

Comprehensive assessment for controlling factor of total Hg level in skipjack tuna from Western North Pacific Ocean

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Background

- Migratory pelagic marine species such as tuna are particularly significant source of MeHg to human (Sunderland, 2007; Yasutake et al., 2003).
- Even among the same species, considerable geographical variation have been reported (Hisamichi et al., 2010; Sunderland, 2007).
- Although age/size dependent increase of Hg has commonly observed, this trend can not be explained solely by bioenergetic process (Trudel & Rasmussen, 2006). To assess diet composition is critical point for mass balance modelling (Ferriss & Essington, 2014).
- In this study, we discuss about controlling factor of THg level in skipjack tuna considering size, trophic position, regional variation, and stable isotope signatures.

Conclusions

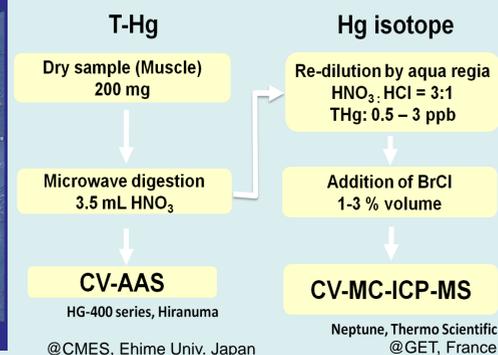
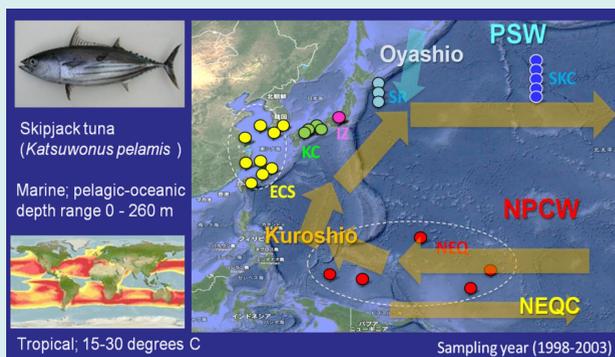
PSW (Polar Subarctic Water): Small size, low THg, high $\Delta^{199}\text{Hg}$ (foraging diet from shallower water).

Kuroshio: Large size, high THg, low $\Delta^{199}\text{Hg}$ (foraging diet from deeper water).

NEQ (Near Equator): Large size, low THg, intermediate $\Delta^{199}\text{Hg}$.

Implications: Variation of $\Delta^{199}\text{Hg}$ possibly the function of foraging depth. Accepting this assumption, body size dependent increase of THg is attributed to the foraging from deep water having higher MeHg concentration. Further study for depth vs isotope signature will be important.

Materials and Methods



Accuracy of measurements were confirmed by measurement of ERM-CE464 (n=8; IRMM, Belgium), DOLT-4 (n=20; NRC, Canada) and oyster tissue (n=35, SRM 1566b, NIST, USA), with the average recovery being 103, 108, and 110%, respectively. The $\delta^{202}\text{Hg}$ of overall measurements of UM-Almaden, ETH-Fluka, and ERM-CE464 were $-0.57 \pm 0.12\text{‰}$ (n=30), $-1.44 \pm 0.24\text{‰}$ (n=46), and $0.59 \pm 0.11\text{‰}$ (n=8), respectively. The $\Delta^{199}\text{Hg}$ of ERM-CE464 was $2.38 \pm 0.02\text{‰}$.

Results & Discussion

Regional variation

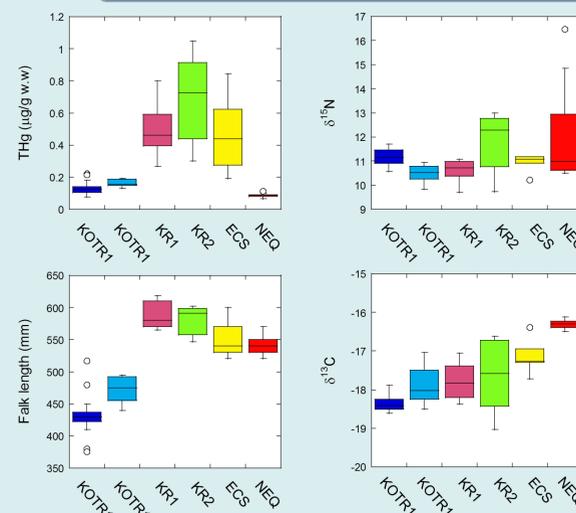


Fig. 1 Regional variation of (a) THg, (b) fork length, (c) $\delta^{15}\text{N}$ and (d) $\delta^{13}\text{C}$.

Body size dependence

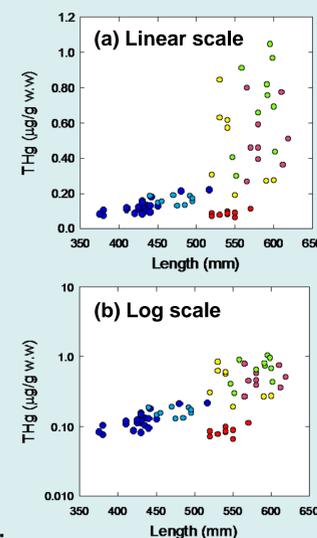


Fig. 2 Relationship between THg and fork length. (a) is linear and (b) is log scales.

$$\text{THg}(\mu\text{g/g-wet}) = 0.0251 \times [\text{body length (cm)}] - 0.971$$

$$R = 0.6568, p < 0.001$$

If the bioenergetics factors are the essence of the body size dependence, proper normalization is needed to consider the factors affecting THg other than size. However, the separate correlation plots for 6 region (Table 1) indicated that only KOTR1 showed significant correlation ($p < 0.001$, $R = 0.7833$), suggesting that body size is, at least in same region, not the primarily factor controlling THg.

Correlation among variables

Table 1. The Pearson's correlation coefficient and statistical significance of stepwise linear-regression analyses. The bold faces indicate p value < 0.05. THg_{corr} represents normalized THg by body length.

	Body length vs.						$\delta^{15}\text{N}$ vs.		$\delta^{13}\text{C}$ vs.		$\delta^{202}\text{Hg}$ vs.		$\Delta^{199}\text{Hg}$ vs.	
	THg	$\delta^{15}\text{N}$	$\delta^{13}\text{C}$	$\delta^{202}\text{Hg}$	$\Delta^{199}\text{Hg}$	THg	$\delta^{15}\text{N}$	THg	$\delta^{13}\text{C}$	THg	$\delta^{15}\text{N}$	THg	$\delta^{13}\text{C}$	
Overall	n	68	58	58	68	58	58	58	58	68	68	68	68	
	R	0.6568	0.1508	0.4821	0.3796	-0.9053	0.1190	0.0354	0.0680	-0.2227	0.3204	0.0415	-0.7514	
	p	0.0076	0.6314	0.0873	0.1990	<0.0001	0.7060	0.9113	0.8301	0.4731	0.2887	0.8959	0.0006	
KOTR1	n	23	21	21	23	23	21	21	23	23	23	23	23	
	R	0.7833	0.1194	-0.5408	0.3592	-0.5899	0.0978	0.0299	-0.2180	0.4154	0.1464	-0.3031	-0.5898	
	p	0.0007	0.7078	0.0563	0.2371	0.0311	0.7593	0.9256	0.4876	0.1634	0.6446	0.3260	0.0312	
KOTR2	n	9	8	8	9	9	8	8	8	9	9	9	9	
	R	-0.0508	-0.1221	0.1419	0.0854	-0.6404	0.6623	0.5943	-0.7305	-0.6760	0.6076	0.2588	-0.1164	
	p	0.8768	0.7106	0.6663	0.7943	0.0336	0.0314	0.0581	0.0148	0.0273	0.0461	0.4248	0.7219	
KRI	n	9	7	7	9	9	7	7	7	9	9	9	9	
	R	0.0926	0.2563	0.4257	0.0563	-0.2170	0.5449	0.3943	0.8855	0.6913	-0.5226	-0.4072	-0.6842	
	p	0.7772	0.4401	0.1970	0.8635	0.5048	0.0950	0.2329	0.0018	0.0292	0.0938	0.2014	0.0209	
KR2	n	10	9	9	10	10	9	9	9	10	10	10	10	
	R	0.4650	0.2525	-0.2929	0.6059	-0.4022	0.5702	0.5397	-0.3587	0.2147	0.8164	0.6700	-0.6581	
	p	0.1353	0.4364	0.3649	0.0426	0.2022	0.0642	0.0822	0.2637	0.5093	0.0021	0.0214	0.0245	
ECS	n	8	6	6	8	8	6	6	6	8	8	8	8	
	R	-0.5428	0.0316	0.7188	-0.5708	0.0526	-0.0143	-0.0531	-0.4340	-0.4988	0.3717	0.4228	-0.5949	
	p	0.0870	0.9252	0.0308	0.0703	0.8732	0.9661	0.8747	0.2023	0.1429	0.2525	0.1906	0.0578	
NEQ	n	9	7	7	9	7	7	7	7	9	9	9	9	
	R	0.5614	0.5745	0.8205	-0.6313	-0.0034	0.5959	N/A	0.1903	N/A	-0.7605	N/A	0.2316	
	p	0.0691	0.0772	0.0062	0.0368	0.9918	0.0659	N/A	0.5667	N/A	0.0076	N/A	0.4762	

Mercury stable isotope

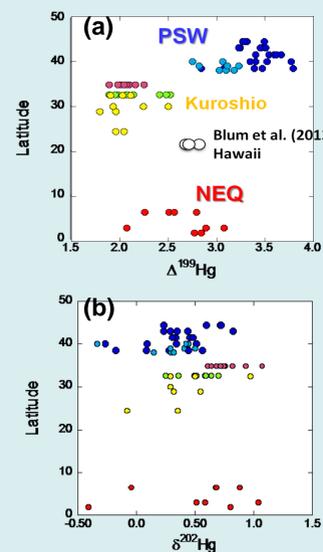


Fig. 3 Latitudinal change of (a) $\Delta^{199}\text{Hg}$ and (b) $\delta^{202}\text{Hg}$.
Fig. 4 Relationship between $\delta^{202}\text{Hg}$ and $\Delta^{199}\text{Hg}$.

The $\delta^{202}\text{Hg}$ and $\Delta^{199}\text{Hg}$ of overall samples were ranged from -0.41 to 1.04‰, 1.90 to 3.77‰, respectively (Fig. 3). These variations are much larger than Blum et al. (2013), which measured same species in Hawaii, showing $\delta^{202}\text{Hg}$ and $\Delta^{199}\text{Hg}$: $0.63 \pm 0.08\text{‰}$ and $2.71 \pm 0.17\text{‰}$, respectively (2SD, n = 3). This indicates isotope signal is variable depending on the region even among the same species.

Interpretation of MIF signature I

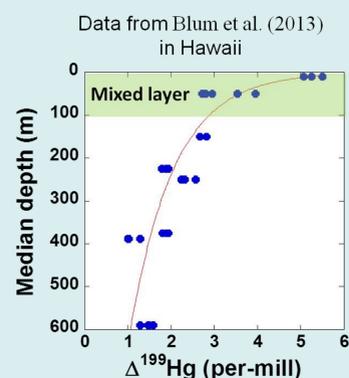
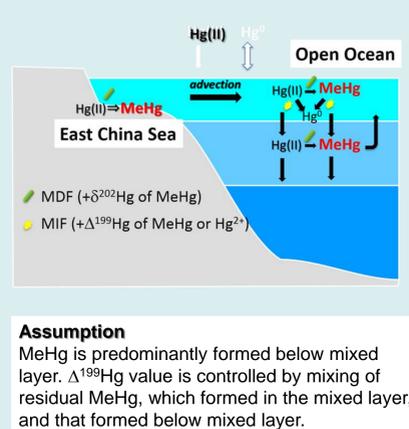


Fig. 5 Proposed model for MIF vs. migration depth trend by Blum et al. (2013)



Interpretation of MIF signature II

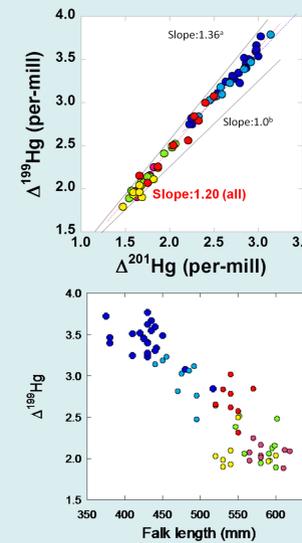


Fig. 6 Relationship between $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$.

- a Bergquist and Blum (2007) MMHg photo-degradation experiment.
- b Bergquist and Blum (2007) Hg^{II} photo-reduction experiment.

Overall slope being 1.20 is same to the other data of marine biota (e.g., Blum et al. 2013).

Fig. 7 Relationship between $\Delta^{199}\text{Hg}$ and fork length

Clear negative correlation
=> Smaller individuals (PSW) forage diet from shallower water, while bigger individuals (Kuroshio, NEQ) are from deeper water.
Increasing size increase thermal inertia and heat production via metabolism. (Barkley et al., 1978)

MIF signature vs THg

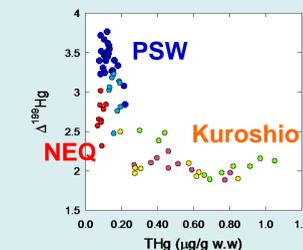


Fig. 8 Relationship between $\Delta^{199}\text{Hg}$ and THg.

PSW
High $\Delta^{199}\text{Hg}$ (shallow migration)
Low THg

Kuroshio
Low $\Delta^{199}\text{Hg}$ (deep migration)
High THg

NEQ
Intermediate $\Delta^{199}\text{Hg}$
Low THg (low MeHg background)

References

Barkley, R.A. et al. (1978) *Fishery Bull.* 76, 653-662.; Bergquist, B.A. & Blum, J.D. (2007) *Science* 318, 417-420.; Blum, J.D. et al., (2013) *Nature Geosci.* 6, 79-884.; Ferriss, B.E. & Essington, T.E. (2014) *Ecol. Model.* 278, 18-28.; Hisamichi, Y. et al. (2010) *Environ. Sci. Tech.* 44, 5971-5978.; Munson, A. et al. (2015) *Global Biogeochem. Cycles* 29, 656-676.; Sunderland, E. (2007) *Environ. Health Persp.* 115, 235.; Trudel, M. & Rasmussen, J.B. (2006) *Can. J. Fish. Aquat. Sci.* 63, 1890-1902.

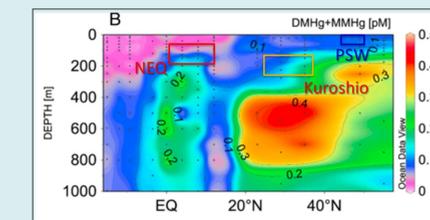


Fig. 9 Transect of Hg species along the CUVAR P16N section (150-175W) by Munson et al. (2015). Depth range was tentatively estimated by $\Delta^{199}\text{Hg}$ value and depth- $\Delta^{199}\text{Hg}$ by Blum et al. (2013).