sample collection, filtration, processing and analytical methods can be compared. 41 These intercomparison efforts are of crucial importance to merge or link data from different 42 origins together. The occupation of the Bermuda Atlantic Time Series (BATS) station 43 (31°45.92'N, 64°04.95'W) by the Netherlands GEOTRACES (GA02; 13 June 2010) as well 44 as US GEOTRACES (GA03; 19-21 November 2011) Atlantic sections expeditions provided 45 an intercomparison opportunity to confirm the compatibility of the 2 different sampling 46 systems as well as different analytical techniques used. In order to compare the data from 47 different sampling systems or analytical techniques, a new statistical approach was developed 48 to include the analytical uncertainty. Furthermore, modifications to an existing multi-element 49 technique (Biller and Bruland, 2012) were made that increased the number of elements 50 analysed and that allows the time consuming extractions to be done shipboard. Overall, we 51 show excellent agreement between data generated by different sampling systems and 52 analytical techniques. However, while both sampling systems are capable of collecting 53 uncontaminated samples for all the GEOTRACES key elements, some apparent outliers are 54 present. The intercomparison between the different analytical techniques also reveals subtle 55 differences that would have gone unnoticed if only reference samples would have been used, 56 underlining the importance and power of intercomparison stations. The results show cross-57 over station data can be used to assess consistency between datasets if these stations have 58 been carefully planned and analysed in combination with (internal) seawater reference 59 samples to assure intra-dataset consistency.

60

62 Introduction

63 The GEOTRACES program is an international study which aims to improve the 64 understanding of the global marine biogeochemical cycles and large-scale distribution of 65 trace elements and their isotopes. As GEOTRACES is an international effort with scientists from approximately 30 nations participating, the emerging data sets are produced using 66 67 differing sampling systems and analytical methods. These datasets are merged in the 68 Intermediate Data Product (The GEOTRACES group, in press). To assure consistency 69 between datasets, the international program is emphasising the concurrent analysis of 70 reference samples from the GEOTRACES reference sample program for concentrations of 9 71 key dissolved trace elements, as well as the planning of cross-over or baseline stations where 72 all aspects of sample collection, filtration, processing and analytical methods can be 73 compared. The intercomparison of data for the Intermediate Data Product has been carried 74 out by the GEOTRACES Standards and Intercalibration committee, which shares the results 75 only with contributors to encourage participation. This paper, independently of the findings 76 of that committee, describes the comparison of 2 sampling systems and four analytical 77 techniques, as well as statistical methods to compare trace metal profiles. 78 The US and Netherlands designed new trace metal clean sampling systems specifically for 79 their respective GEOTRACES programs to obtain accurate values of the distributions of key 80 trace metal elements and isotopes along full depth sections. These are the Titan sampling 81 system of the Netherlands (De Baar et al., 2008) that was recently upgraded with novel 82 PVDF samplers (de Baar et al., 2012) as described in the companion article (Rijkenberg et 83 al., submitted; this issue) and the US GEOTRACES carousel (Cutter and Bruland, 2012). 84 Both systems utilise a conducting cable with a CTD instrument as part of the sampling 85 system and can collect 24 samples, but they differ in the materials used and the procedures of

86 recovery and deployment. For example, the whole Titan system with its 24.4 litre PVDF

87 samplers goes into its own cleanroom van for sub-sampling whereas the 12 litre, Teflon-88 coated, GO-FLO sampling bottles are removed from the US GEOTRACES carousel and are 89 brought into a dedicated cleanroom van for filtration and sub-sampling (for more details see 90 (Cutter and Bruland, 2012; De Baar et al., 2008; Rijkenberg et al., submitted; this issue)). 91 Besides the development of new sampling equipment, considerable progress has also been 92 made in the development of new multi-element methods (e.g. Biller and Bruland, 2012; Lee 93 et al., 2011; Milne et al., 2010; Sohrin et al., 2008) using chelating resins for off-line 94 extraction, with subsequent detection with a high-resolution, magnetic sector, inductively coupled plasma mass spectrometer (ICP-MS). Most of the data presented in this paper has 95 96 been analysed using a modified version of the Biller and Bruland (2012) method that will be 97 described here.

98 With these sampling and processing systems, datasets of unprecedented resolution are 99 being produced that are linked together in a global database (The GEOTRACES group, in 100 press). Therefore, we must be confident the sampling systems as well as different analytical 101 techniques obtain comparable and uncontaminated samples. The occupation of the Bermuda 102 Atlantic Time Series (BATS) station by the Netherlands GEOTRACES during a meridional 103 section of the western Atlantic in June 2010 (GA02) and the US GEOTRACES during a 104 zonal section of the North Atlantic in November 2011 (GA03) provided an opportunity to 105 collect dissolved trace metal samples from two occupations of the same 'cross-over' station 106 with different sampling systems, which were analysed for a suite of trace metals by the same 107 analyst with the same analytical method. For Al, Fe and Mn, shipboard dissolved 108 concentration data values are available from the June 2010 Netherlands BATS occupation 109 allowing further intercomparison with shipboard data. Additionally, for Cd, Fe and Zn, 110 concentration data is available from the US occupation using a double spike multi-collector ICP-MS method (Conway et al., 2013). More data is available from the US occupation as 111

112	well as historic occupations of the BATS station, but for the sake of conciseness not all data
113	and methods can be described and compared in this paper and we contrast shipboard methods
114	and 2 recent ICP-MS methods that apply standard addition versus isotope dilution double
115	spiking as the calibration method.
116	Besides comparing the results of the different sampling systems and analytical techniques
117	(Table 1), this paper also aims to describe the modifications made to the method of Biller and
118	Bruland (2012). In order to compare the data from different sampling systems or analytical
119	techniques, a new statistical approach had to be developed to include the analytical
120	uncertainty that is described in this paper. In the companion paper the new PVDF samplers
121	are described (Rijkenberg et al., submitted; this issue).
122	
123	Methods
124	For the Titan sampling system and the US GEOTRACES carousel the reader is referred to
125	the respective papers (Cutter and Bruland, 2012; De Baar et al., 2008) and the companion
126	paper on the PVDF samplers (Rijkenberg et al., submitted; this issue). The only differences in
127	sample treatment are the filtration and acidification. Seawater samples were acidified to a
128	concentration of 0.024 M hydrochloric acid (HCl) which results in a pH of 1.7 to 1.8 with
129	either Baseline® HCl (Seastar Chemicals Inc.) for the samples from the Netherlands cruises,
130	or Q-HCl (produced at UCSC as 6 N Q-HCl with a sub-boiling quartz still) for samples from
131	the US cruise analysed with the multi element standard addition ICP-MS method. Seawater
132	samples from the US cruise analysed with the double spike ICP-MS method were acidified to
133	a concentration of 0.012 M HCl (which results in a pH of 2.0) with Aristar Ultra HCl (VWR
134	International). Any difference between these acids is accounted for in the blank
135	determination. On the Netherlands GEOTRACES cruise, the samples were filtered using 0.2
136	μm pore size Sartobran capsule filters, whereas on the US GEOTRACES cruise 0.2 μm pore

- 137 size Acropak capsule filters were used. The use of these different brands of filter cartridge
- 138 does not result in significant differences in trace metal concentrations (Cutter and Bruland,

139 2012). All samples were stored in LDPE bottles that were acid washed according to the

140 'Sample and Sample-handling Protocols for GEOTRACES cruises'

141 (http://www.geotraces.org).

- 142
- 143 Double spike ICP-MS and shipboard methods

144 The details of the double spike ICP-MS method for Fe, Zn and Cd by Thermo Neptune

145 multicollector ICP-MS are described by Conway et al. (2013). Accuracy of the double spike

146 method for dissolved concentrations has been previously demonstrated by agreement with the

147 consensus values for SAFe D1, D2 and S reference standards Conway et al. (2013), which

148 were analysed within the same ICP-MS analytical sessions at the Center for Elemental Mass

- 149 Spectrometry (University of South Carolina) as the samples presented in this
- 150 intercomparison. Procedural blanks for this technique were determined as 5 pmol kg⁻¹ (Fe), 1

151 pmol kg⁻¹ for (Zn) and 35 fmol kg⁻¹ for Cd (Conway et al., 2013). The details on the

- 152 shipboard methods of the Netherlands occupation for Fe, Mn and Al were described by
- 153 Rijkenberg et al. (2014), Middag et al. (2011a) and Middag et al. (in press), respectively.
- 154 Typical precisions and detection limits for shipboard Al were 3.3% for multiple
- 155 measurements of a 6.5 nM sample (n=40) and a detection limit of 0.05 nM (defined as three
- 156 times the standard deviation of the lowest concentration observed) (Middag et al., in press).
- 157 For shipboard Mn these were 5.1% for a 0.45 nM sample (n=37) and a detection limit of 0.01
- 158 nM. For shipboard Fe the precision was 4% for replicate measurements of a 0.94 nM
- 159 reference sample (n=29) and a limit of detection of 0.01 nM (Rijkenberg et al., 2014).
- 160

162 *Reagents*

163 This method includes the analysis of yttrium (Y), lanthanum (La), titanium (Ti) and 164 gallium (Ga), in addition to manganese (Mn), iron (Fe), cobalt (Co), nickel (Ni), copper (Cu), 165 zinc (Zn), cadmium (Cd) and lead (Pb) that were determined in the original method by Biller 166 and Bruland (2012). Moreover, we introduce a new 'element dilution' approach that can be 167 used for extractions performed at sea that is less labour intensive than the gravimetrical 168 method described by Biller and Bruland (2012) as the weighing of the samples has been 169 excluded. The extraction of the samples is the process where the trace metals of interest are 170 separated from the original seawater matrix to remove interfering ions, as well as 171 concentrating the samples via the use of a chelating column (Nobias PA1 chelating resin in 172 this method). This pre-concentration is necessary due to the low concentrations of trace 173 metals in the open ocean and the high background salt matrix of seawater. The reagents used for the trace metal analyses are the same as those described by Biller 174 175 and Bruland (2012), with the exception of a Lutetium (Lu)-Indium (In) spike and the use of 176 Hydrofluoric Acid (HF) as described below. A 4000 nM stock of Lu and In was made in 177 0.024 M Nitric Acid (HNO₃) (triple quarts distilled) from dilutions of the respective 1000 μ g g⁻¹ SPEX standards. A 2.4 M and 1.2 M working solution of HF were made by diluting 178 179 concentrated HF (50%, Optima grade, Fisher Scientific). The HF addition was found to 180 increase the recovery of notably Ti (see Modified Multi-element standard addition ICP-MS 181 *Method*). A short outline of the method is given below that mainly focusses on the differences 182 with the method described by Biller and Bruland (2012). For more details on the method and 183 method verification in general, the reader is referred to Biller and Bruland (2012). The 184 method uses Nobias-chelate PA1 resin to extract the metals from the seawater. The metals are subsequently eluted with 1 M nitric acid and the extracts are run on a sector field ICP-MS. 185

The 1 M nitric acid used in this method contains 0.6 mM HF in addition to the 10 ng g⁻¹ Rh in
the original method.

188

189 *Calibration*

190 A multi-element stock standard with natural isotopic abundances of Mn, Fe, Co, Ni, Cu, 191 Zn, Cd, Y, La, Ti, Ga and Pb, in 0.024 M HNO₃ was used to make standard additions to 192 natural seawater with low concentrations of metals for calibration. Five standard additions 193 were used for calibration in this method, and every sample and standard were spiked with Lu 194 and In to obtain a concentration of 5 nM for both elements. In addition to the seawater 195 standard additions, 5 standards were made up in the acid used to elute the metals from the 196 pre-concentration columns (elution acid standards). These standards (also spiked with Lu-In) 197 and the extracted seawater standards were analysed on each run.

198 The resin recovery for the modified multi-element ICP-MS method was monitored as 199 described by Biller and Bruland (2012). Briefly, the slope of the relationship between the 200 ICP-MS signal and metal standard additions to seawater is compared with the slope of the 201 relationship between the ICP-MS signal and metal additions to elution acid that was not 202 passed through the extraction system. This was done for every set of samples that were 203 extracted and analysed by ICP-MS. The recovery for all elements with the exception of Ga 204 and Ti is quantitative (recovery >98%) and agrees with previous results for this resin (Biller 205 and Bruland, 2012; Sohrin et al., 2008).

Lutetium and In are two trace metals, naturally existing at sub pM concentrations (Alibo et al., 1999; Alibo and Nozaki, 1999; Amakawa et al., 1996; Greaves et al., 1991), that are quantitatively concentrated by the extraction procedure (recovery >98%). The 5 nM spike overwhelms the natural concentration in the samples. This Lu-In spiking, when done to an exact volume (see *preconcentration and analysis*) of seawater, serves as an internal standard

211 or, analogous to isotope dilution, as an 'element dilution' calibration technique. This is in addition to the standard additions to a seawater sample: the standard additions serve to 212 213 determine the calibration sensitivity for each metal (i.e. the signal per unit of metal) while the 214 Lu-In spike represents the concentration factor of each sample due to the extraction method. 215 The Lu: 'metal of interest' or In: 'metal of interest' ratio does not change after addition of 216 the spike to the sample and will be the same in the sample and the eluent when recoveries are 217 quantitative. Additionally, after addition of the Lu-In spike to the seawater samples, any 218 evaporation of the seawater samples or the eluants after extraction is irrelevant as that does 219 not change the ratios. If the recovery of a metal is not quantitative, the method will only work 220 if the recovery is stable, i.e. does not vary between samples and standards. In this case the 221 calibration sensitivity will be relatively low, but the non-quantitative recovery will be 222 accounted for in this calibration sensitivity.

223 By dividing the ICP-MS signal for every metal by the signal for Lu, the signal is 224 normalized with respect to the concentration factor as the Lu signal is proportional to the 225 amount of sample loaded and the amount of elution acid used. Similarly, the signal for every 226 metal can be (separately) divided by the In signal to have 2 independent determinations of the 227 concentration factor with the added benefit that a 'failed' extraction shows up in large 228 deviations of the resulting concentrations or Lu/In ratios. The Lu and In signals were 229 determined in both medium and low resolution, and were applied appropriately to metals measured in medium resolution (⁴⁷Ti, ⁵⁵Mn, ⁵⁶Fe, ⁵⁹Co, ⁶⁰Ni, ⁶³Cu, ⁶⁶Zn, ⁶⁹Ga) or low 230 resolution (⁸⁹Y, ¹¹¹Cd, ¹³⁹La and ²⁰⁸Pb). The Rh spike to the elution acid served to guickly 231 232 highlight any aspiration problems or drifts in sensitivity on the ICP-MS for the original 233 gravimetrical method. The data presented here was determined by the gravimetrical method 234 based upon the sensitivity of the standard additions to the seawater. The Lu-In element

dilution method was tested versus the gravimetrical method (see *Modified Multi-element standard addition ICP-MS Method*).

237

238 *Pre-concentration and analysis*

239 The off-line pre-concentration set-up and procedures are as described by Biller and 240 Bruland (2012) with some small but essential differences. Most importantly, to 40 ml of 241 seawater, 20 µL of the 1.2 M HF was added (resulting in a 0.6 mM fluoride ion 242 concentration), followed by a 50 µL addition of the Lu-In spike. This 40 ml was taken using a 243 10 ml pipette from a slightly larger volume of seawater. This seawater was UV oxidised to 244 destroy organic metal binding ligands for dissolved Co and Cu concentration determinations 245 (this was found to be particularly important for obtaining accurate dissolved Co 246 concentrations and significant for obtaining accurate dissolved Cu concentrations (Biller and 247 Bruland, 2012)). The overall imprecision of the pipetting the 40 ml of water was determined 248 gravimetrically and found to be $\sim 0.9\%$. The accuracy of the pipette was monitored and no 249 drift was observed over the course of 2 months. 250 Samples were capped and vigorously shaken (5 seconds) and left overnight (14 hours) to 251 homogenise after the spiking. Immediately prior to extraction, the pH of the samples was 252 adjusted to 5.8 +/- 0.2 with \sim 500 µL of a 3.7 M NH₄Ac solution. About 35 ml of the sample was loaded onto the columns in ~45 minutes (the exact amount of the sample loaded was 253 254 gravimetrically determined (Biller and Bruland, 2012)). The leftover 5 ml was used to retro-255 actively verify that the correct pH was achieved. The columns were rinsed with a 0.05 M 256 NH₄Ac rinse/conditioning solution for 1 and 2 minutes, respectively, before and after sample 257 loading to condition the columns at the correct pH, and remove interfering sea salt before elution. Samples were eluted with ~ 2 ml of the elution solution in ~ 5 minutes (the exact 258 259 amount of the elution acid was gravimetrically determined), resulting in a concentration

factor close to 17.5. Using 8 columns in this off-line manifold allowed the simultaneousextraction of 8 samples.

262 Samples were analysed on a Thermo Element XR Magnetic Sector ICP-MS at the Institute 263 of Marine Science at UCSC with nickel sample and skimmer cones (Spectron) and an ESI-PC3 Peltier cooled spray chamber (4 °C). The sample gas rate was 0.75 mL min⁻¹, and the 264 265 additional gas rate was optimised every run (usually around 0.2 mL min⁻¹). Sample was 266 pumped into the instrument at a rate of 120 µL min⁻¹. Molybdenum (⁹⁵Mo) was measured to 267 correct for Molybdenum oxide interferences on the Cd signal (Biller and Bruland, 2012) that 268 became significant at low surface water concentrations of dissolved Cd. 269 Unused (i.e. not passed through the extraction set-up) elution acid was used to track the 270 baseline associated with the acid and ICP-MS instrument and was subtracted from the raw 271 signals. An elution acid standard (elution acid with added metals) served as a drift standard to 272 track the sensitivity of the ICP-MS during a run even though the Rh (gravimetrical method) 273 and the Lu-In spikes ('element dilution' method) are used to correct for any change in 274 sensitivity. System blanks were determined by performing the normal extraction procedure, 275 but loading a low metal concentration seawater sample (the same as used for the standard 276 additions) onto the column for only 30 seconds versus ~45 minutes for an ordinary sample. 277 This way the column is loaded with sea salts, but only minimally with metals. A system blank 278 was determined for every column, but consistency between the blanks indicated no noticeable 279 difference between columns. A reagent blank was determined by adding double the normal 280 amount of HCl, buffer, HF and Lu-In spike to a sample of the same seawater used for the 281 standard additions. The resulting concentration was compared to the 'zero addition' point of 282 the calibration that only had normal amounts of reagents added. No noticeable reagent blank 283 could be determined for the metals except a small Pb blank of 1.2 pmol kg⁻¹. Blanks and limit of detection (LOD, defined as three times the standard deviation of the system blank) are 284

shown in Table 2. SAFe and GEOTRACES reference samples were analysed in every run(Table 3).

287

288 **Comparing profiles with analytical uncertainty**

289 To compare datasets, traditionally paired two-tailed t-tests or 95% confidence intervals 290 (CI's; expressed as 2 x standard deviation) have been used as well as the root-mean-square 291 (RMS) deviations of the reported values from the mean values (Bowie et al., 2003; Landing 292 et al., 1995). The t-tests have the disadvantage that analytical uncertainty is ignored, for 293 example Bowie et al. (2003) reported significant differences (at the 0.05 significance level) 294 between different analytical techniques based on paired two-tailed t-tests, whereas 95% CI's 295 where often overlapping. In addition, a t-test assumes a normal distribution of the data which 296 is rarely the case for an oceanographic profile. Indeed, the distribution of the data is generally 297 skewed towards higher or lower concentrations in the shape of the vertical profile of the 298 chemical species. Solely looking at 95% CI's or RMS deviations, on the other hand, does not 299 give a quantitative measure of comparability for an entire profile. 300 In order to objectively and reproducibly assess the comparability of analytical methods or

sampling systems while acknowledging there is analytical uncertainty, another approached is needed. Here we describe a relatively straight forward test that can be done in a spreadsheet or statistical program. This approach, however, is limited to comparing analytical techniques based on results from the same samples or subsamples thereof. It is less suited to compare oceanographic samples collected at different occasions since samples will have been collected at (slightly) different depths for which another test was developed (see *Different samples approach*).

308

309 Same sample approach

310 To compare analytical results from the same samples (or sub-samples thereof) first a 311 standard paired two-tailed t-test (α =0.05) was done as described by Bowie et al. (2003). As 312 an example, in case of an oceanographic profile of 24 samples analysed by two different 313 analytical techniques in triplicate, the test compares the 24 analytical result pairs. If this does 314 not result in a statistical difference, the results of the two methods are not significantly 315 different regardless of analytical uncertainty. As mentioned above, this paired two-tailed t-316 test assumes a normal distribution of the data. This constitutes a potential caveat for this 317 approach. A statistical program such as Sigmaplot 13.0 as used in this paper, will test for 318 normality of the data when performing a t-test and, if assumptions are violated, an alternative 319 non parametric test such as a Wilcoxon signed-rank test can be performed instead. The test 320 outlined here is for the comparison of 2 datasets, for multiple datasets the t-test and signed 321 rank test can be replaced by an ANOVA or a Friedman Repeated Measures Analysis of 322 Variance on Ranks, with a Tukey post hoc test to identify differences. If the t-test or signed-323 rank test does give a significant difference, the results of the two methods are subsequently 324 assessed with consideration of the analytical uncertainty. For this, the absolute difference 325 between results of method 1 and method 2 was calculated for each sample. Subsequently the 326 standard deviation of the difference was calculated using equation 1:

327
$$\sigma_d = \sqrt{\sigma_1^2 + \sigma_1^2}$$
 (eq. 1)

Where σ_d is the standard deviation of the difference between the analytical method 1 and analytical method 2, and σ_1 and σ_2 are the standard deviations of the results for method 1 and method 2 respectively. The latter standard deviation is calculated based on repeat measurements of subsamples of the same sample and can also be referred to as the standard error of the mean. If no standard deviation can be calculated for one of the methods (e.g. because no multiple subsamples were analysed) the analytical uncertainty needs to be assessed by other means. Notably for ICP-MS methods, often a single sample is analysed by

335 the instrument a number of times and the average and standard deviation of those measurements is reported. This standard deviation divided by n^{0.5}, were n is the number of 336 337 measurements (instrument scans), is the 'internal' standard error for the measurement 338 (instrument variability). It is known that this 'internal' standard error of one sample analysis 339 approximates the standard deviation of replicate analysis of subsamples (standard error of the 340 mean). However, this does not hold in case of techniques, such as the standard addition ICP-341 MS method, where a considerable amount of variation is not due to instrument variability, but 342 due to variation in extractions on resin, difference in sample bottles or other non-instrument 343 related variability. This shows in an 'internal' standard error for the measurement of one 344 subsample that is smaller than the standard deviation of replicate analysis of multiple 345 subsamples (standard error of the mean). The best determination of the standard deviation 346 (standard error of the mean) is to analyse subsamples of the same samples an infinite number 347 of times. The best approximation of this in a practical sense is probably the analyses of 348 references samples/standards that are analysed with every run. In this paper standard 349 deviation will be denoted with the symbol σ and is based on repeat measurements of 350 separately extracted subsamples of the same sample (standard error of the mean). 351 In case of the double spike ICP-MS method where no multiple subsamples of the same 352 seawater samples were analysed for the whole profile, the analytical uncertainty was 353 estimated to be 2% (Conway et al., 2013) and these estimates were used as the σ . The σ_d and 354 absolute difference was calculated for every pair of results; in the case of an oceanographic 355 profile of 24 samples analysed by two different analytical techniques, this results in 24 356 absolute differences and 24 associated standard deviations of the difference. Subsequently a 357 one-tailed paired t-test or signed-rank test can be performed to assess whether σ_d is larger than the absolute difference for the entire profile. If this is the case, it is concluded the 358 359 analytical results of the two methods are not statistically different within uncertainty as the

uncertainty is larger than the difference. Otherwise the conclusion is that the two methods donot give comparable results.

362 This analytical uncertainty assessment can be repeated using a 2σ -based analytical 363 uncertainty or 95% CI's, but for this paper the more conservative approach of 1σ was chosen. 364 Assuming a normal distribution, the 95% CI denotes the interval that contains the actual 365 mean with 95% probability. This can be calculated by multiplying the standard error (standard error= $\sigma x n^{-0.5}$; note this is *not* equivalent to the standard error of the mean defined 366 above) with the critical t-value for the applicable number of degrees of freedom and chosen 367 368 significance level. For large numbers of replicates (n>20) the critical t-value at α =0.05 is ~2, 369 making the 95% CI about 2 standard errors whereas for n=3 the critical t-value equals 4.3. In 370 other words, the 95% CI is a measure of certainty about the mean reported value and this certainty increases with a larger number of replicates (smaller 95% CI). Additionally, for a 371 372 normal distribution and a large number of replicates, 95% of the results lie within $\sim 2 \sigma$ of the 373 mean, which confusingly is sometimes also referred to as the 95% CI. This number of 2σ is a 374 measure of the spread around the mean and can therefore be considered more appropriate as a 375 measure of analytical uncertainty. However, commonly samples are analysed in triplicate for 376 shipboard flow injection methods, or single sample extracts are analysed for ICP-MS 377 methods, making the estimation of the 95% CI less reliable or impossible due to the lack of 378 degrees of freedom. Similarly, the assumption of 2σ encompassing 95% of the variation 379 around the mean is not valid with a small numbers of replicates and we opted to use 1 σ as 380 the most reliable measure of analytical uncertainty.

Additional information can be gained by dividing the difference between result pairs (either positive or negative) by the σ_d after John and Adkins (2010; their equation 18), resulting in values that are basically individual t-values for each pair of results. Subsequently the additive inverse (-t) of each t-value is added to the list of t-values, giving a collection of t-

385 values with an average that equals zero. The standard deviation of this collection of t-values 386 can be used to differentiate between intra-lab and inter-lab variability. If the standard 387 deviation is ≤ 1 , all differences can be related to intra-lab variability (i.e. the analytical 388 uncertainties). If the standard deviation is ≥ 1 , the number indicates how much greater the 389 actual observed differences are than expected based on intra-lab variability, e.g. if the 390 standard deviation is 1.3 or 4, it means the inter-lab variability is 30% or 300% greater than 391 expected based on intra-lab variability, respectively. A standard deviation greater or smaller 392 than 1, however, does not indicate statistical significance at the 95% significance level as this 393 test just predicts differences based on the reported analytical uncertainty and does not test for 394 differences. For the latter, the above or below described tests are to be used.

395

396 Different samples approach

397 As stated above, the above approach is limited to comparing results from the same 398 samples or subsamples thereof and is not suited to compare oceanographic samples collected 399 by different sampling systems or at different times, as samples will have been collected at (slightly) different depths and/or locations. Therefore a script was developed in the freeware 400 401 program R (R Core Team, 2013) that interpolates oceanographic profiles and the analytical 402 uncertainty and subsequently tests the shape of two profiles within a prescribed depth 403 interval. The interpolation allows comparing a sampled depth on one profile with an 404 interpolated value in the other profile. Besides, errors in interpolation on mean concentrations 405 and standard deviation of each profile are used to determine the standard deviation of the 406 interpolated values assuming a 95%-significance confidence interval. This interpolation is 407 based on depth, but since water masses might occupy different depths between sampling 408 events, conservative tracers such as potential or neutral density, salinity or potential 409 temperature might be better suited. However, for the surface ocean, salinities, temperatures

410 and densities observed in the US BATS occupation were not observed in the NL BATS 411 occupation, making interpolation based on these parameters unfeasible. For the sake of clarity 412 we used depth, as in the deep ocean were comparison is feasible (see section Intercomparison 413 between the different sampling systems), depth and density matched very well between the 414 two occupations. Given that thickness of water masses increases with depth (contracting at 415 the bottom of the ocean), using depth makes comparison between two profiles easier, notably 416 in graphs. After interpolation, the shape of the two profiles is compared simultaneously using 417 two statistical approaches. The first approach consists of comparing samples between each 418 other across depth using a simple T-based distribution relying on a pooled variance 419 assumption as follows:

420
$$t_{z} = \frac{m_{1} - m_{2}}{\sqrt{\left(n_{1}\sigma_{2}^{2} + n_{2}\sigma_{2}^{2}\right)\frac{1}{n_{1}} + \frac{1}{n_{2}}}} \sim T_{n_{1} + n_{2} - 1}$$

421 Where $\overline{m_i}$ and σ_i represent respectively the mean concentration at a given *z* depth of the 422 profile *i* and the standard deviation relative to the number of samples, n_i , analysed at this 423 depth. This gives a *p*-value for the comparison at each individual depth that is compared and 424 thus gives a *p*-value profile, a profile of *p*-values which provides insight into where the 425 largest discrepancies and the best agreements are along the depth profile.

426

The second approach aims at comparing the shape of the two profiles within a depth interval using a Fisher-based test. This test consists in computing both the cumulative sum of the Welch's t_z^2 (Welch, 1947) and the cumulative sum of the randomly generated Fisher distribution F_{1,n_1+n_2-2} . A chosen number of random simulations (usually 100,000) of the Fisher F_{1,n_1+n_2-2} distribution are used to estimate the distribution of the cumulative Fisher law. This distribution is then used to compute an 'integrated' *p*-value which represents an
objective metric to determine how far the two profiles are consistent between each other
within the depth interval. The script is provided in the supplementary material with this
paper.

The use of both these approaches ensures a good understanding of the profile's consistency.
The 'integrated' *p*-value allows quantifying the match between profiles with one metric, i.e.
whether or not there is a significant difference, while the T-based *p*-value profiles ensures to
decipher the reason for a good or poor match between profiles.

440 That said, the computation of the 'integrated' *p*-value is sensitive to the number of samples 441 and to the resolution of sampling over depth, i.e. if the sampling resolution is low in one 442 profile, the interpolation will inherently miss fine scale gradients if present. The 'integrated' p-value estimated from the F or a χ^2 test can differ by about 10 to 20% due to the 443 444 sensitivity on the number of samples on the computed confidence intervals. Therefore this 445 approach cannot be considered alone without a proper analysis of profiles miss-match based 446 on the *p*-value profiles. In the following, the 'integrated' *p*-value will be noted as '*i*-*p*-value' 447 and reports the analysis that is based on computation over the profiles (deeper than 1000 m 448 depth, see Intercomparison between the different sampling systems).

449

450 **Results and Discussion**

451 Modified Multi-element standard addition ICP-MS Method

452 The recovery for Mn, Fe, Co, Ni, Cu, Zn, Cd, Y, La, and Pb was quantitative (>98%) at pH

453 5.8. Two elements demonstrated non-quantitative recovery; Ga (75 %; σ =4%) and Ti (50%

- 454 $\sigma=5\%$). The recovery was assessed over a range of pH and this showed most metals were
- 455 quantitatively recovered over a pH range of 4.5 to 9, but the recovery of Mn is not
- 456 quantitative below pH 5.5 and Ga has its optimum recovery around a pH of 5.5. The recovery

457 of Ti, zirconium and scandium benefited from the addition of fluoride (see supplementary 458 material for details). The highest recovery for Ti (75% with 0.6 mM HF) was at pH 4.5 and 459 the recovery of Ti was relatively constant around 50% at a pH of 5.8, but multiple analyses of 460 reference samples indicated modest precision (σ ~10%).

461 The recovery of zirconium and scandium was unstable, resulting in concentrations with 462 low accuracy. Further development needs to be carried out for these elements and they are 463 currently not included in the method. The fluoride addition had a perceived positive effect on 464 the analysis of Fe and Ga. This was mainly visible in the increased 'smoothness' of oceanic 465 profiles (i.e. rather than meandering the profiles showed uniform trends with increasing 466 depth) and the reproducibility of reference samples. Before addition of HF, occasionally 467 values were measured for the reference samples that were a lot lower than expected (outside 468 the consensus range), whereas after addition of HF this was no longer observed. The 469 improvement of the visual appearance of the profiles and more consistent reference samples 470 results might possibly be due to hindering of precipitation after pH adjustment in the presence 471 of HF, but increased operator experience cannot be excluded either. The effect of fluoride 472 addition to the seawater on the recovery was assessed at concentrations up to 10 mM fluoride 473 and it appeared there is an optimum of about 0.6 mM fluoride for the recovery of Ti. It is 474 hypothesised that the addition of fluoride has an advantageous effect for the elements that 475 form hydrolysis complexes in seawater. Potentially, the formation of metal-fluoride 476 complexes diminishes precipitation and wall adsorption after pH adjustment to pH 5.8, 477 leading to an improved recovery. Alternatively, the formation of metal-fluoride complexes 478 results in an increased lability of the metal-water complexes, leading to a greater chance for 479 the resin to form a complex with the metal. For example in the case of Fe, at the analytical 480 loading pH (5.8) the Fe(OH)₂⁺ complex would dominate (Glasby and Schulz, 1999) and this is relatively slow with respect to water exchange (Hudson et al., 1992). The increased lability 481

of the metal-water complexes could increase the recovery on the resin as the chelating resin is competing for the metals with the complex forming agents in the seawater. The only exception was Al; recovery of Al at pH 5.8 was about 25% without fluoride and after the fluoride addition this decreased to near 0%, indicating the Al-fluoride complex is stronger than the Al-resin complex or kinetically inert on the time scale of its passage through the resin. Overall, the pH of 5.8 +/- 0.2 and an addition of 0.6 mM fluoride to the samples resulted in the best recovery for most elements.

489 SAFe and GEOTRACES reference samples were analysed in every run and the results are 490 in line (1σ intervals overlap) with consensus values for the metals that have those (Table 3). 491 Currently there are no consensus values for Ti on reference samples, but Croot (2011) 492 published results for his analysis of Ti on the reference samples (Table 3). The results for the 493 higher concentrations agree reasonably well (1σ intervals overlap), but there appears to be 494 more discrepancy for the lower concentrations, most notably for SAFe S. However, it should 495 be noted that for one of the SAFe S samples (SAFE S 474) analysed by Croot (2011) the 496 result were in range with our results for SAFe S. This indicates the analysis of Ti is 497 challenging and more method development needs to be done in order to attain reliable data on 498 the distribution of Ti in the open ocean. In this case the method should benefit from using 499 isotope dilution to account for the non-quantitative and potentially variable recovery on the 500 resin. For Y, La and Ga there are currently no consensus values either, and the obtained 501 vertical profiles at BATS are compared to published data below (Intercomparison between 502 the different sampling systems) to assess the method's accuracy. There is data on reference 503 samples for La (Hatje et al., 2014; van de Flierdt et al., 2012) that agrees (1σ intervals 504 overlap) with the here reported values (Table 3). 505

505 To accurately measure the sub-pM concentrations of Lu and In the concentration factor 506 would need to be increased, i.e. much more seawater would need to be pre-concentrated on

507 the resin. Therefore we opted instead to use these two elements as an internal standard (see 508 calibration). This 'element dilution' calibration technique was assessed by comparing the 509 resulting values versus the gravimetrical method as described by Biller and Bruland (2012). 510 No discernible difference was observed between the 'element dilution' and gravimetrical 511 method (Fig. 1), either with a two-tailed paired t-test (p=0.20) or the script for profile 512 comparison that includes analytical uncertainty (p=0.97). The benefit of the element dilution 513 calibration technique is that the gravimetrical determination of the pre-concentration factor is 514 no longer necessary, making it possible to do the time consuming extractions at sea. The 515 results presented in this paper are based on extractions done in the shore based lab, but 516 meanwhile the extractions for the subsequent Netherlands Mediterranean GEOTRACES 517 expedition were carried out aboard RV Pelagia by the Middag group and analysed at the 518 University of California Santa Cruz during a visit to the Bruland lab. These results are written 519 up separately, but the results for the GEOTRACES reference samples that were extracted 520 shipboard are included in Table 4 and are not distinguishable from the results for the here 521 reported samples (P>0.05 in all cases; two-tailed homoscedastic t-test). The good agreement 522 between the reference samples and the comparison of the gravimetrical and the element 523 dilution method shows the off-line extractions are no longer required to be carried out in the 524 home laboratory and can also be done at sea, where gravimetric measurements are not 525 possible. After returning with the extracted samples to the home laboratory all samples can be 526 run in 1 long ICP-MS run (150 hours for the Mediterranean data where a total of roughly 527 2100 profile samples, reference samples, standards and blanks were analysed) creating the 528 possibility to produce the data shortly after return. We believe this is a significant 529 improvement over the shore-based extractions that often take many months to complete. 530 Taking back-up seawater samples to return to the shore laboratory is always recommended in 531 case (i) some extractions fail, or (ii) there is inadvertent contamination of some samples, or

(iii) if a future interest arises in other trace elements that require different extractiontechniques.

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- 535

536 Intercomparison between the different sampling systems

537 The BATS intercomparison or cross-over station (31°45.92'N, 64°04.95'W) was used to 538 compare the sampling systems used by the Netherlands and US GEOTRACES programs for 539 12 dissolved trace elements. The data of all trace elements are attached in the supplementary 540 material. Since a different number of samples were collected at different depths between the 541 occupations, the R script is used to interpolate the profiles and assess the differences. Only 542 data below the main thermocline was used (deeper than 1000 m) to avoid temporal variation 543 between the two occupations that were over a year apart and had a negative and positive sea 544 surface height anomaly during the Netherlands and US occupation, respectively, indicative of 545 an up- and down-welling eddy (this was determined using the Colorado center for 546 astrodynamics research SSH data viewer at http://eddy.colorado.edu/ccar/ssh/nrt global grid viewer). 547 548 Of the first row transition metal elements, Mn, Fe, Co, Ni, Cu, Zn (in order of increasing 549 atomic number (25-30)) are all essential for every living organism (de Baar and La Roche, 550 2003). Within the group 12 elements, the second row element Cd is akin to the first row 551 element Zn, and thus Cd may substitute for Zn in phytoplankton, while an intrinsic Cd-552 carbonic anhydrase also is found in certain diatoms (Xu et al., 2008). Next, transition metal 553 group 3 is represented by second row Y and third row La, and transition metal group 4 by its 554 first row Ti that serves as a geochemical source tracer of mineral dust (Dammshaeuser et al.,

- 555 2011) as well as the construction material of the Netherlands sampling frame. The non-
- transition group 13 is represented by Al and Ga with similar ocean chemistry (Orians and

557 Bruland, 1988), and Al is the construction material of the US sampling frame. Aluminum is 558 discussed last as it is not measured with the ICP-MS method. Finally, group 14 is represented 559 by Pb, a decadal time scale tracer at BATS (Lee et al., 2011) due to the earlier use of, and 560 subsequent phasing out, of Pb in gasoline for automobiles.

561 Dissolved Mn is an element that in the open ocean usually shows a strong surface 562 enrichment due to atmospheric input (Baker et al., 2006; Landing and Bruland, 1980) and 563 photo reduction of Mn oxides (Sunda et al., 1983). Below the surface layer, Mn 564 concentrations are generally low and quite uniform due to (microbially mediated) removal by 565 scavenging (Landing and Bruland, 1980; Statham et al., 1998). The Mn profile at BATS 566 largely follows the classical profile (Fig. 2a). There are some discrepancies in the deep water 567 at the low concentrations that are visible in the p-value profile, but overall the results are not 568 statistically different within analytical uncertainty (*i-p*-value=0.32).

569 Dissolved Fe is arguably one of the most analytically challenging elements to determine in 570 the open ocean and considerable effort has been put in intercalibration in the past (e.g. 571 (Johnson et al., 2007)). Dissolved Fe displays characteristics of both a nutrient-type element 572 and a scavenged-type element in the open ocean which is also visible at the BATS station 573 (Fig. 2b). Surface concentrations are elevated, most likely due to recent dust input, but decrease to ~0.1 nmol kg⁻¹ around 100 m depth due to biological uptake. Dissolved Fe 574 concentrations are relatively constant close to 0.5 nmol kg⁻¹ in the deep ocean, indicating 575 576 equilibrium between input and scavenging removal. The profiles for dissolved Fe (Fig. 2b) 577 are not statistically different within analytical uncertainty (*i-p*-value=0.50). 578 Dissolved Co, like Fe, displays characteristics of both a scavenged-type as well as a 579 nutrient-type element. The latter is due to the nutritional requirement of phytoplankton for Co 580 (e.g. Saito and Goepfert, 2008). For Co, the UV oxidation step is of utmost importance, as 581 can be seen by the difference between the data from samples with and without UV oxidation

(Fig. 2c). Only one replicate was analysed for the UV oxidised samples and thus the analytical uncertainty was conservatively estimated at 2%. The Fisher law in the R script cannot be run when only one replicate measurement is available for both profiles. To allow comparison, two replicates for each data point were assumed (note that increasing the number of replicates makes the test harsher). With this assumed error, the profiles are not statistically different within analytical uncertainty (*i-p*-value=0.93).

588 Dissolved Ni is another nutrient-type element (Fig. 2d) as it is used in the breakdown of 589 urea (Price and Morel, 1991) and by nitrogen fixers (e.g.(Ho, 2013)). The general agreement 590 between the data of the Netherlands and US occupation is good, but the data from the US 591 occupation showed four outliers (3000, 2500, 1800 and 874 m depth). Outliers were defined 592 as described by Middag et al. (2011a), where a linear regression was determined between the 593 data above and below the suspect data point and a deviation of more than 10% from the 594 predicted value was used as the rejection criterion. The samples contaminated for Ni were 595 also contaminated for Cu and Zn at a roughly constant ratio of Ni:Cu:Zn but were not 596 contaminated for these metals in profiles analysed by a different PI in the US GEOTRACES 597 program (Jingfeng Wu, pers. com.), so it does not appear to be a problem with the GO-FLO 598 samplers. Additionally, the Zn samples analysed with the double spike ICP-MS method were 599 not contaminated either. When excluding those four outliers, the profiles are not statistically 600 different within analytical uncertainty (*i-p*-value=0.77) for Ni, further confirming that the US 601 sampling system is not the source of contamination. Either these four subsampling bottles 602 were contaminated for Ni, Cu and Zn or the contamination was introduced during subsequent 603 sample handling. The samples of these four coinciding outliers have been excluded from the 604 profiles for Ni, Cu and Zn.

Dissolved Cu has a hybrid type profile (Fig. 2e) with slightly depleted surface

606 concentrations as there is a biological requirement for Cu (e.g. Annett et al., 2008; Peers and

607 Price, 2006), and relatively uniform deep concentrations. For Cu, the UV oxidation step is 608 also of importance as can be seen by the difference between the data from samples with and 609 without UV oxidation (Fig. 2e). As for Co, only one replicate was analysed for the UV 610 oxidised samples and thus the error was conservatively estimated at 2% and the number of 611 replicate analyses was set at 2. With this assumed error, the profiles are not statistically 612 different within analytical uncertainty (*i-p*-value=0.46). There are two additional outliers (6 613 in total) for Cu at 666 and 500 m depth in the upper ocean of the US profile (not considered 614 in the comparison), implying a contamination issue for Cu of unknown origin. This is further 615 corroborated by the relatively modest agreement between the 2 profiles and further research 616 is required to resolve this, but currently any discrepancies do not seem related to the sampling 617 systems.

618 Dissolved Zn is a nutrient type element due to its biological requirement by 619 phytoplankton, most notably in the carbonic anhydrase enzyme to acquire CO₂ for 620 photosynthesis (Morel et al., 1994). Dissolved Zn is one of the most contamination prone 621 elements and each profile has one outlier (Fig. 2f; US at 2100 m and NL at 4525 m), in 622 addition to the four excluded samples from the US occupation. This demonstrates that the 623 utmost care needs to be taken during sample handling. The profiles for dissolved Zn (Fig. 2f) 624 are not statistically different within analytical uncertainty (*i-p*-value=0.08). Towards the 625 seafloor (deeper than 4200 m), the US samples trend to lower values, whereas the 626 Netherlands remain similar, resulting in lower p-values in the p-value profile (see 627 supplementary material). These decreasing p-values near the seafloor are also seen for Mn, 628 Fe and Co, indicating natural variation. This variation might be related to the thick benthic 629 nepheloid layer that has been observed at the BATS station during the US occupation (Lam et 630 al., 2014).

Dissolved Cd displays nutrient type behaviour in the ocean. This was a long standing
subject of debate as Cd was not previously considered a bio-essential metal (Lane and Morel,
2000). However, biological roles for Cd have now been identified, e.g. Cd can substitute Zn
in carbonic anhydrase in some phytoplankton species (Xu and Morel, 2013). The profiles for
dissolved Cd (Fig. 3a) are not statistically different within analytical uncertainty (*i-p*value=0.57).

637 Dissolved yttrium has a narrow range of concentrations at BATS, ranging between ~125 and 165 pmol kg⁻¹, with the highest concentrations in the deep waters (Fig. 3b). The sparse 638 639 amount of data that exists for Y suggests that it has somewhat of a nutrient-type distribution 640 in the oceans with slightly higher concentrations in the older Pacific waters (Alibo et al., 641 1999). Yttrium has also been reported to have higher concentrations in the Mediterranean due 642 to the high dust input in the latter region (Alibo et al., 1999). The profiles for dissolved Y 643 (Fig. 3b) are not statistically different within analytical uncertainty (*i-p*-value=0.34). 644 The concentrations of dissolved Y have been reported to be tightly correlated with 645 dissolved La (Zhang et al., 1994). At BATS, Y correlates with La with a correlation coefficient, R², 0.84 (n=60; not shown). The relatively modest correlation between Y and La 646 647 is caused by a La maximum around 1500 m depth (Fig. 3c) that is absent in the Y profile, 648 indicating a source to the upper North Atlantic Deep Water for La without accompanying Y. 649 Lanthanum correlates strongly with Si with a R² of 0.97 (n=60; not shown) over the entire 650 water column. The profiles for dissolved La (Fig. 3c) are not statistically different within 651 analytical uncertainty (p=0.42). As part of a GEOTRACES intercalibration effort on rare 652 earth elements (REEs), Pahnke et al. (2012) reported La at several depths at the BATS station 653 from the US GEOTRACES Intercalibration cruise in 2008 and this data agrees with the here 654 reported data within analytical uncertainty (*i-p*-value=0.43). We included La as a typical example of a light REE and Lu works well as a heavy REE, but for the latter the 655

656 concentration factor needs to be increased. A recent paper has shown that this method does657 work well for the suite of REE's (Hatje et al., 2014).

658 The recovery for Ti was not quantitative at ~50% recovery, resulting in relatively poor 659 precision and accuracy. The concentrations were calculated based upon the standard addition 660 sensitivities assuming that all samples and standard additions had the same 50% recovery. 661 Nevertheless, the vertical profiles of both occupations agree quite well (Fig. 3e), and are not 662 statistically different within analytical uncertainty (*i-p*-value=0.80). Due to the low accuracy 663 it might not be suitable to look at the actual concentrations, but the observed trends should 664 represent actual variation in the ocean. However, one thing that is worth noting is that the 665 data of the Netherlands occupation is not significantly higher than the data from the US 666 occupation, indicating there is no noticeable contamination of Ti from the Netherlands 667 titanium frame.

Dissolved Ga is a scavenged type element with an atmospheric source to the surface ocean 668 669 (Orians and Bruland, 1988; Shiller, 1998). This is reflected at BATS in the elevated surface 670 concentrations, followed by a decrease with depth and relatively constant concentrations in 671 the deep waters, indicating equilibrium between input and scavenging removal (Fig. 3e). The 672 agreement between the two occupations is remarkably good considering the low overall 673 precision of the Ga measurements related to the non-quantitative recovery of Ga on the resin 674 and the profiles are not statistically different within analytical uncertainty (*i-p*-value=0.29). 675 The profile shape is also consistent with previous data from the region (Shiller, 1998). 676 Whereas these results are encouraging, non-quantitative recovery is not ideal and is likely to 677 result in significant variability between different measurement runs and hence, relatively 678 modest accuracy. Using isotope dilution should solve this issue. The results for Ga produced 679 with the current method should mainly be used to assess general trends rather than putting emphasis on the actual concentrations or small differences. 680

681 Dissolved Pb is an element with a strong anthropogenic atmospheric input related to the 682 use of leaded gasoline that peaked in the mid-1970's. Since leaded gasoline is no longer used, 683 the anthropogenic input to the surface oceans has been reduced and Pb concentrations in the 684 surface and subsurface Atlantic Ocean have been declining (Lee et al., 2011), due to scavenging removal of Pb. Subsurface concentrations of Pb were about 2 pmol kg⁻¹ higher 685 686 during the NL 2010 occupation than almost a year and a half later during the US occupation 687 of the same station. Below 2000 m, the profiles (Fig. 3f) are not statistically different within 688 analytical uncertainty (*i-p*-value=0.24).

689 Dissolved aluminium is not part of the ICP-MS multi-element method due to the poor 690 recovery of Al on the resin with the current method, especially in the presence of HF. 691 However, it is included in the discussion here as it is the material of construction for the US 692 sampling system and shipboard results are available. The result for Al by flow injection (FI) 693 method from the NL 2010 occupation and the previous US 2008 occupation with the same 694 US sampling system (Fig. 4) are not statistically different within analytical uncertainty (*i-p-*695 value=0.16) and the accompanying reference samples are within 1 standard deviation from 696 each other as well as the consensus values (Table 3), verifying the accuracy of the Al data. It 697 is worth noting that the data of the Netherlands occupation is not significantly lower than the 698 data from the US occupation, indicating there is no noticeable contamination of Al from the 699 US aluminium frame. The shipboard data for the US 2011 occupation (Measures et al., in 700 press) was significantly higher (*i-p*-value<0.01) than the current data and the 2008 701 occupation. This was also noted by Middag et al. (in press) and the reason for the 702 significantly higher values reported by Measures et al. (in press) in comparison to the 2008 703 and current data is currently unknown (see Middag et al. (in press) for further details).

704

705 Intercomparison between methods

Multi-element standard addition ICP-MS versus double spike ICP-MS

706

707 The data for Fe, Zn and Cd from the US occupation of the BATS profile as well as 2 708 additional stations (station KN204-1-12 at 29.7°N, 56.8°W and station KN204-1-20 at 709 22.3°N, 35.7°W) from the two different ICP-MS methods were compared with both the t-test 710 and the R-script. The samples were drawn from the same samplers at the same station and 711 cast. These data are attached in the supplementary material and are discussed in 712 oceanographic context elsewhere ((Conway and John, 2014a; Conway and John, 2014b; 713 Conway and John, 2015; Rijkenberg et al., 2014). The results of the standard addition ICP-714 MS method were triplicate extractions and measurements of the same sample. The standard 715 deviation of the triplicate results was used as the analytical uncertainty. The double spike 716 method is designed to obtain isotope ratios and uses more volume and samples were typically 717 extracted once and analysed twice. A 2% error was used as the analytical uncertainty 718 (Conway et al., 2013). 719 For Fe at the BATS station, the results (Fig. 5a) are not statistically different within

720 analytical uncertainty when using the R-script (*i-p*-value=0.09), but are significantly different 721 when considering the analytical uncertainty using the t-test approach (Table 5). When 722 comparing the 2 additional stations, no significant differences were observed using either test 723 for station 12, but the t-test approach reported a significant difference for station 20 (Table 5). 724 Thus the two ICP-MS methods produce comparable results, but the outcomes of the different 725 tests differ. The different test results between the t-test and the R script are related to the 726 manner of testing, where the R script is more reliable due to the simultaneous testing of both 727 differences and analytical uncertainty as detailed for Zn below.

For Zn at the BATS station, the results (Fig. 5b) are not statistically different using the ttest approach (p=0.21), but the results are different within analytical uncertainty when using the R-script (*i-p*-value=0.03) (Table 5). The p-value profile shows the main discrepancy is in

731 the upper 200m where the Zn concentrations are low (Fig. 5b and supplementary material). 732 When excluding these data points (first 8 of 37), the results are not statistically different 733 within analytical uncertainty (*i-p*-value=0.19). This implies there is an issue with accuracy for 734 the low Zn numbers, but small differences due to low level contamination cannot be excluded 735 either. The differing results between the t-test and R script, like for Fe, are related to the 736 manner of testing. The initial t-test does not consider analytical error and will give an 737 insignificant difference if the 2 profiles are 'snaking' around each other even if individual 738 differences are relatively large, hence the passing of the test for Zn. The secondary test is 739 only performed if the initial test fails (e.g. in case of Fe) and only considers the analytical 740 error and with respect to the absolute difference without regard for the directionality of the 741 difference. In contrast, the R script considers both factors simultaneously. When comparing 742 the 2 additional stations for Zn, no significant differences were observed using either test 743 (Table 5).

744 The low *p*-values in the p-value profile in the surface (Fig. 5b) were also observed for Fe 745 (Fig. 5a and supplementary material) and might be attributed to lower analytical 746 reproducibility for the lower concentrations, for example because of a greater contribution 747 from analytical blanks or lower signal/noise ratio during analysis, or to low level 748 contamination of the original samples, or to a combination of both factors. For the standard 749 addition ICP-MS method, the standard deviations where higher for the lowest concentrations 750 of Fe and Zn, confirming the lower reproducibility for the lower concentrations. Low level 751 contamination might arise from the used sample bottles or during filtrations, e.g. due to 752 metals leaching from the filter or particulates caught on the filter. More research is needed to 753 verify or exclude the latter hypothesis.

For Cd at the BATS station the results (Fig. 5c) are not statistically different within
analytical uncertainty when using the R-script (*i-p*-value=0.30), but are significantly different

756 within analytical uncertainty when using the t-test approach (Table 5). However, when 757 excluding the most deviation data points (more than 30% or 10 pmol/kg; 6 out of 37 data 758 points) the σ_d is significantly larger than the absolute difference for the entire profile and thus 759 are not different within analytical uncertainty (Table 5). When comparing the 2 additional 760 stations, no significant differences were observed using either test (Table 5). Thus the two 761 ICP-MS methods produce very comparable results, but some individual data points do differ. 762 The R script is less sensitive to these individual 'outliers' than the t-test approach. 763 Overall, the two ICP-MS based methods give results that are not statistically different for 764 Fe, Zn and Cd. The p-value profiles (Fig. 5 and supplementary material) show that for the 765 low surface ocean concentrations (closer to the LOD), there is more deviation between the 766 two methods. Additionally, the intercomparison shows the manner of testing is very 767 important and the R-script gives results that are more reliable due to the simultaneous testing 768 of the difference in concentration as well as the analytical uncertainty. The agreement 769 between profiles does differ between the different stations (Table 5) indicating that a single 770 cross-over station might not be representative. Multiple cross-over stations between transects 771 will often be unpractical, but repeat sampling at a cross-over station and independent analyses 772 (e.g. in different runs some weeks apart) could provide more representative intercomparisons. 773

115

*Multi-element standard addition ICP-MS versus shipboard flow injection methods*The metals Mn and Fe were also analysed shipboard using flow injection methods. Both
methods are based on chemiluminescence produced by the reaction between hydrogen
peroxide and luminol which is catalyzed by Fe or Mn (see Klunder et al. (2011) and Middag
et al. (2011a) for details). The vertical profiles, at first sight, demonstrate agreement between
the shipboard methods and the standard addition ICP-MS method (Fig. 2a, 2b), but for both
metals there is a small discrepancy. For Mn, the profiles are not significantly different within

781 analytical uncertainty using the R script (*i-p*-value=0.05) but the t-test approach does give a 782 significant difference within analytical uncertainty. The shipboard data is about ~0.03 nmol 783 kg⁻¹ lower at all depths. A similar offset was observed for the analysed reference samples 784 (Table 3), but results for both methods were within the consensus range. For the shipboard 785 data the blank was undeterminable, so no blank was subtracted. For the ICP-MS data a small blank of 0.04 nmol kg⁻¹ was determined and subtracted from the data. Perhaps this blank is an 786 787 underestimation, but that does not seem very likely since Mn is not a contamination sensitive 788 element. Adjusting either dataset for the observed offset makes them undistinguishable when 789 considering the analytical error using either test (*i-p*-value=0.83 and *p*=0.91 for the R script 790 and t-test approach, respectively), implying that this indeed is an absolute offset. However, 791 given that the offset is just insignificant based on the R-script, the deviation between the 792 profiles roughly equals the analytical uncertainty.

793 For Fe, the profiles are also not significantly different within analytical uncertainty using 794 the R-script (*i-p*-value=0.06), but the t-test approach does give a significant difference within 795 analytical uncertainty. The results for reference samples for both methods were within the 796 consensus range, but showed a similar discrepancy as observed for the profiles with the ICP-797 MS data being ~5% higher. Since the same relative offset is observed at high and low 798 concentrations, this discrepancy appears related to a difference in calibration. For the sake of 799 argument we adjusted the shipboard data towards the ICP-MS data based on the $\sim 5\%$ 800 difference observed in the reference samples. This results in the profiles being no longer 801 distinguishable using the t-test approach either (*i-p*-value=0.18 and *p*=0.82 for the R script 802 and t-test approach, respectively). As for Mn the deviation between the profiles is just smaller 803 than the overall analytical uncertainty.

804 It should be realised that the consensus values for the reference samples have a range and 805 therefore have limited suitability to be able distinguish small differences. Similarly, when

806 comparing to previously published data, there is inherent variation due to discrepancies in the 807 sampling depths as well as truly natural spatial and temporal variations that can mask small 808 differences between methods. Both the ICP-MS and shipboard data included reference 809 sample analysis and these agreed within the consensus range. When looking in detail, it is 810 noticed that there appears to be a small discrepancy between the vertical profiles (ICP-MS vs 811 shipboard) of dissolved Fe and Mn that is also represented between the reference samples, 812 but usually this would have been ignored as both were within the range of reproducibility and 813 accuracy. The small difference observed between the shipboard and ICP-MS methods is just 814 insignificant, but does warrant monitoring.

815

816 Conclusion

817 The intercomparison of the different Netherlands and US sampling system by analysing 818 samples from the BATS cross-over station with the same method and analyst demonstrated 819 that both sampling systems are capable of collecting uncontaminated samples for all the 820 GEOTRACES key elements. Nevertheless, at all times throughout the procedure, utmost care 821 must be taken to avoid inadvertent contamination. All results for reference samples are in 822 agreement with the current consensus values for those elements that have a consensus value 823 available. Moreover, the data reported here for trace metal profiles obtained with different 824 analytical methods are consistent, demonstrating comparibility between techniques. This 825 implies the analytical ICP-MS methods and the shipboard FI methods are functioning as 826 expected and produce accurate data that can be reliably compared between cruises, with the 827 caveat of Ga and Ti profiles, for which we did not yet achieve quantitative recovery. 828 Despite the strong consistency between collection and analytical techniques presented here, ultimately it is realised that all measurements have an uncertainty associated with them. 829 830 Comparison of results from the exact same station i.e. measuring sub-samples from the same

sample with different techniques allowed us to discern small differences between the
shipboard measurements and the ICP-MS measurements. We were fortunate enough to run
sub-samples from the same sample on the different methods, but these comparisons should
also be possible from cross-over station as long as that station is sampled at common depths
and in a relatively short timeframe to minimize natural variation, highlighting the importance
of cross-over stations to the GEOTRACES program.

837

838 *Comments and recommendations*

839 For the design of future crossover stations it is advisable to optimise the coordination of 840 sampling depths, or preferably common neutral density horizons. This would greatly benefit 841 the intercomparison, but might not aid intercomparison in the surface ocean if large variations 842 occur between occupations. For comparisons between analytical techniques or sampling 843 systems the deep ocean is simply best suited to make these comparisons. Comparison in the 844 surface ocean is inherently less accurate due to the larger interpolation necessary and the fact 845 that some processes (such as sinking and remineralising particles) are a function of depth as 846 well as density.

847 In order to merge or link data from different cruises, investigators and nations into a 848 coherent global data set, cross-over stations will be an invaluable tool. This has been done for 849 example by the CARINA program for ocean CO₂ and ancillary measurements (Olsen et al., 850 2009). It should be realised though, that one cross-over station in a long section is not a 851 definitive answer to resolve potential differences. It will only work if every data set is also 852 validated by regular measurements of the international reference samples to ensure 853 consistency within the international community and merged database. 854 A remark here with respect to the reference samples is that these should not only be 855 analysed, but the results also need to be submitted to the reference sample coordinator (K.W.

Bruland; <u>bruland@ucsc.edu</u>) in order to have the most representative updated consensus values available to the community. Furthermore, since the consensus values are not certified or constant, no corrections should be made to datasets based on analyses of reference standards; the values should just be part of the metadata. Internal seawater reference samples are most useful to identify any intra-dataset variability (i.e. variation between analyses performed on different days). The routine use of an internal seawater reference samples to be analysed far more frequently than the international reference samples is recommended,

simply due to the scarcity of the international reference samples.

864

863

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866

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- 1038
- 1039

1040 Tables

1041 Table 1 overview of the comparisons made in this paper

Comparison	Parameters		
Netherlands sampling system vs US sampling	Mn, Fe, Co, Ni, Cu, Zn, Cd, Y, La, Ti,		
system	Ga, Pb via ICP-MS (same analyst)		
Netherlands sampling system vs US sampling	Al via flow injection (different		
system	analysts)		
Standard addition ICP-MS vs double spike	Fe, Zn and Cd		
isotope dilution ICP-MS			
Standard addition ICP-MS vs shipboard flow	Fe and Mn		
injection			

1042

- 1043 Table 2 Blanks and Limit of Detection (LOD) defined as three times the standard deviation of
- 1044 the blank (average for the 8 different columns).

element	Unit	Blank	LOD
Y	pmol kg ⁻¹	0.36	0.21
Cd	pmol kg ⁻¹	0.11	0.10
La	pmol kg ⁻¹	0.06	0.08
Pb	pmol kg ⁻¹	0.94	0.16
Ti	pmol kg ⁻¹	4.22	3.97
Mn	nmol kg ⁻¹	0.03	0.02
Fe	nmol kg ⁻¹	0.01	0.02
Со	pmol kg ⁻¹	0.52	0.22
Ni	nmol kg ⁻¹	0.07	0.01
Cu	nmol kg ⁻¹	0.01	0.01
Zn	nmol kg ⁻¹	0.03	0.02
Ga	pmol kg ⁻¹	1.57	1.19

Metal Unit	Safe S1 91		Safe D2 492		Geotraces S 138		Geotraces D 146	
	Meas.	Cons.	Meas.	Cons.	Meas.	Cons.	Meas.	Cons.
Y	62±1		198±2		130±2		138±2	
pmol kg ⁻¹								
Cd	1.3±0.3	1.1±0.3	977±15	986±23	2.2±0.3	2.1±0.6	262±5	272±6
pmol kg ⁻¹								
La	4.5±0.1		30.6±0.5		13.3±0.3	14.7±1.1^	22.0±0.5	23.6±1.4^
pmol kg ⁻¹								
Pb	48.3±1.1	48.0±2.2	27.7±0.5	27.7±1.5	28.7±0.5	28.7±0.8	42.1±1.3	42.3±1.3
pmol kg ⁻¹								
Ti .	23±8	48±14*	140 ± 32	127±33*	66±19	75±12*	221±61	202±18*
pmol kg ⁻¹								
Mn	0.82±0.02	0.79±0.06	0.36 ± 0.02		1.53±0.04		$0.20{\pm}0.01$	
nmol kg ⁻¹			0.00.001	0.35±0.06	1.47.0.02	1.50±0.11		0.21±0.03
Mn SB			0.33±0.01		1.47±0.03		0.18±0.01	
nmol kg ⁻¹	0.10.0.01		0.0-0.04			0.546.0.046		1.00.010
Fe	0.10±0.01	0.093 ± 0.008	0.97±0.04		0.53±0.04	0.546±0.046	$0.94{\pm}0.04$	1.00 ± 0.10
nmol kg ⁻¹			0.00+0.02	0.933±0.023				
Fe SB			0.89±0.03					
nmol kg ⁻¹	2.0+0.2	4.0+1.5	46.0+1	45.0+2.0	22.4+1	21.0 1.2	(10:00	(45)10
Co	3.9±0.3	4.9±1.5	46.9±1	45.8±3.0	32.4±1	31.8±1.3	64.0±0.9	64.5±1.8
pmol kg ⁻¹ Ni	2.21±0.09	2.29±0.08	8.50±0.17	8.59±0.28	2.05±0.06	2.09±0.05	4.04±0.07	3.97±0.07
nmol kg ⁻¹	2.21±0.09	2.29±0.08	8.30±0.17	0.39±0.20	2.03±0.00	2.09±0.03	4.04±0.07	3.9/±0.0/
Cu	0.5±0.02	0.53±0.05	2.2±0.1	2.32±0.16	0.8±0.1	0.83±0.08	1.5±0.1	1.55±0.13
nmol kg ⁻¹	0.3±0.02	0.55 ± 0.05	2.2±0.1	2.32±0.10	0.8±0.1	0.85±0.08	1.5±0.1	1.55±0.15
Zn	0.07±0.03	0.068±0.010	7.50±0.20	7.47±0.27	0.04±0.02	0.042±0.008	1.76±0.06	1.73±0.09
nmol kg ⁻¹	0.07±0.05	0.000±0.010	7.50±0.20	7.47±0.27	0.04±0.02	0.042±0.000	1.70±0.00	1.75±0.07
Ga	11.1±0.8		7.0±0.4		41.4±2		29.0±1.5	
pmol kg ⁻¹					···· -			
Al SB					27.4±0.3		17.5±0.3	
nmol kg ⁻¹								
NL						27.5±0.2		17.7±0.2
Al SB US					27.4±0.5	1	17.8±0.3	
nmol kg ⁻¹								

1047 denoted one σ standard deviation.

1048 ^ Not actual consensus data but results reported by van de Flierdt et al. (2014)

1049 *Not actual consensus data but results reported by Croot (2011)

1050 SB denotes shipboard data

1051 Table 4 SAFe reference samples comparison of the results in Table 3 with the Mediterranean

Metal Unit	Safe S1 564	2-tailed paired t-test vs results accompanying BATS	Safe D2 54	2-tailed paired t-test vs results accompanying BATS
Y pmol kg ⁻¹	63±1	0.16	197±2	0.71
Cd pmol kg ⁻¹	1.5±0.1	0.21	976±12	0.91
La pmol kg ⁻¹	4.7±0.1	0.07	31.4±0.4	0.07
Pb pmol kg ⁻¹	49.1±0.2	0.2	27.4±0.2	0.31
Ti pmol kg ⁻¹	16±1	0.14	116±7	0.20
Mn nmol kg ⁻¹	0.81±0.01	0.60	0.38±0.01	0.09
Fe nmol kg ⁻¹	0.10±0.02	0.73	0.94±0.01	0.06
Ni nmol kg ⁻¹	2.28±0.02	0.17	8.62±0.18	0.40
Zn nmol kg ⁻¹	0.08±0.02	0.80	7.44±0.07	0.53
Ga pmol kg ⁻¹	11.2±0.9	0.91	8.5±1.2	0.06

1052 GEOTRACES expedition where shipboard extractions were performed (n=4)

1054Table 5 Intercomparison results between subsamples of the same profiles for 3 stations in the1055North Atlantic Ocean using Double spike isotope dilution ICP-MS versus standard addition1056ICP-MS. Both statistical testing approaches were used (R script and t-test approach). Please1057not that for the initial t-test a p>0.05 implies no significant differences between the profiles,1058whereas for the secondary t-test a p<0.05 implies the two methods are not statistically1059different within uncertainty.

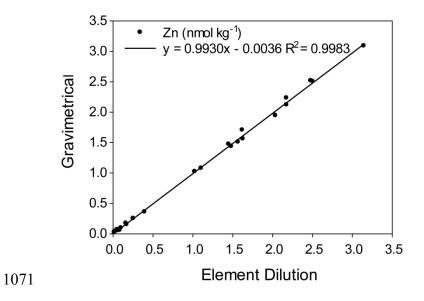
	Double Spike					
	В	ATS	KN204-1-12		KN204-1-20	
	t-test <i>p-</i> <i>value</i>	R script <i>i-p-value</i>	t-test <i>p-value</i>	R script <i>i-p-value</i>	t-test <i>p-value</i>	R script <i>i-p-value</i>
Standard addition Fe	0.99*	0.09	0.28#	0.16	0.99*	0.07
Standard addition Zn	0.22#	0.03	0.10#	0.11	0.47#	0.08
Standard addition Cd	0.11*	0.30	0.02\$	0.75	0.41#	0.10

1060 # initial t-test gives no significant difference (p>0.05)

- absolute difference between data points and the analytical results of the two methods are
- 1063 statistically different when considering uncertainty)
- 1064 \$T-test result when considering the analytical uncertainty < 0.05 (σ_d is greater than absolute
- 1065 difference than data points and the analytical results of the two methods are not statistically
- 1066 different within uncertainty)
- 1067 For the R script, a i-p-value > 0.05 means the analytical results of the two methods are not
- 1068 statistically different within uncertainty
- 1069

^{1061 *} T-test result when considering the analytical uncertainty > 0.05 (σ_d not greater than

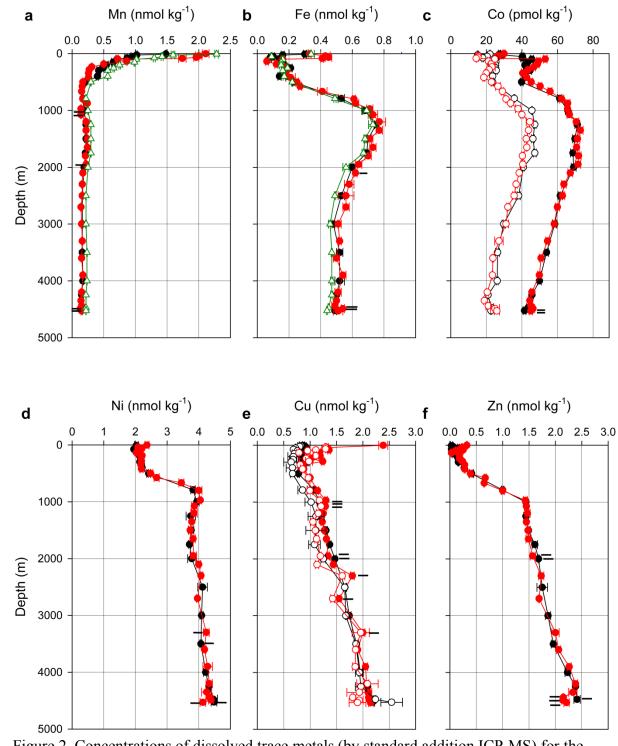
1070 Figures



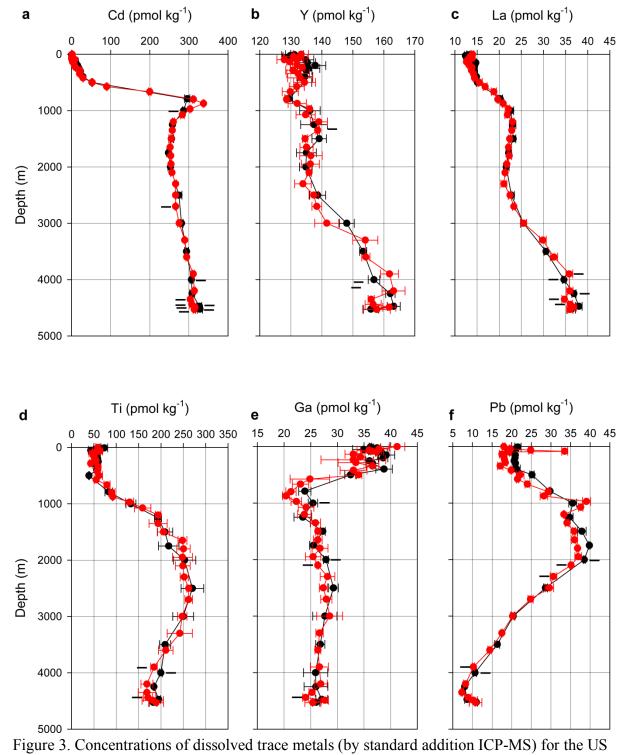
1072 Figure 1. Results of the element dilution calibration vs. the gravimetrical calibration for the 24

1073 samples of the Netherlands BATS occupation for dissolved Zn concentration, by standard

addition ICP-MS.

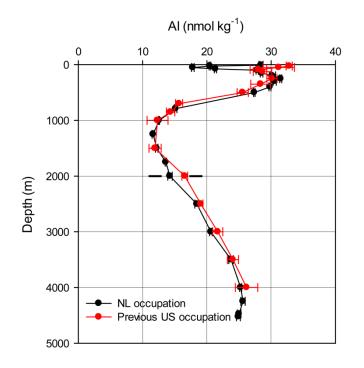


- 1081 Co and Cu. On each graph, a black horizontal bar indicates a significant difference (p < 0.05)
- 1082 between samples at individual depths (deeper than 1000 m) between the US and Netherlands'
- 1083 profiles (UV oxidised in the case of Co and Cu) observed in the *p*-value profile (see *Different*
- 1084 *samples approach* for calculation and the supplementary material for the *p*-value profiles).
- 1085 Note that the integrated *p*-value (*i-p*-value) reported in the text is based on the profiles deeper
- 1086 than 1000 m depth and does not equal the average of the *p*-values at individual depths (see
- 1087 *Different samples approach*).



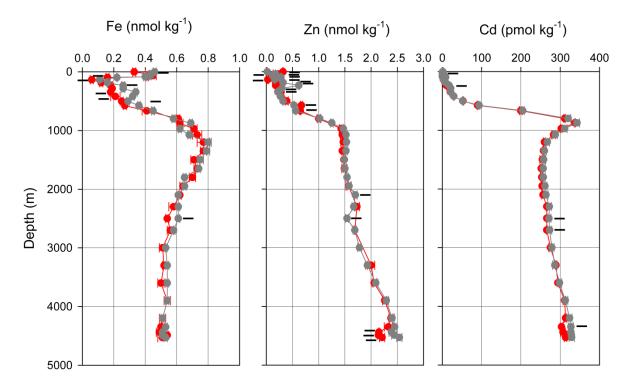
5000figure 3. Concentrations of dissolved trace metals (by standard addition ICP-MS) for the US1089(red circles) and Netherlands (black circles) occupations of the BATS station for A) Cd (pmol1090(red circles) and Netherlands (black circles) occupations of the BATS station for A) Cd (pmol1091kg⁻¹); B) Y (pmol kg⁻¹); C) La (pmol kg⁻¹); D) Ti (pmol kg⁻¹) E) Ga (pmol kg⁻¹); F) Pb (pmol1092kg⁻¹). On each graph, a black horizontal bar indicates a significant difference (p<0.05)1093between samples at individual depths (deeper than 1000 m) between the US and Netherlands'

profiles observed in the *p*-value profile (see *Different samples approach* for calculation and the supplementary material for the *p*-value profiles). Note that the integrated *p*-value (*i-p*value) reported in the text is based on the profiles deeper than 1000 m depth, except for Pb for which the test was based on depths deeper than 2000 m, and this *i-p*-value does not equal the average of the *p*-values at individual depths (see *Different samples approach*).



1100 Figure 4. Concentrations of Al (nmol kg⁻¹) from shipboard analysis during the Netherlands 1101 (black circles) and a previous (2008) US (red circles) occupation of the BATS station. The 1102 US data was analysed by Matt Brown from the University of California Santa Cruz (UCSC) 1103 and is the mean and standard deviation (error bars) of different samples filtered with different 1104 filtration systems as part of the initial Intercalibration efforts of the GEOTRACES program. 1105 The NL data is the mean and standard deviation (error bars) of triplicate analysis of the same 1106 sample. A black horizontal line denotes a significant difference between samples at 1107 individual depths (in the 1000 to 4000 m depth range) between the two profiles observed in 1108 the *p*-value profile (see *Different samples approach* for calculation and the supplementary 1109 material for the *p*-value profile). Note that the integrated *p*-value (*i*-*p*-value) reported in the

1110 text is based on the profiles between 1000 m and 4000 m depth. The excellent agreement is 1111 firm evidence by ocean crossover station approach of the accuracy of the measurements of 1112 dissolved Al in seawater by both laboratories (UCSC and NIOZ). This data was previously 1113 presented in a different form by Cutter and Bruland (2012).



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1115 Figure 5. Concentrations of trace metals for the standard addition ICP-MS method (red 1116 circles) and double spike isotope dilution ICP-MS method (grey circles) analyses at the US BATS station occupation for A) Cd (pmol kg⁻¹); B) Zn (nmol kg⁻¹); C) Fe (nmol kg⁻¹). A 1117 1118 black horizontal bar indicates a significant difference (p < 0.05) between samples at individual 1119 depths (whole water column) between the US and Netherlands' profiles observed in the *p*-1120 value profile (see *Different samples approach* for calculation and the supplementary material 1121 for the *p*-value profiles). Note that the integrated *p*-value (*i*-*p*-value) reported in the text is 1122 based on the entire profile as these samples were collected simultaneously. 1123

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