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Metal bioaccumulation and detoxification processes in cephalopods: a review

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Abstract

In recent decades, cephalopods have been shown to have very high capacities to accumulate most trace elements, regardless of whether they are essential (e.g., Cu and Zn) or non-essential (e.g., Ag and Cd). Among the different pathways of exposure to trace elements, the trophic pathway appears to be the major route of assimilation for numerous metals, including Cd, Co, Hg and Zn. Once assimilated, trace elements are distributed in the organism, accumulating in storage organs. The digestive gland is the main organ in which many trace elements accumulate, independent of the exposure pathway. For example, this organ can present Cd concentrations reaching hundreds to thousands of ppm for some species, even though this represents only a small proportion of the total mass of the animal. Such a specific organotropism towards the digestive gland, regardless of the exposure pathway, poses the question of the detoxification processes evolved by cephalopods in order to sustain high concentrations of both essential and non-essential trace elements. This paper reviews our current knowledge of trace element bioaccumulation in cephalopods, differences in pharmaco-dynamics between organs and tissues, and the detoxification processes used by cephalopods to counteract trace element toxicity. We have focussed on bioaccumulation within the digestive gland by investigating the subcellular locations of trace elements and their protein ligands.

1. Introduction

The increase in anthropogenic sources and emission of pollutants induces a strong contamination pressure on living organisms. Coastal and oceanic ecosystems are considered to be the final receptacle of released contaminants, especially in littoral areas that are close to wastewater discharges. Hence, persistent organic pollutants (POPs), oils (including Polycyclic Aromatic Hydrocarbons), trace elements, radionuclides, litter and debris (including plastics), and pharmaceuticals, found in most coastal areas worldwide, could result in deleterious effects, directly or indirectly affecting living resources (e.g. Williams, 1996). Among these pollutants, metallic trace elements and their potential impacts on marine biota have been extensively studied for more than 40 years (e.g. Islam and Tanaka, 2004). Nevertheless, contamination of shallow water is still a concern, as the major proportion of trace elements is trapped in coastal sediments where they can be continuously released into the water column. Moreover, the development of new metal compounds, such as nanoparticles (e.g. Baun et al., 2008), has contributed to a recent increase in the release of various trace elements into the environment (Salles et al., 2013).

Many studies have reported the trace element concentrations in marine organisms of economic interest, such as fish (McGeer et al., 2011), bivalves and gastropods (Naimo, 1995). Less data is available regarding cephalopods, despite these organisms being distributed and extensively targeted by fisheries worldwide, particularly during the past decades, in which cephalopod landings have increased concomitantly with a decrease in the abundance of small pelagic fish (Jereb et al., 2015). Their massive consumption has led to the assessment of chemical contaminants in edible species, in order to evaluate risks for human health (Nakahara et al., 1982; Rjeibi et al., 2015; Storelli, 2008; Storelli et al., 2005; Storelli et al., 2010). Since then, most research has focused on the analysis of trace element concentrations (Figure 1), and to a lesser extent, on that of organic compounds in cephalopods sampled in the field (Kawano et al., 1986; Son et al., 2015; Tanabe et al., 1984). The values recorded in cephalopod tissues have revealed the exceptional capacity of these organisms to accumulate and concentrate metals compared to other marine taxa (Figure 1), whatever their geographical origin, including contaminated (e.g., along the coast of industrialised countries) or pristine areas (e.g., in the southern Indian Ocean) (Bustamante et al., 2000; Kojadinovic et al., 2011; Martin and Flegal, 1975; Nakahara et al., 1979; Schipp and Hevert, 1978). Moreover, trace element concentrations also show large variation depending on the age, sex and maturation state of individuals, in addition to variation among species and those with different life-history traits, including diet or trophic level (Chouvelon et al., 2011; Miramand et al., 2006; Ueno et al., 2003; Won et al., 2010). Nevertheless, the high concentrations recorded in cephalopods, such as $1003 \mu\text{g}\cdot\text{g}^{-1}$ wet weight of Cd in the digestive gland of *Illex argentinus* (Dorneless et al., 2007), suggests that these organisms efficiently accumulate trace elements from the environment and retain them in their tissues. To the

best of our knowledge, the effects of such contamination on cephalopods remain poorly studied (Le Pabic et al., 2015a; Raimundo et al., 2010a; Semedo et al., 2012), even though the risks for human health appear to be limited when only the muscular parts of the organism are consumed (Rjeibi et al., 2015; Storelli, 2008; Storelli et al., 2006).

Overall, compared to bivalves, a limited number of studies have been devoted to understanding the physiological mechanisms involved in the high bioaccumulation and bioconcentration capacities for trace elements in cephalopods (Bustamante et al., 2006a; Bustamante et al., 1998b; Bustamante et al., 2002a; Costa et al., 2014; Miramand and Bentley, 1992; Miramand and Guary, 1981; Nakahara et al., 1979; Raimundo et al., 2010a, b; Raimundo et al., 2008; Rocca, 1969; Tanaka et al., 1983; Ueda et al., 1985). These studies have demonstrated strong organotropism of a number of trace elements towards the digestive gland (Ag, Cd, Cu, Zn and ^{210}Po) and the branchial hearts and their appendages (Co, Fe, V, ^{241}Am , ^{137}Cs and ^{237}Pu). These organs generally show the highest concentrations among the tissues and organs analysed. This specific organ distribution suggests that these organs play a key role in the detoxification and storage of trace elements (Bustamante et al., 1998a; Bustamante et al., 2002a; Finger and Smith, 1987; Martin and Flegal, 1975; Miramand and Bentley, 1992; Smith et al., 1984), however, our understanding of the underlying physiological mechanisms needs to be improved.

The present review focuses on the metabolism of trace elements in cephalopods, including the different bioaccumulation pathways, the distribution of metals at the tissue and subcellular levels, and the detoxification processes available to counteract the potentially toxic effects of the elevated concentrations recorded in cephalopod tissues.

2. Metal bioaccumulation processes

Explaining why such high concentrations of trace elements are observed in cephalopods requests to evaluate the sources of exposure, and the relative contribution of different uptake pathways in bioaccumulation. The environmental compartments in which trace elements are dispersed, entrapped or accumulated, such as the water column, sediment and prey, are all potential sources of contamination for cephalopods. Therefore, bioaccumulation results from the absorption or assimilation of bioavailable elements from the seawater and sediment (pore water), in addition to those ingested through the food. In addition, the high amount of metals recorded in tissues suggests efficient retention and/or poor elimination of accumulated trace elements.

Most cephalopod species are characterised by their “grow fast, die young” life cycle, represented by their exceptional growth rates, similar to those recorded in vertebrates (O'Dor and Webber, 1986). This trait requires a strict carnivorous regime, high prey consumption rate and efficient assimilation of

nutrients. Hence, assimilation of trace elements through ingestion of contaminated prey most likely constitutes a major source, contributing to the high trace element concentrations found in these organisms (Bustamante et al., 1998a; Martin and Flegal, 1975). As a first argument, the interspecific comparison of Cd concentrations in cephalopods sampled within the same area was higher in octopus species than pelagic species (Bustamante et al., 1998a; Lahaye et al., 2005). This difference could be attributed to the contrasting levels of Cd in benthic prey (crustaceans and bivalves), which provides the benthic cephalopods with a much higher Cd concentration than the fish that constitute the preferential prey of pelagic species (Cossa and Lassus, 1989) (Figure 1). In addition, studies have demonstrated that the retention of accumulated Cd in the cuttlefish *Sepia officinalis* differed depending on its uptake route. Thus, the biological half-life ($T_{b1/2}$) of dissolved ^{109}Cd is approximately 2 months, whereas dietary accumulated metal is retained in tissues for an indeterminate period (Bustamante et al., 2002b), making food the main source of Cd uptake. A similar contribution of the dietary pathway has been suggested in the loliginid squid *Sepioteuthis lessoniana* (Koyama et al., 2000).

Based on concentrations reported in cephalopods worldwide (Figure 2A and Table S2), ommastrephid squid displayed mean Cd levels as high as octopods, but extreme values ranging between 413–1003 $\mu\text{g.g}^{-1}$ wet weight were reported in the digestive gland (approximately 1000–2500 $\mu\text{g.g}^{-1}$ dry weight) of *Illex argentinus* (Dorneles et al., 2007; Gerpe et al., 2000). Again, the trophic regime of this oceanic ommastrephid could explain these high values, as this species consume Cd-rich pelagic crustaceans (amphipods and euphausiids), and display an important cannibalism or predation on other cephalopod species (Santos and Haimovici, 1997). In addition, the differences in bioaccumulation capacities compared to loliginids could result from their contrasting digestive physiologies, as discussed below.

Another trace element considered to be mainly accumulated through trophic pathways in marine ecosystems is Hg (for review: Fitzgerald et al., 2007). Studies in the literature have reported that Hg concentrations are positively correlated with the age and size of cephalopods (Barghigiani et al., 2000; Monteiro et al., 1992; Pierce et al., 2008; Rossi et al., 1993; Storelli and Marcotrigiano, 1999), showing its bioaccumulation over time. In the Bay of Biscay, the Hg concentrations in European squid *Loligo vulgaris*, veined squid *L. forbesi* and the common cuttlefish *Sepia officinalis* have been linked to changes in their trophic niche, as assessed by the stable isotope signature (Chouvelon et al., 2011). Hence, Hg levels in squid are significantly increased, concomitant to a drastic rise in $\delta^{15}\text{N}$, which is probably associated with a change in the preference of prey type, from crustaceans in young squid in favour of fish in larger individuals. However, the correlation of prey type with size is less evident in cuttlefish with a more opportunistic food regime that includes crustaceans (such as crabs) in adults (Blanc et al., 1998). These observations are also consistent with a study that demonstrated that the dietary pathway contributes to more than 70% of inorganic Hg bioaccumulation in the common cuttlefish (Lacoue-Labarthe et al., 2009b). Additionally, in fish of higher trophic levels than

crustaceans, Hg is mainly present as methylmercury (MeHg) (Bloom, 1992; Chauvelon et al., 2008). This organic form of Hg is thought to be very efficiently assimilated and retained by predators (Pierce et al., 2008), resulting in the biomagnification of Hg along food webs. These properties of MeHg also explain the higher Hg concentrations in adult cephalopods at higher trophic levels. Nevertheless, this association between contamination and age/size remains controversial in octopus (Barghigiani et al., 1991; Raimundo et al., 2004; Raimundo et al., 2010c; Seixas et al., 2005) and sub-adult and adult cuttlefish from the coast of Portugal (Raimundo et al., 2014), in which no relationship between Hg concentration and size has been found. Moreover, benthic cephalopods generally display significantly higher Hg concentrations than pelagic cephalopods (Bustamante et al., 2006b and Figure 2B). This suggests that processes other than trophic exposure contribute to Hg bioaccumulation in cephalopods. The transfer of methylated Hg in sediment and accumulation via the skin and gills of benthic and burying species could explain why these species generally display higher Hg concentrations than pelagic cephalopods. Certainly, the toxicokinetics of Hg bioaccumulation, considering both inorganic and organic forms and different sources (sediment, food or water), remains to be clarified, in order to understand differences in Hg levels observed in the field.

These observations underlie the potential predominance of this trophic source as the contamination pathway, however, experimental studies have highlighted that its relative contribution in metal uptake is trace element-dependent. As mentioned above, toxicokinetic data and application of biodynamic models to estimate the relative contribution of the different pathways of exposure (Landrum et al., 1992; Thomann, 1981) has allowed for determination of the major trophic pathways that contribute to Cd, Co, inorganic Hg and Zn accumulation in squid and cuttlefish fed with fish larvae and brine shrimp, respectively (Bustamante et al., 2002b; Koyama et al., 2000; Lacoue-Labarthe et al., 2009b). Nevertheless, the assimilation of trace elements could be affected by: (1) the prey type, as the proportion of accessible trace elements in the prey may vary (Pouil et al., 2016); (2) the ontogenic stage, as the digestive gland of juveniles is not fully functional compared to adults (Nixon and Mangold, 1998; Yim and Boucaud-Camou, 1980); and (3) environmental conditions, such as temperature or acidity, that may affect the physicochemical forms of trace elements and the metabolism of cephalopods. Each of these factors will, in turn, affect the assimilation efficiency of trace elements by individuals.

In contrast to the elements mainly accumulated from the food pathway, diet only appears to be a minor contributor for Ag compared to direct absorption of waterborne elements (Bustamante et al., 2004). Furthermore, the biological half-life of this element is relatively short in cephalopods. The consequence of the water pathway being the main contributor for Ag accumulation, combined with its elevated depuration rate, is that Ag concentrations in the common cuttlefish fluctuate according to environmental contamination conditions during its life cycle. The highest Ag concentrations have been reported to occur in juveniles occupying the coastal nursery grounds, whereas the lowest

concentrations were found in offshore overwintering individuals (Miramand et al., 2006). The tissues that are in direct contact with seawater, such as the gills and skin, are considered to be the main sites of metal absorption. Indeed, following exposure to dissolved ^{110m}Ag , the gills of *S. officinalis* contained 25% of the radiotracer content of the whole body, which decreased to 3% after 6 days in clean water conditions due to its depuration and transfer to other tissues, especially the digestive gland (Bustamante et al., 2004). At the same time, the fraction of Ag in the digestive gland increased from 5 to 61%, implying very efficient translocation of Ag from interface tissues (e.g., gills and epithelium) to this organ for detoxification and storage. Furthermore, it is noteworthy that the experimentally-determined biological Ag half-life does not exceed 2 weeks (Bustamante et al., 2004), suggesting the effective elimination of the metal. This corroborates the decreased Ag concentration recorded in wild animals when their exposure to dissolved metals is reduced during their offshore wintering migration (Miramand et al., 2006).

The dominance of the trophic pathway for the accumulation of trace elements, such as Cd or Zn, do not exclude the significant uptake of dissolved trace elements in seawater (Lacoue-Labarthe et al., 2011; Le Pabic et al., 2015b). Thus, an embryo surrounded by a protective capsule could accumulate dissolved metals that can pass through the eggshell during egg development (for review Lacoue-Labarthe et al., 2016), inducing high exposure to an environmental contaminant. Moreover, the bioaccumulation efficiency from this uptake route could be affected by environmental factors. The salinity, and also water pH, is known to vary in coastal areas, and are expected to decrease due to climate change (Borges and Frankignoulle, 1999; Caldeira and Wickett, 2005), which can influence chemical speciation, and therefore, the bioavailability of metals (often expressed as the free form fraction) (Millero et al., 2009; Stockdale et al., 2016). Moreover, changes to the metabolism of cephalopods induced by variations in temperature, salinity or hypercarbia affect their physiology (i.e., ionoregulation), and thus, the metal bioaccumulation capacity (Lacoue-Labarthe et al., 2012; Lacoue-Labarthe et al., 2009a; Lacoue-Labarthe et al., 2011).

The sediment pathway concerns only nectobenthic and benthic species, such as octopus, cuttlefish or sepiolids that display burying behaviours in sand or mud (Poirier et al., 2004). Nevertheless, experimental data has demonstrated that the contribution of contaminated sediment as a source of Cd, Co and Zn in cephalopod tissues is low (Bustamante et al., 2004; Bustamante et al., 2002b). However, as discussed above, the bioaccumulation of organic Hg, methylated by microorganisms on the seafloor, could significantly contribute to the contamination of cephalopod nectobenthic and benthic species (Bustamante et al., 2006b). To date, no quantitative data are available on this transfer pathway, preventing evaluation of its contribution to whole-body metal concentrations.

The final bioaccumulation pathway, which must be noted due to its potential for causing contamination during early life stages, is maternal transfer. By monitoring the Cu content of egg

compartments during development, Declair et al. (1970) showed progressive transfer of this metal from the yolk to the embryo, suggesting that the supply of this essential trace element for the developing organism is derived from an initial pool of Cu that is incorporated into the vitelline reserve. The hypothesis of maternal transfer of essential elements is also supported by the presence of higher concentrations of Cu, Fe, Mn and Zn compared to those of non-essential elements (e.g., Ag, Cd and Pb) recorded in the eggs of wild *S. officinalis* (Miramand et al., 2006; Rosa et al., 2015; Villanueva and Bustamante, 2006). The transfer of essential metals from the female to its offspring has been demonstrated experimentally using radiotracers, showing that ^{75}Se and ^{65}Zn initially assimilated from radiolabelled food by female common cuttlefish were subsequently transferred to the embryos via incorporation in the vitellus (Lacoue-Labarthe et al., 2008b). Interestingly, this maternal transfer is not restricted to metabolically essential trace elements, as $^{110\text{m}}\text{Ag}$ has been also found in cuttlefish eggs under the same conditions. In the freshly laid eggs, Ag was found to be equally distributed between the eggshell and vitellus, and was also detected in newly hatched juveniles (Bustamante et al., 2004; Lacoue-Labarthe et al., 2008a). Similarly, As is also likely to be maternally transferred, considering the high concentrations recorded in both the gonads of females (Bustamante et al., 2008; Kojadinovic et al., 2011) and in hatchlings compared to the concentration recorded in juveniles (Le Pabic et al., 2015b). However, to the best of our knowledge, this transfer has not been demonstrated experimentally.

Considering these pathways, the taxonomy, ontogenic stage, geographical area, life history strategy and traits, including the diet and trophic position, all appear to drive the concentration of metals recorded in cephalopods. A combination of both experimental and field studies needs to be considered to explain the importance of these different factors. The toxicokinetic approach (e.g., the use of radiotracers allowing at working at environmental concentrations) involves the use of biokinetic models in order to investigate bioaccumulation processes, including the pathways and effects of biotic and abiotic factors, using the same metal concentrations found in seawater. Determining the concentration of metals in wild specimens remains essential for validating these predictive models and assessing large-scale variations at the population or community levels. The high capacity of cephalopods to concentrate metals raises the question of the detoxification strategy evolved by these organisms to avoid negative impacts of contaminants on individuals and the population. This assumes highly efficient storage of metals by these organisms through unique mechanisms at the tissue and subcellular levels, in order to retain these trace elements in a non-metabolically-available form, thereby limiting their toxicity.

3. Tissue distribution

The digestive gland of cephalopods is a complex and polyvalent organ, performing various digestive and excretory functions (Boucaud-Camou and Yim, 1980; Boucher-Rodoni et al., 1987). It is also considered to be the major storage, and consequently, key detoxifying organ, as evidenced by numerous studies highlighting its ability to concentrate trace elements (Bustamante et al., 2003; Bustamante et al., 1998b; Bustamante et al., 2000; Declair et al., 1978; Miramand and Bentley, 1992; Raimundo et al., 2005; Smith et al., 1984; Ueda et al., 1985), except Hg (inorganic and organic) and As, for which the digestive gland contributes less to the total body burden (Bustamante et al., 2008; Bustamante et al., 2006b; Kojadinovic et al., 2011). Miramand and Bentley (1992) measured the concentration of 11 metals in the digestive gland, branchial hearts, kidney and cuttlebone of two nectobenthic species of cephalopods from the English Channel, the common cuttlefish *S. officinalis* and the curled octopus *Eledone cirrhosa*. Although the digestive gland represents only 6 to 10% of the whole animal tissue, it contains > 80% of the total body burden of Ag, Cd and Co, and 40 to 80% of the other measured elements. More recently, Raimundo et al. (2010b) also highlighted the key storage role of the digestive gland in *Octopus vulgaris* from the Portuguese coast. The concentrations of Cd, Cu, Pb and Zn were found to be three or four orders of magnitude higher in the digestive gland when compared to the gills, kidney and gonads. The unique organotropism of trace elements of this organ has raised the question of the role of the digestive gland in terms of the accumulation and detoxification of trace elements, which are stored in a non-toxic form. These observations are supported by evidence of trace elements transfer, monitored using gamma emitting radiotracers (e.g., ^{110m}Ag , ^{109}Cd , ^{57}Co and ^{65}Zn), from interface tissues (skin and gills) that are directly exposed to the waterborne contamination to the digestive gland in *S. officinalis* (Bustamante, 1998; Bustamante et al., 2004; Bustamante et al., 2002b). Nevertheless, this mechanism of trace element accumulation is not without adverse consequences, as DNA damage in *O. vulgaris*, such as broken strands, were found to be higher in the digestive gland compared to the other tissues analysed (Raimundo et al., 2010a), meaning that these potential detoxification mechanisms could be bypassed.

Branchial hearts are another predominant site for the storage of metallic elements such as Co (including ^{60}Co), Ni and V, and of transuranic elements such as ^{241}Am and $^{239-240}\text{Pu}$ (Guary and Fowler, 1982; Guary et al., 1981; Miramand and Bentley, 1992; Miramand and Guary, 1980, 1981; Nakahara et al., 1982; Ueda et al., 1985), even more than the digestive gland itself. This is particularly the case for Co, the concentration of which has been found to be about three-times higher in branchial hearts than in the digestive gland of *O. vulgaris* (Ueda et al., 1979). The branchial hearts were found to contain up to 50% of the whole-body radioactivity in *O. vulgaris* exposed to ^{60}Co , even though this organ only represents 0.2% of the total body weight of the octopus (Nakahara et al., 1982). Other trace elements, such as Cd and Cu, are also concentrated in the branchial hearts, but to a significantly less extent than the digestive gland (Miramand and Bentley, 1992). Bustamante et al. (2002) showed that Cd in the branchial heart was derived from waterborne Cd, whereas Cd from food was completely

contained within the digestive gland. The involvement of the branchial hearts in the hemocyanin metabolism of coleoid cephalopods is consistent with the presence of Cu within this tissue (Beuerlein et al., 1998; Mangold, 1989), Cu being the respiratory pigment of hemocyanin. Moreover, following exposure to dissolved ^{109}Cd and ^{65}Zn radiotracers, the branchial hearts and appendages of *S. officinalis* exhibit high Cd levels following its transfer from the skin, gills and muscles (Bustamante et al., 2002). Considering the short biological half-life of dissolved Cd accumulated in the whole organism (approximately 2 months), and the physiological circulatory and excretory functions of the haemolymph, the waterborne-derived Cd transported by the blood appears to be excreted through the branchial hearts and appendages. A similar mechanism of transfer was also observed for ^{60}Co in *O. vulgaris* (Nakahara et al., 1979).

While the highest Hg concentrations have been recorded in the digestive gland, Hg was found to be mainly stored in the muscle in terms of body burden distribution. Bustamante et al. (2006b) reported that 70 to 90% of the total Hg body burden was stored in the muscular tissue of different cephalopod species collected from the North-east Atlantic Ocean. More than 80% of dissolved inorganic ^{203}Hg accumulated in juvenile cuttlefish was also found to be stored in muscle (Lacoue-Labarthe et al., 2009b). This unique distribution could be firstly explained by the fact that these tissues represent more than 70% of the total body weight. Secondly, in the digestive gland, the main fraction of Hg is the inorganic form, in contrast to the large proportion of organic mercury (MeHg) found in muscle tissue. This suggests that demethylation of organic Hg could occur in this organ, and/or that MeHg is translocated from the digestive gland to muscle due to its high affinity for the sulphhydryl groups of muscular proteins rather than fat tissue (Bloom, 1992; Bustamante et al., 2006b). However, the muscle storage capacity appears to be limited in most contaminated octopuses collected along the Portuguese coast, in which Hg and MeHg concentrations reached a threshold, whereas Hg concentrations were increased in the digestive gland (Raimundo et al., 2010c).

In summary, trace elements are found to be highly concentrated in specific cephalopod tissues. Although the toxicity of contaminants in cephalopods is relatively unknown, the absence of acute effects caused by high metal concentrations, as reported previously, imply that biochemical processes occur at both the tissue and cellular levels to store these elements in non-chemically active forms, thereby limiting their toxic interactions with molecular and cellular compounds, or to a lesser extent, allowing their elimination.

4. Mechanisms of trace element detoxification at the cellular level

4.1. Metal-rich deposits in branchial hearts

As mentioned above, metals such as Co, Cu, V and transuranic elements are abundantly or mainly located in the branchial hearts. This tissue is mainly composed of ovoid cells, also called rhogocytes or

polyhedral cells, hemocytes derived from circulating blood, and the adhesive hemocytes in the wall of the branchial hearts. Rhogocytes are involved in haemolymph detoxification by eliminating endogenous or exogenous substances, such as ferritin, colloidal gold or bacteria (Beuerlein et al., 2002; Beuerlein and Schipp, 1998). These rhogocytes can contain high amounts of numerous pigmented granules composed of metabolic intracellular waste and Fe (III). These structures, called adenochroms (Miramand and Guary, 1981), could be considered to be natural bonders for different metals such as Co or V and transuranic elements, and are believed to participate in their detoxification (Figure 3). Finally, rhogocytes are thought to be involved in hemocyanin metabolism (Beuerlein et al., 2004; Ruth et al., 1999), explaining the elevated Cu levels in branchial hearts.

4.2. Metal-rich deposits in digestive gland cells

4.2.1. Histological insights

Several studies have attempted to unravel the physiological mechanisms involved in the storage capacity of the digestive gland, especially through the use of histological investigations.

Histological observations of the digestive gland structure of cuttlefish revealed that it is composed of numerous tubules with a clearly-defined lumen (Boucaud-Camou and Yim, 1980). As first described by Cuénot (1907), three different cell types make up the digestive gland, including basal cells, digestive cells and excretory cells (Figure 3). Basal cells, with their typical triangular shape, are situated between the basal lamina of tubules and the digestive cells. The numerous digestive cells, derived from the differentiation of basal cells, are involved in intracellular digestion, containing vacuoles that are generally coloured pale yellow to light brown. These vacuoles were originally named “boules” by Cuénot (1907), and probably contain residual peptides and lipids from intracellular digestion (Boucaud-Camou and Yim, 1980). The content of these vacuoles is excreted into the lumen of the digestive tubules, suggesting that these digestive cells potentially play a role in the excretion of digestive waste as well as contaminants. These digestive cells can differentiate into excretory cells, which are more rarely observed according to Costa et al. (2014), located close to the apical lamina, and mainly composed of a single large vacuole. These excretory cells are thought to play an important role in the excretion of salts.

Histological analyses by Martoja and Marcaillou (1993) showed the presence of metal-rich spherules in the basal cells of the digestive gland of *S. officinalis*. In these spherules, they noted the presence of Cu and S in particularly high amounts, and other numerous elements in the descending order: Zn > Ag and Ca > Mg > Sr. Based on several indirect analyses, these authors hypothesised that these spherules contain metallothionein-like proteins (MTLP; cysteine-rich proteins to which several metals have a high affinity) as a trace element ligand embedded within a polymer matrix. Surprisingly they were

unable to find an association between Cd, well-known for its high affinity to MTLP, and such spherules, calling into question their composition and role in trapping metals.

More recently, Costa et al. (2014) provided a more detailed structural and chemical description of the basal cells in the digestive gland, strengthening previous results from Martoja and Marcaillou (1993) in the same species. Thus, calcic spherules rich in Cu and Pb (the latter in a dichromate form) were identified in the basal cells. They were found to be composed of a complex organic protein matrix, originating from an extension of the rough endoplasmic reticulum, also suggested to contain MTLP. Therefore, it is suggested that trace elements initially bound to soluble proteins were trapped in solid proteo-mineral structures. Therefore, the basal cells containing these spherules are likely to be involved in trace element detoxification, allowing their storage in a non-chemically-active form. It is noteworthy that these typical metal-rich structures have not been consistently observed in the digestive gland of cephalopods. Neither Boucaud-Camou and Boucher-Rodoni (1983) nor Bustamante (1998) observed these granule-like structures in the digestive gland cells of *S. officinalis* or *Todarodes sagittatus*, respectively. However, the variety of histochemical protocols utilised could explain these contrasting observations, such as the use of unsuitable fixatives, as suggested by Martoja and Marcaillou (1993).

Another plausible role for the basal cells in the maintenance of Cu homeostasis has been proposed. Martoja and Marcaillou (1993) highlighted structural similarities in shape and cytological properties between the basal cells (previously called “replacement” or “pyramid” cells) of the digestive glands of *S. officinalis* and *Nautilus pompilius*, involved in hemocyanin synthesis (Ruth et al., 1999; Ruth et al., 1988). This particular triangular shape is found in the digestive gland cells of both species, and the rough endoplasmic reticulum is also highly developed in both cases, and extended in these spherule shapes. Thus, a particular function of this special cell type in Cu regulation cannot be excluded, considering the large amount of Cu found in the spherules.

Finally, the digestive cells contain two other structures which can participate in trace element detoxification: (1) the “boules” structures, corresponding to typical vacuoles (e.g., heterolysosomes), probably involved in intracellular digestion and enzyme secretion (Boucaud-Camou, 1972); and (2) the brown bodies, which are large vacuoles containing cellular debris, thought to have an excretory role, previously observed to contain Fe deposits in *S. officinalis* (Costa et al., 2014). As lysosomes are known to accumulate trace elements from the cytosol, considered to be a method of their elimination in invertebrates (Dallinger, 1993; Marigomez et al., 1996; Moore, 1990), a similar role for the “boules” of digestive cells can be expected. The subcellular distribution of trace elements in the digestive gland of *S. officinalis* showed that, among others, 44, 30 and 31% of accumulated Ag, Fe and Zn, respectively, were located within the lysosomal fraction (Tanaka et al., 1983). Similarly, Bustamante et al. (2006a) noticed that more than 75% of the Ag burden was in the lysosome and

mitochondrial fractions. Additionally, Raimundo et al. (2008) showed a positive correlation between metals contained in lysosomes in *O. vulgaris* and the total quantity of metals found in the whole digestive gland. Interestingly, the structure of the digestive gland in Loliginidae is quite different from the other coleoid cephalopods (at least from cuttlefish and ommastrephid squid), as their lysosomal system is comparatively less developed, also lacking the “boules” structures (Boucaud-Camou and Boucher-Rodoni, 1983). Thus, it is expected that such characteristics could limit the metal detoxification/storage performance in Loliginidae, considering that they often exhibit a lower metal concentration compared to other cephalopods from the same area. For instance, Bustamante et al. (2002a) related the lowest Cd bioaccumulation capacity of loliginid squid (Figure 2A) to the absence of boules.

4.2.2. Subcellular partitioning

Overall, only few studies have investigated the subcellular distribution of metals in the digestive gland (Bustamante et al., 2006a; Bustamante et al., 2002a; Craig and Overnell, 2003; Finger and Smith, 1987; Raimundo et al., 2008, 2010d; Rocca, 1969; Tanaka et al., 1983), most of which have considered only the insoluble (including organelles, nuclei and cellular debris) and the soluble (cytosol) subcellular fractions, separated by successive centrifugation steps. Only few metals, Cd, Cu and Zn, have been extensively studied in these works, with Ag investigated to a lesser extent, for which the subcellular distribution varies considerably between species.

In the digestive gland of the common cuttlefish *S. officinalis*, Cd, Co and Cu are mostly found in the cytosol, meaning that these metals are associated with soluble proteins, whereas Ag, Fe, Mn, Pb and Zn are mainly found in the organelle fraction, suggesting that these metals could be bound to or sequestered within cell structures (Bustamante et al., 2006a). These results are consistent with observations in the digestive gland of other cephalopod species, at least with respect to Cd, which has been shown to be mainly (50 to 96%) distributed in the cytosolic fraction of the digestive gland cells of *E. cirrhosa*, *S. officinalis*, *S. orbignyana*, *S. elegans*, *Illex coindetti*, *N. gouldi*, *T. sagittatus*, *O. vulgaris* and *L. vulgaris*, in ascending order (Bustamante et al., 2002a; Finger and Smith, 1987; Raimundo et al., 2008, 2010d). In contrast to this, Cd was mainly found to be associated with the insoluble fraction, composed of organelles, nuclei and cell debris, in the Japanese squid *T. pacificus* (Tanaka et al., 1983). This might be explained by: (1) varying contributions of the trace element uptake pathways (waterborne, sediment and trophic routes) among species, and/or (2) differences in the trace element concentrations in the digestive gland (Bustamante et al., 2006a; Bustamante et al., 2002a). According to a study on *O. ocellatus*, differences in the contribution of exposure routes does not appear to have an effect on subcellular metal distribution (Koyagani et al., 1982). However, Bustamante et al. (2002a) showed that when the Cd concentration in the digestive gland of

cephalopods increased, this metal was continuously transferred from the soluble to the insoluble compartment. Additionally, the subcellular partitioning of Cd investigated in seven cephalopod species displayed a large range of total Cd levels in the digestive gland. The results highlighted a negative correlation between the percentage of Cd found in the soluble fraction and the Cd concentration. This suggests that a concentration threshold probably exists, as Cd is mostly accumulated within insoluble parts of digestive gland cells. The literature reports the precipitation of trace elements into insoluble granules in various invertebrates as a common trace element storage and detoxification mechanism (e.g. George et al., 1982). Thus, the subcellular distribution strengthened the hypothesised role of the basal cell spherules, described above, in metal detoxification. From these observations, it is difficult to find one common mechanism that can be applied to every cephalopod species. Additionally, the observed subcellular distribution cannot be explained by the biological role of specific trace elements, as both essential and non-essential elements are found in the soluble and insoluble fractions in varying proportions. However, it is noteworthy that subcellular fractioning is an accurate procedure from an experimental point of view, as it requires a high degree of precision to ensure that the fraction obtained corresponds to the fraction required. Consequently, it is possible to observe an overlap between several fractions, and their identification using techniques such as enzyme activity measurements or ultrastructural observations, is not always performed.

The relationship between non-essential trace element concentrations in the digestive gland and those found in organelles are also of concern for the toxicological risk to the organism. Regarding Cd and Pb levels in the digestive gland of wild *O. vulgaris*, increased metal concentrations are associated with enhanced Cd and Pb concentrations within the mitochondria and nuclei, respectively (Raimundo et al., 2008). The binding of these metals to key cell structures raises the question of the potential toxic effects of these elements on cellular respiration and nucleus function. Indeed, in the same species, elevated DNA damage has been observed in the digestive gland compared to other tissues (Raimundo et al., 2010a). Nevertheless, DNA damage can be induced by genotoxicants other than trace elements, also known to show high accumulation in this organ. Further studies are required to assess the potential impact of metals on cellular and physiological function, particularly information regarding their toxicology in cephalopods, which is especially lacking.

Cephalopods are a major component of the diet of numerous predators such as fish, marine mammals and seabirds (Clarke, 1996; Croxall and Prince, 1996; Klages, 1996), therefore, they constitute a potentially important source of trace elements (Bustamante et al., 1998a; Muirhead and Furness, 1988). Evaluation of the trophically available metals (TAM) could be one possible tool to estimate contaminant transfer from cephalopods to predators (Wallace et al., 2003; Wallace and Luoma, 2003). This could be performed by assessing the metals contained in the soluble fraction, suggested to contain the efficiently assimilated metals. Thus, the most concerning data would be for Cd, which seems to be more abundant in the soluble fraction of the digestive gland. This is especially worrying due to the

deleterious effects induced by this metal in vertebrates (Fleischer et al., 1974). For example, a grey-headed albatross from Kerguelen Islands was found to have ingested 11.2 mg of Cd per kg of body weight per week from three squid, which is approximately 500-times higher than the weekly tolerable intake for humans (Bustamante et al., 1998a).

4.2.3. Metal-binding proteins

As described above, when trace elements are not bound to organelles, they are often found in the soluble fraction, where they are associated with cytosolic proteins, thereby inhibiting the toxic interactions of metallic ions with sensitive binding sites (proteins, molecules or cell structures). The formation of complexes of trace elements with specific proteins has been subject of several studies in various cephalopods species, including cuttlefish (Bustamante et al., 2006a; Bustamante et al., 2002a; Declair et al., 1978), squid (Castillo and Maita, 1991; Finger and Smith, 1987; Tanaka et al., 1983), and octopus (Bustamante et al., 2002a; Nakahara et al., 1979; Raimundo et al., 2010b; Raimundo et al., 2008, 2010d; Rocca, 1969; Ueda et al., 1985). The targeted tissues included the digestive gland as the preferential organ for trace element accumulation, in addition to the branchial hearts, branchial gland, kidneys, blood and muscle. As for other studies that have investigated trace element detoxification in cephalopods, only few trace elements are usually targeted, the most common of which are Cd, Cu and Zn. Even with these metals, the results obtained vary significantly among species, making it difficult to conclude a single or general rule.

In the digestive gland of the common cuttlefish *S. officinalis* and the *Ommastrephes bartrami*, *Nototodarus gouldi* and *T. pacificus* squid, soluble Cd is mostly bound to high molecular weight (HMW) proteins (> 70 kDa) (Bustamante et al., 2006a; Castillo and Maita, 1991; Finger and Smith, 1987; Tanaka et al., 1983), whereas 53% of Cd in the squid *Onychoteuthis borealijaponica* is bound to low molecular weight (LMW) proteins (> 3 kDa) (Castillo and Maita, 1991) (Table 1). Similarly, Raimundo et al. (2010d) also showed a preferential association of Cd to LMW protein complexes in the digestive gland of *O. vulgaris*. Such differences could be explained by: (1) differences in protocol for the chromatographic analysis, in particular, whether not a reducing agent was used (Table 1); (2) inter-specific variations in detoxification processes using soluble proteins; and (3) the contrasting magnitude of metal exposure between the species evaluated. More precisely, when Cd is present in the digestive gland at low concentrations it is bound to only one type of protein (HMW in squid and cuttlefish, LMW in octopus), whereas two proteins fractions are identified when the total Cd concentration exceeds 40 $\mu\text{g}\cdot\text{g}^{-1}$, regardless of the species (Table 1). As suggested by Raimundo et al. (2010d), the Cd level in the digestive gland drives a transient detoxification strategy, based on the formation of Cd complexes with HMW proteins when Cd levels are low, which changes to binding to

LMW proteins when the Cd concentration exceeds a certain threshold. In this case, Cd could be trapped within insoluble structures, as described above.

Concerning the nature of these metal-binding proteins, several authors have suggest LMW proteins such as MTLP (Bustamante et al., 2002a; Declair et al., 1978; Finger and Smith, 1987; Raimundo et al., 2010b; Tanaka et al., 1983). Metallothioneins (MTs) are well-known LMW (about 6–7 kDa), heat stable and cysteine-rich proteins. Furthermore, metals like Ag, Cd, Cu, Hg and Zn have a high affinity for MTs due to their high thiol group content (Viarengo and Nott, 1993). Thus, MTs are thought to play a key role in the homeostasis of essential metals, in addition to an important role in the trace element tolerance of organisms (Amiard et al., 2006). Indeed, their unique affinity for trace elements gives them the ability to scavenge not only essential, but also non-essential trace elements, with the affinity suggested as follows: Hg > Cu, Ag, >> Cd > Pb > Zn > Co (Vasak, 1991). It has been widely demonstrated that exposure of organisms to trace elements could lead to the induction of MTs through Zn-sensitive inhibitor mediation of MT genes (see Amiard et al., 2006). To the best of our knowledge, structural characterisation of MTs from cephalopods has not yet been performed. Nevertheless, several authors have confirmed the presence of proteins of a similar weight that bind trace elements, using MTs from other species (rat, rabbit and horse). It has been shown that proteins with a molecular weight close to that of MTs from other species mainly bind Cu in *Todarodes pacificus* (Tanaka et al., 1983) and *Sepia officinalis* (Bustamante et al., 2006a), bind Cd in *Ommastrephes borealijaponica*, *O. bartrami* and *T. pacificus* squid (Castillo and Maita, 1991; Tanaka et al., 1983), and Zn in *O. bartrami* (Castillo and Maita, 1991).

The quantification of MTs in the digestive gland through the measurement of thiol groups has highlighted the wide variation in MT concentrations among cephalopod species, ranging from 700 to 3500 $\mu\text{g}\cdot\text{g}^{-1}$ wet weight (Bustamante et al., 2002a; Raimundo et al., 2010b). However, these levels are comparable to those found in other molluscs, such as the mussel *Mytilus galloprovincialis* (Pavičić et al., 1993) and gastropod *Littorina littorea* (Bebiano et al., 1992). Interestingly, the lowest levels of Cd (and also the highest percentage of soluble Cd) were found to be associated with the highest levels of MTs (Bustamante et al., 2002a), suggesting: (1) MTs are probably not the main pathway of detoxification for Cd in cephalopods; and (2) other detoxification mechanisms occur beyond a particular concentration threshold, potentially involving heavier soluble proteins or binding to insoluble structures.

Regarding essential metals, Le Pabic et al. (2015b) demonstrated a positive correlation between the whole organism Cu and Zn concentrations and MTs in juvenile cuttlefish, which is consistent with the essential role that MTs may play in the homeostasis of essential trace elements in cephalopods. However, based on several indirect methods, Craig and Overnell (2003) showed there was no evidence of MT involvement in Cu or Zn regulation in the eggs, hatchlings or adult tissues (digestive gland,

muscle and eyes) of the veined squid *L. forbesi*. Nevertheless, taking into account the small amount of lysosomes contained within the digestive gland of Loliginidae, combined with the histological observations that MTLPs are mainly located within lysosomes (Martoja and Marcaillou, 1993), these observations may be in accordance with those previously reported. This raises the question of the relative importance of MTs in the homeostasis of essential elements in cephalopods, and further investigations are required in order to elucidate their role.

Additionally, Castillo and Maita (1991) purified a protein fraction of approximately 16 kDa from the digestive gland cells of the squid *O. bartrami*. They demonstrated the occurrence of three differently charged proteins that bind Cd, Cu and Zn within a similar weight fraction. Amino acid characterisation of these proteins found that the cysteine content appeared to be lower than the detection limit, suggesting that this fraction contained only a very low amount of MTs. Nevertheless, these authors highlighted the high content of aspartic and glutamic acids in these proteins, which can also bind trace elements (Viarengo et al., 1988). When combined, aspartic and glutamic acids constitute 21–29% of the total molar ratio of these proteins, which is comparable to the high cysteine content (25–30%) found in MTs. The authors suggested that trace elements must be bound to dicarboxylic amino acids instead of thiolates.

Conclusion

This paper attempted to review the existing knowledge regarding the high capacity of cephalopods to concentrate metallic trace elements in their tissues, to assist in unravelling the physiological mechanisms involved in metal homeostasis. It also highlighted the lack of knowledge in this area, which requires investigation in order to obtain an overall understanding of the ecotoxicology of these exceptional organisms. It appears that the detoxification mechanisms include both dynamic and transient processes, allowing the storage and/or elimination of metals depending on the exposure conditions. However, such mechanisms need to be further explored, taking into account the physiology, particularly the digestive physiology, and the life traits of each species, in order to explain the differences in bioaccumulation capacity observed among cephalopod species. In addition, the trophically available metals (TAM) fractions that could be transferred from cephalopods to predators (Wallace et al., 2003; Wallace and Luoma, 2003) might be driven by the detoxification / storage strategies in the prey. As cephalopods are a major component of the diet of numerous predators such as fish, marine mammals and seabirds (Clarke, 1996; Croxall and Prince, 1996; Klages, 1996), unravelling their metal metabolism is therefore essential for perceiving their key role in trace elements transfer and/or biomagnification in marine food web (Bustamante et al., 1998a; Muirhead and Furness, 1988).

Furthermore, little is known about the toxicological risks of such bioaccumulation capacities on the physiological and ecological performance and fitness of cephalopods. As recently shown, the cephalopod population is increasing globally, despite growing anthropogenic pressure and changes in climatic conditions occurring in oceans (Doubleday et al., 2016), suggesting an apparent tolerance to marine environment contamination. Nevertheless, considering the growing interest in cephalopods by fisheries as food resources, and their key role in marine trophic webs, this work highlights the need to gain a better understanding of the ecotoxicology of metals, and more broadly, of others pollutants in cephalopods. Further works would have to combine physiological, toxicological and biogeographical approaches to address the gaps of knowledge reported in this paper.

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Table 1. Concentrations ($\mu\text{g}\cdot\text{g}^{-1}$ dw) and proportion in the cytosol (%) of Cd in the digestive gland of 4 cephalopod species as well as the molecular weight of proteins associated with Cd. Main and minor peaks refer to the fractions to which associated Cd concentrations were the highest and lowest (but detectable), respectively, following chromatographic analysis.

Species	[Cd] $\mu\text{g}\cdot\text{g}^{-1}$ dw	% in cytosol	Main peak	Minor peak	Protein weight (kDa)	Reducing agent	Reference
<i>Nototodarus gouldi</i>	10-30 40-80	70 %	HMW	IMW	70 kDa > 70 kDa + 20 kDa	Mercaptoethanol ^a	Finger and Smith (1987)
<i>Ommastrephes bartrami</i>	211	NA	IMW	LMW	16 kDa (10-30 kDa) > 70 kDa	Mercaptoethanol no	Castillo and Maita (1991)
<i>Onychoteuthis borealijaponica</i>	NA	NA	LMW		3 kDa	Mercaptoethanol	Castillo et al. (1990)
<i>Todarodes pacificus</i>	38-82	30 %	HMW	IMW	70 kDa + 11-16 kDa	no	Tanaka et al. (1983)
<i>Sepia officinalis</i>	15	53 %	HMW		70 kDa	no	Bustamante et al. (2006a)
<i>Octopus vulgaris</i>	10-30 57-252	93 %	LMW	HMW	6-11 kDa 6-11 kDa + 130-140 kDa	no no	Raimundo et al. (2010d) Raimundo et al. (2008)

HMW: high molecular weight; IMW: intermediate molecular weight; LMW: low molecular weight. NA: not available.

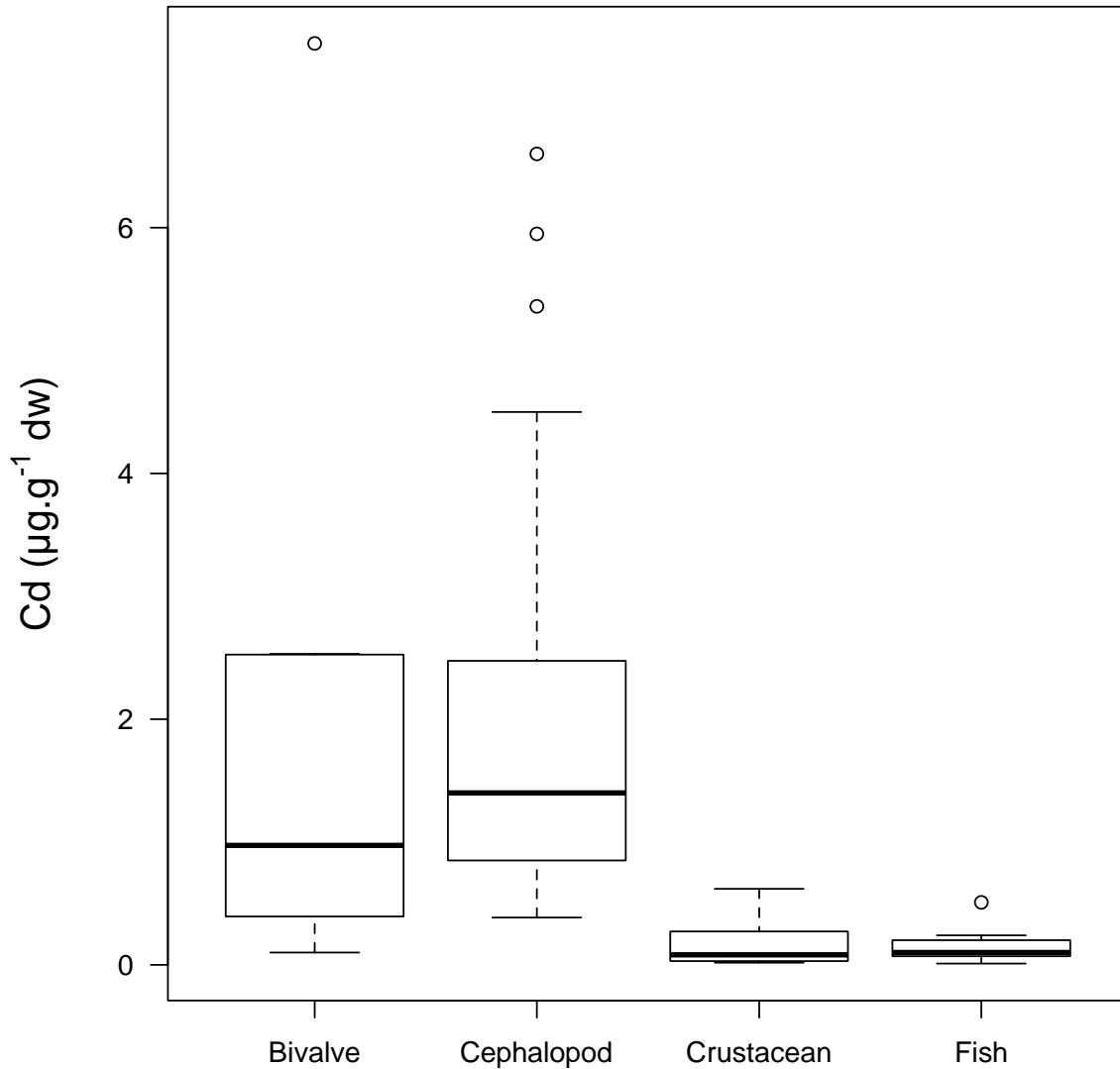


Figure 1. Whole body (or soft parts for Bivalves) concentrations of Cd ($\mu\text{g}\cdot\text{g}^{-1}$ dry weight) in different order of Fish, in Crustacean Decapods, in different Families of Bivalves and Cephalopods collected in the Bay of Biscay and the English Channel. Please note that these data are not exhaustive and illustrate the capacity of Cd bioaccumulation of cephalopod compared to others marine organisms. Per: Perciforme; Clup: Clupeiforme; Gad: Gadiforme; Myc: Myctophiforme; Ang: Anguilliforme; Dec: Decapod; Ven: Veneridae; Tel: Tellinidae; Myt: Mytilidae; Ost: Ostreidae; Pec: Pectinidae; Oct: Octopodidae; Sep: Sepiidae; Loli: Loliginidae; Om: Ommastrephidae; His: Histioteuthidae; Cra: Cranchiidae. Data extracted from the literature are detailed in Table S1.

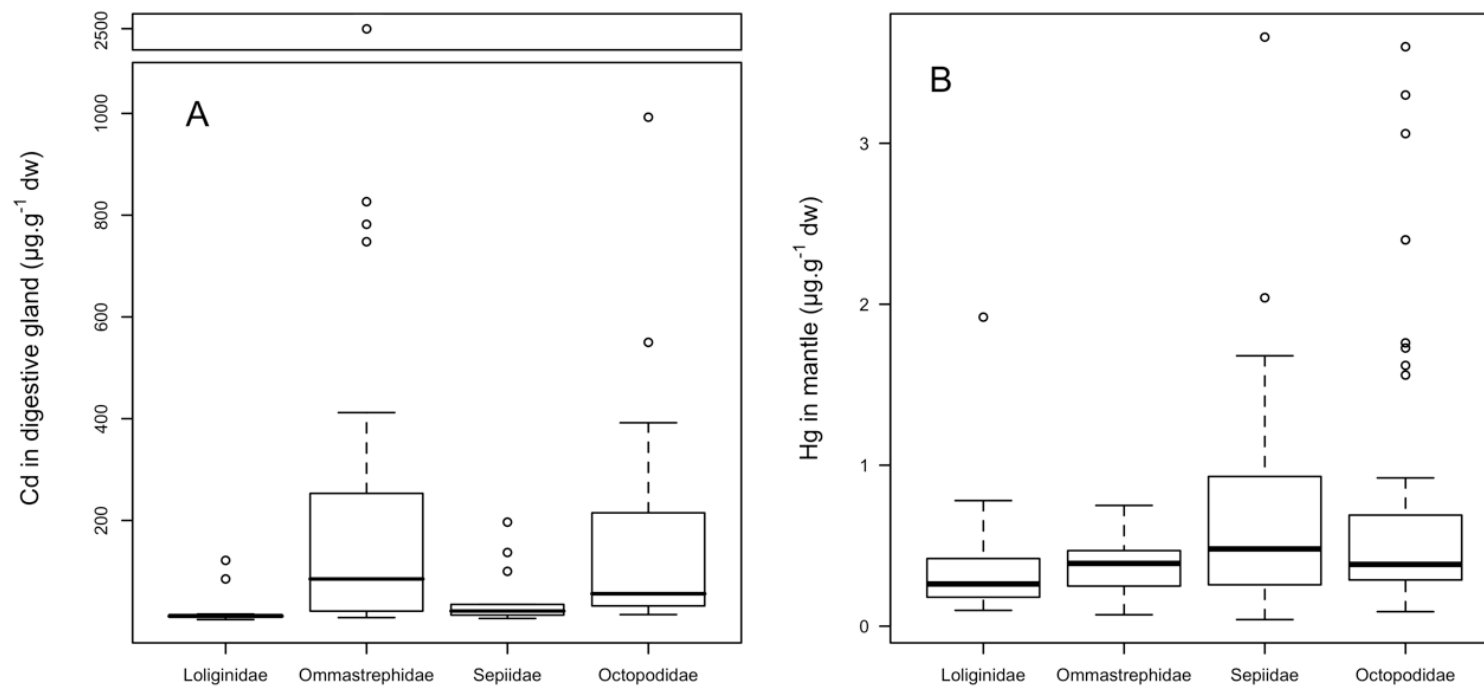


Figure 2. Cd (A) and Hg (B) concentrations (in $\mu\text{g.g}^{-1}$ dry weight) in the digestive gland and muscle, respectively, of cephalopods from the main four families (i.e. Loliginidae and Ommastrephidae squids, Sepiidae and Octopodidae) collected worldwide, with taxonomy discrimination. Data extracted from the literature are detailed in Table S2 and S3.

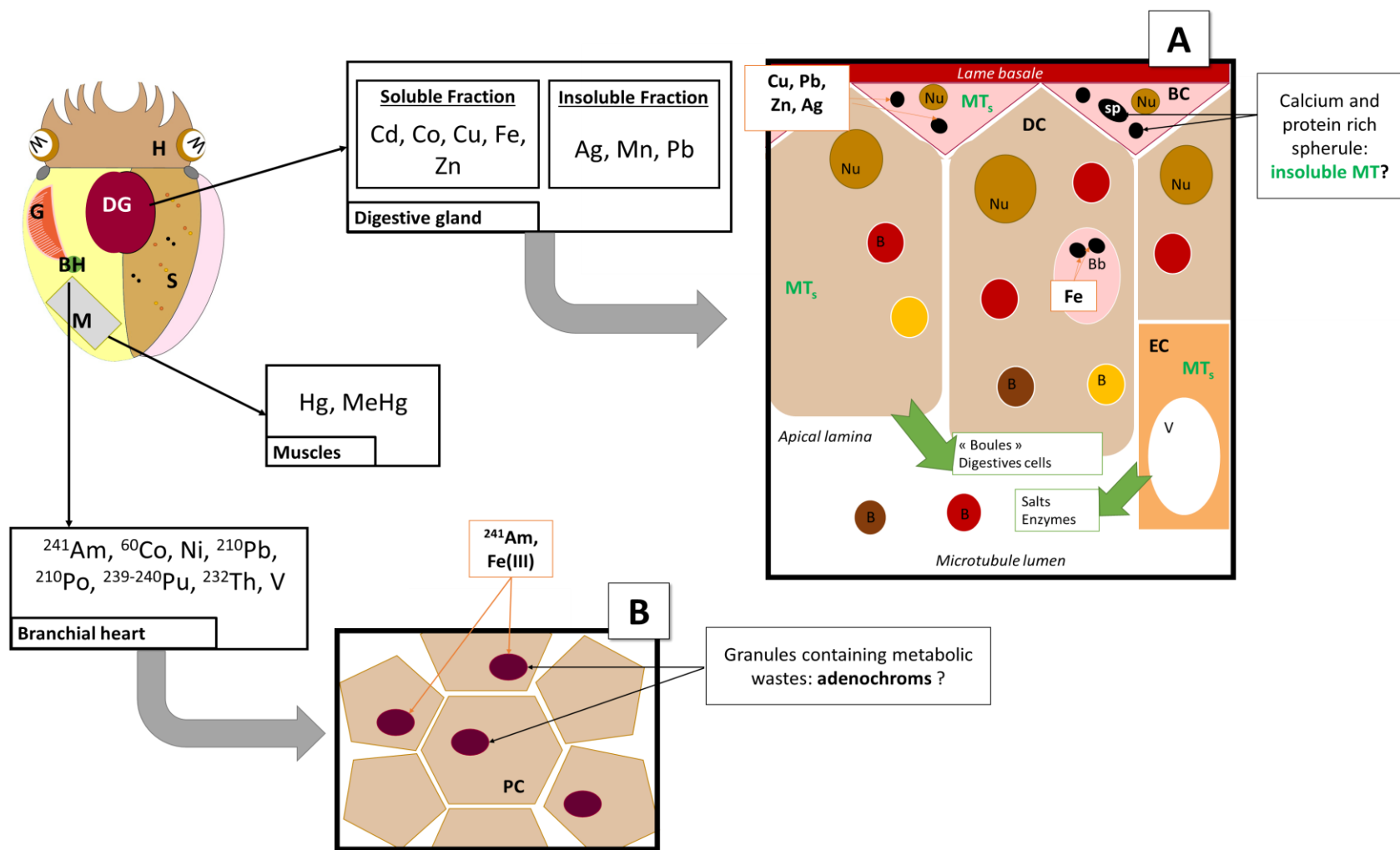


Figure 3. Simplified representation of metal bioaccumulation and organotropism in the common cuttlefish, *S. officinalis*. (A) Subcellular schematic view of the digestive gland. B: “boules” structures, heterolysosomes; Bb: Brown bodies; BC: Basal cells; DC: Digestive cells; EC: Excretory cells; MTs: Soluble metallothioneins; Nu: Nucleus; V: vacuole. (B) Subcellular schematic view of branchial hearts. PC: Polyhedral cells.

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