

Stable isotope analysis as a critical tool on the advancement of mercury and selenium interaction in biota: New insights on mercury demethylation mechanism

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Stable isotope analysis as a critical tool on the advancement of mercury and selenium interaction in biota **New insights on mercury demethylation mechanism**^[1]

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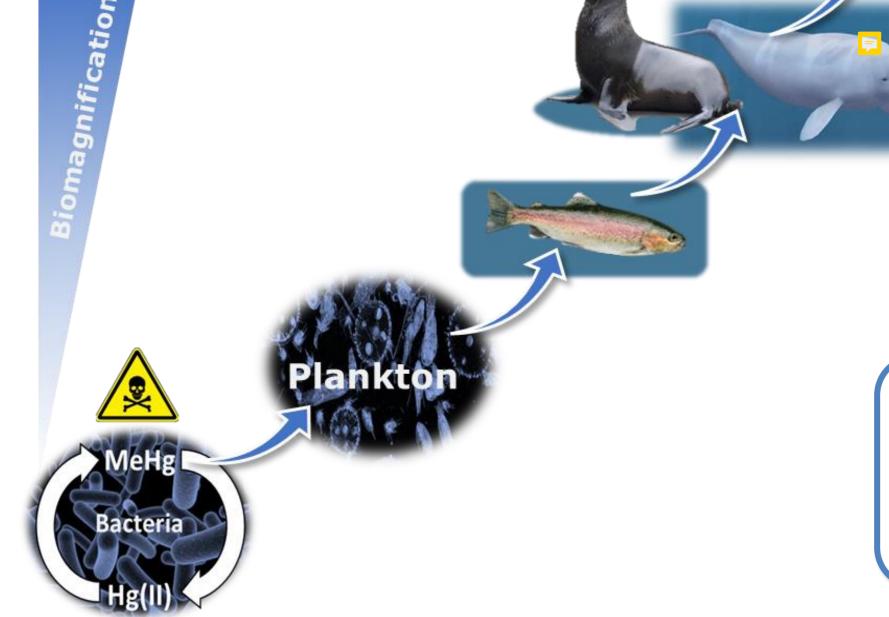
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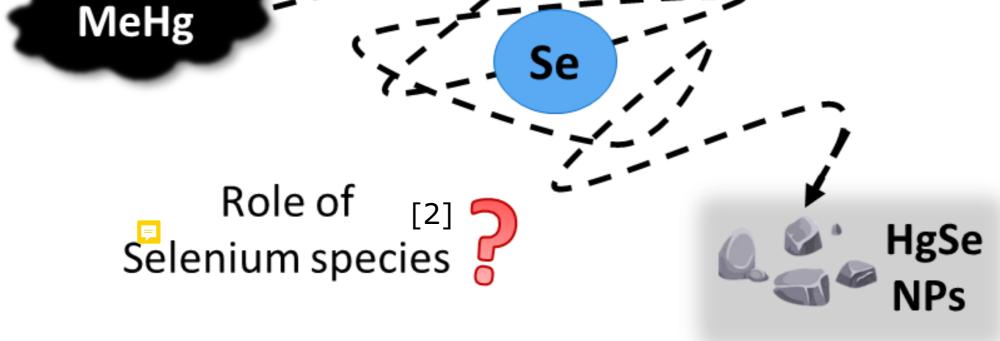
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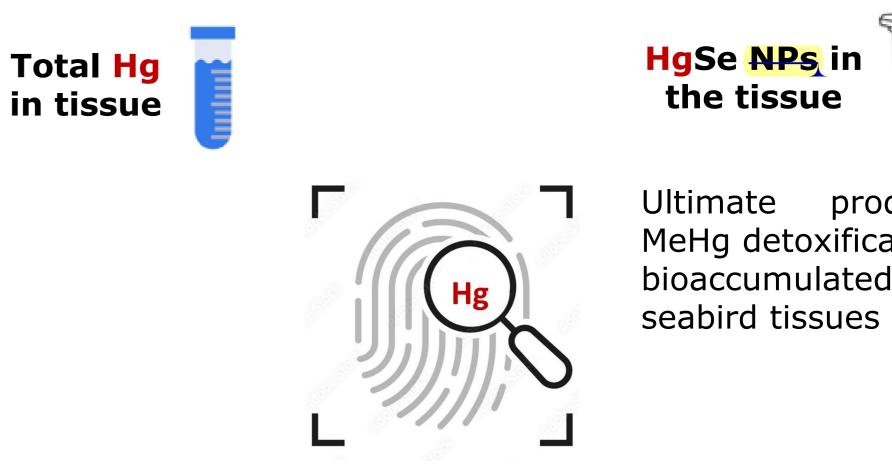
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Background Mercury (Hg) it is a globally distributed **Isotopic characterization** pollutant Hg fate and its detoxification mechanism Δ Its more neurotoxic chemical form methylmercury δ^{202} Hg (MDF) - insights into Hg internal transformations in biota remains unclear Seabird /:\ (MeHg) bioaccumulates and biomagnifies through the aquatic food chain Δ^{199} Hg (MIF) – insights into Hg food sources

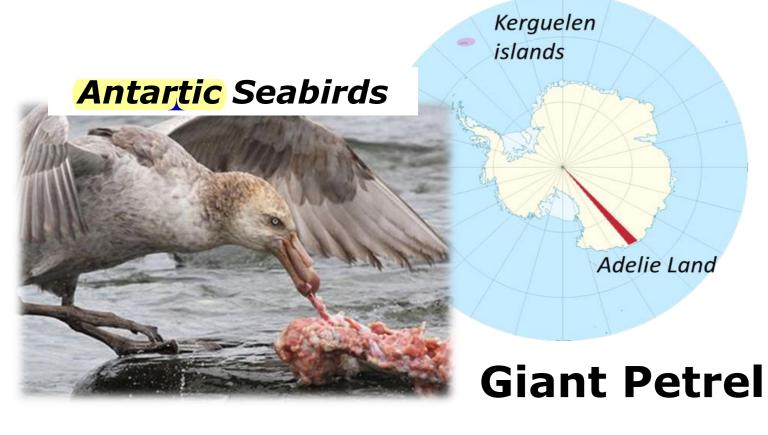




Simultaneous information about the (pollution) sources and the ل ج ک ک fate on living organisms of Hg can be obtained by investigating Hg isotopes mass dependent (MDF) and mass independent fractionation (MIF)

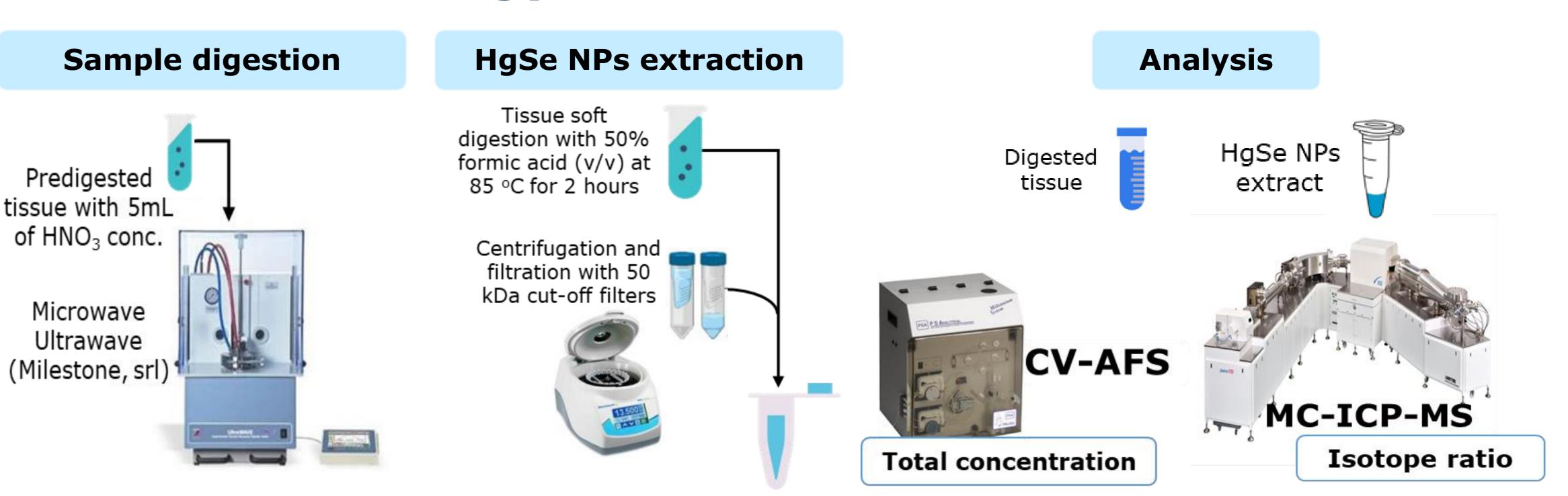


product of MeHg detoxification bioaccumulated in these

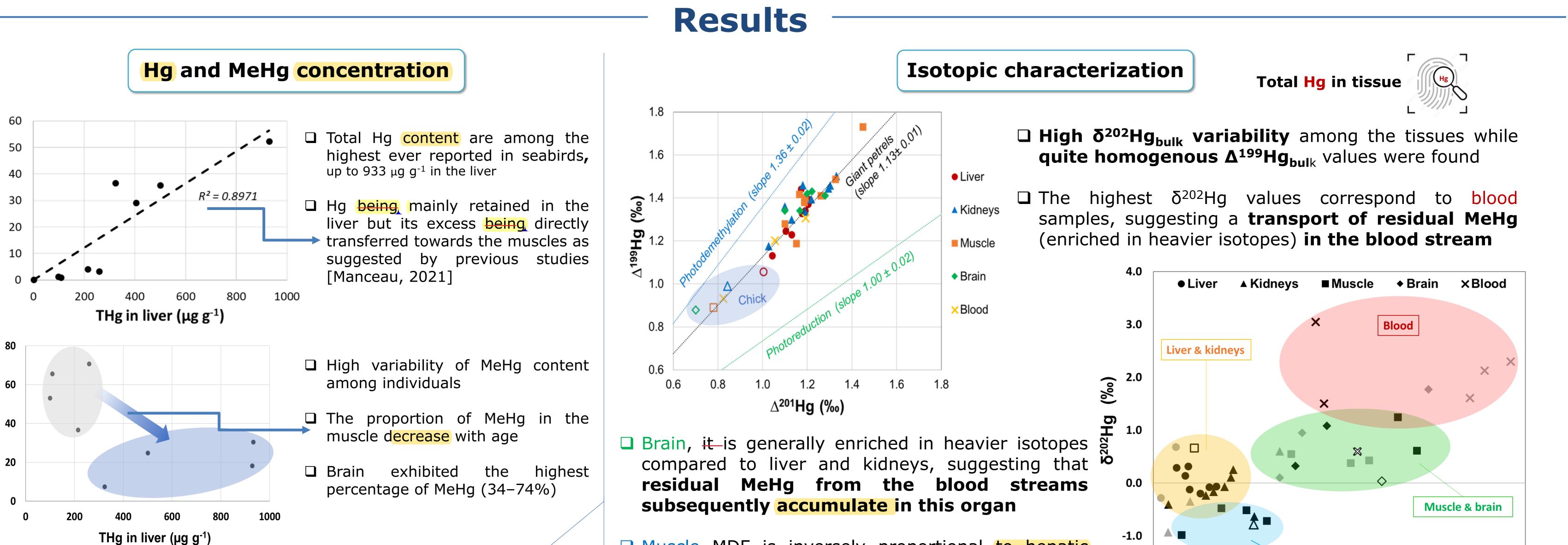


- **Top scavengers** of the Southern Ocean
- High Hg and Se bioaccumulation in tissues (up to 445 mg(Se) g^{-1} and **1 mg(Hg) g^{-1}**)

Methodology



• Samples from 5 key tissues of **11 dead individuals** (liver, kidneys, muscle, and brain) – 41 tissues





THg in muscle (µg g⁻¹)

(%)

MeHg

1. First time HgSe NPs species-specific (Hg) isotopic characterization (δ^{202} Hg, Δ^{199} Hg) in seabirds and the largest characterization of this compound in biota

2. No isotopic fractionation appears to be the HgSe NPs induced during biomineralization step from the precursordemethylated species

3. A shift of 2–3 ∞ δ 202Hg could be associated to the cleavage of the C-Hg bond. Because the difference btw Hg isotopic composition of MeHg (estimated) and demethylated species.

References

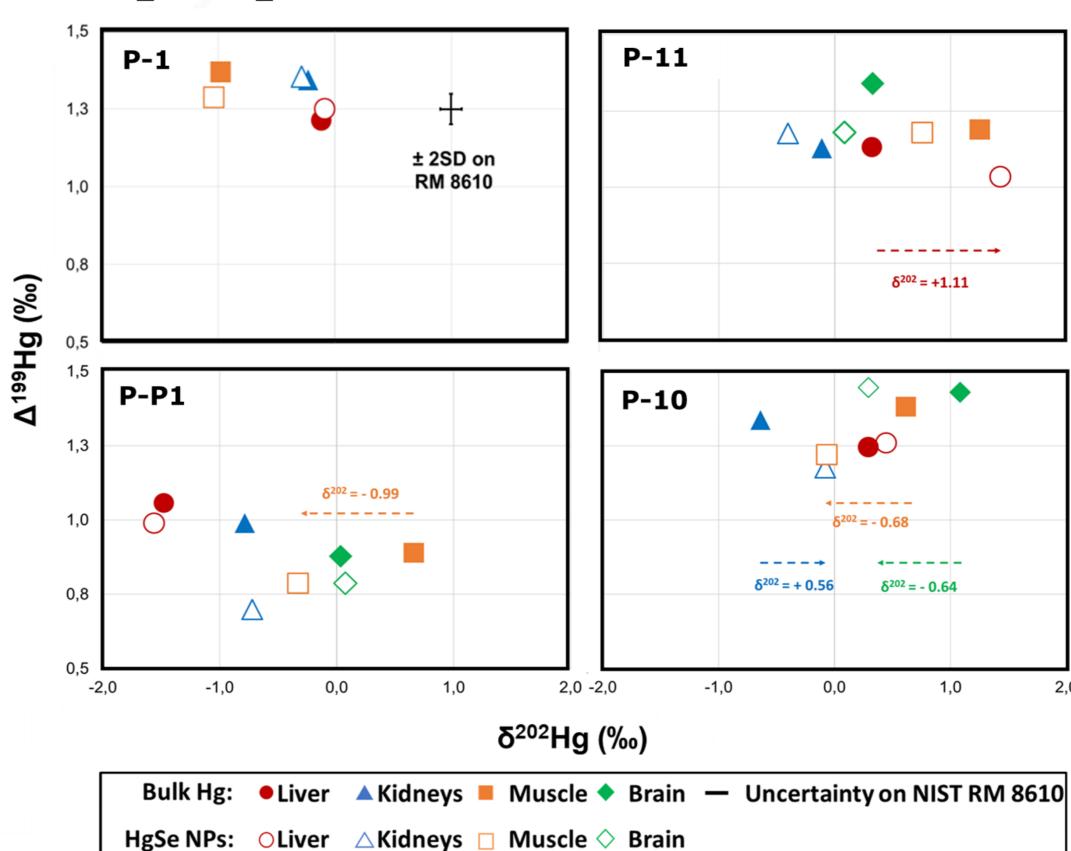
^[1] S. Queipo-Abad et. al., Journal of Hazardous Materials, 2022, 127922, 1-8.

^[2] K. El Hanafi et. al., Environmental Science & Technology, 2022, 5, 3288-3298.

^[3] A. Manceau et. al., Environmental Science & Technology, 2021, 55, 1515-1526.

□ Muscle MDF is inversely proportional to hepatic THg content. Hence, for older individuals, demethylation occurs effectively in this tissue





Muscle for individuals with high hepatic Hg concentration -2.0 80 20 60 100 MeHg (%)

□ HgSe NPs in liver (P-11) and kidneys (P-10) are enriched in heavier isotopes in the bulk, to which comparison are unexpected results considering HgSe NPs the final product of MeHg breakdown

□ Muscle and brain's HgSe NPs (P-10) are enriched in lighter isotopes compared with the bulk, in agreement with the higher fraction of Hg present as MeHg in those tissues

□ A match for both MDF and MIF, was observed in liver, kidneys, and muscle between $\delta^{202}Hg_{bulk}$ and $\delta^{202}Hg_{HqSe}$ of exhibiting the highest THg individuals hepatic concentrations (P-1), suggesting that **no isotopic fractionation is induced** during the biomineralization step from the precursor (demethylated) species.



Demethylation

(-) MDF

Hg:Se (1:4)^[3]

HgSe

MeHg

P-01 2.10 ‰

P-02 1.98 ‰

P-03 2.94 ‰

Estimated values

P-01 0.18 %

P-02 0.74 ‰

P-03 -0.32 %

Measured values