The coupling of reversed-phase HPLC with ICP-MS offers an attractive orthogonal approach to size-exclusion HPLC-ICP-MS for the characterisation of complex bioinorganic systems. An interface between microbore RP-HPLC and ICP-MS using a direct injection nebulizer which allows the introduction of mobile phases up to 50% methanol is described. Its applications for the analysis of cobalamins and metallothioneins are illustrated.

The complexity of many naturally existing classes of bioinorganic species makes the resolution of size-exclusion chromatography insufficient for a more detailed characterisation (within a given class) of the metal complexes present. It is well known that some species with essentially similar molecular masses may show some structural differences which may be crucial for their role in life processes. Therefore, despite SEC-ICP-MS being the primary technique for the detection of metal species in natural samples [1], the compounds detected by this technique should be further examined by chromatographic or electrophoretic techniques with complementary separation mechanisms.

Whereas in the case of SEC the coupling with ICP-MS is readily achieved owing to the compatibility of the mobile phase composition and flow-rates used, the use of ICP-MS as a detector in RP-HPLC with gradient elution using mobile phases containing more than 20% of an organic solvent have been relatively scarce because of the low tolerance of the plasma to the solvent vapour [2–12]. Only two applications concerned metalloproteins [3,12]. Although the ICP mass spectrometer used in this work can tolerate the introduction of 50% or more aqueous methanol in the FIA mode (100 µL injection into an aqueous stream of 1 mL min⁻¹), experiments showed that concentrations of methanol above 15% in the mobile phase pumped continuously at 1 mL min⁻¹ lead to the extinction of the plasma within a few minutes rendering the acquisition of a chromatogram impossible [12].

High levels of organic solvents in the mobile phase can be coped with by using a refrigerated spray chamber [5–7] or a nebulizer with a desolvation unit [8–11]. An elegant way to overcome the incompatibility of the mobile phase composition with the ICP is a post-column dilution of the effluent to bring the concentration of methanol to a level tolerated by the plasma. An interface between microbore HPLC and ICP-MS based on this principle and examples of its applications are presented below.

### Description of the interface

The interface developed is shown in figure 1. It is based on the use of a direct injection nebulizer (DIN). The DIN is a microconcentric nebulizer with no spray chamber; it nebulizes the liquid sample directly into the central channel of the ICP torch [13]. Compared with conventional nebulizers it offers 100% sample introduction efficiency into the ICP, accommodation of low sample introduction flow rates (10 – 100 µL min⁻¹), an extremely small internal volume (< 2 µL) and fast sample washout with minimal memory effects.

The use of the DIN eliminates the basic problem of the conventional nebulizers to couple micro or narrowbore chromatography with ICP-MS as the use of a microbore column (flow rate of 40 µL min⁻¹) would require a makeup flow of 20 times higher to match the working range of a cross-flow or concentric nebulizer (0.6 – 1 mL). This would lead to a
considerable dilution and high post-column dead volume to which the contribution of the spray chamber should be taken into account. In the case of the DIN the dilution factor is 2.5 which, on one hand, is sufficient to decrease the concentration of the organic modifier so that it does not make the plasma unstable, and, on the other hand, does not contribute much to the post-column dispersion and the loss of the number of the theoretical plates. Under the optimized conditions the sensitivity of the DIN-ICP-MS for the nuclide was a factor of 2 lower than that obtained with the cross-flow nebulizer and corresponded to the manufacturer’s (CETAC) specification. The relative standard deviation (for signals not limited by counting statistics) was 1 – 2%.

Species selective analysis of cobalamin analogues

Cobalamins are coenzymatically active forms of vitamin B\textsubscript{12} (cyanocobalamin) which is a water-soluble vitamin and a nutrient essential for all cells [14,15]. They belong to the group of corrinoids which are composed of a corrin nucleus with a cobalt atom in its centre. The cobalt atom is coordinately linked to a nitrogen of a pendant 5,6-dimethylbenzimidazole group. The molecule also contains ribose with an \(\alpha\)-glycosidic linkage. Depending on the substituent of the cobalt atom (upper axial ligand) different forms of vitamin B\textsubscript{12}: cyanocobalamin (CN), adenosylcobalamin (coenzyme B\textsubscript{12}, 5′-desoxyadenosyl), methylcobalamin (CH\textsubscript{3}), hydroxocobalamin (OH) and aquocobalamin (H\textsubscript{2}O), exist. Cobalamins should be distinguished from their potentially harmful analogues devoid of enzymatic activity such as e.g., cobinamides that lack the nucleotide moiety (5,6-dimethylbenzimidazole). The monitoring of the intermediate and finished products of cobalamin drugs is essential for process and quality control. Analysis of multivitamin preparations for parental nutrition is required to evaluate the stability (aging) of corrinoid compounds to light exposure and temperature, and their interactions with other vitamins and trace elements.

A chromatogram obtained by microbore HPLC-DIN-ICP-MS under the conditions optimized above is shown in figure 2a. Calibration curves are linear over three decades and the analytical precision varies between 2 – 5%.

ICP-MS is an element selective detector which detects only Co which forms 5% of the total molecule. Nevertheless the detection limits obtained by microbore HPLC-DIN-ICP-MS are about 100 times lower than those obtained with UV detection and are the lowest ever reported for on-line detection of cobalamins. The value of 20 ng mL\textsuperscript{-1} corresponds to an ADL of 0.1 ng of a cobalamin and matches the detection limits normally determined off-line by radioisotope dilution. Microbore HPLC-DIN-ICP-MS is an attractive technique for the evaluation of the serum cobalamin status.

Figure 2b shows the application of the method developed to the analysis of the status of a hydroxocobalamin preparation at detection levels not accessible by UV or ESI-MS. Two signals, in addition to the peak of Co\textsuperscript{2+} (dead volume); 2 - hydroxocobalamin; 3 - adenosylcobalamin. Conditions: pH 4.0, injected volume 5 \(\mu\)L, each standard at 50 ng. The gradient profile is indicated in % CH\textsubscript{3}OH by the dotted line.
Sequestration of metal ions (e.g. Cd\textsuperscript{2+}) in stable, intracellular macromolecular complexes is a major mechanism by which cells resist the cytotoxic effects of the metal. Metallothioneins (MTs) [a group of non-enzymatic low molecular mass proteins (6 – 7 kDa)] represent a class of the most common ligands. Classic techniques for the determination of MTs such as metal-saturation assays, polarography via sulphhydryl groups and immunoassays fail to provide information on the original metal composition and do not allow one to identify and to quantify the individual MT isoforms. The complexity of the polymorphism of mammalian metallothioneins and the variety of potentially bound metals (Cd, Zn, Cu, Hg) requires an approach coupling a high resolution separation technique able to distinguish between the proteins with a single aminoacid heterogeneity, and an element-selective detection technique able to determine the metals bound \[16,17\].

The developed coupling of microbore reversed-phase chromatography with ICP-MS was applied to a study of rabbit liver MT preparations isolated by size-exclusion chromatography and purified by anion-exchange chromatography to give the fractions of the two major isoforms: MT–1 and MT–2.

Chromatograms obtained under the optimum conditions are shown in figure 3. In contrast to the size-exclusion chromatograms of these MT–1 and MT–2 preparations which show one peak, the reversed-phase chromatograms are more complex. For MT–2 (Fig. 3a) the major peak apparently consists of three poorly resolved peaks and is accompanied by a noisy baseline and some minor peaks. One of these peaks can be identified according to its retention time as the Cd–MT–2 complex. HPLC-ICP-MS does not allow the attribution of the small peaks to labile Cd–MT–2 and Cd–MT–2 complexes or to different MT–2 sub-isoforms with microheterogeneities. A small copper peak (with an abundance corresponding to that calculated on the basis of the ratio of total Cd and Cu) is present at the retention time of the major peak. The identity of the retention time suggests the presence of a mixed Cd–Cu–MT–2 complex.

Figure 3b shows an even more complex HPLC – DIN – ICP-MS chromatogram of the other major MT isoform – MT–1. Again it is not possible to state whether true sub-isoforms are involved or artefacts of differently metallated species of one isoform. The distribution of Cu among these isoform does not follow that of Cd. Whether the early eluting peaks contain pure copper or mixed Cd-Cu complexes remains to be elucidated. At this pH (6.0) Zn complexes are too labile to be separated.

Conclusions

The coupling of reversed-phase chromatography with ICP-MS offers an attractive approach to examine the purity of a metal species isolated by size-exclusion chromatography. The use of an interface based on the post-column dilution of the methanolic mobile phase with water followed by direct injection nebulization allows the use of microbore columns and the introduction of up to 50% methanol into an ICP. This is sufficient for the majority of studies of speciation and of the metallothioneins not only in terms of metals bound (Cd, Cu) but also in terms of iso- and sub-isoforms.

References