

### 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**<sup>[1]</sup>

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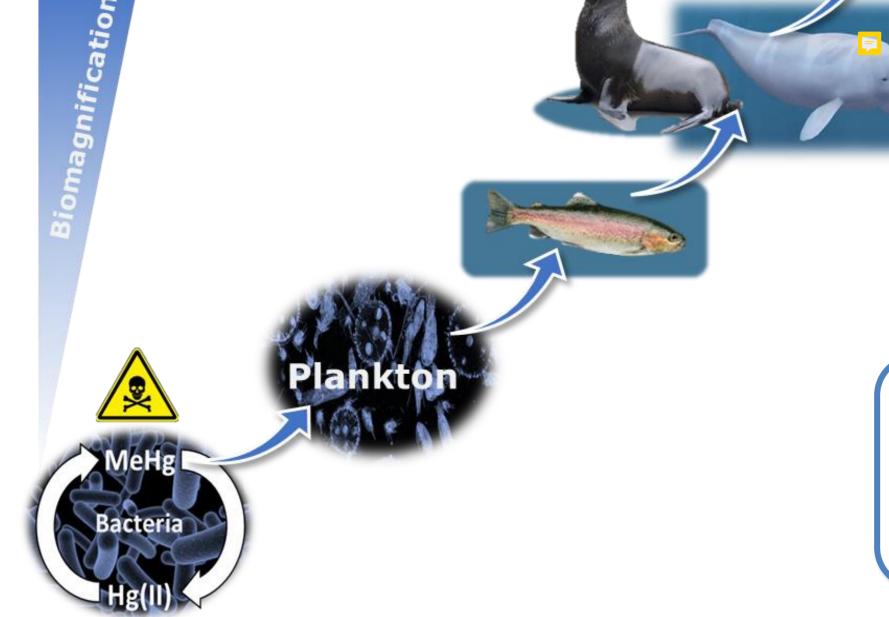
1-Université de Pau et des Pays de l'Adour, E2S UPPA, CNRS, IPREM, Institut des Sciences Analytiques et de Physico-chimie pour l'Environnement et les matériaux, Pau, France. 2-Advanced Isotopic Analysis Research Company, Pau, France.

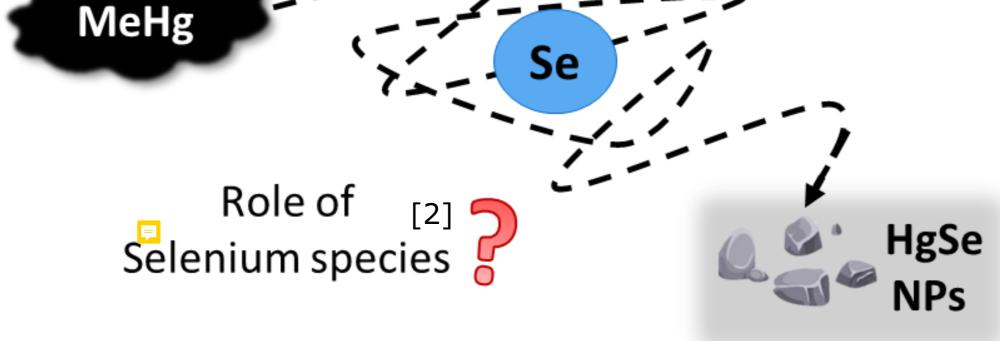
3-PS Analytical, Orpington, Kent, UK.

4-Centre d'Etudes Biologiques de Chizé, UMR 7372 CNRS - La Rochelle Université, Villiers-en-Bois, France.

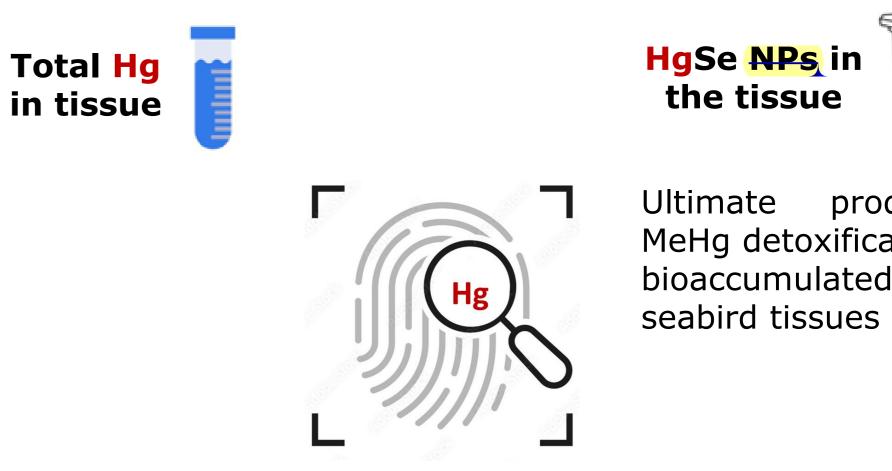
5-Institut Universitaire de France (IUF), Paris, France.

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

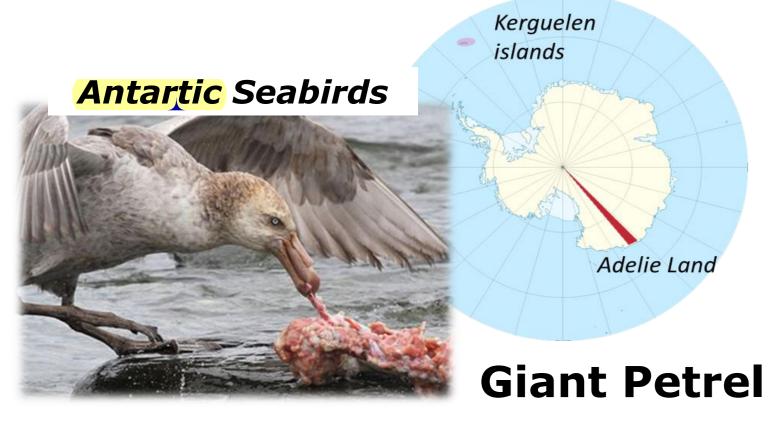




Simultaneous information about the (pollution) sources and the ل ج ک ک fate on living organisms of Hg can be obtained by investigating Hg isotopes mass dependent (MDF) and mass independent fractionation (MIF)

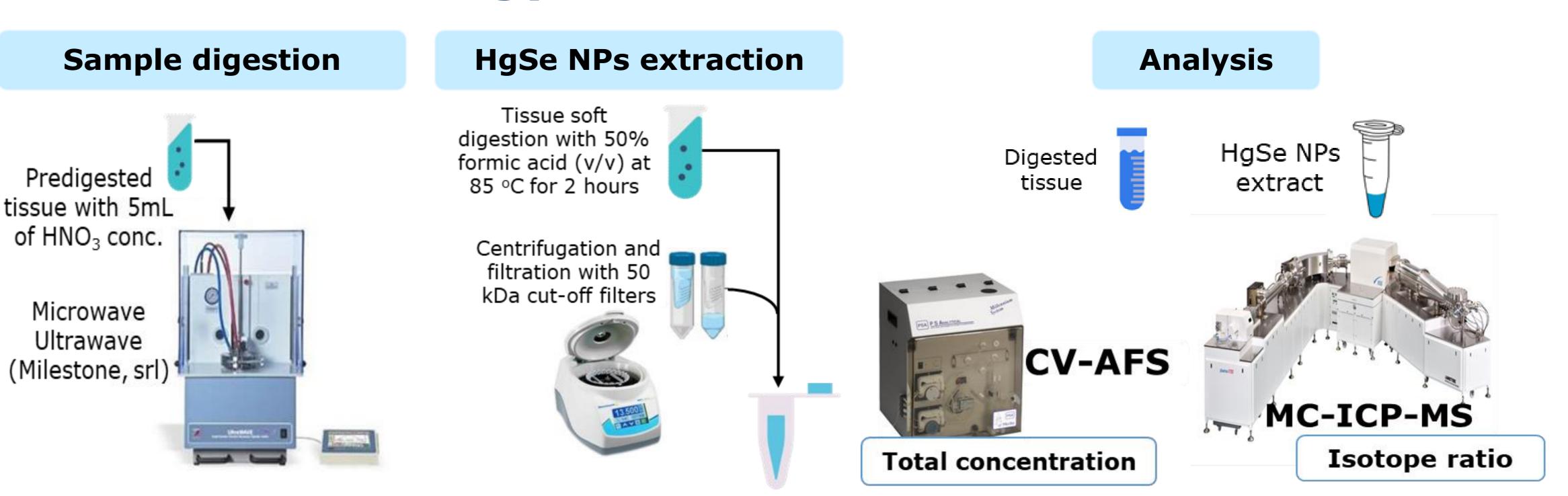


product of MeHg detoxification bioaccumulated in these

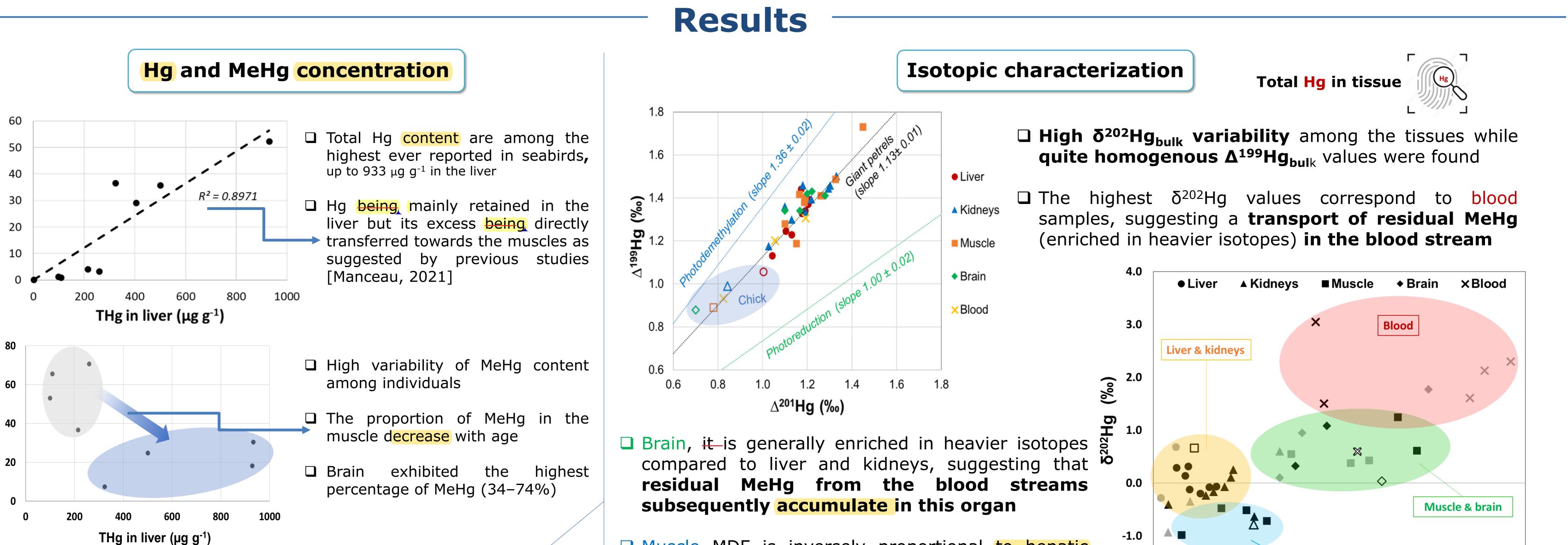


- **Top scavengers** of the Southern Ocean
- High Hg and Se bioaccumulation in tissues (up to 445 mg(Se)  $g^{-1}$  and **1 mg(Hg) g^{-1}**)

# Methodology



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





THg in muscle (µg g<sup>-1</sup>)

(%)

MeHg

1. First time HgSe NPs species-specific (Hg) isotopic characterization ( $\delta^{202}$ Hg,  $\Delta^{199}$ 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  $\infty$   $\delta$  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

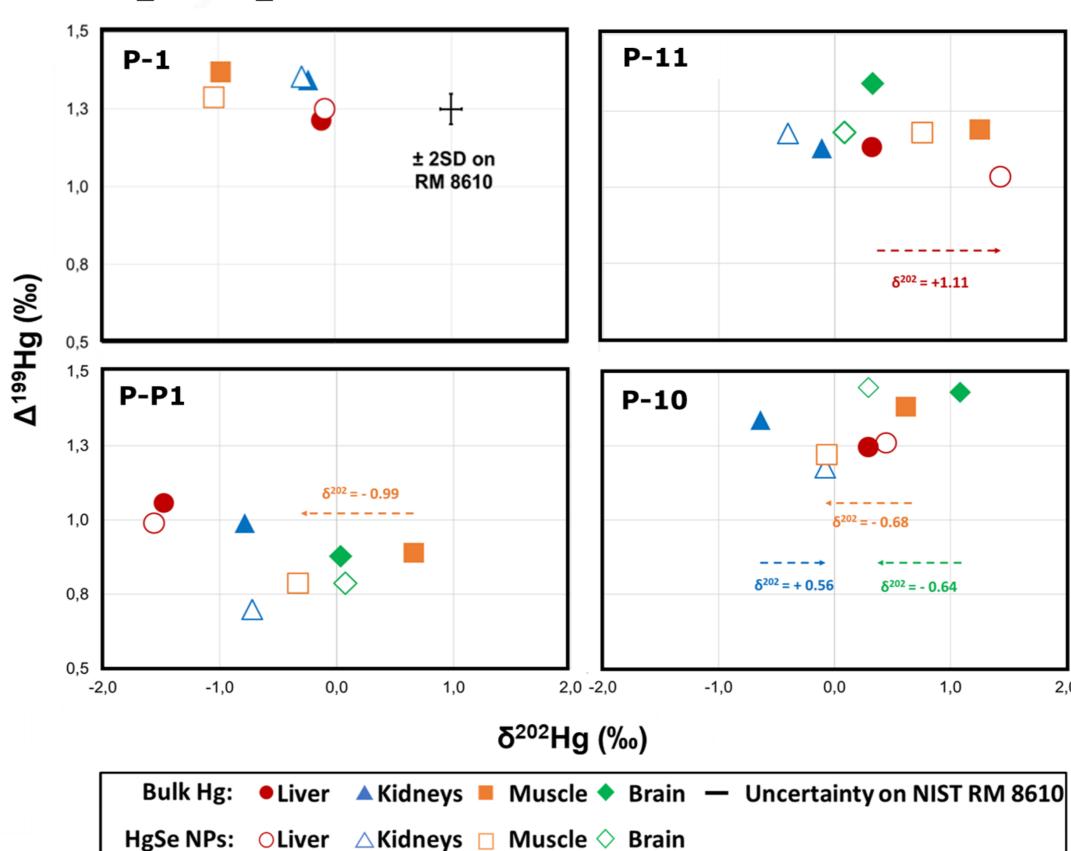
<sup>[1]</sup> S. Queipo-Abad et. al., Journal of Hazardous Materials, 2022, 127922, 1-8.

<sup>[2]</sup> K. El Hanafi et. al., Environmental Science & Technology, 2022, 5, 3288-3298.

<sup>[3]</sup> 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)<sup>[3]</sup>

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