Bioaccumulation of mercury in fishes of Jagadishpur Reservoir, Nepal

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Abstract
Stable isotope ratios of nitrogen (δ¹⁵N, ‰) and carbon (δ¹³C, ‰), accompanied by stomach contents were used to assess the food chain and trophic transfer of mercury in fifty-three marketable sized fish belonging to five species (Channa punctatus, Mystus vittatus, Puntius sophore and Xenentodon cancila) from the Jagadishpur Reservoir, Nepal. The highest total Hg concentration was found in X. cancila with an average of 800.42(±279.36) µg/kg exceeding the international marketing limit (500 mg/kg), a carnivorous species. However, for some individuals of N. nandus, total Hg concentrations in other fish species in the present study were significantly lower than that limit. The fish community had at least two trophic levels (Δ¹⁵N > 5.6), C. punctatus with the highest and M. vittatus the lowest signatures of δ¹⁵N, which was also supported by the stomach content analysis. There was neither correlation between total Hg and δ¹⁵N nor connectivity in food resource utilization (based on δ¹³C), indicating no biomagnification among these fish species. In addition, Hg concentrations were not significantly correlated to total fish length in any of the species. Fish species in the present study have low Hg content accompanied by low biomagnification through the studied fish community.

Keywords: Fish diet, Stable isotopes of carbon, Stable isotopes of nitrogen, Total mercury, Trophic position

Introduction
Mercury is pervasive and causes toxic threats to society, mainly through the consumption of aquatic food (e.g., fish). Its concentrations have increased in nature, primarily through human activities (Swain et al., 2007). Methylmercury (MeHg) is the main form of Hg that has neurotoxic effects on humans and is easily bioaccumulated through the food web (Freije & Awadh, 2009; Watras et al., 1998), and top predator fish have maximum values in freshwaters (Lockhart et al., 2005). Therefore, in many fish species, total Hg concentrations are positively correlated to body size and age of fish (Donald et al., 2015; Taylor et al., 2014). Although global Hg biomagnification remains unexplained entirely (Lavoie et al., 2013), many other factors are controlling the Hg dynamics in freshwaters bodies; for example, reduced pH enhances Hg uptake by fish (Watras et al., 1998), dissolved organic carbon has a positive relationship with Hg in fish (Belger & Forberg, 2006; Lavoie et al., 2013; Watras et al., 1998), size of the lake has negative relations to Hg in fish (Bodaly et al., 1993), extensive logging activities in the watershed increase Hg concentrations in aquatic biota (Garcia & Carignan, 2000), increment in total phosphorus decreases Hg magnification (Lavoie et al., 2013).

Stable isotope analysis is used to examine the trophic transfer of Hg through the food webs of freshwater ecosystems (Lavoie et al., 2013). Nitrogen isotope ratios (δ¹⁵N= ¹⁵N/¹⁴N) are adequate to determine the trophic position of an organism within the food web in an ecosystem (Post, 2002), whereas carbon isotope ratios (δ¹³C = ¹³C/¹²C) are useful to find out the origin of different food sources (Campbell et al., 2006). However, the reservoir fisheries with human inputs (e.g., feed) may show abnormally deviated δ¹⁵N enrichment than natural water bodies (Wang et al., 2019); thus, trophic levels should cautiously be interpreted.

In Nepal, some studies related to Hg accumulation in fish fillets in lakes (Sharma et al., 2013; Thapa et al., 2014) and rivers (Pandey et al., 2017) had been conducted. These stud-
ies focused mainly on Hg concentrations in fish fillets, and their potential health implications through fish consumption. However, similar studies are lacking from the lowlands of Nepal, particularly from reservoirs. Therefore, the study was carried out to provide baseline data for Hg contamination in fish from Nepal's lowland region using five major fish species, supposedly representing different trophic levels.

Materials and Methods

Study area and sampling
The Jagadishpur Reservoir, a Ramsar Site, is considered a popular tourism hub in Nepal. It is situated at a latitude of N 27° 37.174' and a longitude of E 083° 05.833' at an elevation of 197 m. above sea level (Fig. 1) with a surface area of 225 ha.

Forty-three fish species were recorded by IUCN (2015), of which the five primary fish species Garal (*Channa punctatus*), Tengerkanti (*Mystus vittatus*), Dhoke (*Nandus nandus*), Sidre (*Puntius sophore*), and Kauwa (*Xenentodon cancila*) were studied at present. Fifty-three fish were collected during April 2012 using locally available fishing gears (by local fishers) and identified on-site before they were transported to the laboratory and stored in a frozen condition for further analyses. The total length was measured in centimeters (cm) before dissection. Fish muscle sampling was performed following the methods described in Sharma et al. (2008, 2009). Briefly, the axial muscle was collected from each fish after skin removal from the dorsolateral side, between the dorsal fin and lateral line. The muscle samples were kept in the freezer until further analyses in the laboratories.

Analytical procedures
The muscle was used to analyze total mercury (THg), δ¹⁵N, and δ¹³C following the method described by Sharma et al. (2013). Trophic positions of the fish species were determined based on δ¹⁵N and δ¹³C values, as Post (2002) and Campbell et al. (2006) explained. The δ¹⁵N indicates the relative trophic positions of the organisms whereas δ¹³C gives information on the carbon source of the diet (Sharma et al., 2008).

The methods for Hg analysis can be found in Pandey et al. (2017) as the muscle samples from this study were analyzed.
together in the same batch. In brief, THg were analyzed by cold vapor atomic spectrometry using a direct mercury analyzer (Hydra II, Teledyne Leeman Labs, Hudson, NH, USA) after calcination in an O2 stream and amalgamation on an Au trap. The detection limits for the analyses were 0.01 mg/kg (3 times of SD for ten measurements of blanks). The standard materials used for the analytical accuracy and precision measurements were GSS-9 (n=4) and Tort-2 (n=3). Precisions were within 5.7 % and 1.7 %, and the recovery was within 91-107 % and 104-107 % for GSS-9 and Tort-2, respectively. Each sample was measured twice, and if the relative standard deviation (RSD) were >10%, the sample was measured again. The RSD for all samples was < 6 %.

The analytical procedures for the stable isotopes of nitrogen and carbon were followed after Pandey et al. (2017). In brief, an aliquot of each sample was freeze-dried and was then homogenized using a ceramic grinder. The sample was measured for stable isotopes of carbon and nitrogen (δ13C and δ15N) with an isotope ratio mass spectrometer (Finnigan MAT Delta V Advantage, Thermo Fisher Scientific, Waltham, MA, USA). Isotopic data are presented in units of per mil (%). Stable isotopes (δ13C and δ15N) were determined as

\[ \delta^{13}C_{\text{Sample}} = \left( \frac{R_{\text{sample}}}{R_{\text{standard}}} - 1 \right) \times 1000 \]

where \( R \) is the ratio of heavy to light isotope (13N/14N or 13C/12C) in the sample, standards are atmospheric air for N and Pee Dee Belemnite for C. The overall analytical precision was ± 0.2 ‰, including sample preparation and analysis.

Stomach contents were preserved in 70 % alcohol and analyzed later under a dissecting microscope. Mean volume percentages of prey in the stomach contents were measured in all fish species, categorized into six items: worms, detritus, aquatic plants, fish insects, and unidentified following Pandey et al. (2017). 14 out of 53 analyzed fish had empty stomachs.

**Statistical analysis**

The comparison of THg in fish muscle was performed by a One-Sample t-test. Correlation and regression analyses were carried out to test relationships among THg concentrations, δ15N, δ13C, and total length. The relationship between THg concentrations and exposure time was explained by the regression of log-THg against the total length of fish. The biomagnification rate of Hg among the fish community was determined with the regression of log-THg against δ15N. Trophic positions of the fish species in the food chain were identified with a simple food web structure based on the relationship between nitrogen isotope (δ15N) and carbon isotope (δ13C) of the fish species. All the relationships were considered statistically significant at \( p \leq 0.05 \).

**Results and Discussion**

The simple food web structure based on δ15N and δ13C showed that Channa punctatus occupied the highest trophic position, and M. vittatus the lowest (Fig. 2). However, our study lacks baseline isotopic data of the prey species; therefore, the figure gives an idea of the relative trophic positions of fish only. Nevertheless, this indicates a food chain of at least two trophic levels (Δδ15N > 5.6) in the fish community of Jagadishpur Reservoir, since δ15N increases approximately by three ‰ for each trophic level (Vander Zanden & Rasmussen, 2001). All the individuals from the lowest trophic position (as indicated by the stomach contents and the δ15N signatures), M. vittatus had aquatic plants as their dominant diet. The stomach contents of C. punctatus, the species at the higher trophic position based on δ15N signatures, were dominated by animal foods (e.g., fish and insects; Fig. 3). Similar results were reported previously from the commercial fish species in Lake Phewa, Pokhara, Nepal (Sharma et al., 2013).

THg concentrations in fish samples were within the range of 65.13 to 1263.54 mg/kg (w/w) in the present study (Table 1). Channa punctatus had the lowest concentration, whereas carnivorous species X. cancila had the highest concentrations of THg. The higher THg concentrations in X. cancila is probably a result of bioaccumulation of mercury through diet as its food was dominated by fish, worms, and insects (Fig. 3) which was similar to the findings observed by Deribe et al. (2014) on predator fishes in Ethiopian Rift Valley Lakes.

Nandus nandus had the second-highest THg concentrations possibly due to fish as one of the main components in its diet, which was also reported as having mid-range THg concentration (90 mg/kg) among other fish species in the polluted rivers in Bangladesh (Shoeb et al., 2017). In the remaining species, THg concentrations were relatively low, which might be because of the consumption of food from lower trophic levels as also mentioned by Campbell et al. (2003) and Desta et al. (2007) for some fishes in Lake Victoria and Lake Awassa, respectively.
Mean volume (%) of diet composition (stomach contents) of C. punctatus, M. vittatus, N. nandus, P. sophore, and X. cancila.

However, the lowest value of Hg in C. punctatus in the present study despite having some animal diet in the stomachs (Fig. 3) and highest $\delta^{15}$N values (indicating higher trophic position, Fig. 2) does not follow the general patterns of explanation and needs further investigation. Sometimes, fish species from different trophic levels may have similar mercury concentrations due to unknown reasons. Furthermore, low Hg concentrations in predator fishes of reservoirs could be due to the short food webs and limited magnification factor compared to natural water bodies (Xu et al., 2018). Although overall average concentrations of total Hg in fishes from the present study had significantly lower values ($t = -3.87, p < 0.001$) compared to the international marketing limit (500 mg/kg; WHO, 2008), X. cancila had significantly exceeded this limit ($t = 2.85, p = 0.015$; 86% individuals exceeded). Furthermore, 45% of individuals exceeded the marketing limits, although no statistically higher Hg concentrations were observed in N. nandus. Compared to the present study, lower Hg in fishes was reported by Wang et al. (2019) in an artificial Changshou Lake and Xu et al. (2018) in Three Gorges Reservoirs in China.

In the present study, mercury concentration was not significantly correlated to total fish length in any species (Table 2). This could be due to the disproportionation between fish growth with Hg bioaccumulation in eutrophic water bodies, as explained by Wang et al. (2019). Furthermore, a possible shift in the diet of old individuals could be a cause for the weak relationship as discovered in a fish species in the Nepalese rivers (Pandey et al., 2017) and some African lakes (Tadiso et al., 2011). This phenomenon is also explained by Desta et al. (2006) as changing the habit of diet by age, where fishes consume zooplankton in the younger stage and aquatic plants when mature. Water management practices, particularly in reservoirs, profoundly influence Hg bioaccumulation to a great extent (Willacker et al., 2016). Moreover, watershed characteristics and in-lake processes (e.g., food web structure) are also considered important phenomena for controlling Hg concentrations in a fish community. However, these explanations need further study in the fish community from this reservoir. Although a significant positive relationship was found between log-THg and $\delta^{15}$N for N. nandus ($p = 0.038, R = 0.47$) and X. cancila ($p = 0.041, R = 0.78$), no significant relationship was found in separate fish species (Table 2) as well as pooled data (Fig. 4).

Table 1 Mean (± SD) and range of length (cm), THg concentrations (mg/kg, ww), $\delta^{13}$C (‰) and $\delta^{15}$N (‰) of five different fish species sampled from Jagadishpur Reservoir, Nepal.

| Fish species | n  | Length (cm) (Mean±SD) (Range) | Total Hg (mg/kg, ww) (Mean±SD) (Range) | $\delta^{15}$N (‰) (Mean±SD) (Range) | $\delta^{13}$C (‰) (Mean±SD) (Range) |
|--------------|----|------------------------------|----------------------------------------|--------------------------------------|--------------------------------------|
| C. punctatus | 6  | 15.03±0.74 (14.20-16.20)     | 144.53±31.29 (108.19-200.95)           | 11.29±1.18 (9.85-12.79)              | -23.83±1.77 (-26.82 to -21.98)       |
| M. vittatus  | 3  | 12.10±0.76 (11.30-12.80)     | 165.91±40.64 (126.37-207.56)           | 9.33±2.16 (7.09-11.40)               | -23.59±3.14 (-26.76 to -20.49)       |
| N. nandus    | 20 | 14.00±1.25 (11.80-15.80)     | 467.75±142.68 (143.17-698.73)          | 9.38±1.03 (7.44-11.14)              | -23.02±2.70 (-27.47 to -16.30)      |
| P. sophore   | 17 | 10.09±1.17 (8.00-11.90)      | 162.46±72.89 (77.06-368.16)            | 10.20±1.44 (8.07-12.61)             | -23.86±1.89 (-25.27 to -19.46)      |
| X. cancila   | 7  | 26.55±2.93 (20.5-29.3)       | 800.42±279.36 (429.04-1263.54)         | 10.71±0.70 (9.83-11.77)             | -23.28±2.57 (-26.33 to -19.43)      |
Table 2 Regression of mercury concentrations (log-THg) against length (cm), and δ¹⁵N were analyzed for individual fish species. Also included in the table are a regression of δ¹⁵N and δ¹³C against the length of different species. The sample size (n), intercept, slope, R² and p-values are given for each regression. Significant results for p < 0.05 at α = 0.05 (95 % confidence interval) are written in bold.

| Species     | Regression       | (n) | Intercept | Slope | S.E. of Slope | R   | p-value |
|-------------|------------------|-----|-----------|-------|---------------|-----|---------|
| All         | log-THg vs. δ¹⁵N | 53  | 2.60      | -0.02 | 0.03          | 0.07| 0.64    |
| C. punctatus | log-THg vs. Length | 6   | 1.89      | 0.02  | 0.06          | 0.15| 0.78    |
|             | log-THg vs. δ¹⁵N | 6   | 2.01      | 0.01  | 0.04          | 0.17| 0.75    |
|             | δ¹⁵N vs. Length   | 6   | 15.14     | -0.26 | 0.78          | 0.16| 0.76    |
|             | δ¹³C vs. Length   | 6   | 5.03      | -1.92 | 0.71          | 0.81| 0.05    |
| M. vittatus  | log-THg vs. Length | 3   | 0.49      | 0.14  | 0.01          | 0.99| 0.06    |
|             | log-THg vs. δ¹⁵N | 3   | 2.67      | -0.05 | 0.01          | 0.99| 0.06    |
|             | δ¹⁵N vs. Length   | 3   | 4.31      | -2.82 | 0.51          | 0.98| 0.16    |
|             | δ¹³C vs. Length   | 3   | -4.25     | 1.57  | 3.85          | 0.38| 0.75    |
| N. nandus    | log-THg vs. Length | 20  | 2.09      | 0.04  | 0.03          | 0.31| 0.19    |
|             | log-THg vs. δ¹⁵N | 20  | 1.97      | 0.07  | 0.03          | 0.47| 0.038   |
|             | δ¹⁵N vs. Length   | 20  | 8.58      | 0.06  | 0.19          | 0.07| 0.77    |
|             | δ¹³C vs. Length   | 20  | -4.47     | 1.55  | 0.36          | 0.72| <0.001  |
| P. sapphire  | log-THg vs. Length | 17  | 2.43      | -0.03 | 0.04          | 0.17| 0.52    |
|             | log-THg vs. δ¹⁵N | 17  | 1.88      | 0.03  | 0.03          | 0.24| 0.35    |
|             | δ¹⁵N vs. Length   | 17  | 9.36      | 0.08  | 0.32          | 0.07| 0.79    |
|             | δ¹³C vs. Length   | 17  | -24.18    | 0.03  | 0.42          | 0.02| 0.94    |
| X. cancila   | log-THg vs. Length | 7   | 3.28      | -0.02 | 0.02          | 0.27| 0.55    |
|             | log-THg vs. δ¹⁵N | 7   | 4.76      | -0.18 | 0.06          | 0.78| 0.041   |
|             | δ¹⁵N vs. Length   | 7   | 6.08      | 0.18  | 0.08          | 0.72| 0.07    |
|             | δ¹³C vs. Length   | 7   | -27.94    | 0.18  | 0.39          | 0.19| 0.67    |

No significant relationship of stable isotopes, δ¹⁵N and δ¹³C, with total fish length was found in the studied fish species except for N. nandus, which had a significantly positive relationship between δ¹³C and total length (p < 0.001, R = 0.72; Table 2). Interestingly, N. nandus was the only species having a wide variation in δ¹³C values with both the lowest and highest values of carbon signatures (Table 1), indicating a correspondingly wide range of habitat use and carbon sources in the diet as was the case of T. zilli observed in Lake Ziway (Tadiso et al., 2011). However, all other species seemed to consume the diet from the same locality and thus the same basic carbon sources, regardless of their size. The overall results indicate that the habitat within the fish community is dominated by pelagic carbon sources (Fig. 2) as can be inferred from light (more negative) carbon signatures (Bootsma et al., 1996; Deribe et al., 2014; Hecky & Hesslein, 1995), a similar condition like the one possibly resulting from the eutrophication (Gautam & Bhattarai, 2008), which is described by Eagles-Smith et al. (2008) for the eutrophic Clear Lake.

The slope of the regression equation of log-THg and δ¹⁵N values (as a measure of trophic magnification slope) can be used for the quantitative measurement and comparative studies of biomagnification rates among aquatic systems (Borga et al., 2012; Lavoie et al., 2013). However, there was no relationship between log-THg and δ¹⁵N, indicating that the biomagnification of mercury along the food web of Jagdishpur Reservoir was not clear (Table 2). Another study in the Nepalese lake (Lake Phewa) by Sharma et al. (2013) also observed low biomagnification of THg through the food chain. Furthermore, a similar finding was reported in a tropical lake with a lower rate of biomagnification of Hg, as indicated by the slope of the regression equation (Desta et al., 2006).

Figure 4 The regression between log-THg (mg/kg, ww) and δ¹⁵N (%) for all fish species from Jagdishpur Reservoir. Fish codes used in the figure are C. punctatus (CP), M. vittatus (MV), N. nandus (NN), P. sapphire (PS), and X.
cancila (NC). Lastly, the absence of representative fish species from the food chain in the studied lake might have some impacts hiding the relationship between THg and δ15N as described by Sharma et al. (2013).

Conclusion
The carnivorous fish had higher concentrations of THg compared to omnivorous and herbivorous fishes in Jagadishpur Reservoir; however, none of the fishes statistically exceeded WHO marketing guidelines. Mercury concentrations in the fish community of the Jagadishpur Reservoir did not show significant relationships with the length of fish and their trophic positions, except_N. nandus. The fish community in the present study represented at least two trophic levels (Δ15N > 5.6); C. punctatus at the highest and M. vittatus the lowest based on δ15N signatures, which was also supported by the stomach content analysis. There is a clear indication that the fish community in our samples revealed a low biomagnification rate of mercury, possibly due to no representation from all the trophic levels.

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