

MINI-REVIEW

Involvement of metallothionein and copper in cell proliferation

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Metallothionein is a low-molecular weight, cysteine-rich, metal-binding protein which has been implicated in the detoxification of toxic metals (cadmium, mercury), metabolism of zinc and copper, as well as in the scavenging of free radicals. Recent evidence suggests that the protein may also be involved in cell proliferation. Based on the experiments carried out so far, it is assumed that the fundamental role of metallothionein in cell proliferation may be to detoxify and/or transfer copper ions from the cytoplasm to the nucleus at the G₁/S phase, which in turn participate in some way in nuclear DNA synthesis.

Keywords: cell proliferation, copper, metallothionein

Introduction

Metallothionein (MT) is a metal-binding protein originally isolated from equine renal cortex by Margoshes & Vallee (1957). Later studies revealed that MT occurs throughout the animal kingdom, and is also found in higher plants, eukaryotic micro-organisms and some prokaryotes. In animals, the protein is most abundant in parenchymatous tissues, i.e. liver, kidney, pancreas and intestines (Kägi & Schäffer 1988, Robinson 1989, Kay *et al.* 1991).

The characteristic features of MT are its low molecular weight, 6000–7000, and its unusual amino acid composition: cysteine accounts for 30% of the residues and aromatic amino acids are absent. The cysteine residues along the polypeptide chain are fixed and organized into Cys–X–Cys, Cys–Cys or Cys–X–Y–Cys sequences. All cysteinyl-SH groups are involved in ligation of metal ions. Normally MT binds seven atoms of cadmium or zinc per molecule or 12 atoms of copper. A tetrahedral arrangement of four sulfur atoms surrounds each cadmium or zinc atom, while a trigonal arrangement surrounds the copper atoms. The metals are localized in two

polynuclear clusters in two distinct domains, each containing half of the polypeptide chain. The C-terminal or α -domain contains four cadmium or zinc atoms, whereas the N-terminal or β -domain contains three atoms of the metals. Each domain also has the potential to bind six copper atoms. Interestingly, copper binding occurs preferentially in the β -domain and cadmium or zinc in the α -domain. Moreover, copper in its cuprous form binds more firmly than cadmium, which in turn binds more firmly than zinc. MT from *Neurospora*, which corresponds to the β -domain of vertebrate MT, as well as yeast MT bind *in vivo* only copper (for review, see Armitage *et al.* 1982, Bremner 1987, Kägi & Schäffer 1988, Bremner & Beattie 1990).

A most remarkable feature of MT is its inducibility. It is induced by a wide range of metals (cadmium, zinc, copper, mercury, bismuth), as well as by specific hormones and other agents including glucocorticoids, catecholamines, glucagon, interleukin-1, interleukin-6, dibutyl cAMP, phorbol esters, protein kinase C and protein kinase A (Kägi & Schäffer 1988, Garrett *et al.* 1992). The control of MT synthesis occurs at the transcriptional level. The transcription of the MT gene is mediated through specific DNA sequences in the promoter region and transcription factor(s) of protein nature, e.g. ACE1 in *Saccharomyces cerevisiae* (Thiele 1988), and

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MBF-1 (Imbert *et al.* 1989), p39 (Andersen *et al.* 1990), MEP-1 (Labbe *et al.* 1991) and MREBP (Koizumi *et al.* 1992) in mammalian cells. It has been proposed that the transcription factors bind metals, become activated and then bind to specific sequences on the MT gene, thereby increasing transcription (Palmiter 1987, Searle 1987, Fürst *et al.* 1988, Karin *et al.* 1988).

Although MT was discovered 35 years ago, the function specific to the protein still remains a topic of discussion. As far the protein has been implicated in the detoxification of toxic metals (cadmium, mercury), in the metabolism of zinc and copper, as well as in the scavenging of free radicals (Cousins 1985, Richards 1989, Bremner & Beattie 1990). Accumulating evidence suggests that MT may also be involved in cell proliferation, which is the subject of this article.

MT and the cell cycle

The highly conserved protein in evolutionary terms, its ubiquitous occurrence, and the programmed synthesis of MT during tissue regeneration and development (Andrews *et al.* 1991, Nemer *et al.* 1991) are significant arguments for its involvement in some fundamental cellular processes, e.g. the cell cycle. Ohtake *et al.* (1978) have shown, for the first time, the parallelism of enhanced DNA synthesis with increased MT formation in the liver of rats recovering from partial hepatectomy. Later studies, using immunohistochemical methods, have revealed the presence of MT under normal physiological conditions, especially in actively dividing cells, e.g. in various epithelia (Nishimura *et al.* 1989b, 1990, 1991). The authors conclude that MT immunostaining is related mainly to the turnover rate of a tissue, e.g. the relatively short turnover of the cornea epithelium (3–4.5 days) accounts for stronger MT immunostaining in this tissue than in the epidermis in which keratinocytes are replaced with cells migrating from the basal layer with a turnover of 15–30 days (Nishimura *et al.* 1991). In addition, application of tetradecanoyl phorbol ester, a potent cancer promoting and hyperplasia-inducing agent, to mouse skin results in activation of cell divisions in the basal layer (Nishimura *et al.* 1988). Under these conditions both [³H]thymidine was found to be incorporated into cell nuclei and MT immunofluorescence was observed in the dividing basal cells. The intense MT immunostaining has been also demonstrated in spermatogonia and basal cells of seminal vesicles and the ejaculatory duct in rats (Nishimura *et al.* 1990), as well as in human

uroepithelium displaying dysplastic changes or carcinoma (Bahnson *et al.* 1991).

Immunohistochemical methods have also provided interesting information on the intracellular distribution of MT. For example, in the rat perinatal liver, which may be used as a model for rapidly dividing cells, there is a gradual increase in levels of cytoplasmic MT during gestation, but at day 20, MT is mainly localized in the nuclei and remains so for several days after birth; thereafter, the intranuclear localization of MT decreases with age (Panemangalore *et al.* 1983). The intense immunostaining of MT is also found in the nuclei of fetal and newborn human liver and kidney (Nartey *et al.* 1987), as well as in the nuclei of leptotene spermatocytes (Nishimura *et al.* 1990). Furthermore, in the partially hepatectomized rat liver, which is used as a model for actively growing tissues, MT is localized predominantly in the nuclei, whereas MT is found only in the cytoplasm of the laparotomized rat liver (Nishimura *et al.* 1989a). To get more insight into the relationship between MT and the cell cycle, Tsujikawa *et al.* (1991) stimulated primary cultured adult rat hepatocytes by epidermal growth factor (EGF) and insulin and then examined by indirect immunofluorescence the intracellular localization of MT. The study revealed that MT staining is confined to the cytoplasm in G₁ phase, while in the early S phase the protein is mainly localized in the cell nuclei. However, in the period of maximal DNA synthesis MT staining in nuclei appears to be weaker than in the cytoplasm. The authors pointed out that the stimulation of proliferation by EGF is necessary for localization of MT in the nuclei of primary cultured hepatocytes. Generally, these studies strongly suggest that intracellular changes in the localization of MT are closely related to cell proliferation, especially to early stages (G₁/S) of this process.

Unfortunately, immunohistochemical methods, although very important in examining the intracellular distribution of MT under normal physiological conditions in particular, have not provided any information on the role of MT in the nucleus and cytoplasm during the cell cycle, as well as whether the nuclear and cytoplasmic MT are present as zinc and/or copper-containing proteins. The answers to these questions, at least in part, have been obtained by using biochemical methods. For instance, Ohtake *et al.* (1978) have found the presence of high amounts of Zn-MT in the cytoplasm of liver in rats recovering from partial hepatectomy. High levels of MT associated mainly with zinc and negligible amounts of copper are also present in the cytoplasm

of fetal and postnatal liver of rats (Mason *et al.* 1980), mouse and Chinese hamster (Bakka & Webb 1981), and humans (Riordan & Richards 1980, Klein *et al.* 1991). On the contrary, the cytoplasmic MT from the perinatal liver of calf (Hartman & Weser 1977), Syrian hamster (Bakka & Webb 1981) and guinea-pig (Lui 1987) contains primarily copper. It is worthy of note, however, that both the metal composition and intracellular localization of MT may undergo essential changes during the course of development. The idea is supported by a recent study (Włostowski 1992) which revealed that the cytoplasmic MT in the newborn liver of bank vole is a zinc-containing protein, the concentration of which abruptly decreases between 1 and 5 days following birth, just in the period directly preceding a rapidly growing liver, but most of the lost cytoplasmic MT is recovered in the nuclei as a copper-containing protein. These changes may be explained by a substitution process involving the pool of Zn-MT already present in the cytoplasm and the increasing concentration of the total and nuclear copper during that period (Włostowski 1992). Since copper has a higher binding affinity for MT than zinc (Day *et al.* 1981, Kägi & Schäffer 1988), one may conclude that copper displaces zinc already bound to MT in the cytoplasm and subsequently the complex Cu-MT is translocated to the nuclei. There is some circumstantial evidence in the literature indicating that it may also be the case in the neonatal liver of rats. Thus an abrupt decline in the cytoplasmic Zn-MT at 4–14 days postpartum observed by Wong & Klaassen (1979) coincides with a sharp increase in the total and nuclear copper concentrations (Evans *et al.* 1970, Terao & Owen 1977). In addition, at the 13th day following birth the cytoplasmic MT contains greater amounts of copper than zinc (Holt *et al.* 1987), suggesting an involvement of the substitution process.

Although more study is needed to confirm the assumption, the aforementioned data suggest that MT may function as a copper transfer protein from cytoplasm to nuclei and/or as a copper detoxifying factor in the cell nuclei. Also, it cannot be excluded that the complex Cu-MT, by itself, is involved in some way in the triggering of the S phase as the protein is localized in nuclei only in the early S phase (Tsujikawa *et al.* 1991). However, if MT is a necessary protein for the onset of S phase, then a lack of this protein should prevent the cell from the proliferation. There is some evidence in the literature indicating that it is not the case, at least for *S. cerevisiae* cells. It has been shown that yeast cells lacking their own MT-like copper-binding protein

grow normally under standard laboratory conditions, but are hypersensitive to copper poisoning (Thiele *et al.* 1986). Interestingly, expression of monkey MT complementary DNAs in these cells restored their ability to grow in a high copper medium. Thus it is conceivable that MT plays a similar role in mammalian cells and that the protein is not involved directly in cell proliferation, but indirectly, e.g. through detoxification and/or transfer of copper from cytoplasm to nucleus. This conclusion implies, however, that copper may also play a fundamental role in cell proliferation.

Copper and cell proliferation

The precise role of copper in cell proliferation is as yet unknown, although increasing evidence supports the idea that this trace element may be involved in the cell cycle. For instance, the nuclear copper concentration reaches the highest level (62–71% of the total tissue copper) in the period directly preceding and during a rapidly growing liver in the neonatal bank vole (Włostowski 1992). Likewise, in the human fetal liver, as well as in the newborn rat liver the highest proportion of copper is recovered from the nuclear fraction, whereas over 60% of the total tissue copper is present in the cytosol in the adult liver (Evans *et al.* 1970, Nartey *et al.* 1987). Apart from the perinatal liver exhibiting several-fold higher concentrations of copper than the adult liver (Evans *et al.* 1970, Terao & Owen 1977, Riordan & Richards 1980, Bakka & Webb 1981, Lui 1987, Nartey *et al.* 1987, Klein *et al.* 1991, Włostowski 1992), the cells of malignant tumors display a higher ability to accumulate copper than their normal counterparts (Nederbragt *et al.* 1989). However, if an increase in intracellular copper concentration, especially in the nuclei, is a necessary condition for cell proliferation, then any deficiency of this metal should suppress the process. Indeed, an inhibition of the proliferation, particularly nuclear DNA synthesis, has been found to occur in spleen lymphoid cells (SLC) from copper-deficient rodents (Lukasiewicz & Prohaska 1983, Davis *et al.* 1987, Kramer *et al.* 1988). A more recent study (Padhye *et al.* 1992) has revealed that naphthoquinone thiosemicarbazones—potent chelators of copper—efficiently inhibit DNA synthesis in P388 lymphocyte cells. Although thiosemicarbazones which strongly bind copper may have directly limited the availability of the metal for the cells in the latter case, in the former one the limitation of copper for SLC may have resulted from an accompanied deficiency of ceruloplasmin activity in the blood plasma (Kramer *et al.* 1988) as this

glycoprotein, synthesized in the liver, is thought to be a major copper donor to extrahepatic tissues (Cousins 1985). If this is the case, then one may assume also that the need for greater amounts of copper, e.g. for the proliferation of cells participating in immune response, may account for an elevation of ceruloplasmin and copper levels in the plasma observed during inflammation and infection (Laurin & Klasing 1987, Klasing 1988, Richards & Augustine 1988). Furthermore, it is noteworthy that cytokines (e.g. interleukin-1) released by activated macrophages induce in the liver both ceruloplasmin and MT (Klasing 1988, Barber & Cousins 1988). In addition, cytokines probably induce the synthesis of MT also in lymphocytes (Oberbarnscheidt *et al.* 1988), in which the regulatory mechanism of MT production seems to be similar to that in the liver (Yamada & Koizumi 1991). Thus it cannot be excluded that MT, ceruloplasmin and copper are involved, among other things, in the lymphocyte proliferation during immune response.

Further circumstantial evidence suggesting an involvement of ceruloplasmin in copper delivery to dividing cells comes from studies on copper metabolism during and after pregnancy in mammals (Terao & Owen 1977). For instance, maternal plasma ceruloplasmin and copper concentrations increase during pregnancy but decrease postpartum. Simultaneously, biliary excretion of copper almost stops at term and remains decreased thereafter. In addition, the content of copper in the newborns, especially in the liver, is very high. These data strongly suggest that the mother stores copper for the use of her fetuses and newborn pups, and that ceruloplasmin transfers copper directly (fetuses) or indirectly (through milk) to the rapidly growing offspring. Otherwise their growth and development can be significantly disturbed. This situation exists, e.g. for dams producing milk deficient in copper, which bear an autosomal mutation referred to as toxic milk (Rauch 1983). Thus, as in the case of an inflammatory response, it is reasonable to assume that high amounts of copper being transferred on ceruloplasmin from mother to offspring may also be implicated in the process of cell division.

Concluding remarks

The literature data presented in this paper strongly suggest that MT and copper play important functions in the cell cycle, although their precise role still remains to be established. Nevertheless, based on the experiments carried out to date, one may

conclude that the fundamental role of MT in cell proliferation may be to detoxify and/or transfer copper ions from the cytoplasm to the nucleus at the G₁/S phase, which in turn may participate in some way in the nuclear DNA synthesis as the process is accompanied by very high levels of copper in the cell nuclei and any deficiency of copper suppresses the incorporation of [³H]thymidine. Further studies are urgently needed, however, to verify the conclusion since it is based mainly on indirect inference.

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