

METAL CONCENTRATIONS IN THE INTESTINAL FRACTIONS PREPARED BY DIFFERENTIAL CENTRIFUGATION PROCEDURE

FINAL PROJECT MEETING

Integrated evaluation of aquatic organism responses to metal exposure: gene expression, bioavailability, toxicity and biomarker responses (BIOTOXMET) Zagreb, 17th February 2025

Sara Šariri

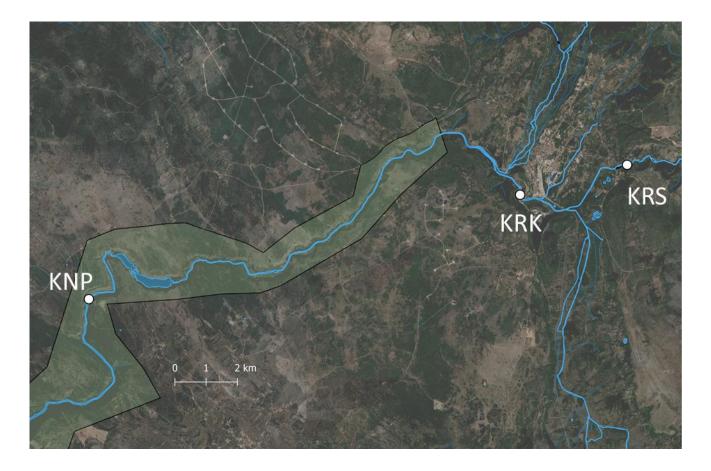
The subcellular distribution of metals

- only a fraction of the total metal concentration in an organism is bioavailable (metabolic processes, toxic effects) while the rest is detoxified
- within cells, metals are distributed across different compartments bioavailable fraction - metals bound to organelles and the cytosol
- the subcellular distribution of metals has been studied in the fish liver, gills, and gonads, while there are no studies on the intestine
- previous studies on the fish intestine have mostly reported total metal concentrations
- the aim of this study for the first time determine the distribution of metals among subcellular fractions of the fish intestine using differential centrifugation procedure

Methods

- sampling in autumn 2021
- Intestinal tissue of 4 individuals from each location





Subcellular fractionation

b

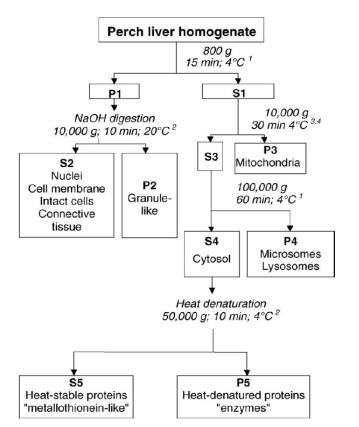


Fig. 1. Differential centrifugation method used to separate the various subcellular fractions. P: pellet, S: supernatant. (1) Durand et al. (1999), (2) Wallace et al. (2003), (3) Takeda and Shimizu (1982), and (4) Olsson and Haux (1986).

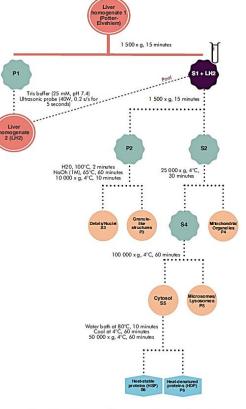


Fig. 2. Schematic illustration of the subcellular partitioning procedure used to separate Yellow Perch (a) gonads and (b) liver into operationally defined fractions. Small tubes represent the 50-µL homogenate fractions. Fractions represented by orange circles were tested with marker enzymes.

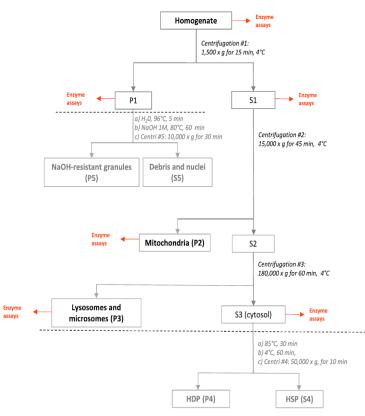
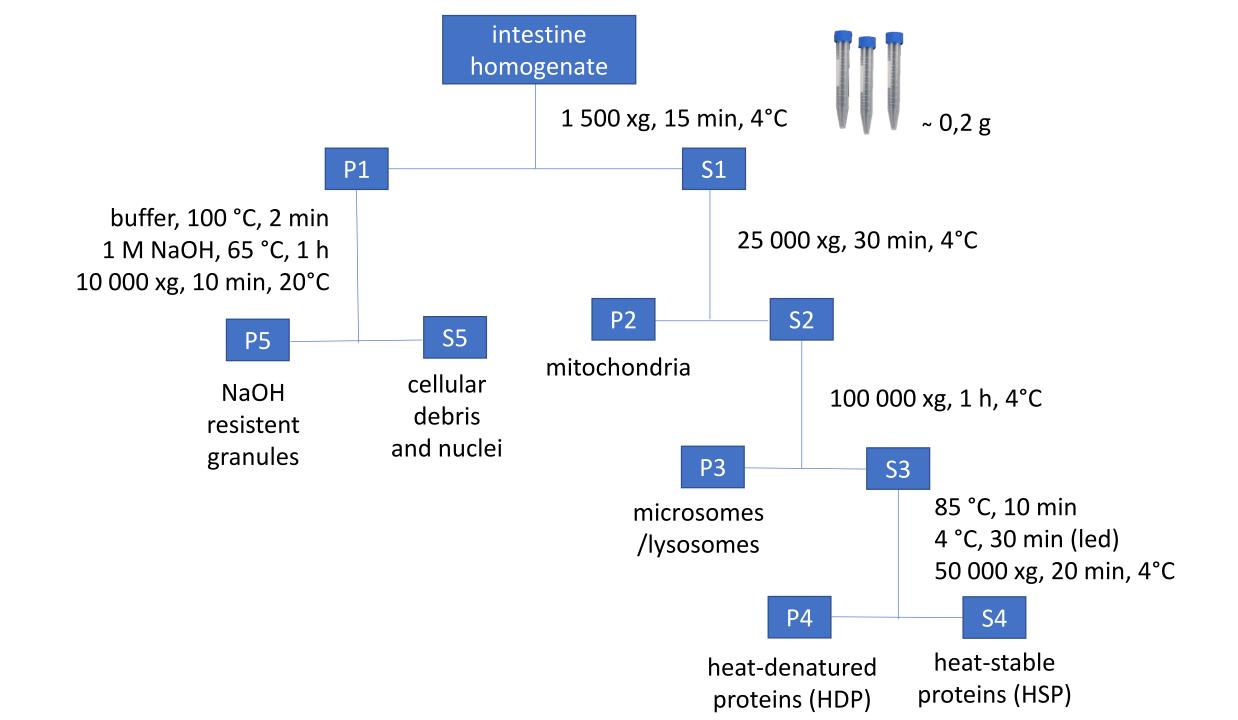


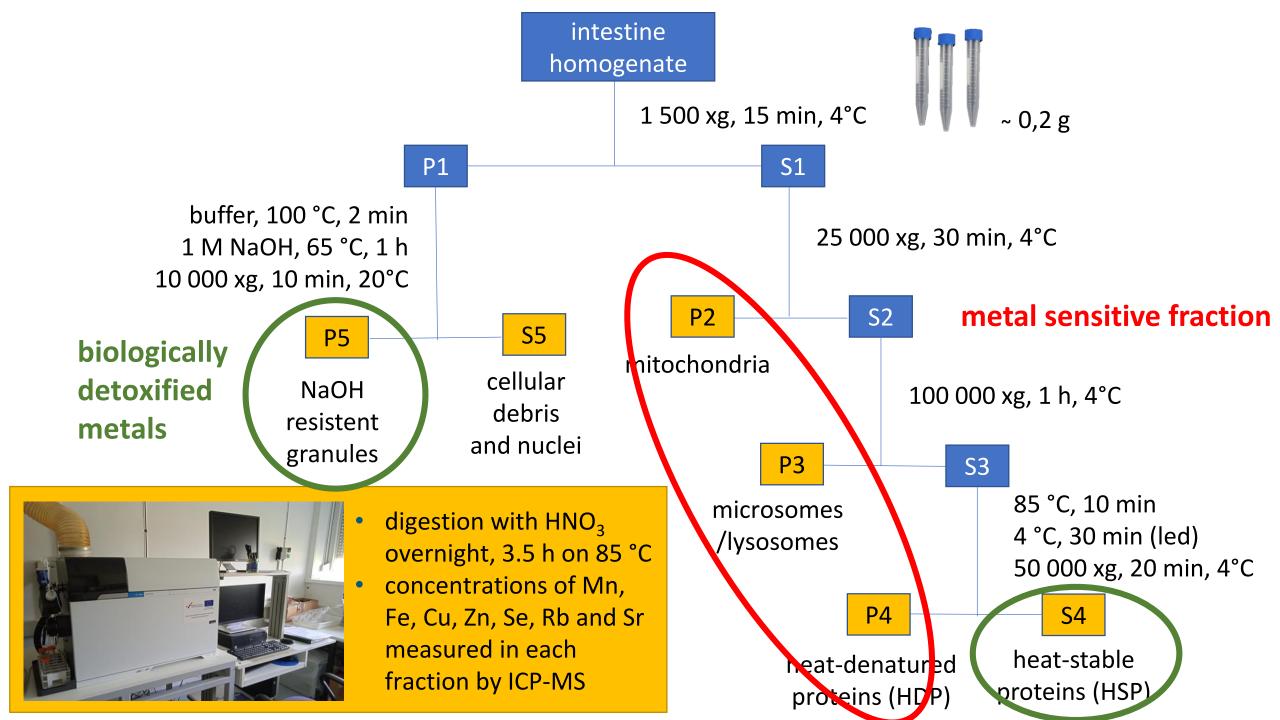
Fig. 1. Flowchart of the generic subcellular fractionation protocol for metal distribution studies. The letter P indicates pellet and S is for supernatant. The acronyms HSP and HDP mean, respectively, "heat-stable proteins" and "heat-denaturable proteins." The mention of "enzyme assays" indicates that marker enzymes can be measured in these fractions to validate the protocols. Fractions in gray and below the dotted lines were not subjected to enzyme assays as they had undergone heat treatments.

Giguere et al. 2006

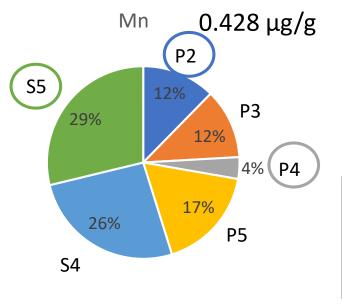
Khadra et al. 2019

Urien et al. 2020





Results







• P2 - mitochondria

• P3 - microsomes/lysosomes

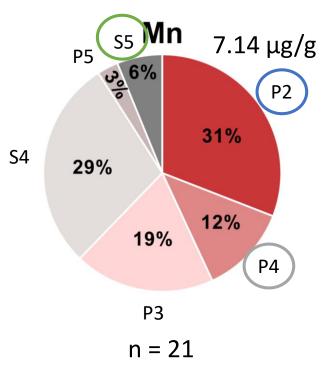
• P5 - NaOH resistent granules

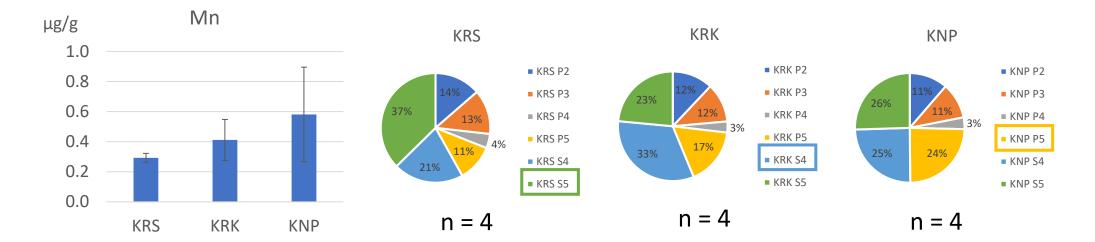
• S4 - heat-stable proteins (HSP)

• S5 - cellular debris and nuclei

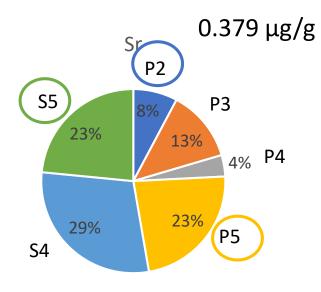
• P4 - heat-denatured proteins (HDP)

Desjardins et al. 2022 (yellow perch, liver)

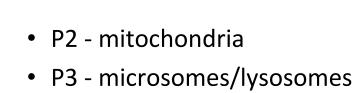




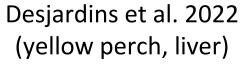
Sr

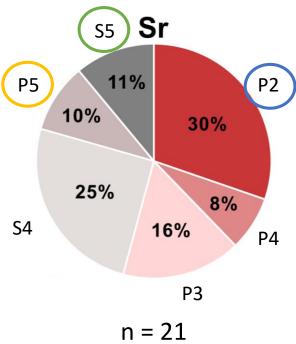


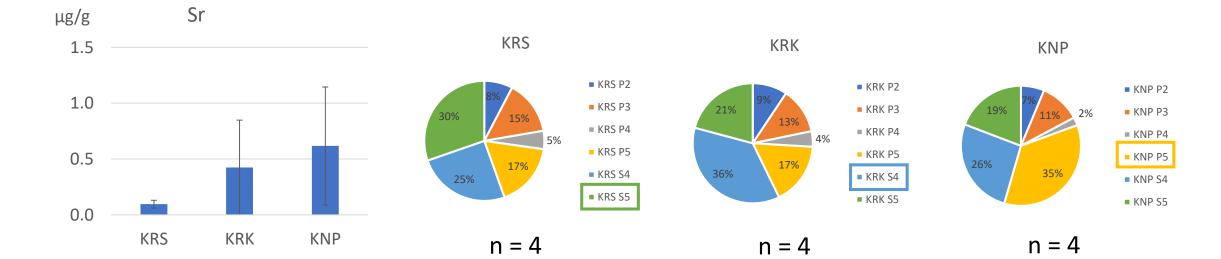
n = 12



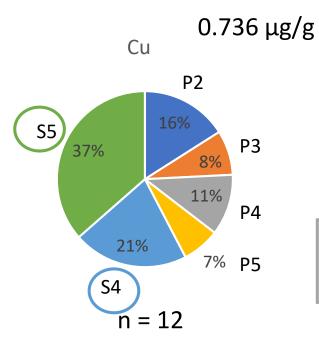
- P4 heat-denatured proteins (HDP)
- P5 NaOH resistent granules
- S4 heat-stable proteins (HSP)
- S5 cellular debris and nuclei

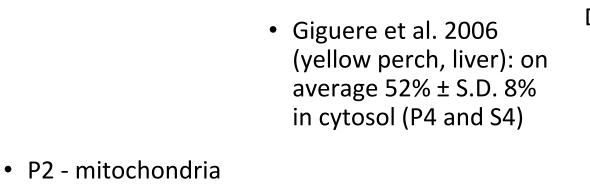




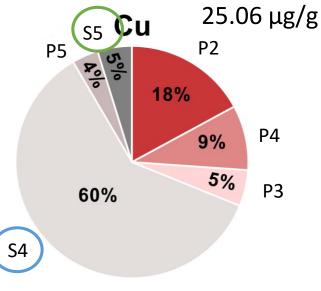




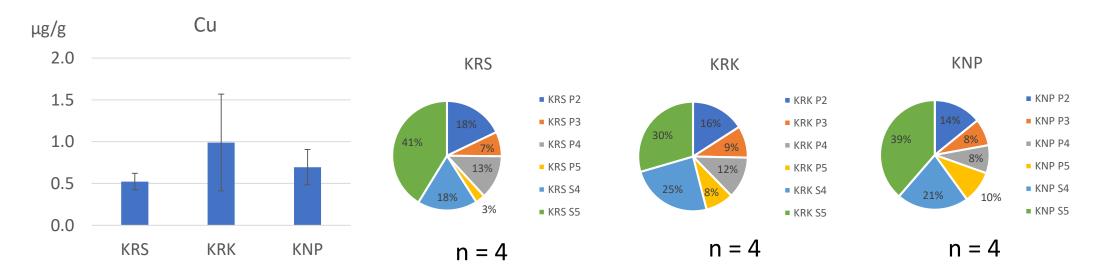




Desjardins et al. 2022 (yellow perch, liver)



n = 21



P3 - microsomes/lysosomes

• P5 - NaOH resistent granules

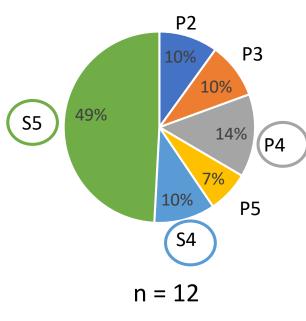
• S4 - heat-stable proteins (HSP)

S5 - cellular debris and nuclei

P4 - heat-denatured proteins (HDP)



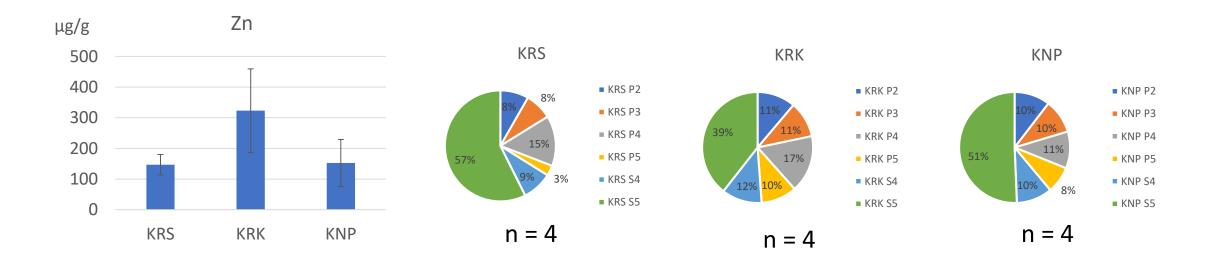
Zn 207.364 µg/g



- P2 mitochondria
- P3 microsomes/lysosomes
- P4 heat-denatured proteins (HDP)
- P5 NaOH resistent granules
- S4 heat-stable proteins (HSP)
- S5 cellular debris and nuclei

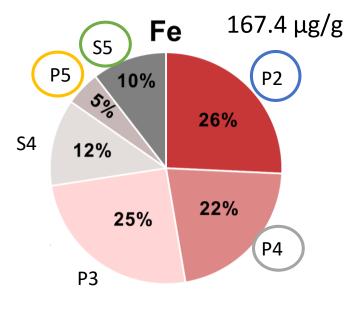
 Giguere et al. 2006 (yellow perch, liver): on average 51% ± S.D. 8% in cytosol (P4 and S4)

n = 48

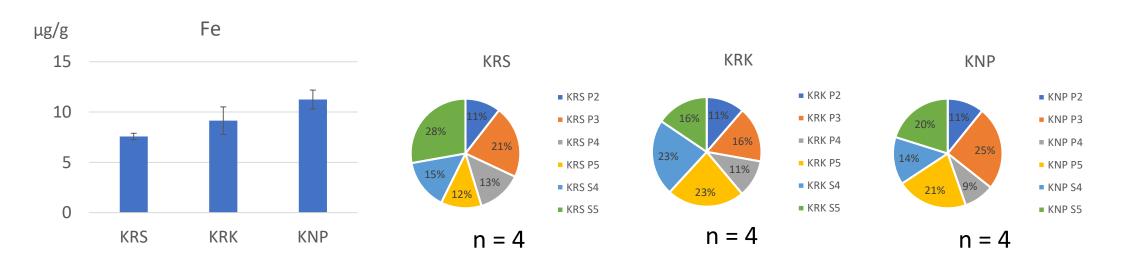


Fe

Desjardins et al. 2022 (yellow perch, liver)



n = 20



• P2 - mitochondria

• P3 - microsomes/lysosomes

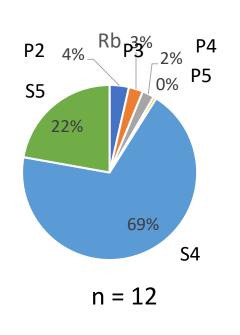
• P5 - NaOH resistent granules

• S4 - heat-stable proteins (HSP)

• S5 - cellular debris and nuclei

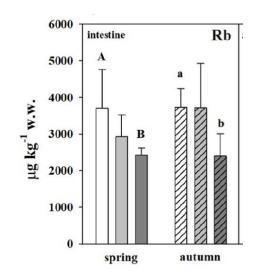
• P4 - heat-denatured proteins (HDP)

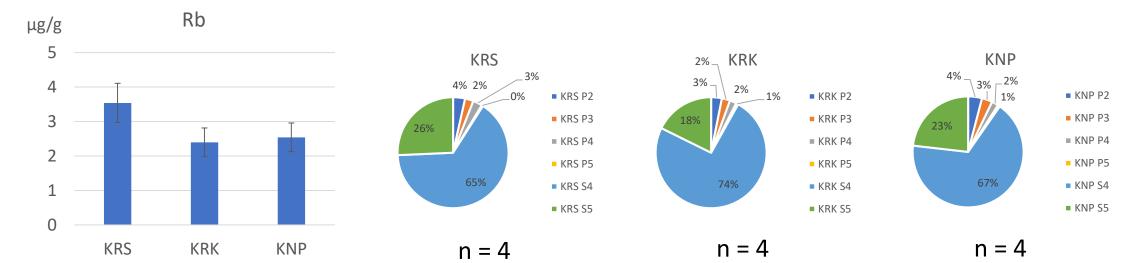
Rb



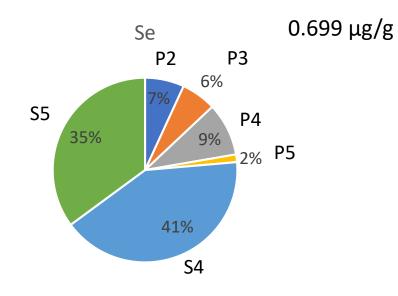
2.826 µg/g

- P2 mitochondria
- P3 microsomes/lysosomes
- P4 heat-denatured proteins (HDP)
- P5 NaOH resistent granules
- S4 heat-stable proteins (HSP)
- S5 cellular debris and nuclei



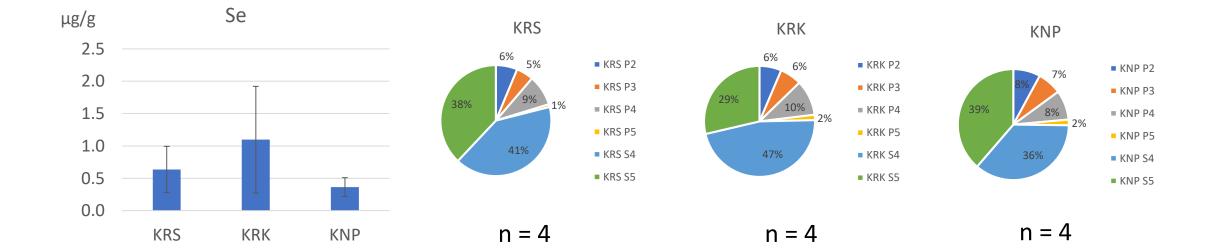


Se



n = 12

- P2 mitochondria
- P3 microsomes/lysosomes
- P4 heat-denatured proteins (HDP)
- P5 NaOH resistent granules
- S4 heat-stable proteins (HSP)
- S5 cellular debris and nuclei



Conclusions

- the first data for the intestinal tissue of brown trout
- no significant differences between sampling sites
- all elements higher percentages in S5 fraction (cellular debris and nuclei) compared to the literature data – tissue specificity, incomplete homogenization



Thank you for your attention!

 Croatian Science Foundation project: Integrated evaluation of aquatic organism responses to metal exposure: gene expression, bioavailability, toxicity and biomarker responses (BIOTOXMET) project number: IP-2020-02-8502