



Using mercury isotopes to understand the bioaccumulation of Hg in the subtropical Pearl River Estuary, South China



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HIGHLIGHTS

- Hg isotopic compositions in wild fish from subtropical south China showed different patterns.
- Small $\Delta^{199}\text{Hg}$ values in fish indicate less photo-degradation of MeHg before entering the food web.
- Significant differences of $\Delta^{199}\text{Hg}$ values among different species of fish show different feeding guilds.

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ABSTRACT

Coastal and estuarine regions are important areas of mercury pollution. Therefore, it is important to properly characterize the sources and bioaccumulation processes of mercury in these regions. Here, we present mercury stable isotopic compositions in 18 species of wild marine fish collected from the Pearl River Estuary (PRE), south China. Our results showed variations in mass-independent fractionation ($\Delta^{199}\text{Hg}$: $+0.05 \pm 0.10\text{‰}$ to $+0.59 \pm 0.30\text{‰}$) with a $\Delta^{199}\text{Hg}/\Delta^{201}\text{Hg}$ of ~ 1.26 , suggesting that aqueous MeHg underwent photo-degradation prior to incorporation into the food chain. For the results, we discovered small but significant differences of $\Delta^{199}\text{Hg}$ values among herbivorous, demersal, and carnivorous fish, indicating that different feeding guilds of fish may have incorporated MeHg with various degrees of photo-demethylation. The consistent mercury isotope compositions between fish feeding habitat and mercury sources in the estuary provide potentially important findings on the transformation and bioaccumulation of this toxic metal in subtropical coastal environments.

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1. Introduction

Mercury (Hg), a global pollutant, is released into the environment from both natural and anthropogenic sources. In an aqueous environment, inorganic Hg (IHg) can form methyl mercury (MeHg), a more toxic form that is readily bioaccumulated and biomagnified in aquatic food chains, posing a significant threat to aquatic ecosystems and human health (Fitzgerald et al., 2007; Blum et al., 2013). Anthropogenic activities have greatly altered the sources and biogeochemical processes of Hg, however, its impact on the

transformation and bioaccumulation in the aqueous environment is less clear (Fitzgerald et al., 2007).

With the recent documentation showing that Hg isotopes exhibit mass-dependent fractionation (MDF, as $\delta^{202}\text{Hg}$ values) and mass-independent fractionation (MIF, as $\Delta^{199}\text{Hg}$ values), Hg isotope geochemistry has greatly increased our understanding of the sources and fates of this toxin (Hintelmann, 2012; Blum et al., 2014; Yin et al., 2010a, 2014). Similar to $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$, Hg isotopes may offer insights into the sources and biogeochemical behavior of Hg in food webs (Bergquist and Blum, 2007; Senn et al., 2010; Jackson et al., 2008; Das et al., 2009; Gantner et al., 2009; Laffont et al., 2009, 2011; Gehrke et al., 2011; Perrot et al., 2010, 2012; Kwon et al., 2012, 2013, 2014; Blum et al., 2013; Li et al., 2014).

The $\delta^{202}\text{Hg}$ values have been applied to link fish and Hg sources.

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For instance, a strong correlation between $\delta^{202}\text{Hg}$ in fish and that in co-located intertidal sediments was found in the San Francisco Estuary, USA (Gehrke et al., 2011). However, the use of fish $\delta^{202}\text{Hg}$ to trace Hg sources was called into question by the occurrence of Hg-MDF during Hg bioaccumulation (Jackson et al., 2008; Das et al., 2009). Higher $\delta^{202}\text{Hg}$ values have been observed in mammal predators [e.g., humans (Laffont et al., 2009, 2011; Sherman et al., 2013), whales (Li et al., 2014), seals (Perrot et al., 2012), and birds (Point et al., 2011)] than in their fish prey. Recent studies observed no Hg-MDF during MeHg bioaccumulation by fish (Kwon et al., 2012, 2013), whereas intestinal MeHg degradation can cause significant MDF in mammals (Sherman et al., 2013). The most intriguing finding is the observation of Hg-MIF in fish (Bergquist and Blum, 2007; Jackson et al., 2008; Das et al., 2009; Gantner et al., 2009; Laffont et al., 2009, 2011; Gehrke et al., 2011; Perrot et al., 2010, 2012; Senn et al., 2010; Kwon et al., 2012, 2013, 2014; Blum et al., 2013; Li et al., 2014). The Hg-MIF in fish was initially explained to be caused during *in vivo* biological processes (Das et al., 2009), whereas such an explanation is not supported by the observation of no Hg-MIF during microbial transformations (Rodríguez-González et al., 2009; Kritee et al., 2013; Jiménez-Moreno et al., 2013), bioaccumulation and trophic transfer (Laffont et al., 2009, 2011; Perrot et al., 2012; Kwon et al., 2012, 2013; Blum et al., 2013). The Hg-MIF in fish was later explained by the aqueous MeHg photo-degradation process prior to bioaccumulation, due to the fact that >90% of Hg in fish is MeHg (Blum et al., 2013). With unperturbed Hg-MIF during bioaccumulation, it is speculated that fish Hg-MIF can provide important information on the photochemical cycling of MeHg in aquatic ecosystems (Bergquist and Blum, 2007; Blum et al., 2013). For instance, higher Hg-MIF ($\Delta^{199}\text{Hg} > 1\text{‰}$) has been documented in pelagic fish than in coastal fish ($\Delta^{199}\text{Hg} < 1\text{‰}$), indicating that MeHg in the open ocean water column is subjected to more photo-degradation than in coastal waters (Gehrke et al., 2011; Senn et al., 2010; Blum et al., 2013; Kwon et al., 2014). A systematic decline in $\Delta^{199}\text{Hg}$ values with the depth of fish feed was observed, suggesting that Hg-MIF can be a good indicator of the feeding depth of fish (Blum et al., 2013).

The Pearl River Estuary (PRE) is a subtropical estuary located in

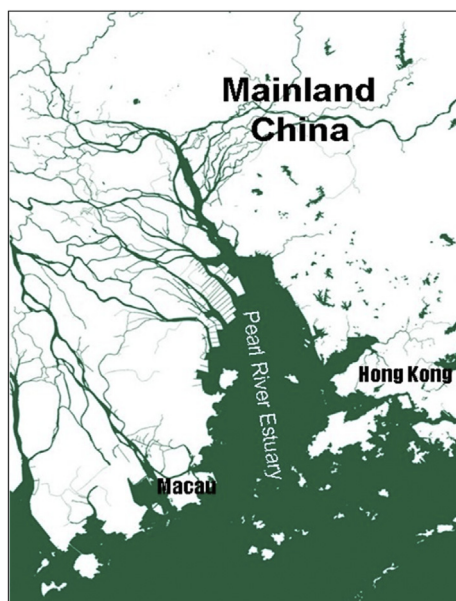


Fig. 1. Map showing the Pearl River Estuary (PRE), South China.

the northern part of the South China Sea (SCS) (Fig. 1). During the last few decades, rapid economic development in the surrounding Pearl River Delta (PRD) region has led to excessive discharges of Hg and other pollutants into the PRE (Fu et al., 2010; Liu et al., 2011, 2012). Our recent study investigated Hg isotopes in surface sediments of the PRE, which indicated that the PRE receives Hg from atmospheric deposition, industrial waste discharges, urban emissions, and watershed runoff, with distinct Hg isotopic signatures (Yin et al., 2015). However, the impact of Hg pollution on the bioaccumulation in the aquatic food web in such a subtropical region is poorly understood. Here, we investigated the Hg isotopic variations in 18 fish species in the PRE. The objectives of this study were (1) to investigate the sources of MeHg in different feeding guilds of fish; (2) to evaluate the capability of Hg isotopes in tracking Hg through aquatic food webs using Hg isotopes; and (3) to evaluate the potential effect of human activities on Hg dynamics in food webs of the PRE.

2. Materials and methods

2.1. Fish sampling

The PRE (Fig. 1) covers an area of ~2000 km², with depths varying from 0 to 30 m. The average distance is 49 km from north to south, and 4–58 km from east to west (Shi et al., 2010; Yin et al., 2015). A total of 39 individual fish from 18 species of fish were collected in the PRE from August to December 2011. Details of the processing of the samples, the total mercury (THg) concentrations (9.5 ± 2.4 to 53 ± 15.2 ng g⁻¹ wet weight), the MeHg concentrations (8.4 ± 1.9 to 50.3 ± 16.0 ng g⁻¹ wet weight), the $\delta^{15}\text{N}$ ($10.2 \pm 0.4\text{‰}$ to $18.7 \pm 0.4\text{‰}$), and the $\delta^{13}\text{C}$ ($-23.4 \pm 0.5\text{‰}$ to $16.9 \pm 0.2\text{‰}$) of the fish samples have been given by Pan et al. (2014). The average fraction of MeHg (F_{MeHg}) in the fish ranged from $87.6 \pm 4.8\%$ to $102.5 \pm 7.8\%$, consistent with the previous finding that the majority of the Hg in fish was MeHg (Fitzgerald et al., 2007; Blum et al., 2013). The analyses of Hg isotopes performed in this study were for THg. Based on THg, $\delta^{15}\text{N}$ data, and dietary information, the fish in the PRE can be categorized into three main groups: herbivorous, demersal, and carnivorous (Fig. 2).

2.2. Hg isotopic composition analysis

About 0.5 g of dried samples (wet/dry ratio: 4.2–4.5) were weighed and digested at 95 °C for 3 h with a 10 mL acid mixture ($\text{HNO}_3:\text{H}_2\text{SO}_4 = 7:3$, v/v). About 500 μL of BrCl was later added to each sample and kept for 12 h to allow the conversion of Hg to Hg(II). Then, 5 mL of a 20 M sodium hydroxide (NaOH) solution were slowly added to each digest to ensure that the acid concentration was <20%. Finally, the digests were diluted to 1–3 ng mL⁻¹ of Hg using Milli-Q water (18.2 M Ω cm). Certified reference material (TORT-2, Lobster *Hepatopancreas*) was prepared in the same way as the fish samples. The THg recovery of TORT-2 was in the range of 94–109% ($n = 6$). A mercury isotope analysis was based on a sample-standard bracketing method validated by Yin et al. (2010b). The bracketing standard solutions (NIST SRM 3133) in a 10% acid

Table 1
Comparisons among the published Hg isotopic values of TORT-2.

$\delta^{202}\text{Hg}$	2sd	$\Delta^{199}\text{Hg}$	2sd	n	References
0.04	0.12	0.72	0.08	13	Tui et al. (2012)
0.10	n.a	0.79	n.a	1	Kwon et al. (2014)
-0.33	0.16	0.82	0.07	3	Masbou et al. (2013)
-0.09	0.12	0.73	0.08	6	This study

n.d. means not available.

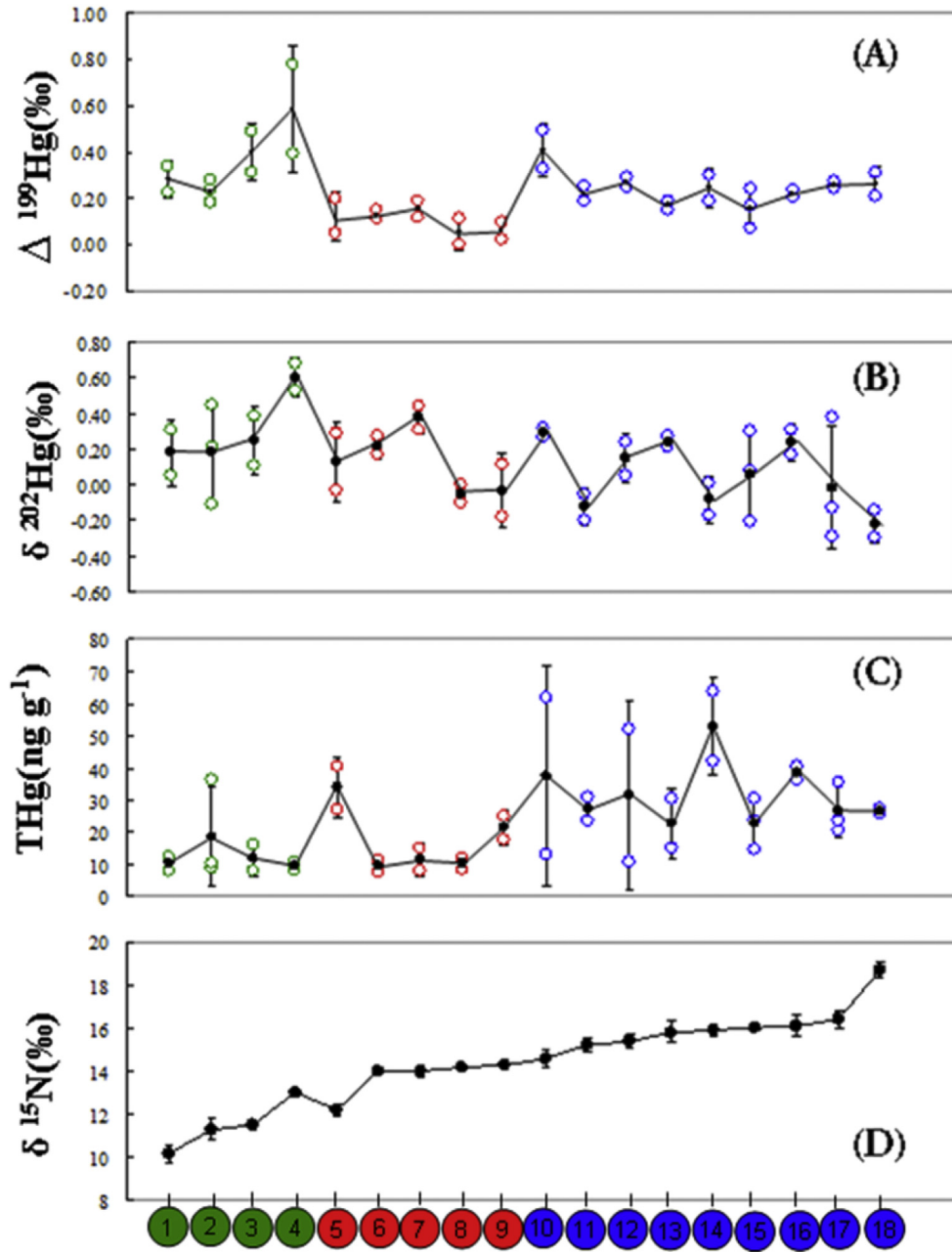


Fig. 2. $\delta^{202}\text{Hg}$ (A), $\Delta^{199}\text{Hg}$ (B), THg (C) and $\delta^{15}\text{N}$ (D) values in herbivorous [1. Flathead grey mullet (n = 2); 2. Western Pacific gizzard shad (n = 3); 3. Taiwanese mullet (n = 2) and 4. Silver pomfret (n = 2)], demersal [5. Shortnose ponyfish (n = 2); 6. Large scale mullet (n = 2); 7. Mottled spinefoot (n = 2); 8. Speckled tonguesole (n = 2) and 9. Largescale tonguesole (n = 2)] and carnivorous fish [10. Belanger’s croaker (n = 2); 11. Donkey croaker (n = 2); 12. Bartail flathead (n = 2); 13. Banded grouper (n = 2); 14. Devil stinger (n = 2); 15. Russell’s Jewfish (n = 3); 16. Striped Catfish (n = 2); 17. Longtooth grouper (n = 3) and 18. Yellow Drum (n = 2)]. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

solution ($\text{HNO}_3:\text{H}_2\text{SO}_4 = 7:3$, v/v) were adjusted to within 10% of the THg concentration in the fish digests. Mercury isotope compositions are reported in terms of per mil deviations from the NIST SRM 3133 Hg standard and expressed in delta notation (Blum and Bergquist, 2007):

$$\delta^{\text{xxx}}\text{Hg}(\text{‰}) = \left\{ \frac{(\text{xxxHg}/^{198}\text{Hg}_{\text{sample}})}{(\text{xxxHg}/^{198}\text{Hg}_{\text{NIST 3133}})} - 1 \right\} \times 1000 \quad (1)$$

and xxx represent the mass of Hg isotopes between 199 and 202. Hg-MIF is reported in $\Delta^{\text{xxx}}\text{Hg}$ notation (deviation from mass dependency in units of permil, ‰) using the following formulas (Blum

and Bergquist, 2007):

$$\Delta^{199}\text{Hg} \approx \delta^{199}\text{Hg} - (\delta^{202}\text{Hg} * 0.2520) \quad (2)$$

$$\Delta^{201}\text{Hg} \approx \delta^{201}\text{Hg} - (\delta^{202}\text{Hg} * 0.7520) \quad (3)$$

Replicate digests were prepared for each fish sample (n = 2) and standard reference material (TORT-2) (n = 6). The UM-Almadén standard solution was measured once in every 10 samples. Data uncertainties in this study corresponded to the larger value of the measurement uncertainty of either the replicate fish digests or UM-Almadén. Measurements of UM-Almadén ($\delta^{202}\text{Hg}$: $-0.49 \pm 0.11\text{‰}$;

$\Delta^{199}\text{Hg}$: $-0.05 \pm 0.08\%$; $\Delta^{201}\text{Hg}$: $-0.05 \pm 0.06\%$; 2σ , $n = 11$) were consistent with previous data (Blum and Bergquist, 2007). As shown in Table 1, the TORT-2 ($\delta^{202}\text{Hg}$: $-0.09 \pm 0.12\%$, $\Delta^{199}\text{Hg}$: $-0.73 \pm 0.08\%$, $\Delta^{201}\text{Hg}$: $-0.55 \pm 0.08\%$, 2σ , $n = 6$) was within the range of errors of previous studies (Masbou et al., 2013; Tsui et al., 2012; Kwon et al., 2014).

3. Results and discussion

3.1. Differences in Hg isotope compositions between sediments and fish in the PRE

The Hg isotopic compositions of the 18 species of fish are summarized in Table S1 of Supporting Information. Various degrees of both $\delta^{202}\text{Hg}$ ($-0.22 \pm 0.10\%$ to $+0.38 \pm 0.10\%$) and $\Delta^{199}\text{Hg}$ ($+0.05 \pm 0.10\%$ to $+0.59 \pm 0.30\%$) (Fig. 2A and B) were observed among different feeding guilds of fish, which may be interpreted as having been caused by the mixing of isotopically distinct sources and isotope fractionation during Hg biogeochemical cycling in sediments (or water column) and during trophic transfer to fish (Bergquist and Blum, 2007; Jackson et al., 2008; Das et al., 2009; Gantner et al., 2009; Laffont et al., 2009, 2011; Gehrke et al., 2011; Senn et al., 2010; Perrot et al., 2010, 2012; Kwon et al., 2012, 2013, 2014; Blum et al., 2013; Li et al., 2014).

In the PRE, Hg is released into the environment mainly in inorganic Hg (IHg) forms (Yin et al., 2015). Inorganic Hg is subsequently converted to MeHg in deep anoxic water and sediments by sulfate- and iron-reducing bacteria (Fitzgerald et al., 2007; Kwon et al., 2014). An apparent increase in MeHg versus $\delta^{13}\text{C}$ was observed in the PRE fish, suggesting that the trophic transfer of MeHg was largely via the benthic pathway. In the PRE, high levels of MeHg in sediments (Shi et al., 2010) and high fractions of MeHg in seawater (~10%) have been reported (Fu et al., 2010), compared to the much lower fractions of MeHg (~1%) in precipitation and river waters (Fu et al., 2010; Liu et al., 2012). This indicates that external sources of MeHg (e.g., riverine and atmospheric inputs of MeHg) are generally limited in the PRE. The Hg isotopic compositions of the PRE sediments have been investigated and characterized to have relative low $\delta^{202}\text{Hg}$ (-1.92% to -0.68% , $n = 27$) and $\Delta^{199}\text{Hg}$ (-0.15 to $+0.16\%$, $n = 27$) values (Yin et al., 2015). The observation of the isotopic differences between fish and sediments is important for understanding the biogeochemical cycling of Hg in the PRE. The positive offset in $\delta^{202}\text{Hg}$ between fish and sediment in this study is well agreed with numerous recent studies. As no Hg-MDF has been demonstrated to occur during the assimilation of Hg by fish (Kwon et al., 2012, 2013), we suggest that fish mainly assimilate Hg with higher $\delta^{202}\text{Hg}$ values. $\delta^{202}\text{Hg}$ reported in this study are for bulk fish Hg, which contains the dominant fraction of MeHg and minor IHg (Table S1). The IHg in the PRE fish may represent the bioavailable IHg in sediments and water column, while MeHg mainly comes from Hg methylation of the bioavailable IHg fractions. Bioavailable IHg species have been reported as being characterized by higher $\delta^{202}\text{Hg}$ values (Yin et al., 2013a). The loss of lighter Hg isotopes during Hg adsorption of sediment particles [containing thiol (Wiederhold et al., 2010) and goethite (Jiska et al., 2012), Hg(II) photo-reduction (Bergquist and Blum, 2007; Zheng and Hintelmann, 2009), microbial Hg(II) reduction (Kritee et al., 2013), and evasion processes (Zheng et al., 2007) can shift the bioavailable IHg to higher $\delta^{202}\text{Hg}$ values. In sediments and bottom water, MeHg is expected to have relative lower $\delta^{202}\text{Hg}$ than the IHg, because preferential methylation of light Hg isotopes has been demonstrated during biotic and abiotic methylation of bioavailable IHg (Rodríguez-González et al., 2009; Jiménez-Moreno et al., 2013). However, once transported to the surface water column via diffusion, advection and resuspension processes, subsequent photo- and

microbial-demethylation of Hg can shift the MeHg toward higher $\delta^{202}\text{Hg}$ values (Bergquist and Blum, 2007; Kritee et al., 2013). Differences in water conditions (e.g., depth and water clarity) may cause significant variations in the isotopic signatures of IHg and MeHg in water column and fish among different regions (Blum et al., 2013). For instance, Masbou et al. (2013) reported higher $\delta^{202}\text{Hg}$ in MeHg than in the IHg fraction, while Jackson et al. (2008) demonstrated higher $\delta^{202}\text{Hg}$ in IHg than in the MeHg fraction. In this study, negative correlations between fish $\delta^{202}\text{Hg}$ and THg (Fig. 3A, $p < 0.05$, ANOVA) were observed, which contrasts with the findings reported in previous literature (Bergquist and Blum, 2007). However, no clear correlation between $\delta^{202}\text{Hg}$ and F_{MeHg} can be found ($p > 0.3$, ANOVA). Given the fact that the fish collection represents diverse feeding habitats in different locations of the PRE, it is still unclear whether the $\delta^{202}\text{Hg}$ variability represents the variability of MeHg with isotopically distinct sources, or a mixing of IHg (with high $\delta^{202}\text{Hg}$) and MeHg (with low $\delta^{202}\text{Hg}$). Hence, more studies are needed in the future to investigate the key processes involved in the subtropical estuary.

Compared to the PRE sediments, small positive $\Delta^{199}\text{Hg}$ values were observed for the fish in the PRE (Fig. 2A). No clear correlations were observed between THg and $\Delta^{199}\text{Hg}$ (Fig. 3B, $p > 0.10$, ANOVA), or between F_{MeHg} and $\Delta^{199}\text{Hg}$ ($p > 0.10$, ANOVA). Insignificant Hg-MIF during bioaccumulation and trophic transfer have been documented in prey-predator (e.g., fish-human, fish-seal) (Laffont et al., 2009, 2011; Point et al., 2011; Perrot et al., 2012; Sherman et al., 2013; Li et al., 2014) and plant studies (e.g., rice and aspen tree) (Demers et al., 2013; Yin et al., 2013b), as well as fish feeding experiments (Kwon et al., 2012, 2013). The Hg-MIF has been understood by the nuclear volume effect (NVE) and the magnetic isotope effect (MIE) (Schauble, 2007; Buchachenko et al., 2007). In laboratory experiments, the NVE was observed during several processes [e.g., elemental Hg volatilization (Estrade et al., 2009), the non-photo reduction of Hg species (Zheng and Hintelmann, 2010), and

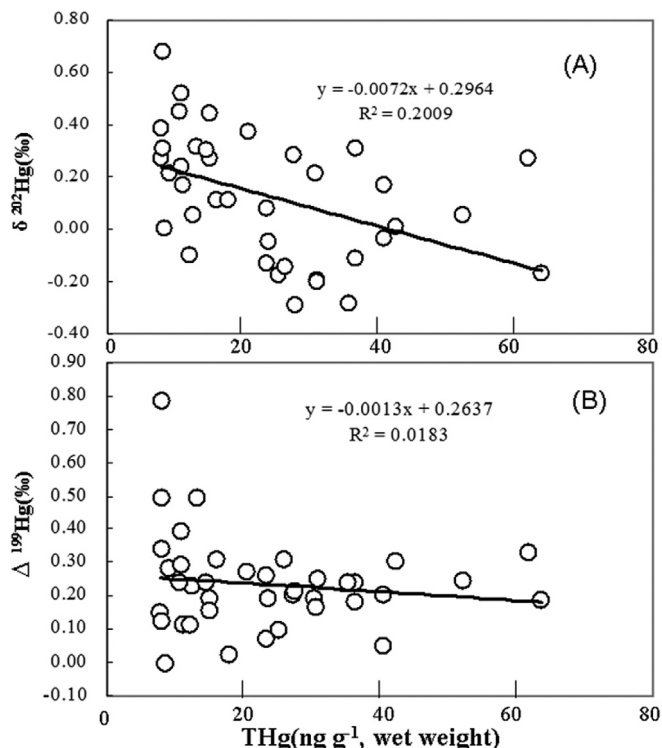


Fig. 3. Correlations of THg to $\delta^{202}\text{Hg}$ (A) and $\Delta^{199}\text{Hg}$ (B) in the PRE fish.

Hg-thiol complexation (Wiederhold et al., 2010)], demonstrating $\Delta^{199}\text{Hg}/\Delta^{201}\text{Hg}$ of ~ 1.6 . These processes may not represent the Hg-MIF in the PRE fish because the linear regression of $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ yielded a slope of 1.26 (Fig. 4, $p < 0.05$, ANOVA), which is consistent with fish in other coastal regions (Gehrke et al., 2011; Senn et al., 2010; Kwon et al., 2014). Laboratory experiments on the aqueous MeHg photo-degradation induced by MIE showed $\Delta^{199}\text{Hg}/\Delta^{201}\text{Hg}$ of ~ 1.3 , which was in accordance with the $\Delta^{199}\text{Hg}/\Delta^{201}\text{Hg}$ observed in the PRE fish (Bergquist and Blum, 2007). Hence, it is thought that the Hg-MIF is probably first generated by the MeHg photo-degradation prior to being transferred to fish.

3.2. Using Hg-MIF to differentiate aquatic trophic structures

Once methylated, MeHg can be released to the water column via diffusion, advection, and resuspension, and subjected to photo-degradation prior to being transferred to fish via food uptake and bioaccumulation (Kwon et al., 2013). Therefore, Hg-MIF in fish can reveal important information on the MeHg exposure pathways and characteristics of water columns (Bergquist and Blum, 2007; Gantner et al., 2009; Gehrke et al., 2011; Senn et al., 2010; Blum et al., 2013; Sherman and Blum, 2013; Kwon et al., 2014). In the present study, both demersal fish and herbivorous fish showed lower THg and $\delta^{15}\text{N}$ values (Table 2 and Fig. 5). A possible explanation for this is the fact that demersal fish mainly inhabit and feed on benthic organisms in bottom waters (close to sediment interface), and herbivorous fish mainly live and feed on algae and seagrass in the photic zone (Table S1). Large differences in $\Delta^{199}\text{Hg}$ values ($p < 0.05$, ANOVA) were observed between demersal fish (mean: $+0.10 \pm 0.07\%$, σ , $n = 10$) and herbivorous fish (mean: $+0.36 \pm 0.18\%$, σ , $n = 10$). The offset of Hg-MIF between the pelagic and benthic organisms was also reported in several arctic lakes (Gantner et al., 2009) and in the North Pacific Subtropical Gyre (Blum et al., 2013). The low Hg-MIF in demersal fish confirms that the source of MeHg in the PRE fish was largely via the benthic pathways. We suggested that demersal fish are mainly exposed to a benthic MeHg source that has not undergone much photochemical breakdown, while herbivorous fish are more likely to represent MeHg that has undergone intensive photo-degradation in surface

waters. Carnivorous fish show intermediate mean $\Delta^{199}\text{Hg}$ value ($+0.24 \pm 0.09\%$, σ , $n = 20$) between that of demersal and herbivorous fish (Table 2). Carnivorous fish with the highest THg and $\delta^{15}\text{N}$ values (Fig. 2) suggest that such fish prey upon herbivorous, demersal fish, and other marine species, which will result in the intermediate Hg-MIF (Fig. 5). This further supports the hypothesis that the Hg-MIF in biological samples is directly constrained by the isotopic signature of the assimilated MeHg, rather than by *in vivo* mass-independent isotope fractionation (Blum et al., 2013; Kwon et al., 2012, 2013).

3.3. Low Hg-MIF in the PRE fish

Fig. 4 shows differences in Hg-MIF among fish from different geographic areas. In comparison with fish in the open ocean (Senn et al., 2010; Blum et al., 2013), much smaller Hg-MIF was observed in fish in the PRE and other coastal areas (Gehrke et al., 2011; Kwon et al., 2014), indicating that MeHg has undergone much less photo-degradation before entering the coastal food web. Differences in water circulation between the pelagic and coastal areas may be responsible for the variability of fish Hg-MIF caused by the MeHg photo-degradation (Senn et al., 2010). The Hg-MIF signatures in fish have been used to represent the degree of MeHg photo-degradation in the water column (Blum et al., 2013; Sherman et al., 2013; Kwon et al., 2014). In the PRE, a low rate of MeHg photo-degradation is expected due to low water clarity in the PRE (generally < 1 m) and to the riverine transport of large volumes of suspended particulate matter (Liu et al., 2012; Yin et al., 2015). We estimated the percentage of MeHg photo-degradation based on the $\Delta^{199}\text{Hg}$ values of the PRE fish. We calculated the expected relationship between $\Delta^{199}\text{Hg}$ and the fraction of Hg^{2+} remaining in the aqueous system according to:

$$10^3 \cdot \ln[(10^{-3} \Delta^{199}\text{Hg}_t + 1)/(10^{-3} \Delta^{199}\text{Hg}_i + 1)] = S \cdot \ln(1-f) \quad (4)$$

Where f represents the fraction of MeHg that was photo-demethylated; $\Delta^{199}\text{Hg}_i$ represents the initial $\Delta^{199}\text{Hg}$ value ($\Delta^{199}\text{Hg}_i = 0$), and $\Delta^{199}\text{Hg}_t$ represents the $\Delta^{199}\text{Hg}$ value at a given time (here we use the fish data). The slope (S) is calculated by

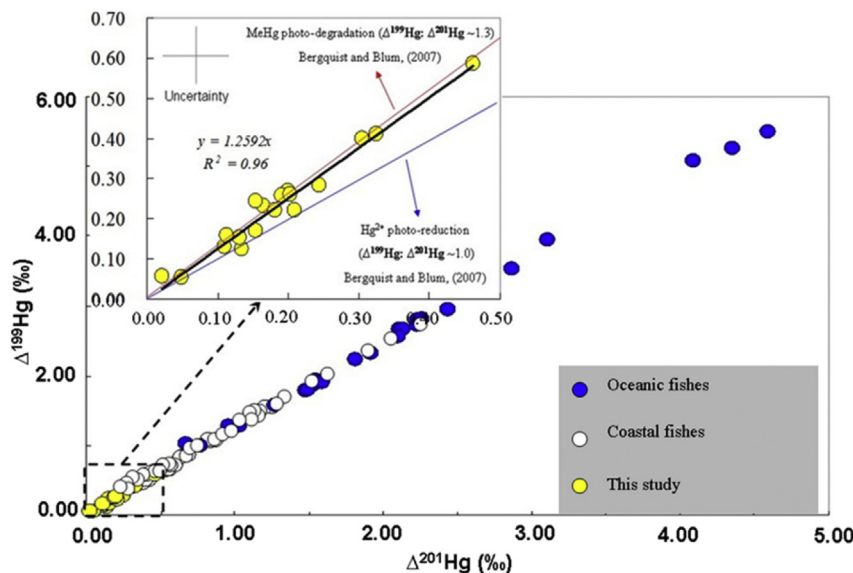


Fig. 4. Mass-independent fractionation of Hg in coastal and oceanic fish. Oceanic fishes were collected from Gulf of Mexico (Senn et al., 2010) and North Pacific Ocean (Blum et al., 2013); Coastal fishes were collected from Terrebonne Bay (Senn et al., 2010), San Francisco Estuary (Gehrke et al., 2011) and the Northeast coast estuaries (Kwon et al., 2014) of United States.

Table 2
Fish species, $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ isotopes, Hg contents and bulk tissue Hg isotopic compositions in different fish groups of the PRE.

Fish groups	n	THg ng g ⁻¹	σ ng g ⁻¹	F _{MeHg} %	σ %	$\delta^{15}\text{N}$ ‰	σ ‰	$\delta^{202}\text{Hg}$ ‰	σ ‰	$\Delta^{199}\text{Hg}$ ‰	σ ‰	$\Delta^{201}\text{Hg}$ ‰	σ ‰
Herbivorous fish	9	75.8	33.3	94.1	5.8	11.48	1.00	0.29	0.25	0.36	0.18	0.30	0.16
Demersal fish	10	77.4	40.4	96.5	5.5	13.71	0.87	0.13	0.20	0.10	0.07	0.10	0.06
Carnivorous fish	20	208.2	314.3	98.6	7.4	16.03	1.05	0.06	0.23	0.24	0.09	0.19	0.08

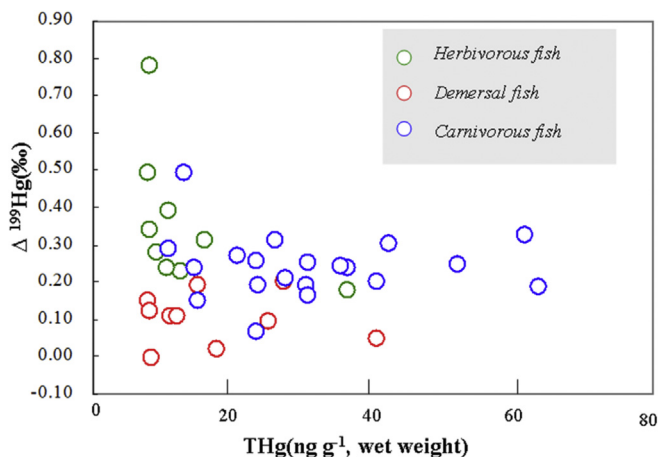


Fig. 5. Utilization of THg and $\Delta^{199}\text{Hg}$ to distinguish Hg sources in the PRE fish. ANOVA tests indicate statistical differences in $\Delta^{199}\text{Hg}$ values between herbivorous and demersal fish ($P < 0.05$), and statistically higher THg values for carnivorous fish than herbivorous and demersal fish ($P < 0.05$). Insignificant correlations can be observed with each group of fish, and the combined data ($P > 0.2$, ANOVA).

plotting $10^3 \cdot \ln\left[\frac{(10^{-3}\Delta^{199}\text{Hg}_t + 1)}{(10^{-3}\Delta^{199}\text{Hg}_i + 1)}\right]$ versus $\ln(1-f)$ for the relevant MeHg photo-degradation (Bergquist and Blum, 2007). For the 10 mg/L DOC photo-reductions of Hg^{2+} experiment, $S = -7.82$. Finally, we estimated that less than 10% of MeHg is photo-degraded prior to entering the food web.

4. Conclusions

Coastal areas such as the PRE have received Hg inputs from natural and anthropogenic sources (Yin et al., 2015). Methylation of Hg in deep anoxic waters and sediments contributes to the dominant sources of Hg in aquatic ecosystems (Kwon et al., 2014). In the surface water column, MeHg can be subjected to photo-degradation prior to being bioaccumulated into aquatic food webs; hence, Hg-MIF can be subsequently coded in fish. The present study observed much lower Hg-MIF in the PRE fish compared to other coastal and oceanic fish, indicating that much less MeHg photo-degradation occurs in the PRE than in coastal and oceanic waters. The environmental risk associated with MeHg in fish can increase due to the low efficiency of MeHg photo-degradation. As a result, Hg-MIF in fish can be used to evaluate the process of MeHg accumulation. This study documented for significant variability in Hg-MIF among herbivorous, demersal, and carnivorous fish species, suggesting that different feeding guilds of fish may have incorporated MeHg with various degrees of photo-demethylation in the water column. Based on these results, we concluded that Hg isotopes (especially Hg-MIF) can be a powerful tool for revealing the exposure pathways and geochemical behaviors of MeHg in coastal food webs.

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Appendix A. Supplementary data

Supplementary data related to this article can be found at <http://dx.doi.org/10.1016/j.chemosphere.2015.12.100>.

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