

Function of Metallothionein in Gene Expression and Signal Transduction: Newly Found Protective Role of Metallothionein

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(Received December 29, 2007)

A heavy-metal-binding protein, metallothionein (MT), is induced by heavy metal overload and reactive oxygen species (ROS). The metals and ROS are rendered harmless by binding to MT or by oxidizing MT, respectively. This is a well-known role of MT. MT is also induced by glucocorticoids and cytokines. Induced MT may increase intracellular free zinc and modulate the activity of transcription factors. Changes in MT levels are thought to help in adaptation to changes in environmental conditions. MT modulates inflammatory reactions, including lipopolysaccharide (LPS)-induced expression of cytokines, nitric oxide production from macrophages in response to LPS, and resistance to LPS/D-galactosamine-induced lethality. In this review, we focus on a newly found protective role of MT, which acts mainly via changes in intracellular zinc levels or modulation of gene expression.

Key words — metallothionein, zinc, metal-responsive element-binding transcription factor-1, nuclear factor- κ B, inflammatory reaction

INTRODUCTION

Metallothionein (MT) is a low-molecular-weight metal-binding protein.^{1,2)} One-third of its amino acid composition is cysteine, and it lacks disulfides, aromatic amino acids, and histidine. In mouse, four *MT* genes have been cloned, *MT-I* to *IV*. *MT-I* and *II* are actively expressed in many cell types in various organs and tissues, as well as in most cultured cells. In contrast, *MT-III* and *IV* show a very restricted cell-type-specific expression pattern.³⁾ We focus on *MT-I* and *II*. MT has high affinity for heavy metals and is usually saturated with essential metals such as zinc. It has been suggested to maintain zinc homeostasis.^{4,5)} It is known to be induced by heavy metals.^{1,2)} Defense against some heavy metals such as cadmium has been considered to be one of its toxicological roles. Heavy-metal-induced transcriptional activation of

MT-I and *II* is mediated by a *cis*-acting DNA element, metal-responsive element (MRE).^{6–8)} MRE-binding transcription factor-1 (MTF-1) is a highly conserved zinc finger transcription factor (TF) that regulates the transcription of target genes, including MT and Zn transporter-1 (ZnT-1), in response to heavy metals.^{9–12)} MT is also induced by reactive oxygen species (ROS),^{13,14)} glucocorticoids (GC),^{15–17)} and some cytokines.^{18–23)} Therefore, the level of MT changes in response to environmental conditions to allow adaptation to the environmental conditions. Some reports show that MT protects cells against factors other than heavy metals.^{24–27)} Here we review; the molecular mechanisms of MT induction, the mechanism of nuclear trafficking of MT, the effect of MT on transcriptional activity, and the physiological role of MT based mainly on our findings.

OVERVIEW OF MT INDUCTION MECHANISMS

MT is induced by heavy metals and ROS. Loss

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of MTF-1 silences endogenous *MT* genes, indicating that MTF-1 is required for both the basal and zinc-induced transcription of *MT* genes.^{28,29)} MTF-1 has a metalloreulatory DNA-binding domain composed of six Cys₂His₂ zinc fingers.^{9–11)} Deletion mutants missing finger 1, fingers 5 and 6, or only finger 6 had severely attenuated metal-induced binding to mouse *MT-I* promoter.³⁰⁾ A study of point mutations of zinc fingers showed that the six fingers are not functionally equivalent in zinc response.³¹⁾ A potential role in MTF-1 zinc sensing contributed by an unusual (non-canonical) peptide RGEYT linker connecting the two N-terminal zinc fingers was reported recently.³²⁾ MTF-1 is also the key factor in response to cadmium and ROS. But the exact mechanisms of *MT* induction are not clear.

Activation of MTF-1 depends only on zinc in a cell-free system; *e.g.*, addition of cadmium to recombinant MTF-1 or cell extract does not activate MTF-1 in electrophoretic mobility shift assay, as shown in a cell-free, MTF-1-dependent transcription system.³³⁾ Transcriptional induction by cadmium, copper, or H₂O₂ additionally requires the presence of zinc-saturated *MT* (Zn-*MT*). This induction is explained by the preferential binding of cadmium or copper to *MT* or its oxidation by H₂O₂; the concomitant release of zinc in turn leads to the activation of MTF-1 (Fig. 1). But this ex-

planation contradicts the result that approximately 100 μM zinc is needed for activation of MTF-1 in cultured cells but only 20 μM of cadmium. Twenty μM cadmium might not release the same amount of zinc as 100 μM zinc. The upstream stimulatory factor (USF) family and the antioxidant response element (ARE), which overlaps the USF-binding site (USF/ARE), are involved in *MT* induction by H₂O₂ and cadmium.¹⁴⁾ A cysteine-rich cluster, from the C-terminus to a serine/threonine-rich transcriptional activation domain, was specifically required for MTF-1 to activate transcription by heavy metals.³⁴⁾ Phosphorylation might be involved in the activation of MTF-1 in response to heavy metals.^{35–37)}

Molecular mechanisms of *MT* induction by cadmium and ROS are still unclear. *MT* was induced under hypoxia, and the induction was mediated by cooperative interactions between MTF-1 and hypoxia-inducible factor-1α.^{38–40)} *MT* expression was suppressed in primary human hepatocellular carcinoma and was mediated through inactivation of CCAAT/enhancer-binding protein α by the phosphatidylinositol 3-kinase signaling cascade.⁴¹⁾ Sp1, a TF, might inhibit *MT* expression via competition with MTF-1–MRE binding.⁴²⁾ The mechanism of MTF-1-mediated *MT* expression might be complicated.

MT synthesis is induced in response to

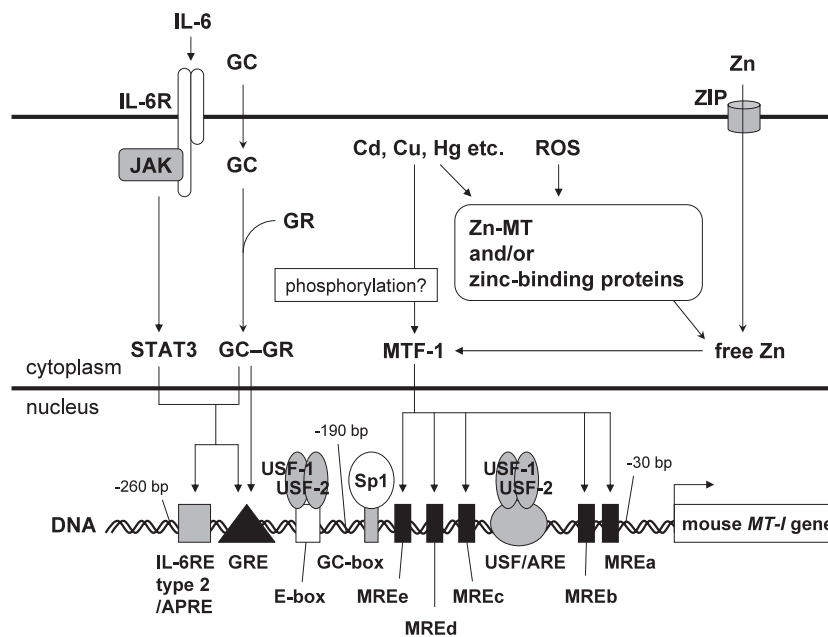


Fig. 1. Proposed Molecular Mechanisms in Mouse *MT-I* Gene Transcription

In response to heavy metals or ROS, zinc is released from Zn-*MT* or Zn-binding protein(s). The released zinc and other zinc brought in by ZIP family zinc importers activates MTF-1-mediated *MT-I* gene transcription. Phosphorylation of MTF-1 might be involved in its activation. In response to IL-6, activated STAT3 and GC-bound GR synergistically activate *MT-I* gene transcription. Sp1 and USF-1/USF-2 constitutively bind to these binding elements.

inflammation-producing compounds.^{20,43–45} We revealed that the MT-inducing factor is interleukin (IL)-6.²² We detected the MT-inducing factor of cultured hepatoma cells in the serum of lipopolysaccharide (LPS)-treated mice, and found that the presence of GC at the physiological level was necessary for the factor's MT-inducing activity. The activity was inhibited by the addition or preadministration of anti-IL-6 antibody in cultured cells and experimental animals.⁴⁶ Studies at the promoter gene level showed that both the type 2 IL-6-responsive element and the GC-responsive element are required for synergistic activation by IL-6 and GC (Fig. 1).²² This synergistic activation required not only the binding of signal transducer and activator 3 (STAT3) and GC receptor (GR) to their responsive elements, but also probably the interaction of STAT3 and GR with each other.⁴⁷ MT induction by some heavy metals (Ce^{3+} , V^{5+} , Mn^{2+}) was completely or partially mediated by IL-6.^{48–50} IL-6 is the acute-phase protein-inducing factor, and MT is a kind of acute-phase protein. Other cytokines—IL-1,¹⁹ tumor necrosis factor (TNF)- α and interferon- γ ,^{18,23} and GC^{15–17}—also induce MT expression. These results suggest that MT modulates the inflammatory response. The putative role of MT in the acute-phase response is discussed below.

NUCLEAR TRAFFICKING OF METALLOTHIONEIN

MT was first reported as a heavy-metal-binding protein in the cytosolic fraction.⁵¹ It was subsequently found in the nucleus.^{52–57} Localization of MT in nuclei was observed in growing primary-cultured hepatocytes.⁵⁴ Its translocation in digitonin-permeabilized BALB/c3T3 cells depended on the presence of ATP and cytosol factor(s) in a similar manner as for classical nuclear localization signal (NLS)-mediated nuclear translocation, even though there is no NLS in MT and the cytosolic factor is different.⁵⁸ MT was localized in the nucleus in response to the generation of a feeble ROS at the S phase.⁵⁹ Fluorescence-labeled *MT-II* diffused into the nucleus of SCC25 cells, where it was selectively and actively retained by nuclear binding factors.⁶⁰ However, the detailed mechanism of the nuclear localization of MT is still unknown. These observations suggest that nuclear MT protects nuclei from oxidation occurring with

progression of the cell cycle. On the other hand, some TFs contain zinc finger domains for interaction with their cognate DNA sequences. The zinc is the essential factor for the DNA-binding, because removing zinc ions complexed in these zinc fingers abrogates DNA-binding and transcriptional activity. Estrogen receptor (ER) in metal-depleted nuclear extracts exhibits reduced DNA-binding.⁶¹ Its binding can be restored with Zn-MT. Hence, apothionine (apo-MT) inhibits DNA-binding by abstracting zinc from functional ER.

FUNCTION OF MT IN MODIFICATION OF TRANSCRIPTIONAL ACTIVITY

We reported that MTF-1 was activated during endotoxemia, and that this activation was mediated by an increase in Zn-MT (Fig. 2).⁶² The intracellular free zinc concentration is tightly regulated. Transcription mediated by Zur and ZntR, Zn-responsive TFs of *Escherichia coli* (*E. coli*), responded to around 10^{-15} M free zinc.⁶³ The maximum volume of a typical *E. coli* cell in exponential phase is about 10^{-15} liter. The lowest possible concentration of free zinc, corresponding to one zinc atom per cell, would be 1×10^{-9} M. Thus, the concentration is $\ll 1$ labile zinc ion per *E. coli* cell. In eukaryotic cell, labile zinc concentration is estimated $\gg 1$ ion/cell.⁶⁴ The increase in Zn-MT must increase the intracellular available zinc pool because Zn-MT is equilibrium to free Zn and apo-MT (Fig. 3). On the other hand, increased MT expression maintains the intracellular zinc level under limited availability of zinc from the environment. MT-conferred MTF-1 activation may be mediated through an increase of the available zinc pool.⁶⁵ Because MT, despite its high metal binding constant ($K_{\text{Zn}} = 3.2 \times 10^{13} \text{ M}^{-1}$ at pH 7.4), can transfer zinc to the apoforms of zinc enzymes that have inherently lower stability constants.⁶⁶ Several results indicate that MT acts as a zinc chelator or donor for zinc-finger-type TFs, including ER, Sp1, and TF IIIA.^{61,67–69}

MT activates the DNA-binding activity of nuclear factor- κB (NF- κB), and causes NF- κB -mediated transactivation (Fig. 2).^{70,71} Activity of the human cytomegalovirus major immediate-early promoter was regulated by intracellular zinc levels. Overexpression of MT upregulates the DNA binding of NF- κB and thus NF- κB -induced activation

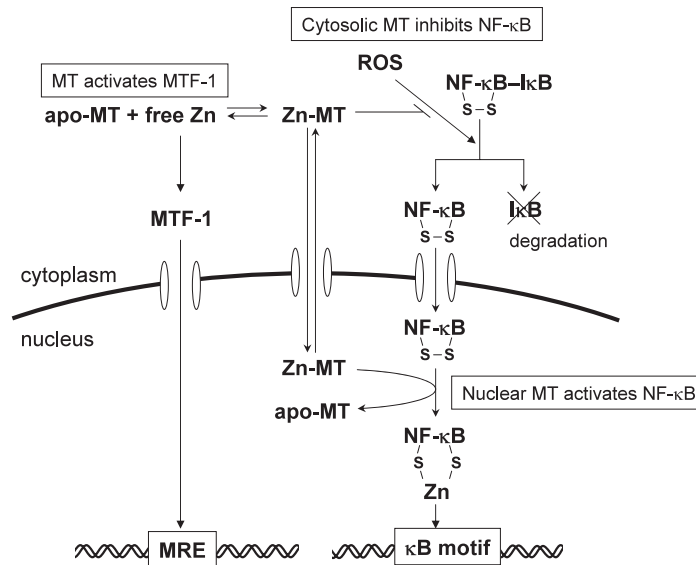


Fig. 2. Proposed Mechanisms of MT-mediated Modification of MTF-1 and NF- κ B activities

MTF-1 activation during endotoxemia is mediated by an increase in Zn-MT. NF- κ B is activated by nuclear Zn-MT. Disulfides in NF- κ B are reduced by Zn-MT. Zn bound to MT is transferred to cysteine residues in NF- κ B. In cytosol, MT scavenges radicals and inhibits ROS-mediated NF- κ B activation.

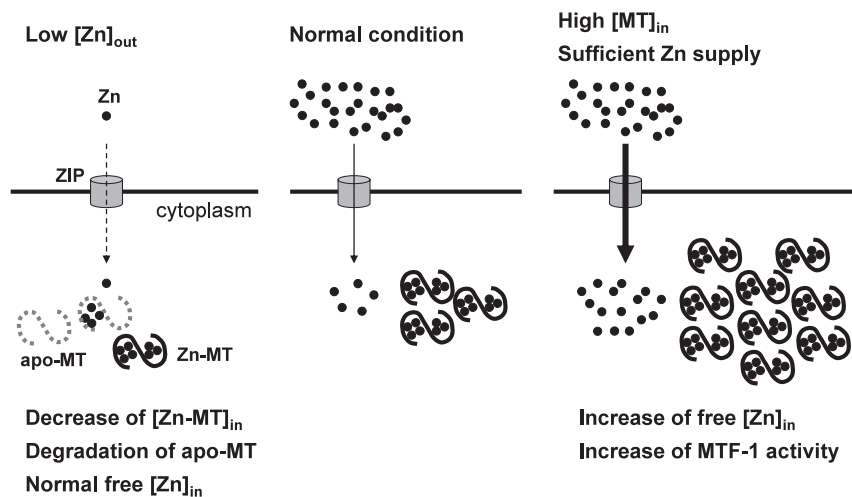


Fig. 3. Proposed Mechanisms of MT-mediated Control of Intracellular Zinc Concentration

When extracellular zinc is restricted, produced MT incorporates zinc. If the incorporated zinc is not sufficient to saturate apo-MT, apo-MT is degraded. During endotoxemia, MT expression is induced and Zn-MT is increased. When zinc supply is sufficient, the intracellular free zinc concentration is increased in response to increase in Zn-MT.

of transcription. NF- κ B is not a zinc-finger-type TF, but zinc is necessary for its activation. In the nucleus, disulfides in NF- κ B are reduced to thiol residues by thioredoxin.⁷²⁾ The cysteine residues in NF- κ B must free in order to bind zinc.⁷³⁾ A specific interaction between MT and the p50 subunit of NF- κ B was reported.⁷⁴⁾ MT may be easily oxidized and so provide zinc immediately. MT acts as a redox factor in the same way as thioredoxin and thus donates zinc. LPS-induced expres-

sion of TNF- α , IL-1 α , and IL-6 mRNAs was decreased in MT-knockout (MT^{-/-}) peritoneal exudate macrophages. These decreased expressions caused inadequate activation of NF- κ B.⁷⁵⁾ Zinc-induced expression of macrophage colony-stimulating factor in MC3T3-E1 and L929 cells required MT-mediated NF- κ B activation.⁷⁶⁾ On the other hand, NF- κ B is a redox-regulated TF, and MT may control intracellular redox status. Cytosolic MT may act as a negative regulator of NF- κ B ac-

tivity.⁷⁷⁾ Whether MT activates or inhibits NF- κ B might depend on the cell type and the stage in the cell cycle.

PHYSIOLOGICAL ROLE OF MT AND ITS PUTATIVE MECHANISM

As mentioned above, MT aids in maintaining zinc homeostasis and protects against toxicity induced by excess zinc and other heavy metals.^{1,2,4,5)} The physiological role of MT has been studied in MT^{-/-} mice. Independent researchers found that MT modulates the intracellular zinc level (Fig. 3) and gene expression, cell function, and acute-phase response. A number of MT-regulated genes were reported.⁷⁸⁻⁸⁰⁾ Some could be regulated by the MT-mediated modification of MTF-1 and NF- κ B activity, as shown in Fig. 2. Some tissue-specific regulation was observed. For example, glucose-6-phosphatase expression in MT^{-/-} mice was down-regulated in liver and upregulated in kidney.⁸⁰⁾ The regulation mechanism is unknown. Further characterization of these genes will determine whether their expressions are primary or secondary effects of an MT deficiency.

MT modulates cell functions. Reduced bactericidal activity was observed in MT^{-/-} peritoneal exudate macrophages.⁸¹⁾ Macrophages play important roles in the immune response and in inflammation. They produce nitric oxide (NO) in response to microbial infection as a bactericidal factor. Levels of expression of inducible NO synthase (iNOS) mRNA and production of iNOS protein in response to LPS stimulation were similar in MT^{+/+} and MT^{-/-} cells. The reduced production of NO in MT^{-/-} macrophages is due mainly to reduced activity of iNOS. Thiol-dependent NO stores increased the tissue NO level during renal ischemia.⁸²⁾ NO can react with the SH group of MT *in vitro*, leading to the release of metals from MT.⁸³⁾ The MT molecule has the potential to trap NO, and might modulate NO-mediated signaling. Zinc homeostasis might be partially regulated by interaction of NO with MT.⁸⁴⁾

The specific role of MT in liver regeneration after partial hepatectomy, chemical injury, and fibrosis was reported.²⁶⁾ Defective liver regeneration after injury in MT^{-/-} mice was observed. On the other hand, MT might be involved in cell proliferation because zinc modulates cell proliferation (Fig. 4). Zinc inhibits protein tyrosine phosphatase

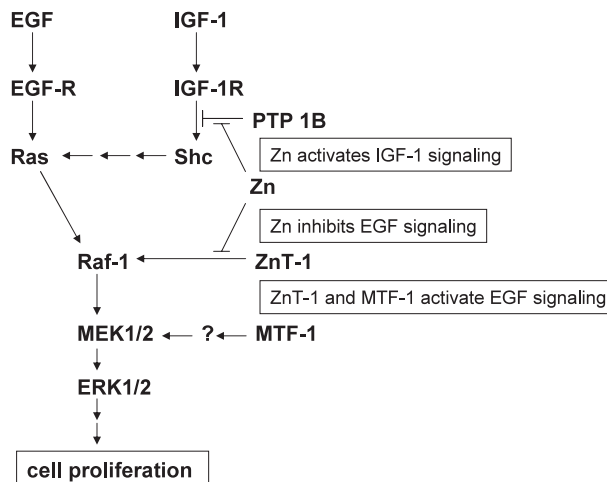


Fig. 4. Relationship between Zinc and ERK Pathway

IGF-1 receptor (IGF-1R) is phosphorylated in response to binding of IGF-1. IGF-1 signaling is inhibited by a phosphatase, PTP 1B. PTP 1B activity is inhibited by zinc. ZnT-1, zinc efflux transporter, activates Raf-1. The activation of Raf-1 is inhibited by zinc treatment. MTF-1, which regulates ZnT-1 expression, activates the ERK pathway.

activity in insulin/insulin-like growth factor (IGF)-1 signaling.⁸⁵⁾ ZnT-1, a zinc exporter whose gene is one of the targets of MTF-1, modulates the activity of the Ras-extracellular-signal-regulated kinase (ERK) cascade.^{86,87)} ZnT-1 binds to the amino-terminal regulatory portion of Raf-1, an upstream kinase in the ERK cascade, and promotes the enzymatic activity. Zn inhibits Raf-1 binding to ZnT-1. Raf-1 plays an important role in signal transduction and cell proliferation. We reported that inhibition of MTF-1 by dominant-negative MTF-1⁸⁸⁾ and MTF-1 siRNA decreased epidermal growth factor-dependent ERK phosphorylation in primary hepatocytes.⁸⁹⁾ Phosphorylation of ERK has been shown to be essential for cell proliferation. MTF-1 mediates MT production, and MT might modulate MTF-1 activity.^{33,62)} This signaling cycle might modulate cell function.

MT modulates inflammation. MT^{-/-} mice are more sensitive to LPS/D-galactosamine (GalN)-induced lethality than are MT^{+/+} mice.^{24,25)} The levels of vital mediators—TNF, NO, and platelet-activating factor—were similar in MT^{+/+} and MT^{-/-} mice, whereas the mRNA levels of the protective protein, α_1 -acid glycoprotein, in response to LPS/GalN were lower in MT^{-/-} mice. In this LPS/GalN model, zinc preadministration protects against lethality.⁹⁰⁾ The effects of an MT deficiency might be dysfunction of zinc-mediated protective gene expression. MT also protects against coagu-

latory and fibrinolytic disturbance and multiple organ damage induced by LPS, at least partly, via inhibition of the expression of proinflammatory proteins.²⁷⁾

CONCLUSIONS

We reviewed the physiological role of MT and showed some roles of MT with their mechanisms. Most of the functions can be explain by changes in intracellular zinc concentration or modulation of gene expression. MT is induced by exposure to xenobiotics and changes in environment. Induced MT modulates intracellular zinc concentration or gene expression and protects cells from the xenobiotics and changes in environment. MT has many other functions as well; for example, it protects against carcinogenesis.⁹¹⁾ These functions might also be based on changes in intracellular zinc level or modulation of gene expression. A rapid increase in intracellular zinc concentration is important for cell signaling in mast cells.⁹²⁾ MT might be involved in a lot of cell signaling systems. Further study of MT will reveal the exact molecular mechanisms.

Acknowledgements We thank our laboratory staff for their helpful advice. This research was partially supported by the Japan Society for the Promotion of Science (Grant-in-Aid for Scientific Research).

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