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Mercury Isotope Fractionation by Internal Demethylation and Biomineralization Reactions in Seabirds: Implications for Environmental Mercury Science

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ABSTRACT: A prerequisite for environmental and toxicological applications of mercury (Hg) stable isotopes in wildlife and humans is quantifying the isotopic fractionation of biological reactions. Here, we measured stable Hg isotope values of relevant tissues of giant petrels (*Macronectes* spp.). Isotopic data were interpreted with published HR-XANES spectroscopic data that document a step-wise transformation of methylmercury (MeHg) to Hg-tetraselenolate (Hg(Sec)₄) and mercury selenide (HgSe) (Sec = selenocysteine). By mathematical inversion of isotopic and spectroscopic data, identical $\delta^{202}\text{Hg}$ values for MeHg (2.69 ± 0.04 ‰), Hg(Sec)₄ (-1.37 ± 0.06 ‰), and HgSe (0.18 ± 0.02 ‰) were determined in 23 tissues of eight birds from the Kerguelen Islands and Adélie Land (Antarctica). Isotopic differences in $\delta^{202}\text{Hg}$ between MeHg and Hg(Sec)₄ (-4.1 ± 0.1 ‰) reflect mass-dependent fractionation from a kinetic isotope effect due to the MeHg \rightarrow Hg(Sec)₄ demethylation reaction. Surprisingly, Hg(Sec)₄ and HgSe differed isotopically in $\delta^{202}\text{Hg}$ ($+1.6 \pm 0.1$ ‰) and mass-independent anomalies (i.e., changes in $\Delta^{199}\text{Hg}$ of ≤ 0.3 ‰), consistent with equilibrium isotope effects of mass-dependent and nuclear volume fractionation from Hg(Sec)₄ \rightarrow HgSe biomineralization. The invariance of species-specific $\delta^{202}\text{Hg}$ values across tissues and individual birds reflects the kinetic lability of Hg-ligand bonds and tissue-specific redistribution of MeHg and inorganic Hg, likely as Hg(Sec)₄. These observations provide fundamental information necessary to improve the interpretation of stable Hg isotope data and provoke a revisitation of processes governing isotopic fractionation in biota and toxicological risk assessment in wildlife.

INTRODUCTION

The efficient biomagnification of methylmercury (MeHg) in aquatic food webs results in elevated dietary exposure of mercury (Hg) to top predators such as seabirds and marine mammals.¹ After the

assimilation of dietary MeHg in the digestive tract, MeHg is circulated in the bloodstream to various tissues and organs (e.g., brain) and forms MeHg-cysteine (MeHg-Cys) complexes with proteins.² At the organism level the toxicological effects of MeHg are controlled, in part, by two central processes: the demethylation of MeHg within cells and the internal exchange and depuration of both MeHg and inorganic Hg degradation products. The biomineralization of MeHg to HgSe granules is accepted as an important internal transformation of Hg in vertebrates,^{3,4} but until recently little was known on the biochemistry of the reaction. Development and application of new analytical approaches, most notably high energy-resolution X-ray absorption near-edge structure (HR-XANES) spectroscopy, identified that MeHg is first demethylated to a Hg-tetraselenolate complex (Hg(Sec)₄; Sec, selenocysteine) associated with selenoprotein P (SelP).⁵ Hg(Sec)₄ bound to SelP, the latter containing at least ten Sec residues in vertebrates,⁶ is hypothesized to be the template for biomineralization of HgSe.⁵ This hypothesis is strengthened by the observed co-existence of Hg(Sec)₄ and HgSe in diverse tissues of seabirds (giant petrels, *Macronectes* spp.),⁷ mammals (long-finned pilot whales, *Globicephala melas*),⁸ and fish (blue marlin, *Makaira* sp.).⁹ The step-wise MeHg → Hg(Sec)₄ → HgSe demethylation and biomineralization reactions appear to be universal in vertebrates.^{4,5,7-10} Notably, proportions of the three dominant chemical forms of Hg (MeHg, Hg(Sec)₄, HgSe) vary considerably between tissues and organisms, the latter reflecting variable physiological efficiencies for MeHg detoxification for reasons yet known. These observations have important toxicological implications for the risk from MeHg to organisms.

Variations in the ratio of naturally occurring stable Hg isotopes provide complementary information on the internal transformations and exchange of Hg within organisms. Isotopic fractionation of Hg due to internal transformations has focused on the kinetic mass-dependent fractionation (MDF) of Hg for the MeHg → Hg(Sec)₄ demethylation reaction. Waterbirds from diverse aquatic systems exhibited a constant fractionation of $\delta^{202}\text{Hg}$ of -2.2 ± 0.1 ‰ for the MeHg → Hg(Sec)₄ demethylation reaction, and a uniform MeHg isotopic composition between tissues, suggesting a rapid internal exchange of MeHg between tissues of the birds.¹¹ MDF due to MeHg demethylation accounts for the decrease in $\delta^{202}\text{Hg}$ of tissues with decreasing fraction of MeHg ($f(\text{MeHg})$), as observed in birds,^{11,12} fish¹³ and mammals,¹⁴⁻¹⁶ and parallel increase in $\delta^{202}\text{Hg}$ of the residual MeHg in keratinized tissues.^{14,17,18}

However, the $\delta^{202}\text{Hg}$ values of tissues of higher trophic level organisms often cannot be explained by the simple mixing of two isotope endmembers (e.g., $\delta^{202}\text{MeHg}$ and $\delta^{202}\text{Hg}(\text{Sec})_4$), as evidenced by significant variability in $\delta^{202}\text{Hg}$ values in tissues with the same $f(\text{MeHg})$ ¹²⁻¹⁴ and the increase in $\delta^{202}\text{Hg}$ values by >1.0 ‰ in the liver of marine mammals as $f(\text{MeHg})$ approaches zero.^{15,16,19} The $\delta^{202}\text{Hg}$ of a tissue ($\delta^{202}\text{Hg}_t$) is the weighted sum of the $\delta^{202}(\text{Sp}_i)_t$ values of the chemical species

$$\delta^{202}\text{Hg}_t = \sum f(\text{Sp}_i)_t \times \delta^{202}\text{Sp}_i \quad (1)$$

and therefore other chemical forms of Hg (e.g., HgSe) may explain behavior in $\delta^{202}\text{Hg}$ in organisms. It was shown recently that the increase in $\delta^{202}\text{Hg}$ in liver and kidneys of long-finned pilot whales at low $f(\text{MeHg})$ fingerprints the transformation of Hg(Sec)₄ to HgSe.⁸

Aside from demethylation, other processes that fractionate Hg isotopes have not been documented within biota, including mass-independent fractionation (MIF; expressed as $\Delta^{199}\text{Hg}$, $\Delta^{200}\text{Hg}$, $\Delta^{201}\text{Hg}$, $\Delta^{204}\text{Hg}$) from kinetic processes or equilibrium isotope effects, the latter driven by both MDF and nuclear volume effect (NVE).²⁰⁻²² A fundamental understanding of the internal processes controlling the isotopic fractionation of Hg in organisms is essential to fully utilize isotopic values for

environmental applications, such as identifying the dietary origins of MeHg,^{11, 23} climate change effects,²⁴ and response to changes in Hg sources.^{25, 26}

To address these science needs, the isotopic fractionation of Hg between the three prominent chemical forms of Hg (MeHg, Hg(Sec)₄, HgSe) was quantified using diverse tissues of giant petrels. We measured the Hg isotopic composition of 23 tissues from eight giant petrels (P1-P8) analyzed previously for chemical speciation by HR-XANES and transmission electron microscopy (Figure 1).⁷ The giant petrels demethylate MeHg with extreme efficiency, with near-complete demethylation of MeHg in the muscle and varying degrees of HgSe biomineralization across diverse tissues (liver > kidneys > muscle ≈ brain).⁷ $\delta^{202}\text{Hg}$ values of each chemical form of Hg ($\delta^{202}\text{MeHg}$, $\delta^{202}\text{Hg}(\text{Sec})_4$, $\delta^{202}\text{HgSe}$) were determined with high precision by the inversion of isotopic and spectroscopic data using linear algebra. The calculated $\delta^{202}\text{MeHg}$ value was compared with values of $\delta^{202}\text{MeHg}$ directly measured on a subset of tissues using an established method.²⁷ The magnitude of isotopic fractionation quantified for the MeHg → Hg(Sec)₄ reaction in the giant petrels contrasts with that quantified for waterbirds and mammals, indicating that fractionation patterns arise from different physiological efficiencies for demethylation. Further, mechanistic evidence is provided for the isotopic fractionation of Hg for the Hg(Sec)₄ → HgSe biomineralization reaction. We discuss the implications of these findings for the toxicological understanding of Hg in organisms and the environmental application of stable Hg isotopes.

MATERIALS AND METHODS

Study animals. Giant petrels are large scavengers that feed primarily on carrion of marine mammals and other seabirds. They remain all year long within the Southern Ocean and in fringing subtropical waters. Being both top consumers and long-lived animals, giant petrels accumulate large amounts of mercury in their tissues. Eight dead specimens were opportunistically collected at the subantarctic Kerguelen Islands and in Adélie Land (Antarctica) and stored at -20°C until dissection. All were adult males, but their exact age was unknown.

Mercury stable isotope measurement. All analyses for Hg stable isotopes were performed at the USGS Mercury Research Lab (Madison, WI). Tissues were digested in concentrated nitric acid (HNO₃) and heated at 80 °C for 8 h. Sample digests were further oxidized using bromine monochloride (BrCl) to 10% total volume and heated for an additional 2 h to ensure the degradation of any organic Hg species. Samples were diluted to 1-5% H⁺ content before isotope analysis. Concentration recoveries for digestions of certified reference material (DOLT-2, dogfish liver) were 105 ± 3% (*n* = 3).

Species-specific isotope measurements were performed on tissues that contained variable *f*(MeHg) using a previously established distillation and anion-exchange resin method (*n* = 9).²⁷ Briefly, samples were distilled to removed matrix interferences, and distillates were preserved to a final acid concentration of 0.5% hydrochloric acid (HCl). After acidification, samples were passed through a pre-treated AG1-X4 resin column.²⁷ MeHg is neutrally charged under these conditions (0.5% HCl) and therefore does not sorb to the AG1-X4 resin, whereas negatively charged inorganic species were sorbed to the AG1-X4 resin. If required, large volumes of eluent were further pre-concentrated using rapid gold amalgamation and collected into an oxidant trap.²⁸ MeHg isotope samples had acceptable recoveries, within 7% of previously measured MeHg concentrations, indicating no process-induced

isotope fractionation. A certified reference material (DOLT-2) was also analyzed with samples and had acceptable recoveries (96 and 106%; $n=2$) after distillation and resin preconcentration.

Total Hg and MeHg isotope samples were analyzed using a multicollector inductively coupled plasma mass spectrometer (MC-ICP-MS, Thermo Scientific). Analysis was conducted using standard-sample bracketing with National Institute of Standards and Technology (NIST) 3133. Standards, samples, and reference materials were introduced to the MC-ICP-MS using stannous chloride reduction in a custom gas-liquid separator²⁹ at a flow rate of 0.85 mL min⁻¹. A thallium standard (Tl, 40.9 g/L) was simultaneously introduced to the gas-liquid separator for mass bias correction using a desolvating nebulizer (Apex, ESI).²⁸ The MC-ICP-MS was tuned for optimal voltage and signal intensity for ²⁰²Hg and ²⁰⁵Tl prior to analysis.

Isotope values are expressed as $\delta^{XXX}\text{Hg}$ and $\Delta^{XXX}\text{Hg}$ in reference to NIST 3133. Isotope values of total Hg of tissues and the MeHg pool are denoted as $\delta^{202}\text{Hg}_f$ and $\delta^{202}\text{MeHg}_f$ for feathers, $\delta^{202}\text{Hg}_m$ and $\delta^{202}\text{MeHg}_m$ for muscle, $\delta^{202}\text{Hg}_k$ and $\delta^{202}\text{MeHg}_k$ for kidneys, $\delta^{202}\text{Hg}_l$ and $\delta^{202}\text{MeHg}_l$ for liver, and $\delta^{202}\text{Hg}_b$ and $\delta^{202}\text{MeHg}_b$ for brain, respectively. A secondary standard (NIST RM 8610, UM Almadén) was run every five samples to ensure accuracy and precision of measurements and isotope values agreed with certified values ($\delta^{202}\text{Hg} = -0.55 \pm 0.06 \text{ ‰}$, $\Delta^{199}\text{Hg} = -0.01 \pm 0.04 \text{ ‰}$, $\Delta^{200}\text{Hg} = 0.01 \pm 0.04 \text{ ‰}$, $\Delta^{201}\text{Hg} = -0.03 \pm 0.03 \text{ ‰}$, $\Delta^{204}\text{Hg} = -0.01 \pm 0.06 \text{ ‰}$, 2SD; $n=14$). Certified reference material (DOLT-2) was comparable for total Hg and species-specific isotope measurements to literature values (Table S4).²⁷ Furthermore, a comparison of total Hg and species-specific isotope measurements of three of the petrel feathers (P2, P4, and P5), which contained exclusively MeHg, shows excellent agreement (Figure S7), with differences in values between total Hg and MeHg $\leq 0.16 \text{ ‰}$, $\leq 0.03 \text{ ‰}$, $\leq 0.07 \text{ ‰}$, $\leq 0.03 \text{ ‰}$, and $\leq 0.05 \text{ ‰}$ for $\delta^{202}\text{Hg}$, $\Delta^{199}\text{Hg}$, $\Delta^{201}\text{Hg}$, $\Delta^{200}\text{Hg}$, and $\Delta^{204}\text{Hg}$, respectively. The aforementioned differences were an assessment of the precision of total Hg and species-specific isotope measurements. Hg stable isotope data generated during this study are available in the U.S. Geological Survey data repository.³⁰

Data analysis. Species-specific $\delta^{202}\text{Sp}_i$ values were calculated using the proportion of each species ($f(\text{Sp}_i)_t$) quantified by HR-XANES spectroscopy⁷ and the weighted average $\delta^{202}\text{Hg}_t$ values as an inverse problem, which can be solved using linear algebra.³¹ Equation 1 can be represented as follows:

$$\underbrace{\begin{bmatrix} f(\text{MeHg})_1 & f(\text{Hg}(\text{Sec})_4)_1 & \dots & f(\text{HgSe})_1 \\ f(\text{MeHg})_2 & f(\text{Hg}(\text{Sec})_4)_2 & \dots & f(\text{HgSe})_2 \\ \dots & \dots & \dots & \dots \\ f(\text{MeHg})_n & f(\text{Hg}(\text{Sec})_4)_n & \dots & f(\text{HgSe})_n \end{bmatrix}}_{\mathbf{A}} \underbrace{\begin{pmatrix} \delta^{202}\text{Sp}_1 \\ \delta^{202}\text{Sp}_2 \\ \dots \\ \delta^{202}\text{Sp}_m \end{pmatrix}}_{\mathbf{x}} = \underbrace{\begin{pmatrix} \delta^{202}\text{Hg}_1 \\ \delta^{202}\text{Hg}_2 \\ \dots \\ \delta^{202}\text{Hg}_n \end{pmatrix}}_{\mathbf{b}} \quad (2)$$

where \mathbf{A} is a $n \times m$ matrix of the spectroscopic $f(\text{Sp}_i)_t$ values, n is the number of tissue samples (22-23), and m is the number of Hg species (3). \mathbf{x} and \mathbf{b} are two column vectors associated with the unknown $\delta^{202}\text{Sp}_i$ values and the measured $\delta^{202}\text{Hg}_t$ values, respectively. \mathbf{x} can be obtained by minimizing the functional $\varphi(\mathbf{x})$, or residual, that quantifies the difference between the predicted (\mathbf{Ax}) from the observed (\mathbf{b}) data:

$$\varphi(\mathbf{x}) = \|\mathbf{Ax} - \mathbf{b}\|^2 = (\mathbf{Ax} - \mathbf{b})^T \mathbf{C}_d^{-1} (\mathbf{Ax} - \mathbf{b}) \quad (3)$$

where $(\mathbf{Ax}-\mathbf{b})^T$ denotes the transpose of the vector $(\mathbf{Ax}-\mathbf{b})$, and \mathbf{C}_d^{-1} is the inverse matrix of the squared standard deviations of the isotope measurements. Here, the total number of unknowns is 3 ($\delta^{202}\text{MeHg}$, $\delta^{202}\text{Hg}(\text{Sec})_4$, $\delta^{202}\text{HgSe}$) for 22-23 equations (i.e., tissues), therefore the inverse problem is overdetermined and the objective functional has a unique minimum. In our previous study on long-finned pilot whales⁸ the linear system had 92 unknowns for 89 equations. Therefore, the inverse problem was underdetermined and its resolution required the addition of a regularization term to Eq. (3).

As isotope measurements are independent, the input covariance \mathbf{C}_d matrix is diagonal and contains the experimental uncertainties on $\delta^{202}\text{Hg}_t$. The unknown vector \mathbf{x} is obtained at the minimum of φ , which is equivalent to zero in its first derivative:

$$\partial_{\mathbf{x}}\varphi(\mathbf{x}) = \mathbf{A}^T\mathbf{C}_d^{-1}\mathbf{Ax} - \mathbf{A}^T\mathbf{C}_d^{-1}\mathbf{b} = 0 \quad (4)$$

giving

$$\underbrace{\mathbf{A}^T\mathbf{C}_d^{-1}\mathbf{A}}_{\mathbf{B}}\mathbf{x} = \underbrace{\mathbf{A}^T\mathbf{C}_d^{-1}\mathbf{b}}_{\mathbf{y}} \quad (5)$$

where \mathbf{B} is a full ranked matrix and \mathbf{y} a vector. Eq. 5 was solved with classical linear system resolution methods. In addition to providing \mathbf{x} , classical inverse problem theory provides an uncertainty assessment of the solution through the output covariance matrix \mathbf{C}_m :

$$\mathbf{C}_m = (\mathbf{A}^T\mathbf{C}_d^{-1}\mathbf{A})^{-1} \quad (6)$$

Its diagonal elements are the variances of the \mathbf{x} values, which is the square of the standard deviations of the species-specific $\delta^{202}\text{Sp}_i$ values, as evaluated from the uncertainties of the isotopic measurements ($\delta^{202}\text{Hg}_t$).

The linear inverse problem theory allows obtaining, in the least-squares sense, the optimal solution ($\mathbf{x}_i = \delta^{202}\text{Sp}_i$) to the set of equations (\mathbf{Ax}) and its standard deviation. However, data are affected also by uncertainties on the species proportions ($f(\text{Sp}_i)_t$) obtained from the HR-XANES fits. The classical inverse problem treatment does not include uncertainties on the elements from the \mathbf{A} matrix that contains the $f(\text{Sp}_i)_t$ values. This uncertainty was assessed by introducing a statistical treatment to the numerical solutions obtained from the inverse analysis. Uncertainties on the finite $f(\text{Sp}_i)_t$ values can be added to the \mathbf{A} matrix by constructing the $\tilde{\mathbf{A}}$ matrix:

$$\underbrace{(\mathbf{A} + \mathbf{C})}_{\tilde{\mathbf{A}}}\mathbf{x} = \mathbf{b} \quad (7)$$

where \mathbf{C} contains the variability on the \mathbf{A} elements (i.e, precision on $f(\text{Sp}_i)_t$). For example, the element (i,j) of the matrix $\tilde{\mathbf{A}}$ is $\tilde{\mathbf{A}}(i, j) = \mathbf{A}(i, j) \pm \mathbf{C}(i, j)$. An ensemble of matrices $\tilde{\mathbf{A}}_e$ can be constructed from a uniform random sampling of all $\tilde{\mathbf{A}}$ matrices spanned by the $\mathbf{C}(i,j)$ values. Each $\tilde{\mathbf{A}}_e$ matrix is treated by the above-described inverse algorithm, yielding the new solution \mathbf{x}_e . Considering a sufficiently large ensemble of N $\tilde{\mathbf{A}}_e$ matrices, a mean solution $\bar{\mathbf{x}}$ can be computed from all solutions $\mathbf{X} = (\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_n)$.

The covariance matrix of $\bar{\mathbf{x}}$ can be computed as $\tilde{\mathbf{C}}_m = \frac{1}{N-1}\mathbf{M}\mathbf{M}^T$, where the matrix \mathbf{M} is built from the perturbation of the mean solution $\mathbf{M} = (\mathbf{x}_1 - \bar{\mathbf{x}}, \mathbf{x}_2 - \bar{\mathbf{x}}, \dots, \mathbf{x}_N - \bar{\mathbf{x}})$. The diagonal elements of

\tilde{C}_m represent the variances and their square roots represent the standard deviations of $\bar{\mathbf{X}}$, which now include both the isotopic and spectroscopic uncertainties. $\bar{\mathbf{X}}$ and \tilde{C}_m from the giant petrel data were converged after a few hundreds of random $\mathbf{C}(i,j)$ draws. Values reported herein correspond to $N = 1000$. The Python code of the inversion algorithm is available in the Supporting Information and the GRICAD GitLab's site.³²

RESULTS

The tissues and feathers of giant petrels exhibit large variability in $\delta^{202}\text{Hg}_t$ values that follow a hierarchy across all individuals based on tissue type: feathers ($\delta^{202}\text{Hg}_f = 2.74 \pm 0.16 \text{ ‰}$, $n = 5$, mean \pm standard deviation) > brain ($\delta^{202}\text{Hg}_b = 1.09 \pm 0.77 \text{ ‰}$, $n = 3$) > kidneys ($\delta^{202}\text{Hg}_k = 0.01 \pm 0.25 \text{ ‰}$, $n = 5$) = liver ($\delta^{202}\text{Hg}_l = 0.04 \pm 0.11 \text{ ‰}$, $n = 5$) > muscle ($\delta^{202}\text{Hg}_m = -0.80 \pm 0.23 \text{ ‰}$, $n = 5$) (Figures 2 and 3, Table S1). $\delta^{202}\text{Hg}_t$ for liver tissues are within the range of literature data for seabirds ($-0.64 \text{ ‰} < \delta^{202}\text{Hg}_l < 0.68 \text{ ‰}$),¹² whereas $\delta^{202}\text{Hg}$ for feathers are modestly higher than those measured previously for seabirds ($1.10 \text{ ‰} < \delta^{202}\text{Hg}_f < 2.35 \text{ ‰}$).^{12, 33} Notably, muscle tissues exhibit considerably lower $\delta^{202}\text{Hg}_t$ values than liver tissues (on average 0.84 ‰), counter to observations to other birds and mammals,^{11, 13-16, 18} and were as much as approximately -3.5 ‰ lower compared to feathers (Figure 2, Table S1). To the best of our knowledge, differences in $\delta^{202}\text{Hg}$ of this magnitude are the largest reported between tissues in vertebrates. The relationship between $\delta^{202}\text{Hg}_t$ and $f(\text{MeHg})$ (Figure 3) shows a general trend of lower $\delta^{202}\text{Hg}_t$ values in tissues with decreasing $f(\text{MeHg})$ yet pronounced variability in $\delta^{202}\text{Hg}_t$ values in tissues with $f(\text{MeHg}) \leq 0.1$ (as great as 1.4 ‰). Under the premise that Hg in the petrel tissues reflects an integrated isotope signature of environmental sources, we interpret this variability to indicate that $\delta^{202}\text{Hg}_t$ values cannot solely be explained by isotopic fractionation due to the demethylation of MeHg to a single inorganic Hg species (e.g., $\text{Hg}(\text{Sec})_4$). This observation is dissimilar to those in waterbirds with modest levels of demethylation,¹¹ but similar to those in liver and kidneys of long-finned pilot whales with high levels of demethylation and formation of HgSe.^{15, 16}

The analysis of tissues by HR-XANES spectroscopy quantified Hg as three primary species (MeHg, $\text{Hg}(\text{Sec})_4$, HgSe) that vary in relative abundance (Figure 1, Table S2).⁷ The petrel feathers contain exclusively MeHg, muscle and kidneys tissues contain mixtures of $\text{Hg}(\text{Sec})_4$ and HgSe with or without minor MeHg, and the liver tissues contain almost exclusively HgSe ($0.90 \leq f(\text{HgSe}) \leq 1.00$). Of the three brain tissues, Hg is present as a mixture of MeHg, $\text{Hg}(\text{Sec})_4$, and HgSe with varying amounts of MeHg ($0.13 \leq f(\text{MeHg}) \leq 0.83$). Taken together, the isotopic and spectroscopic results suggest that each Hg species has a unique $\delta^{202}\text{Sp}_i$ value and the $\delta^{202}\text{Hg}_t$ of tissues (Figure 2) are the weighted sum of $\delta^{202}\text{MeHg}$, $\delta^{202}\text{Hg}(\text{Sec})_4$, and $\delta^{202}\text{HgSe}$. This hypothesis was tested by solving for $\delta^{202}\text{Sp}_i$ values the following 23 linear equations using an inversion algorithm⁸:

$$\delta^{202}\text{Hg}_t = f(\text{MeHg}) \times \delta^{202}\text{MeHg} + f(\text{Hg}(\text{Sec})_4) \times \delta^{202}\text{Hg}(\text{Sec})_4 + f(\text{HgSe}) \times \delta^{202}\text{HgSe} \quad (8)$$

where $\delta^{202}\text{Hg}_t$ is the measured isotopic value (Table S1) and the Hg species fractions ($f(\text{MeHg})$, $f(\text{Hg}(\text{Sec})_4)$, $f(\text{HgSe})$) are the proportions obtained by HR-XANES for each tissue (Figure 1, Table S2). Three species-specific $\delta^{202}\text{Hg}$ values proved sufficient to quantitatively describe the whole data set within experimental uncertainties (complete details on the inversion procedure are provided in the Methods section). Species-specific $\delta^{202}\text{Hg}$ are $\delta^{202}\text{MeHg} = 2.68 \pm 0.04 \text{ ‰}$, $\delta^{202}\text{Hg}(\text{Sec})_4 = -1.27 \pm 0.06$

‰, and $\delta^{202}\text{HgSe} = 0.19 \pm 0.02$ ‰ (value \pm standard deviation). With twenty-three equations and only three unknowns, the mathematical system is overdetermined.

To verify this mathematical approach, $\delta^{202}\text{Hg}_t$ values of tissues were recalculated (termed $\delta^{202}\text{Hg}_{calc}$) using the established $\delta^{202}\text{Sp}_i$ values and $f(\text{Sp}_i)_t$ determined by HR-XANES (Eq. 8) (Table S2). Figure 4 shows excellent agreement between $\delta^{202}\text{Hg}_{calc}$ and $\delta^{202}\text{Hg}_t$ with a slope near unity (slope = 0.97), indicating that the computational procedure provides robust isotopic solutions with uncertainties comparable to the precision of isotopic measurements (see Methods section) and is resistant to random and systematic errors in the independent spectroscopic and isotopic data sets. A single tissue was identified as an outlier (kidneys of individual P5 with $\delta^{202}\text{Hg}_k = 0.26$ ‰ and $\delta^{202}\text{Hg}_{calc} = -0.63$ ‰) and was excluded from the data set to provide an improved estimate of $\delta^{202}\text{Sp}_i$ values. After recalculation ($n=22$, Eq. 8), the determined $\delta^{202}\text{Sp}_i$ values were only marginally different ($\delta^{202}\text{MeHg} = 2.69 \pm 0.04$ ‰, $\delta^{202}\text{Hg}(\text{Sec})_4 = -1.37 \pm 0.06$ ‰, and $\delta^{202}\text{HgSe} = 0.18 \pm 0.02$ ‰), once again highlighting the robustness of the linear algebra approach. The recalculated $\delta^{202}\text{Sp}_i$ values were further used in all formal analyses in this study.

To independently verify the $\delta^{202}\text{Sp}_i$ values, MeHg was isolated from a subset of the tissues ($n=9$; Table S3),²⁷ including at least one sample of each tissue type, for direct $\delta^{202}\text{MeHg}$ measurement. Of the measured $\delta^{202}\text{MeHg}$ values, five were in good agreement (2.58 – 2.87 ‰) and within analytical precision (0.16 ‰, see Methods section) with the calculated $\delta^{202}\text{MeHg}$ value (2.69 ± 0.04 ‰), whereas one was modestly higher (feathers for P5, $\delta^{202}\text{MeHg}_f = 3.05$ ‰) and three lower than the calculated $\delta^{202}\text{MeHg}$ value (muscle and liver for P3, $\delta^{202}\text{MeHg}_m = 2.07$ ‰ and $\delta^{202}\text{MeHg}_l = 1.87$ ‰; muscle for P4, $\delta^{202}\text{MeHg}_m = 0.90$ ‰). Notably, instances where measured $\delta^{202}\text{MeHg}$ values deviated from the calculated $\delta^{202}\text{MeHg}$ value were of muscle and liver tissues with low MeHg ($f(\text{MeHg}) = 0.07$) or MeHg below the spectroscopic detection limit ($f(\text{MeHg}) < 0.03$). Thus, discrepancies between calculated and measured $\delta^{202}\text{MeHg}$ values can, in part, be attributed to low MeHg levels of tissues being negligible in the computation of $\delta^{202}\text{Sp}_i$ values. The feather with a modestly higher measured $\delta^{202}\text{MeHg}$ value compared to that calculated (3.05 ± 0.03 ‰ versus 2.69 ± 0.04 ‰, respectively) likely reflects subtle variability in $\delta^{202}\text{Hg}$ typical of seabirds with diverse dietary sources and feeding ranges.^{11, 12, 19} Taken together, the aforementioned assessments highlight the precision and accuracy of calculated $\delta^{202}\text{Sp}_i$ values by the inversion of isotopic and spectroscopic data.

The large variability in $\delta^{202}\text{Hg}_t$ values in the petrel tissues (≤ 3.5 ‰) is therefore due to the coexistence of MeHg and two inorganic species, $\text{Hg}(\text{Sec})_4$ and HgSe , with contrasted $\delta^{202}\text{Hg}$ values. Accordingly, the feathers with exclusively MeHg exhibit an average $\delta^{202}\text{Hg}_t$ value weighted by measurement uncertainties of 2.71 ± 0.06 ‰, consistent with $\delta^{202}\text{MeHg} = 2.69 \pm 0.04$ ‰, and the liver tissues with nearly exclusively HgSe exhibit a weighted average $\delta^{202}\text{Hg}_t$ values of 0.04 ± 0.06 ‰ close to $\delta^{202}\text{HgSe} = 0.18 \pm 0.02$ ‰ (Figure 1, Table S2). The $\delta^{202}\text{Hg}_t$ values of muscle tissues are up to 1.2 ‰ lower than those of kidney tissues due to the high proportion of Hg as isotopically distinct $\text{Hg}(\text{Sec})_4$ ($0.40 \leq f(\text{Hg}(\text{Sec})_4) \leq 0.72$) with the same proportion of MeHg. Further, the kidneys contain intermediate proportions of $\text{Hg}(\text{Sec})_4$ and HgSe , yet a similar mean $\delta^{202}\text{Hg}$ value than the liver ($\delta^{202}\text{Hg}_k = 0.01 \pm 0.25$ ‰). The reason is the presence of small proportions of MeHg in the kidneys ($0 \leq f(\text{MeHg}) \leq 0.12$) that are not present in the liver. These observations underscore that the chemical speciation of Hg determines tissue-specific differences in $\delta^{202}\text{Hg}$ values within and between petrels.

The ratio of MIF anomalies ($\Delta^{199}\text{Hg}/\Delta^{201}\text{Hg}$) of tissues shows a $\Delta^{199}\text{Hg}/\Delta^{201}\text{Hg}$ with a slope of 1.31 (Figure S1), which supports that photo-chemical processes prior to food web uptake are primarily responsible for $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values within petrel tissues. Surprisingly, a careful analysis of the data identifies that $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values in liver and kidney tissues were significantly lower than values of the MeHg pool as determined by feather values (predominantly MeHg) and direct MeHg isotope measurements from muscle, liver, or kidneys (t-test, p -value <0.001 , $n=20$; Figure S2). There was an observed hierarchy in $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values across the five petrels, which we interpret to reflect subtle differences in the isotopic signature of Hg from environmental sources. Despite this subtle variability, liver and kidney tissue $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values were 1.43 ± 0.08 and 1.27 ± 0.06 ‰ (average \pm standard deviation, $n=10$) compared to 1.68 ± 0.06 and 1.45 ± 0.06 ‰ for MeHg ($n=8$), respectively. These differences greatly exceed the analytical precision of ≤ 0.03 and ≤ 0.07 ‰ for $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$, respectively (Figure S7, Table S5). Conversely, the differences in $\Delta^{200}\text{Hg}$ and $\Delta^{204}\text{Hg}$ values between the liver and kidney tissues (-0.02 ± 0.02 ‰ and 0.05 ± 0.04 ‰, respectively) and MeHg pool (0.00 ± 0.02 ‰ and 0.01 ± 0.04 ‰, respectively) were negligible and within the analytical precision (≤ 0.03 and ≤ 0.05 ‰ for $\Delta^{200}\text{Hg}$ and $\Delta^{204}\text{Hg}$ measurements, respectively; Figure S7).

Previous HR-XANES and species-specific isotope measurements of birds from diverse aquatic systems demonstrate that the $\text{MeHg} \rightarrow \text{Hg}(\text{Sec})_4$ demethylation reaction does not induce MIF.¹¹ Therefore, we evaluated if the $\text{Hg}(\text{Sec})_4 \rightarrow \text{HgSe}$ biomineralization reaction could be responsible for the observed MIF. Complete details of this analysis are provided in the Supporting Information and summarized in Table S4. For each petrel (P1-P5), “initial” $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values were assigned using measurements of MeHg, either of feathers or MeHg extracts of tissues, as no petrel tissues contained exclusively $\text{Hg}(\text{Sec})_4$ (Figure 1, Table S2). Next, differences in $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ were calculated for each individual between the “initial” MeHg pool and HgSe, the latter determined using liver tissues that contain nearly exclusively HgSe. Differences in $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ between MeHg and HgSe range from -0.10 to -0.32 ‰ and from -0.06 to -0.21 ‰ in the five individuals, respectively. The ratio of the difference in $\Delta^{199}\text{Hg}$ to the difference in $\Delta^{201}\text{Hg}$ (Diff. $\Delta^{199}\text{Hg}/\text{Diff. } \Delta^{201}\text{Hg}$) between MeHg and HgSe was 1.6 ± 0.3 (average \pm standard deviation, $n=5$) and is compared graphically in Figure S4 to $\Delta^{199}\text{Hg}/\Delta^{201}\text{Hg}$ ratios quantified previously for dark nuclear volume effect (NVE, 1.54 ± 0.22 ;²¹ 1.59 ± 0.05)²² and photochemical processes (1.0 to 1.35).³⁴ Consistency in the fractionation of $\Delta^{199}\text{Hg}$ relative to $\Delta^{201}\text{Hg}$ between MeHg and HgSe in the petrel and dark NVEs supports that equilibrium isotope effects underlie the isotopic fractionation for the $\text{Hg}(\text{Sec})_4 \rightarrow \text{HgSe}$ biomineralization reaction.

DISCUSSION

We demonstrate that the $\delta^{202}\text{Hg}$ values of diverse tissues of giant petrels can be described with a high degree of confidence as a weighted average of three species-specific $\delta^{202}\text{Sp}_i$ values corresponding to those of MeHg, $\text{Hg}(\text{Sec})_4$, and HgSe identified by HR-XANES.⁷ The quantification of $\delta^{202}\text{Sp}_i$ values is accomplished using the novel inversion of isotopic and spectroscopic data.⁸ Observed differences in species-specific $\delta^{202}\text{Sp}_i$ values are interpreted to reflect the isotopic fractionation of Hg of two internal transformations, the $\text{MeHg} \rightarrow \text{Hg}(\text{Sec})_4$ demethylation reaction (-4.1 ± 0.1 ‰) and $\text{Hg}(\text{Sec})_4 \rightarrow \text{HgSe}$ biomineralization reaction ($+1.6 \pm 0.1$ ‰). The latter transformation that induced fractionation of

$\delta^{202}\text{Hg}$ was accompanied by modest decreases in $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values in the liver and kidneys tissues (magnitude of changes of $\leq 0.3\text{‰}$ and $\leq 0.2\text{‰}$, respectively). The species-specific $\delta^{202}\text{Sp}_i$ values of MeHg, $\text{Hg}(\text{Sec})_4$, and HgSe were consistent across diverse tissues of all eight petrels, indicating that total $\delta^{202}\text{Hg}$ tissue values arise from varying mixtures of Hg species and that both MeHg and inorganic Hg redistribute internally.¹⁵ Therefore, the large negative fractionation of $\delta^{202}\text{Hg}$ due to MeHg to $\text{Hg}(\text{Sec})_4$ demethylation and subsequent positive fractionation in $\delta^{202}\text{Hg}$ values due to $\text{Hg}(\text{Sec})_4 \rightarrow \text{HgSe}$ biomineralization can explain the variability of $\delta^{202}\text{Hg}$ observed in and across biological tissues of vertebrates,^{14, 16, 19, 24} highlighting the importance of chemical speciation of Hg on stable Hg isotope values. Below we discuss the results that support the aforementioned interpretations and detail the implications of these findings for the toxicological understanding of Hg in high trophic level organisms and environmental application of stable Hg isotopes.

Isotope Fractionation Due to Internal Demethylation and Biomineralization Reactions. The lower $\delta^{202}\text{Hg}$ in $\text{Hg}(\text{Sec})_4$ is attributed to MDF from a chemical kinetic isotope effect of Hg^{35} during enzymatic demethylation of MeHg³⁶ and complexation to SelP⁵, a process documented in other waterbirds¹¹ and long-finned pilot whales⁸. The underlying reason for the greater $\delta^{202}\text{Hg}$ in HgSe is less clear, but several lines of evidence support that changes in the Hg bonding environment are responsible for it. First, the $\text{Hg}(\text{Sec})_4 \rightarrow \text{HgSe}$ transformation is not a one-step reaction but proceeds via a multi-step reaction mechanism $\text{Hg}(\text{Sec})_4 \rightarrow \text{Hg}_x(\text{Se},\text{Sec})_y \rightarrow \text{HgSe}$.^{5, 7, 9} The multinuclear intermediate $\text{Hg}_x(\text{Se},\text{Sec})_y$ species, which is spectroscopically and isotopically indistinguishable from $\text{Hg}(\text{Sec})_4$ when disordered,¹¹ undergoes biomineralization to nanocrystalline HgSe⁷ as the ratio of x to y increases from 1:4 to 1:1 and Sec (selenocysteine) residues are replaced by inorganic Se ligands. Importantly, the isotopic fractionation of Hg attributed to the $\text{Hg}(\text{Sec})_4 \rightarrow \text{HgSe}$ transformation is observed in the petrels concurrent with HR-XANES and electron microscopy evidence for an increase in crystallinity of disordered $\text{Hg}_x(\text{Se},\text{Sec})_y$ to HgSe.⁷

Second, the observed fractionation of $\delta^{202}\text{Hg}$ between $\text{Hg}(\text{Sec})_4$ and HgSe is accompanied by significant decreases in $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ (Figure S2). The differences in $\Delta^{199}\text{Hg}$ relative to $\Delta^{201}\text{Hg}$ between MeHg and HgSe in the petrels (1.6 ± 0.3), which indicate the isotope fractionation process, are consistent with the measured and theoretical ratios for NVE (mean of 1.54 – 1.59 and 1.65, respectively).^{21, 22} Notably, the magnitude of isotope differences attributed to $\text{Hg}(\text{Sec})_4 \rightarrow \text{HgSe}$ biomineralization ($\delta^{202}\text{Hg} = 1.6\text{‰}$; $\Delta^{199}\text{Hg} \leq 0.3\text{‰}$; $\Delta^{201}\text{Hg} \leq 0.2\text{‰}$) in giant petrels are greater than those observed previously in laboratory experiments between aqueous $\text{Hg}(\text{II})$ complexes and a bidentate thiol complex ($\leq 0.6\text{‰}$, $\leq 0.1\text{‰}$, and $\leq 0.05\text{‰}$, respectively),²¹ perhaps due to the multi-dentate nature of the Hg-Se complexes and biomineralization of disordered $\text{Hg}_x(\text{Se},\text{Sec})_y$ to compact, crystalline HgSe. Although other processes could explain these observations, the directionality and relative magnitude of isotopic fractionation in the tissues are remarkably similar to laboratory experiments and theoretical calculations of equilibrium isotope effects from MDF and NVE^{21, 22} and inconsistent with other phenomena (e.g., MIF by kinetic magnetic isotope effects).³⁴ Thus, we assert that equilibrium isotope effects of MDF and NVE are likely responsible for the isotopic fractionation between $\text{Hg}(\text{Sec})_4$ and HgSe. This is the first study *in natura* documenting evidence for NVE of Hg due to dark biotic transformation.

Isotopic and spectroscopic measurements of other seabirds, fish, and mammals support our interpretations. In long-finned pilot whales (*G. melas*), the analysis of HgSe nanoparticles from the

liver show $\delta^{202}\text{Hg}$ values increase from $-0.85 \pm 0.06 \text{ ‰}$ at 2.5 years of age to $-0.09 \pm 0.06 \text{ ‰}$ at 29 years of age as the particles increase in size, in agreement with the isotopic evolution of Hg with a progressive increase of the ratio of x to y of $\text{Hg}_x(\text{Se},\text{Sec})_y$ towards HgSe. Further, the magnitude of isotopic fractionation of $\delta^{202}\text{Hg}$ between $\text{Hg}(\text{Sec})_4$ and HgSe is comparable in the long-finned pilot whales ($+1.4 \text{ ‰}$)⁸ and giant petrels ($+1.6 \text{ ‰}$). In regards to MIF anomalies, $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values of long-finned pilot whales of Li et al.¹⁶ are significantly lower ($\leq -0.25 \text{ ‰}$) in the liver compared to the muscle of adult whales (Figure S5), which was not identified by the authors of the study, and consistent with HgSe biomineralization primarily in the adult whale liver.⁸ A separate study of beluga whale liver tissues (*Delphinapterus leucas*) shows increases in $\delta^{202}\text{Hg}$ concomitant with decreases in $\Delta^{199}\text{Hg}$ in older individuals not explained by ecological factors (e.g., no differences in $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ which are indicative of shifts in foraging behavior). Beyond whales, further indirect evidence for the occurrence of $\text{Hg}(\text{Sec})_4$ is provided in the literature by the deviation from a linear correlation between $\delta^{202}\text{Hg}$ and $f(\text{MeHg})$ in marine fish (*Brosme brosme*),¹³ other seabirds (*Pachyptila desolata*, *Procellaria aequinoctialis*),¹² and seals (*Phoca sibirica*).^{14, 19} $\delta^{202}\text{Hg}$ differed by as much as 1.2 ‰ between liver and muscle tissues of giant petrels in this study and 1.6 ‰ in two other giant petrels with identical low $f(\text{MeHg})$ (Figure S6),¹² which are supporting evidence for a mixture of isotopically distinct HgSe and $\text{Hg}(\text{Sec})_4$. Thus, the isotopic fractionation due to $\text{Hg}(\text{Sec})_4 \rightarrow \text{HgSe}$ biomineralization appears to occur in diverse vertebrates.

The magnitude of measured isotopic fractionation of $\delta^{202}\text{Hg}$ for the $\text{MeHg} \rightarrow \text{Hg}(\text{Sec})_4$ demethylation reaction is significantly greater in giant petrels ($-4.1 \pm 0.1 \text{ ‰}$) than other waterbirds ($-2.2 \pm 0.1 \text{ ‰}$)¹¹ and long-finned pilot whales ($-2.6 \pm 0.1 \text{ ‰}$).⁸ This observation is likely explained by physiological differences between organisms in the rate of demethylation relative to new dietary MeHg intake and MeHg depuration, rather than mechanistic differences in the demethylation reaction. The giant petrels demethylate MeHg with extreme efficiency, as indicated by (1) predominantly inorganic Hg in the muscle, kidneys, and liver, and (2) a vast isotopic difference (-4.1 ‰) between reactant (i.e., MeHg) and product (i.e., $\text{Hg}(\text{Sec})_4$). In contrast, waterbirds from diverse aquatic environments with modest levels of demethylation, including a south polar skua (*Stercorarius maccormicki*) from the same geographic area as the giant petrels, exhibited an isotopic difference between MeHg and $\text{Hg}(\text{Sec})_4$ of -2.2 ‰ .¹¹ We propose that organisms that exhibit faster $\text{MeHg} \rightarrow \text{Hg}(\text{Sec})_4$ demethylation relative to dietary MeHg replenishment and MeHg depuration via keratinized tissues² will behave increasingly like a closed isotope system rather than an open isotope system with infinite reactant reservoirs (i.e., MeHg), resulting in a residual MeHg pool with a greater positive $\delta^{202}\text{Hg}$ offset and expansion in the measured isotopic fractionation between MeHg and $\text{Hg}(\text{Sec})_4$. This assessment is supported by $\delta^{202}\text{Hg}$ values of feathers in this study exceeding all previous observations of Antarctic seabirds^{12, 33} and consistent with observations in Arctic seabirds.¹⁹ Physiological controls on demethylation efficiency in birds and other animals are unknown but may depend on the availability of Se and rate of transcription of the SelP gene.^{37, 38}

Internal Redistribution of MeHg and Inorganic Hg. The speciation and isotopic results provide a new understanding of the metabolic routing of Hg between tissues (Figure 5). The relative invariance of $\delta^{202}\text{MeHg}$ between tissues and across individual petrels can be explained by (1) the continuous supply of new dietary MeHg with similar isotopic composition, and its cellular uptake from, and efflux to, the blood,³⁹ and (2) the highly labile nature of the Hg-SR bond,⁴⁰ that results in rapid internal cycling of MeHg with a greater $\delta^{202}\text{Hg}$ value (i.e., kinetic isotope effect).

The observed invariance of $\delta^{202}\text{Hg}(\text{Sec})_4$ and $\delta^{202}\text{HgSe}$ can be explained by (1) the formation of $\text{Hg}(\text{Sec})_4$ from MeHg of uniform isotopic composition and (2) the internal redistribution of inorganic Hg, perhaps as $\text{Hg}(\text{Sec})_4$. Mechanistic studies in nature linking tissue-specific transformations of Hg and internal redistribution in the circulatory system are lacking and needed. Notwithstanding, HgSe granules should be immobile because they are aggregated.⁷ SelP is expressed and secreted in plasma primarily by the liver and secondarily by extrahepatic tissues (e.g., heart and brain)^{37, 38, 41-45} and is taken up into cells through the apolipoprotein E Receptor-2 (ApoER2) to supply Se for selenoprotein synthesis.⁴⁶⁻⁴⁹ It is likely that MeHg is demethylated to $\text{Hg}(\text{Sec})_4$ by SelP and $\text{Hg}(\text{Sec})_4$ bound to SelP is redistributed to other tissues, including muscle,^{12-17, 19} via the circulatory system. Under this scheme, the biomineralization of HgSe within hepatic and extrahepatic tissues occurs from $\text{Hg}(\text{Sec})_4$ of uniform isotopic composition, and $\text{Hg}(\text{Sec})_4$ forms from isotopically uniform MeHg. Previous observation of Hg-bound SelP in the plasma of Inuits supports this interpretation,⁵⁰ as does isotopic differences between the blood of long-finned pilot whales compared to tissue MeHg values.¹⁵ Further research is needed to establish linkages between Hg transformations and Se homeostasis, which, along with evidence for the highly labile nature of the Hg-ligand complexes in biological systems,⁴⁰ will fuel the advancement of toxicokinetic models on Hg.

Implications for Environmental Application of Hg Stable Isotopes. Stable Hg isotopes are frequently used tools in the examination of Hg bioaccumulation in high trophic level organisms, often focusing on pressing environmental questions in Hg science, and will undoubtedly aid in improving our understanding of Hg cycling and monitoring the effectiveness of global Hg mitigation strategies (i.e. Minamata Convention on Mercury).^{25, 26} Our findings demonstrate that stable Hg isotope values in top predators are highly dependent on the chemical speciation of Hg and can reflect both kinetic and equilibrium isotope effects during internal transformations, which need to be considered if stable Hg isotopes are used for source apportionment. The accurate interpretation of stable Hg isotopes in birds and mammals necessitates a quantitative understanding of the chemical speciation of Hg across all tissue types, as internal demethylation and biomineralization transformations can exert dramatic, bidirectional influence on $\delta^{202}\text{Hg}$ values of hepatic and non-hepatic tissues (e.g., muscle) and more subtle shifts in $\Delta^{199}\text{Hg}$ in primarily liver tissues. Across organisms, differences in the physiological efficiency of the $\text{MeHg} \rightarrow \text{Hg}(\text{Sec})_4$ demethylation reaction will influence the $\delta^{202}\text{Hg}$ value of the residual MeHg pool across all tissues and measured isotopic fractionation for the $\text{MeHg} \rightarrow \text{Hg}(\text{Sec})_4$ reaction. This likely explains the observed range in isotopic $\delta^{202}\text{Hg}$ offset between dietary MeHg and MeHg in wildlife and humans (from +0.3 to +2.0 ‰) from previous work.^{17, 18, 51, 52} The findings further provoke a revisitation of the notion that internal transformations of Hg cannot induce MIF, as evidenced in modest MIF from $\text{Hg}(\text{Sec})_4 \rightarrow \text{HgSe}$ biomineralization in seabirds and whales.

Our results demonstrate that the speciation of Hg should be quantified, and not assumed, irrespective of tissue type,^{5, 7, 9, 11} particularly in organisms that have different extents of internal

demethylation. When the tissue-specific fractions of MeHg, Hg(Sec)₄, and HgSe are known ($f(\text{Sp}_i)_t$), as in this study, their isotopic $\delta^{202}\text{Hg}$ values ($\delta^{202}\text{Sp}_i$) resulting from demethylation and biomineralization reactions can be obtained mathematically by linear algebra (see SI). In this case, the linear system is well-posed and the $\delta^{202}\text{Sp}_i$ solution is unique. If only $f(\text{MeHg})_t$ is known, typically from chemical analysis, the linear system is modestly ill-posed and unique meaningful $\delta^{202}\text{Sp}_i$ values still can be obtained by regularization techniques, as demonstrated for long-finned pilot whales.^{8, 53}

Often the goal of Hg stable isotope applications is to connect Hg within the food web back to specific sources or processes, which becomes demonstrably more difficult when multiple Hg chemical species are present and contribute to the total Hg isotope pool. Our observations have important implications for Hg isotope source assessments in marine animals, especially baseline studies targeting climate change and the effectiveness of the Minamata Convention on Mercury, and suggest that interpretations can be skewed if Hg speciation is not considered. With these considerations in mind, the advancements presented here and elsewhere on new analytical and mathematical tools^{5, 7-9, 11} in Hg science will propel an improved mechanistic understanding on Hg transformation and toxicology in biota and facilitate applications of isotopic and spectroscopic approaches at the organism and ecosystem level.

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ASSOCIATED CONTENT

Supporting Information

Description of mass-independent fractionation analysis, supplementary figures S1-S7 (PDF).

Python code of the inversion software (Inv3-202Hg) (txt)

ReadMe of Inv3-202Hg (txt)

Input files of Inv3-202Hg (tx)

Input dataset (csv)

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Notes

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LEGENDS TO FIGURES

Figure 1. Ternary plot of the proportions of MeHg, Hg(Sec)₄, and HgSe in the liver (L, black), kidneys (K, blue), muscle (M, purple), and feathers (F, red) of five individuals (P1-P5), and the brain (B, green) of three other individuals (P6-P8), of giant petrels, as obtained by HR-XANES.⁷

Figure 2. Bar charts of $\delta^{202}\text{Hg}_t$ values of feathers (red), kidneys (blue), liver (black), and muscle (purple) tissues of five giant petrels (P1-P5) and brain tissues (green) of three additional giant petrels (P6-P8). Error bars present $\pm 1\text{SD}$ of isotope measurements (Table S2) and analytical precision in $\delta^{202}\text{Hg}$ of certified reference materials is provided in Table S5. The high $\delta^{202}\text{Hg}_t$ values of feathers are due to the exclusive presence of MeHg ($\delta^{202}\text{MeHg} = 2.69 \pm 0.04 \text{ ‰}$). Muscle tissues exhibit lower $\delta^{202}\text{Hg}_t$ values than liver tissues because muscle tissues contain a higher proportion of isotopically distinct $\text{Hg}(\text{Sec})_4$ ($\delta^{202}\text{Hg}(\text{Sec})_4 = -1.37 \pm 0.06 \text{ ‰}$) and the liver tissues contain primarily HgSe ($\delta^{202}\text{HgSe} = 0.18 \pm 0.02 \text{ ‰}$).

Figure 3. Relationship between $\delta^{202}\text{Hg}_t$ and $f(\text{MeHg})$ of giant petrel tissues ($n=23$, 1SD). The solid black line presents the linear regression between $\delta^{202}\text{Hg}_t$ and $f(\text{MeHg})$ excluding the muscle data (slope 2.6, y-intercept of 0.0 ‰, and a Pearson correlation $R^2 = 0.98$). The dashed black line is the linear regression between $\delta^{202}\text{Hg}_t$ and $f(\text{MeHg})$ including the muscle data ($R^2 = 0.91$). Error bars present uncertainties in $f(\text{MeHg})$ from HR-XANES fits⁷ and $\pm 1\text{SD}$ of isotope measurements (Table S2), and analytical precision in $\delta^{202}\text{Hg}$ of certified reference materials is provided in Table S5. Methylmercury is completely demethylated in the liver and almost completely demethylated in muscle and kidneys. The large variability in $\delta^{202}\text{Hg}_t$ for the same $f(\text{MeHg})$ at $f(\text{MeHg}) \leq 0.1$ results from the coexistence of two inorganic species, $\text{Hg}(\text{Sec})_4$ and HgSe, with contrasted species-specific $\delta^{202}\text{Hg}$ values.

Figure 4. Calculated versus measured species-averaged values of $\delta^{202}\text{Hg}$ ($\delta^{202}\text{Hg}_{calc}$ versus $\delta^{202}\text{Hg}_t$, respectively) in the giant petrel tissues ($n=23$). Error bars present $\pm 1\text{SD}$ of isotope values (Table S2). The solid black, dashed gray, and dotted gray lines present the linear fit, 95% confidence intervals of the fit, and prediction intervals of the fit, respectively. The outlier kidney of individual P5 (identified by the black arrow) was omitted in the calculation of the regression line ($\delta^{202}\text{Hg}_{calc} = 0.972 \times \delta^{202}\text{Hg}_t + 0.007$, $R^2 = 0.983$).

Figure 5. Schematic diagram showing the isotopic fractionation of ^{202}Hg due to the $\text{MeHgCys} \rightarrow \text{Hg}(\text{Sec})_4 \rightarrow \text{HgSe}$ demethylation and biomineralization reactions (left panel), and diagrammatic representation of the proportions of MeHgCys, $\text{Hg}(\text{Sec})_4$, and HgSe in the tissues of giant petrels (expressed as the percentage of total Hg), and of the species-specific and species-averaged $\delta^{202}\text{Hg}$ values (right panel).

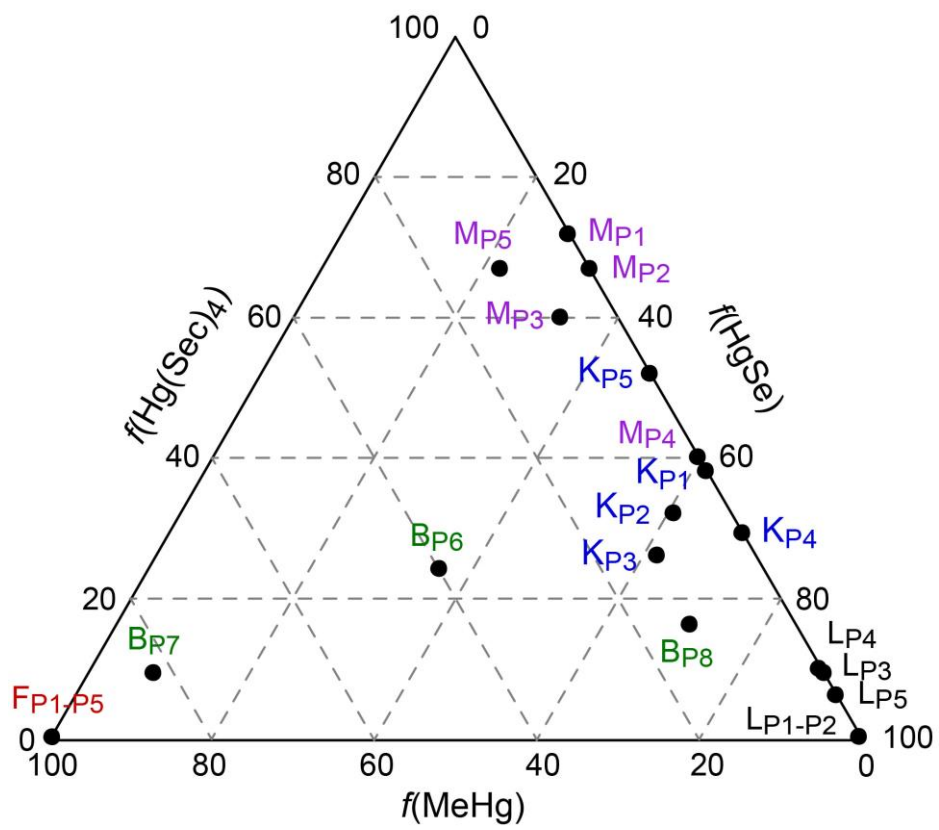


Figure 1

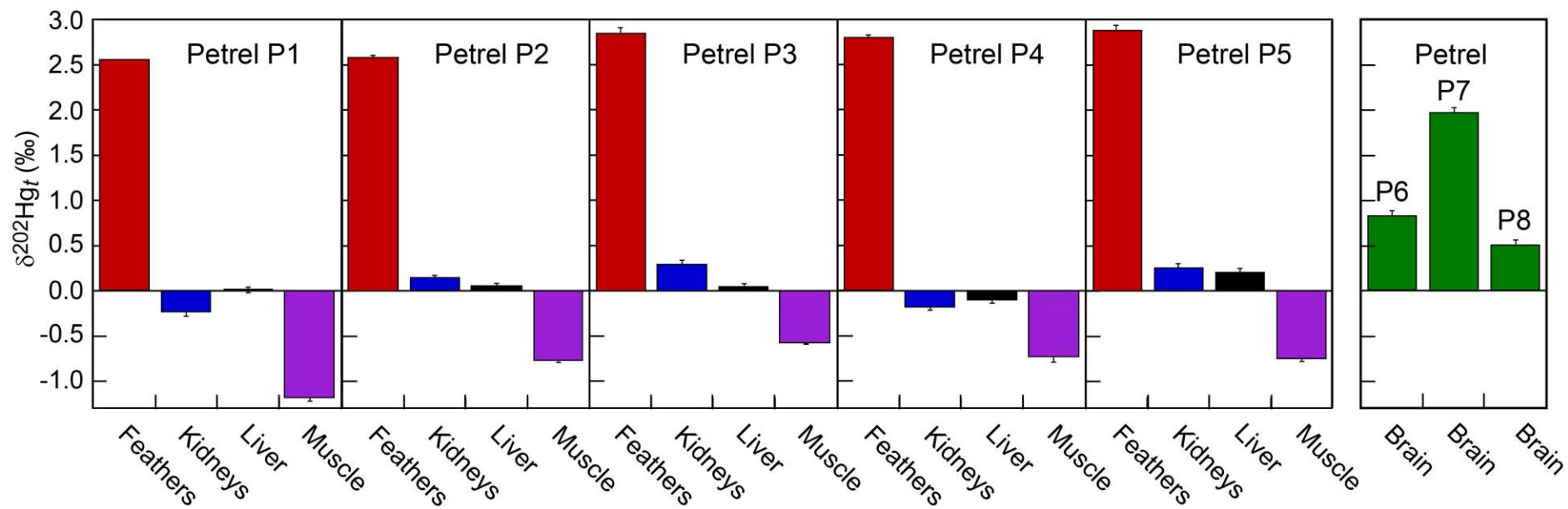


Figure 2

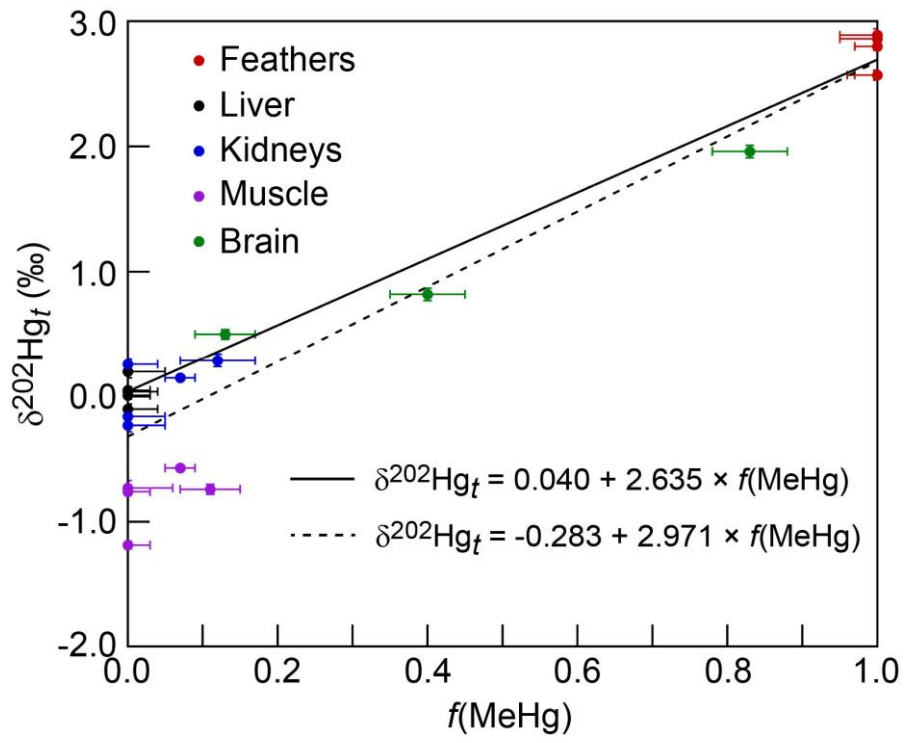


Figure 3

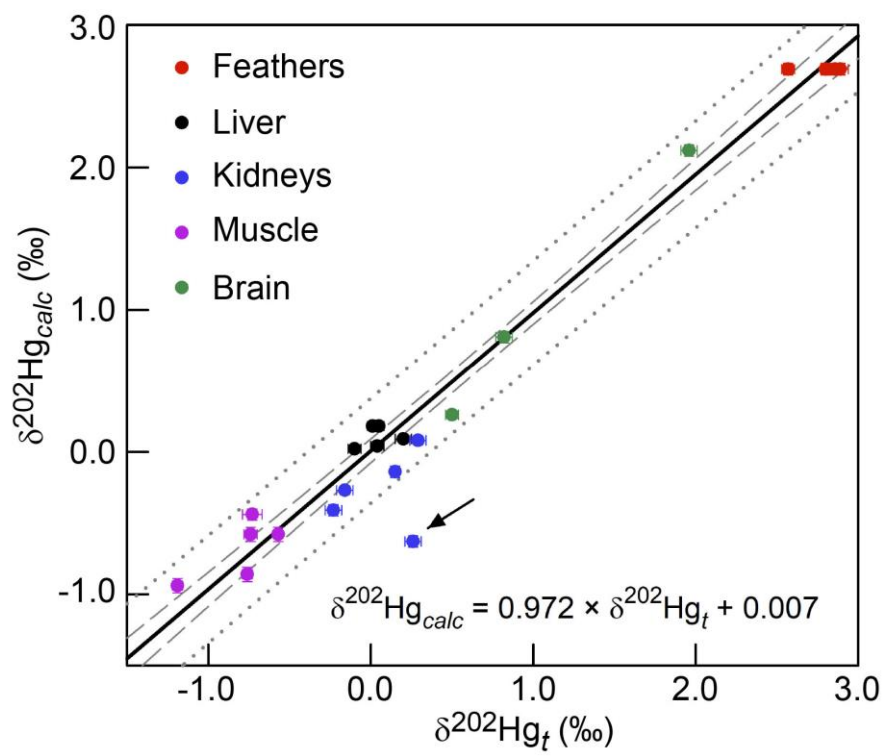
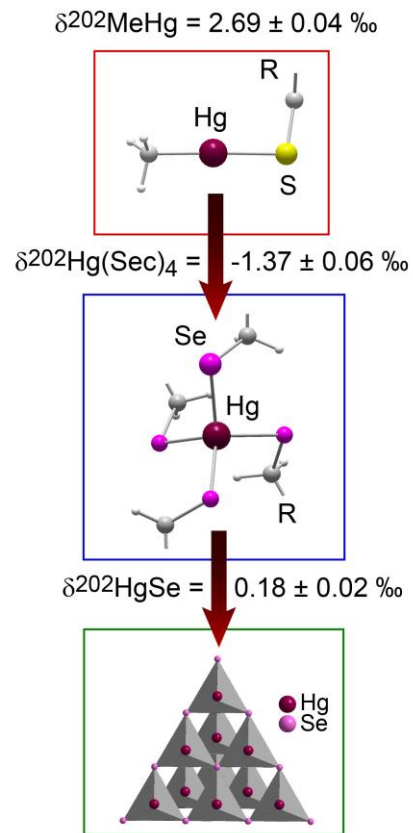


Figure 4

Species-specific fractionation



Species-averaged fractionation

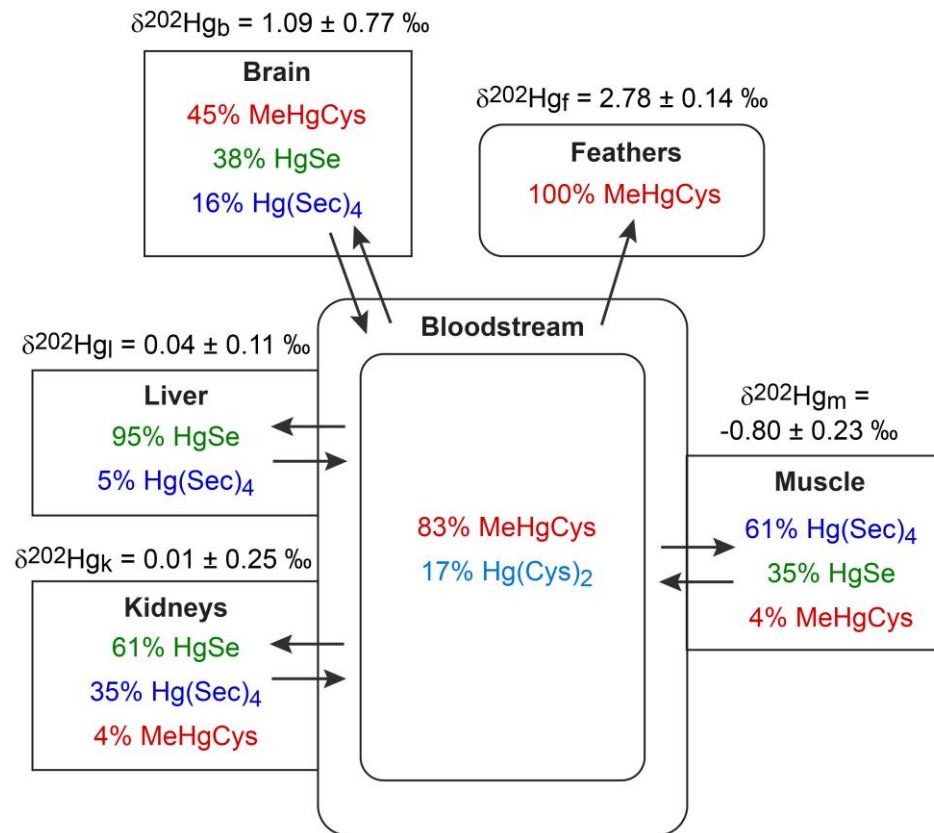


Figure 5

Supporting Information

Mercury isotope fractionation by internal demethylation and biomineralization reactions in seabirds: implications for environmental mercury science

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Supporting information includes 7 Figures, 5 Tables, and 11 pages.

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Mass-Independent Fractionation (MIF) Analysis

The MIF of mercury (Hg) isotopes was evaluated in giant petrels (*Macronectes* spp.) P1-P5 for the mercury-tetraselenolate complex ($\text{Hg}(\text{Sec})_4$) \rightarrow mercury selenide (HgSe) biomineralization reaction. Changes in $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ were calculated for each bird (Table S4). First, “initial” $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values were assigned for each petrel. Previous high energy-resolution X-ray absorption near-edge structure (HR-XANES) spectroscopy and species-specific isotope measurement of birds from diverse aquatic systems demonstrate that the demethylation reaction of methylmercury into mercury tetraselenolate (MeHg) \rightarrow $\text{Hg}(\text{Sec})_4$) does not induce mass independent fractionation (MIF).¹ Therefore, the “initial” $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values were assigned for each individual using isotope values of feathers (which contained exclusively MeHg , Table S2) or the mean of feather values and values of MeHg extracted from tissues. The $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values of the MeHg pool were assumed to be integrated across tissues and be representative of MeHg during the adult life of the bird. The specific tissues used to establish the “initial” $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values are provided in Table S4. Next, $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values of HgSe were assigned for each petrel using isotope values of liver tissues, which contained nearly exclusively mercury selenide ($0.90 \leq f(\text{HgSe}) \leq 1.00$). The $\text{Hg}(\text{Sec})_4 \rightarrow \text{HgSe}$ biomineralization reaction is nearly complete in the liver of the petrels, and therefore difference in

$\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values between the “initial” MeHg and liver were calculated. As shown in Table S4, the $\text{Hg}(\text{Sec})_4 \rightarrow \text{HgSe}$ biomineralization reaction resulted in decreases in $\Delta^{199}\text{Hg}$ values by -0.10 to -0.32 ‰ and $\Delta^{201}\text{Hg}$ values by -0.06 to -0.21 ‰, respectively. Differences in $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values exceeded the analytical precision (Table S5, Figure S7). The ratio of the difference in $\Delta^{199}\text{Hg}$ to difference in $\Delta^{201}\text{Hg}$ ($\Delta^{199}\text{Hg}/\Delta^{201}\text{Hg}$) between MeHg and HgSe, which is interpreted to represent the slope of change in $\Delta^{199}\text{Hg}$ to $\Delta^{201}\text{Hg}$ for the internal $\text{Hg}(\text{Sec})_4 \rightarrow \text{HgSe}$ biomineralization reaction, averaged 1.6 ± 0.3 (average \pm standard deviation, $n=5$; ranged from 1.10 to 1.93) and was in good agreement with the measured ($\sim 1.54 \pm 0.22$) and theoretical (1.65) ratio for nuclear volume effect (NVE).²

All Hg isotope data collected as part of this study is available as part of a supplementary data release.¹⁰

Supplementary Figures

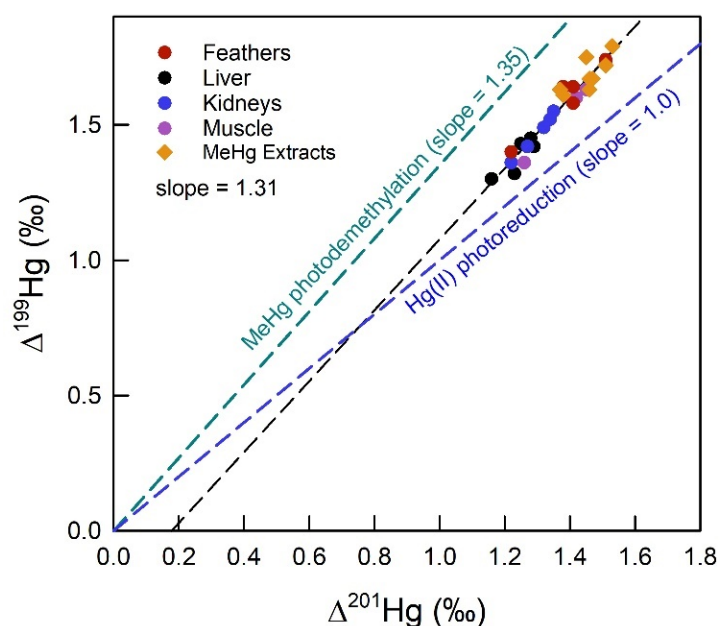


Figure S1. Odd mass-independent fractionation (MIF) isotope values ($\Delta^{199}\text{Hg}$ vs $\Delta^{201}\text{Hg}$) of total Hg digests of giant petrel tissues ($n=20$) and MeHg extracts of selected tissues ($n=9$). The dashed gray line presents the fit of the data. Dashed blue and green lines present empirical slopes for residual Hg(II) photoreduction and residual MeHg photodemethylation processes in water,³ respectively. Analytical precision in $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values of certified reference materials is provided in Table S5. An outlier (P5, muscle, $\Delta^{199}\text{Hg} = 1.98 \pm 0.01$ ‰, $\Delta^{201}\text{Hg} = 1.33 \pm 0.01$ ‰) was identified based on deviation in the $\Delta^{199}\text{Hg}$ versus $\Delta^{201}\text{Hg}$ regression and is neither shown in the figure nor included in the regression.

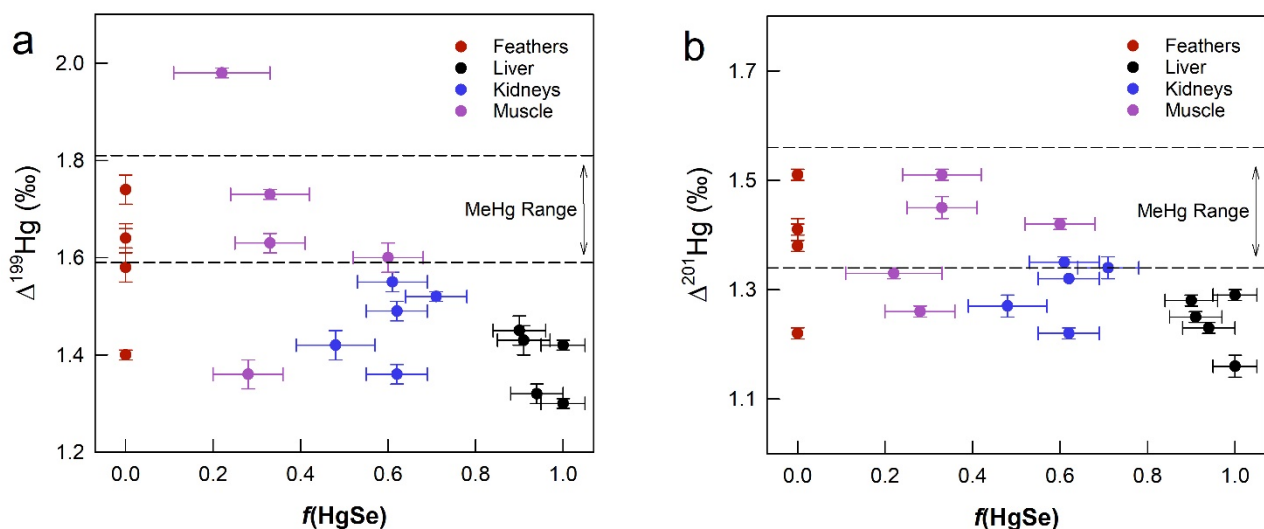


Figure S2. Scatter plot showing the (a) $\Delta^{199}\text{Hg}$ and (b) $\Delta^{201}\text{Hg}$ values versus the fraction of HgSe ($f(\text{HgSe})$) of giant petrel feathers and tissues with respect to the range of MeHg (dashed black lines). Differences in $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values between samples with $f(\text{HgSe})=0$ (feathers, and MeHg extracts, $n=10$) and kidney and liver tissues ($0.48 \leq f(\text{HgSe}) \leq 1.00$; $n=10$) were of statistical significance ($p < 0.001$; student's t-test). Vertical error bars present ± 1 standard deviation (SD) of isotope values and horizontal error bars present uncertainties in $f(\text{HgSe})$. An outlier sample (P5, muscle) was identified based on deviation in the $\Delta^{199}\text{Hg}$ versus $\Delta^{201}\text{Hg}$ regression, but is shown here for completeness. Analytical precision in $\Delta^{199}\text{Hg}$ and $\Delta^{201}\text{Hg}$ values of certified reference materials is provided in Table S5.

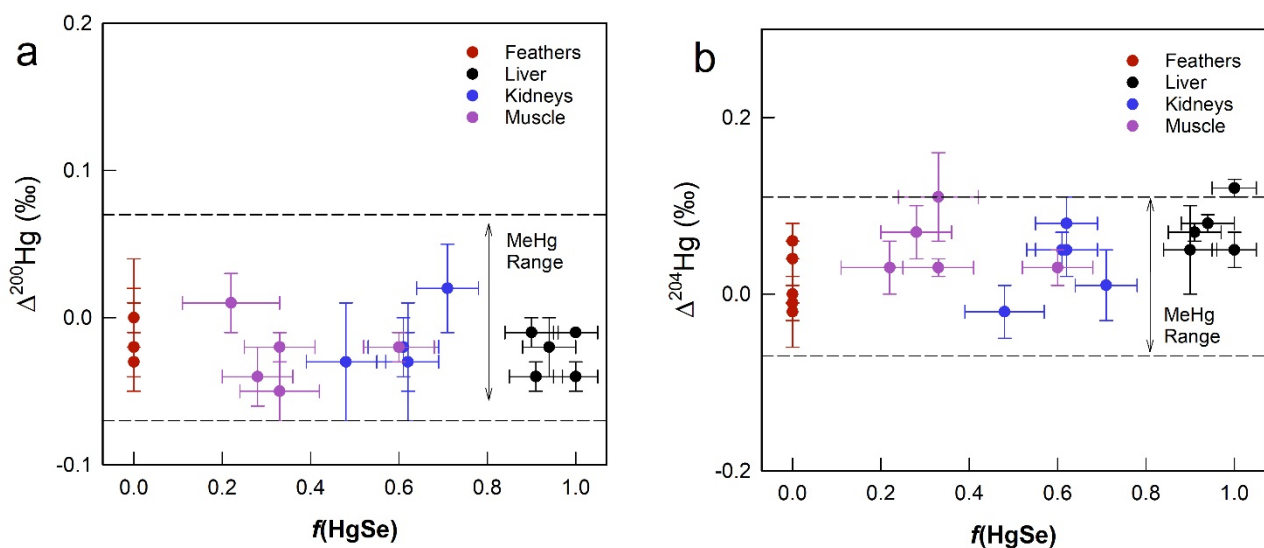


Figure S3. Scatter plot showing that the variability in (a) $\Delta^{200}\text{Hg}$ and (b) $\Delta^{204}\text{Hg}$ versus the fraction of HgSe ($f(\text{HgSe})$) of giant petrel feathers and tissues is within the range of MeHg (dashed black lines). Vertical error bars present ± 1 SD of isotope values and horizontal error bars present uncertainties in $f(\text{HgSe})$. Analytical precision in $\Delta^{200}\text{Hg}$ and $\Delta^{204}\text{Hg}$ values of certified reference materials are provided in Table S5.

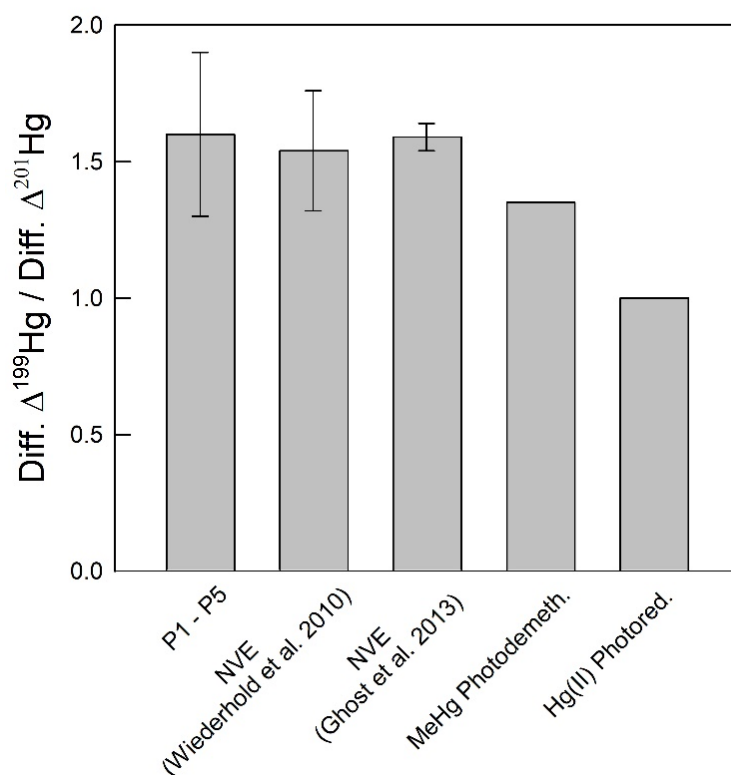


Figure S4. The ratio of the difference in $\Delta^{199}\text{Hg}$ to difference in $\Delta^{201}\text{Hg}$ ($\text{Diff. } \Delta^{199}\text{Hg} / \text{Diff. } \Delta^{201}\text{Hg}$) calculated between the MeHg pool and HgSe for giant petrels P1-P5 (average \pm standard deviation) and $\Delta^{199}\text{Hg} / \Delta^{201}\text{Hg}$ ratios induced by dark nuclear volume effect (NVE),^{2, 4} MeHg photodemethylation, and inorganic Hg(II) photoreduction.³

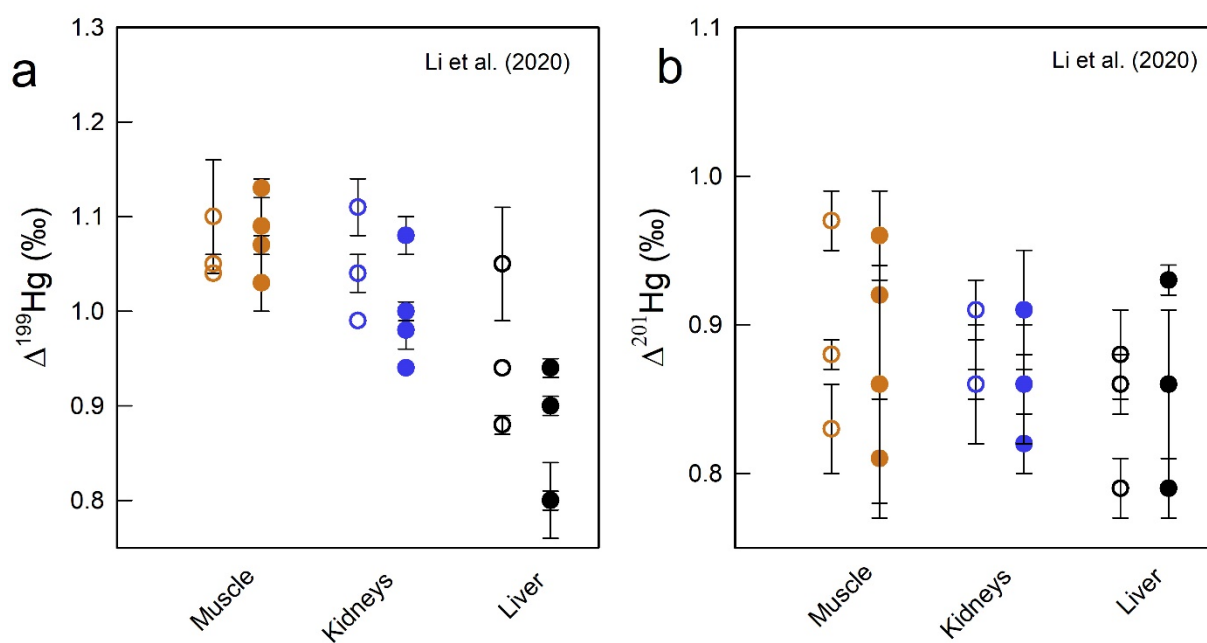


Figure S5. (a) $\Delta^{199}\text{Hg}$ and (b) $\Delta^{201}\text{Hg}$ values of long-finned pilot whale (*Globicephala melas*) muscle, kidneys, and liver from Li et al.⁵. Open symbols are for juvenile whales and solid symbols are for adult whales. Error bars present $\pm 1\text{SD}$ of isotope values.

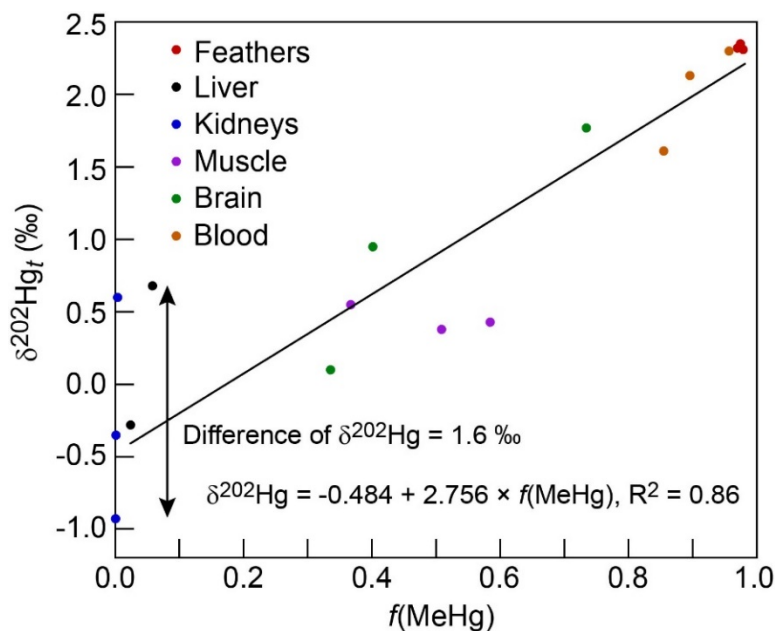


Figure S6. Plot of total Hg isotopes ($\delta^{202}\text{Hg}_t$) versus $f(\text{MeHg})$ for giant petrel tissues analyzed by Renedo et al.⁶ The main difference between these data and data from this study (Figure 3 in the main text) is the higher $\delta^{202}\text{Hg}_t$ and $f(\text{MeHg})$ values of the muscle in the previous study, which indicates that Hg was less demethylated in the Renedo et al.⁶ giant petrel individuals than in the present individuals. The large variability (as great as $\sim 1.6 \text{‰}$) in $\delta^{202}\text{Hg}_t$ of liver and kidneys observed at low $f(\text{MeHg})$ is a strong indication for the coexistence of $\text{Hg}(\text{Sec})_4$ and HgSe in the two tissues.

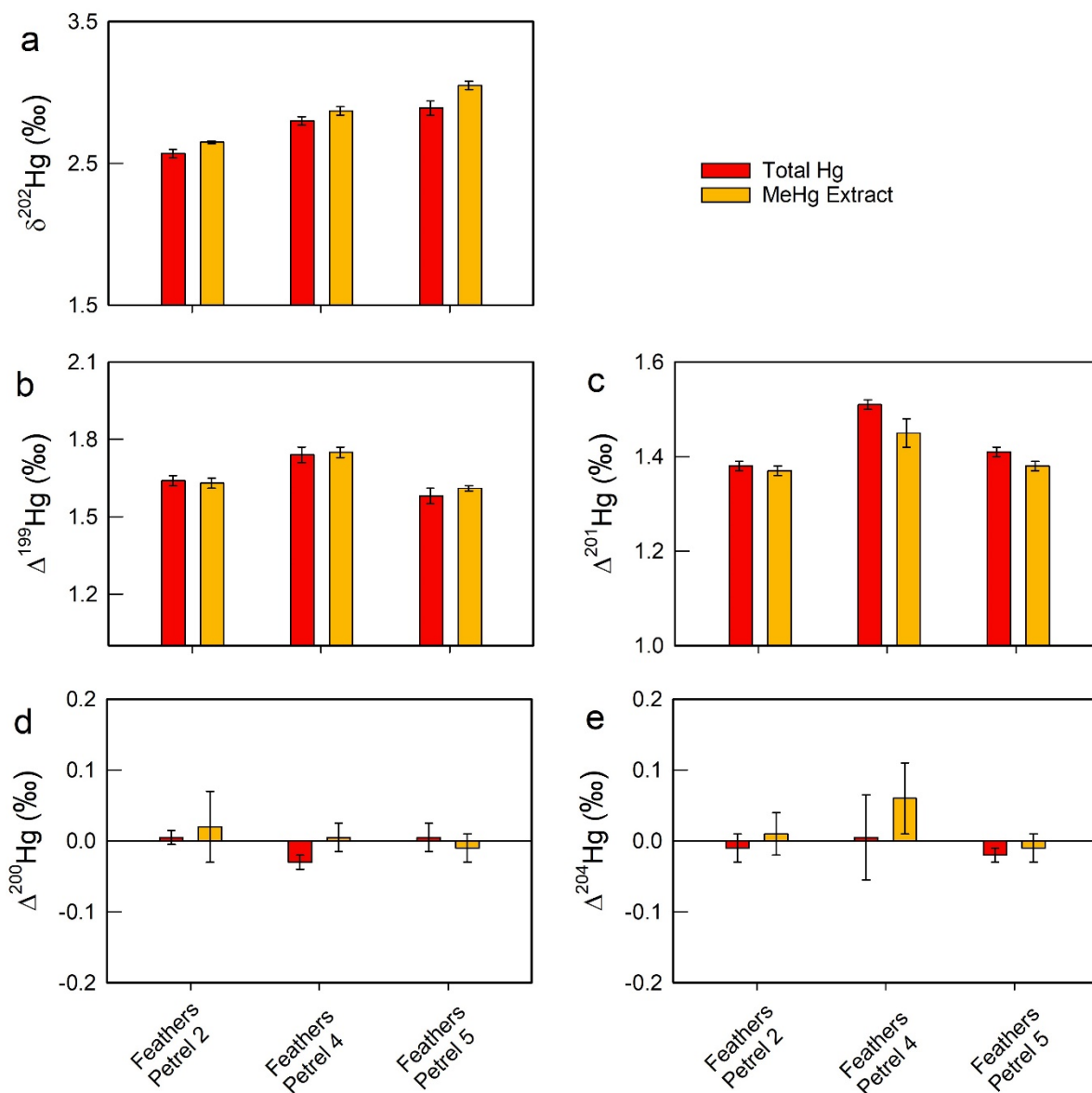


Figure S7. Stable mercury isotope values of (a) $\delta^{202}\text{Hg}$, (b) $\Delta^{199}\text{Hg}$, (c) $\Delta^{201}\text{Hg}$, (d) $\Delta^{200}\text{Hg}$, and (e) $\Delta^{204}\text{Hg}$ of total Hg digests (red bars) and methylmercury (MeHg) extracts (yellow bars) of feathers of three giant petrels (P2, P4, and P5). Error bars present $\pm 1\text{SD}$. Differences in values between total Hg and MeHg were $\leq 0.16\text{‰}$, $\leq 0.03\text{‰}$, $\leq 0.07\text{‰}$, $\leq 0.03\text{‰}$, and $\leq 0.05\text{‰}$ for $\delta^{202}\text{Hg}$, $\Delta^{199}\text{Hg}$, $\Delta^{201}\text{Hg}$, $\Delta^{200}\text{Hg}$, and $\Delta^{204}\text{Hg}$, respectively.

Supplementary Tables

Table S1. Isotopic values (± 1 SD) of total mercury in giant petrel tissues rounded to the nearest hundredth.

ID	Tissue	$\delta^{199}\text{Hg}$	$\pm 1\text{SD}$	$\delta^{200}\text{Hg}$	$\pm 1\text{SD}$	$\delta^{201}\text{Hg}$	$\pm 1\text{SD}$	$\delta^{202}\text{Hg}$	$\pm 1\text{SD}$	$\delta^{204}\text{Hg}$	$\pm 1\text{SD}$	$\Delta^{199}\text{Hg}$	$\pm 1\text{SD}$	$\Delta^{200}\text{Hg}$	$\pm 1\text{SD}$	$\Delta^{201}\text{Hg}$	$\pm 1\text{SD}$	$\Delta^{204}\text{Hg}$	$\pm 1\text{SD}$
P1	Feathers	2.04	0.02	1.29	0.03	3.15	0.04	2.57	0.04	3.90	0.05	1.40	0.01	0.00	0.04	1.22	0.01	0.06	0.02
	Liver	1.30	0.02	-0.04	0.02	1.16	0.02	0.01	0.03	0.13	0.04	1.30	0.01	-0.04	0.01	1.16	0.02	0.12	0.01
	Kidneys	1.30	0.03	-0.14	0.03	1.05	0.03	-0.23	0.05	-0.30	0.05	1.36	0.02	-0.03	0.04	1.22	0.01	0.05	0.03
	Muscle	1.06	0.03	-0.63	0.03	0.37	0.02	-1.19	0.03	-1.70	0.03	1.36	0.03	-0.04	0.02	1.26	0.01	0.07	0.03
P2	Feathers	2.28	0.02	1.29	0.02	3.31	0.02	2.57	0.03	3.83	0.05	1.64	0.02	0.00	0.01	1.38	0.01	-0.01	0.02
	Liver	1.43	0.02	0.02	0.02	1.33	0.03	0.05	0.03	0.13	0.05	1.42	0.01	-0.01	0.00	1.29	0.01	0.05	0.02
	Kidneys	1.59	0.03	0.05	0.03	1.47	0.02	0.15	0.02	0.27	0.04	1.55	0.02	-0.02	0.02	1.35	0.01	0.05	0.02
	Muscle	1.44	0.03	-0.41	0.02	0.88	0.03	-0.76	0.03	-1.11	0.05	1.63	0.02	-0.02	0.01	1.45	0.02	0.03	0.01
P3	Feathers	2.36	0.04	1.41	0.05	3.56	0.05	2.86	0.05	4.30	0.06	1.64	0.03	-0.02	0.03	1.41	0.02	0.04	0.02
	Liver	1.44	0.03	-0.02	0.03	1.28	0.04	0.04	0.04	0.13	0.05	1.43	0.03	-0.04	0.01	1.25	0.01	0.07	0.01
	Kidneys	1.56	0.04	0.11	0.05	1.53	0.04	0.29	0.05	0.50	0.07	1.49	0.02	-0.03	0.02	1.32	0.00	0.08	0.03
	Muscle	1.59	0.02	-0.33	0.03	1.08	0.02	-0.57	0.02	-0.74	0.03	1.73	0.01	-0.05	0.02	1.51	0.01	0.11	0.05
P4	Feathers	2.44	0.04	1.38	0.03	3.62	0.02	2.80	0.03	4.18	0.04	1.74	0.03	-0.03	0.01	1.51	0.01	0.00	0.06
	Liver	1.43	0.04	-0.06	0.03	1.20	0.02	-0.10	0.04	-0.10	0.03	1.45	0.03	-0.01	0.01	1.28	0.01	0.05	0.05
	Kidneys	1.47	0.03	-0.07	0.05	1.22	0.03	-0.16	0.05	-0.23	0.07	1.52	0.01	0.02	0.03	1.34	0.02	0.01	0.04
	Muscle	1.42	0.04	-0.38	0.03	0.87	0.04	-0.73	0.06	-1.06	0.07	1.60	0.03	-0.02	0.01	1.42	0.01	0.03	0.02
P5	Feathers	2.31	0.04	1.45	0.04	3.58	0.04	2.89	0.05	4.30	0.08	1.58	0.03	0.00	0.02	1.41	0.01	-0.02	0.01
	Liver	1.37	0.03	0.08	0.04	1.38	0.04	0.20	0.05	0.38	0.08	1.32	0.02	-0.02	0.02	1.23	0.01	0.08	0.01
	Kidneys	1.48	0.03	0.10	0.06	1.46	0.04	0.26	0.04	0.37	0.06	1.42	0.03	-0.03	0.04	1.27	0.02	-0.02	0.03
	Muscle	1.78	0.02	-0.36	0.02	0.77	0.03	-0.74	0.04	-1.07	0.06	1.97	0.02	0.01	0.01	1.33	0.00	0.03	0.01
P6	Brain	1.74	0.02	0.43	0.03	1.93	0.04	0.82	0.05	1.24	0.07	1.54	0.01	0.02	0.01	1.32	0.01	0.03	0.03
P7	Brain	1.82	0.02	0.98	0.03	2.64	0.04	1.96	0.05	3.00	0.07	1.33	0.02	-0.01	0.01	1.16	0.01	0.06	0.03
P8	Brain	1.64	0.03	0.25	0.03	1.68	0.03	0.50	0.04	0.82	0.07	1.51	0.02	0.00	0.01	1.30	0.00	0.07	0.02

Table S2. Chemical concentrations, fractions of Hg species by HR-XANES, and measured and calculated $\delta^{202}\text{Hg}$ isotopic values ($\pm 1\text{SD}$) of giant petrel tissues rounded to the nearest hundredth^a.

ID	Tissue	[Hg] _{tot} mg/kg ^b dw	f(MeHg)	f(Hg(Sec)4)	f(HgSe)	$\delta^{202}\text{Hg}_r$ ‰	$\delta^{202}\text{Hg}_{\text{calc}}^c$ ‰
P1	Feathers	4.8	1.00 ± 0.05	-	-	2.57 ± 0.04	2.69 ± 0.04
	Liver	226	-	-	1.00 ± 0.05	0.01 ± 0.03	0.18 ± 0.02
	Kidneys	54.5	-	0.38 ± 0.07	0.62 ± 0.07	-0.23 ± 0.05	-0.41 ± 0.04
	Muscle	26.6	-	0.72 ± 0.07	0.28 ± 0.08	-1.19 ± 0.03	-0.94 ± 0.05
P2	Feathers	26.2	1.00 ± 0.05	-	-	2.57 ± 0.03	2.69 ± 0.04
	Liver	804	-	-	1.00 ± 0.05	0.05 ± 0.03	0.18 ± 0.02
	Kidneys	37.7	0.07 ± 0.03	0.32 ± 0.09	0.61 ± 0.08	0.15 ± 0.02	-0.14 ± 0.04
	Muscle	42.0	-	0.67 ± 0.08	0.33 ± 0.08	-0.76 ± 0.03	-0.86 ± 0.05
P3	Feathers	14.9	1.00 ± 0.05	-	-	2.86 ± 0.05	2.69 ± 0.04
	Liver	293	-	0.09 ± 0.06	0.91 ± 0.06	0.04 ± 0.04	0.04 ± 0.02
	Kidneys	40.9	0.12 ± 0.03	0.26 ± 0.09	0.62 ± 0.07	0.29 ± 0.05	0.08 ± 0.03
	Muscle	21.8	0.07 ± 0.03	0.60 ± 0.09	0.33 ± 0.09	-0.57 ± 0.02	-0.57 ± 0.05
P4	Feathers	12.3	1.00 ± 0.05	-	-	2.80 ± 0.03	2.69 ± 0.04
	Liver	1499	-	0.10 ± 0.06	0.90 ± 0.06	-0.10 ± 0.04	0.02 ± 0.02
	Kidneys	414	-	0.29 ± 0.07	0.71 ± 0.07	-0.16 ± 0.05	-0.27 ± 0.03
	Muscle	88.7	-	0.40 ± 0.08	0.60 ± 0.08	-0.73 ± 0.06	-0.44 ± 0.04
P5	Feathers	10.3	1.00 ± 0.05	-	-	2.89 ± 0.05	2.69 ± 0.04
	Liver	170	-	0.06 ± 0.06	0.94 ± 0.06	0.20 ± 0.05	0.09 ± 0.02
	Kidneys	9.84	-	0.52 ± 0.09	0.48 ± 0.09	0.26 ± 0.04	-0.63 ± 0.04
	Muscle	2.87	0.11 ± 0.05	0.67 ± 0.13	0.22 ± 0.11	-0.74 ± 0.04	-0.58 ± 0.05
P6	Brain	2.8	0.40 ± 0.06	0.24 ± 0.15	0.36 ± 0.13	0.82 ± 0.05	0.81 ± 0.04
P7	Brain	1.6	0.83 ± 0.05	0.09 ± 0.09	0.08 ± 0.08	1.96 ± 0.05	2.12 ± 0.04
P8	Brain	13.2	0.13 ± 0.05	0.16 ± 0.10	0.71 ± 0.08	0.50 ± 0.04	0.26 ± 0.03

^aThe inversion calculations were performed on unrounded isotopic values, as reported in the SI Input dataset file.

^bFrom Ref. ⁷.

^cCalculated value using Eq. 2 in main text, fractions of Hg species by HR-XANES, and $\delta^{202}\text{MeHg} = 2.69 \pm 0.04$ ‰, $\delta^{202}\text{Hg}(\text{Sec})4 = -1.37 \pm 0.06$ ‰, and $\delta^{202}\text{HgSe} = 0.18 \pm 0.02$.

Table S3. Isotopic values (± 1 SD) of methylmercury (MeHg) in giant petrel tissues.

ID	Tissue	$\delta^{199}\text{Hg}$	$\pm 1\text{SD}$	$\delta^{200}\text{Hg}$	$\pm 1\text{SD}$	$\delta^{201}\text{Hg}$	$\pm 1\text{SD}$	$\delta^{202}\text{Hg}$	$\pm 1\text{SD}$	$\delta^{204}\text{Hg}$	$\pm 1\text{SD}$	$\Delta^{199}\text{Hg}$	$\pm 1\text{SD}$	$\Delta^{200}\text{Hg}$	$\pm 1\text{SD}$	$\Delta^{201}\text{Hg}$	$\pm 1\text{SD}$	$\Delta^{204}\text{Hg}$	$\pm 1\text{SD}$
P2	Feathers	2.30	0.02	1.35	0.05	3.36	0.02	2.65	0.01	3.98	0.02	1.63	0.02	0.02	0.05	1.37	0.01	0.01	0.03
	Kidneys	2.35	0.02	1.34	0.02	3.49	0.02	2.70	0.03	4.00	0.05	1.67	0.01	-0.02	0.02	1.46	0.01	-0.03	0.01
	Muscle	2.43	0.02	1.40	0.03	3.60	0.03	2.78	0.03	4.19	0.02	1.72	0.02	0.00	0.01	1.51	0.02	0.04	0.05
P3	Liver	2.14	0.02	0.96	0.01	2.87	0.03	1.87	0.03	2.81	0.02	1.67	0.02	0.02	0.01	1.47	0.02	0.02	0.06
	Muscle	2.15	0.01	1.04	0.03	3.02	0.01	2.07	0.03	3.11	0.03	1.63	0.01	0.00	0.02	1.46	0.02	0.01	0.02
P4	Feathers	2.47	0.01	1.44	0.03	3.61	0.05	2.87	0.03	4.35	0.02	1.75	0.02	0.00	0.02	1.45	0.03	0.06	0.05
	Muscle	2.01	0.01	0.43	0.02	2.21	0.01	0.90	0.04	1.30	0.06	1.79	0.02	-0.02	0.01	1.53	0.03	-0.05	0.02
P5	Feathers	2.38	0.02	1.53	0.02	3.68	0.02	3.05	0.03	4.54	0.05	1.61	0.01	-0.01	0.02	1.38	0.01	-0.01	0.02
P8	Brain	2.22	0.02	1.28	0.03	3.28	0.03	2.58	0.04	3.82	0.03	1.57	0.02	-0.01	0.01	1.33	0.02	-0.04	0.04

Table S4. Isotope values used to assess the mass-independent fractionation (MIF) in giant petrels.

ID	$\Delta^{199}\text{Hg}_{\text{initial}}^{\text{a}}$ ‰	Samples for $\Delta^{199}\text{Hg}_{\text{initial}}^{\text{b}}$	$\Delta^{199}\text{HgSe}^{\text{c}}$ ‰	Diff. in $\Delta^{199}\text{Hg}^{\text{d}}$ ‰	$\Delta^{201}\text{Hg}_{\text{initial}}^{\text{a}}$ ‰	Samples for $\Delta^{201}\text{Hg}_{\text{initial}}^{\text{b}}$	$\Delta^{201}\text{HgSe}^{\text{b}}$ ‰	Diff. in $\Delta^{201}\text{Hg}^{\text{d}}$ ‰	Diff. $\Delta^{199}\text{Hg} /$ $\text{Diff. } \Delta^{201}\text{Hg}$
P1	1.40 ± 0.01	$\Delta^{199}\text{Hg}_{\text{f}}$	1.30 ± 0.01	-0.10 ± 0.01	1.22 ± 0.01	$\Delta^{201}\text{Hg}_{\text{f}}$	1.16 ± 0.02	-0.06 ± 0.02	1.67
P2	1.66 ± 0.04	$\Delta^{199}\text{MeHg}_{\text{f}}$, $\Delta^{199}\text{MeHg}_{\text{k}}$, $\Delta^{199}\text{MeHg}_{\text{m}}$	1.42 ± 0.01	-0.24 ± 0.03	1.43 ± 0.07	$\Delta^{201}\text{MeHg}_{\text{f}}$, $\Delta^{201}\text{MeHg}_{\text{k}}$, $\Delta^{201}\text{MeHg}_{\text{m}}$	1.29 ± 0.01	-0.14 ± 0.05	1.71
P3	1.65 ± 0.02	$\Delta^{199}\text{Hg}_{\text{f}}$, $\Delta^{199}\text{MeHg}_{\text{l}}$, $\Delta^{199}\text{MeHg}_{\text{m}}$	1.43 ± 0.03	-0.22 ± 0.02	1.45 ± 0.03	$\Delta^{201}\text{Hg}_{\text{f}}$, $\Delta^{201}\text{MeHg}_{\text{l}}$, $\Delta^{201}\text{MeHg}_{\text{m}}$	1.25 ± 0.01	-0.20 ± 0.02	1.10
P4	1.77 ± 0.03	$\Delta^{199}\text{MeHg}_{\text{f}}$, $\Delta^{199}\text{MeHg}_{\text{m}}$	1.45 ± 0.03	-0.32 ± 0.03	1.49 ± 0.06	$\Delta^{201}\text{MeHg}_{\text{f}}$, $\Delta^{201}\text{MeHg}_{\text{m}}$	1.28 ± 0.01	-0.21 ± 0.04	1.52
P5	1.61 ± 0.01	$\Delta^{199}\text{MeHg}_{\text{f}}$	1.32 ± 0.02	-0.29 ± 0.02	1.38 ± 0.01	$\Delta^{201}\text{MeHg}_{\text{f}}$	1.23 ± 0.01	-0.15 ± 0.01	1.93

^aUncertainties are ± 1 SD of individual measurements or ± 1 SD of mean of multiple measurements.

^b $\Delta^{\text{XXX}}\text{Hg}_{\text{initial}}$ are values of total Hg digests of feathers, and $\Delta^{\text{XXX}}\text{MeHg}$ are values of the MeHg extracted from feathers or tissues. Subscripts identify tissue type (f, feather; k, kidneys; m, muscle; l, liver).

^cValues are of liver tissues that are predominantly HgSe ($f(\text{HgSe}) > 0.90$; see Table S2 and Figure 1 in main text).

^dThe difference between $\Delta^{199}\text{MeHg}$ and $\Delta^{199}\text{HgSe}$ or $\Delta^{201}\text{MeHg}$ and $\Delta^{201}\text{HgSe}$.

Table S5. Measured mercury isotopic values of certified reference materials (CRMs) and standards and references these data were compared against.

CRM	Fraction of Hg	$\delta^{199}\text{Hg}$ ($\pm 2\text{SD}$)	$\delta^{200}\text{Hg}$ ($\pm 2\text{SD}$)	$\delta^{201}\text{Hg}$ ($\pm 2\text{SD}$)	$\delta^{202}\text{Hg}$ ($\pm 2\text{SD}$)	$\delta^{204}\text{Hg}$ ($\pm 2\text{SD}$)	$\Delta^{199}\text{Hg}$ ($\pm 2\text{SD}$)	$\Delta^{200}\text{Hg}$ ($\pm 2\text{SD}$)	$\Delta^{201}\text{Hg}$ ($\pm 2\text{SD}$)	$\Delta^{204}\text{Hg}$ ($\pm 2\text{SD}$)	<i>n</i>	Ref.
DOLT-2	Total Hg	0.60 (0.09)	-0.22 (0.09)	0.25 (0.09)	-0.49 (0.08)	-0.72 (0.10)	0.73 (0.07)	0.03 (0.05)	0.62 (0.03)	0.01 (0.03)	3	Rosera et al. ⁸
DOLT-2	MeHg	0.98 (0.06)	0.10 (0.06)	0.91 (0.08)	0.12 (0.04)	0.11 (0.01)	0.95 (0.05)	0.04 (0.03)	0.82 (0.05)	-0.07 (0.06)	2	Rosera et al. ⁸
UM Almadén ^a	Total Hg	-0.15 (0.04)	-0.26 (0.05)	-0.45 (0.05)	-0.55 (0.06)	-0.83 (0.12)	-0.01 (0.04)	0.01 (0.04)	-0.03 (0.03)	0.01 (0.06)	14	NIST ⁹

^aSecondary Isotope Standard.

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