Effect of cytokines and ultraviolet B radiation on the promoter activity of the metallothionein gene in keratinocytes

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ABSTRACT

Metallothionein (MT) has many functions that are modulated by several factors, including ultraviolet (UV) radiation and cytokines. We thought that these diverse functions of MT might reflect specific regulatory mechanisms of its expression. To understand some of the molecular mechanisms underlying MT expression, we examined the effects of several cytokines and UVB on the promoter activity of the MT gene. First, we introduced the MT promoter construct into the HaCaT keratinocytes and treated them with various concentrations of interleukin-1α (IL-1α) and IL-6. The addition of IL-1α and IL-6 led to an increase in the promoter activity of the MT gene. UVB is known to induce MT expression in epidermal keratinocytes, and IL-6 is a possible mediator of MT induction by UV radiation. Therefore, we investigated whether UVB could induce MT promoter activity. Our results showed, interestingly, that UVB radiation has no or little effect on the promoter activity. This suggested a complex molecular regulation of the MT gene.

Keywords: Sunburn; Cytokine; Epidermis; Ultraviolet Light; Promoter

1. INTRODUCTION

Metallothionein (MT) is a ubiquitously distributed, cysteine-rich, low molecular weight protein having a high binding capacity for metals such as zinc, copper, and cadmium. It plays a role in zinc homeostasis and detoxification of heavy metals [1]. Several studies have shown that MT acts as a reactive oxygen species scavenger [2] and MT induction has protective effects against oxidative stresses such as anticancer drugs and ultraviolet (UV) radiation [3]. MT gene expression is induced not only by heavy metals but also by various stress-inducing agents such as UV [4] and X-ray radiation [5]. We found that β-thujaplicin induced MT expression in keratinocytes, both in vitro and in vivo, and thereby reduced UVB irradiation-induced apoptosis [6]. Furthermore, a number of cytokines, including interferon-α and β, tumor necrosis factor-α, interleukin-1 (IL-1), and IL-6, increase MT expression, suggesting that MT expression also involves additional functions, including immunomodulation, cell growth, and cell differentiation [7].

We thought that these diverse functions of MT might reflect specific regulatory mechanisms of its expression. We have recently cloned the promoter region of the MT gene and performed a functional assay [8]. Thus, to understand some of the molecular mechanism underlying MT expression, we focused on the effects of several cytokines and UVB radiation on the promoter activity of the MT gene.

2. MATERIALS AND METHODS

2.1. Cell Culture

The spontaneously transformed human epidermal keratinocyte cell line, HaCaT, (kindly provided by Dr. Husenig) was cultured in Dulbecco’s modified minimal essential medium supplemented with 10% fetal calf serum, 1% l-glutamine, and 1% antibiotic/antimycotic solution. The cells were maintained at 37°C in a humidified atmosphere containing 5% CO2. The viability of the cells treated with cytokines and UVB radiation for 24 h was approximately 95%, as determined by trypan blue exclusion staining.
2.2. Plasmid Constructs

The vector p5’MT-CAT, containing the 5'-flanking region of the MT-IIA gene, was generated by ligating a HindIII/BamHI fragment of the MT-IIA promoter, spanning from −764 to +79, (American Type Culture Collection, Rockville, MD, USA) to the pBS0CAT reporter construct [8]. The integrity of the reporter constructs was confirmed by direct sequencing.

2.3. UVB Radiation Source

As the UVB source, a bank of 7 fluorescent sunlamps (FL20SE.30; Toshiba Medical Supply, Tokyo, Japan) emitting rays of 275 - 305 nm and peaking at 305 nm was used [9]. The radiation dose was measured using a radiometer (UVR-3036/S; Clinical Supply, Kakamigahara, Japan).

2.4. Transient Transfection Experiments

Transient transfection was performed using Trans IT transfection reagent (PanVera, Madison, WI, USA). Briefly, 60% confluent HaCaT cells were placed in a 60-mm dish and then incubated with 0.5 μg reporter gene and the transfection reagent for 6 h. For monitoring the transfection efficiency, the cells were cotransfected with the RSV-β-galactosidase expression vector. The cells were treated with various concentrations of cytokines for 24 h. The cells were also irradiated with UVB, followed by further incubation with the medium for 24 h. The treated cells were rinsed twice with phosphate buffered saline and then lysed in 200 ml of reporter lysis buffer (Promega, Madison, WI, USA). As positive control, we used the transfected cells that were incubated with 10 μM of cadmium.

2.5. Chloramphenicol Acetyltransferase (CAT) Assays

The CAT activity, an indicator of the promoter activity, was determined by incubation with [14C]-chloramphenicol [8]. The β-galactosidase activity of all the samples were measured, and each CAT activity value was corrected for the β-galactosidase activity in the corresponding cell culture transfected in parallel. CAT activity was quantified by measuring the amount of [14C]-chloramphenicol converted to the monoacetylated form. The promoter activity was expressed as the rate of CAT activity of the sample to that of the positive control (10 μM of cadmium).

3. RESULTS

3.1. Effect of IL-1α, IL-6, and IL-10 (Figure 1)

First, we introduced the MT promoter construct into the HaCaT cells and treated them with various concentrations of IL-6, IL-1, and IL-10. After the addition of IL-1α, the promoter activity of the MT gene increased in a dose-dependent manner and reached the maximum at a concentration of 10 ng/ml. IL-6 induced the promoter activity at an even lower concentration than IL-1α. On the other hand, IL-10 did not activate the promoter.

3.2. Effect of UVB (Figure 2)

Next, we examined the effect of UVB radiation on the promoter activity of the MT gene. We harvested the HaCaT cells 24 h after UVB irradiation. The results showed
that UVB had no or little enhancing effect on the promoter activity at doses ranging from 5 to 30 mJ/cm². Furthermore, we collected the treated cells several times after irradiation, but we could not observe any strong induction of the promoter activity.

4. DISCUSSION

IL-1α has many physiological functions in the immune, metabolic, and hematopoietic systems. Keratinocytes are a major source of IL-1α. As a proinflammatory cytokine, IL-1α is involved in inflammatory and allergic skin diseases such as psoriasis and contact dermatitis. It also plays an important role in many recently defined autoinflammatory diseases [10]. IL-6 acts both as a pro- and anti-inflammatory cytokine. It is also secreted by keratinocytes, and it stimulates the immune response in both normal and abnormal skin conditions [11]. Furthermore, IL-1α and IL-6 are known to increase the MT gene expression and thereby, its protein expression. In this study, we first examined the effects of IL-1α and IL-6 on the MT promoter activity in epidermal keratinocytes. Our results show that both cytokines induced promoter activity in a dose-dependent manner. We also investigated the effect of IL-10 on MT expression, and as expected, we did not find any significant effect of IL-10 on the promoter activity.

The epidermis is the outermost layer of the skin. Keratinocytes form a majority, i.e., about 95%, of the epidermis’ cells and are major targets of solar radiation. UVB benefits humans by catalyzing the production of vitamin D, but it also causes sunburn, photoaging, and skin cancers. UVB is responsible for the production of many cytokines in the keratinocytes [12]. Furthermore, UVB is known to induce MT expression in the epidermal keratinocytes, and IL-6 is a possible mediator of MT induction by UV radiation [13]. Therefore, we investigated whether UVB radiation could induce MT promoter activity. Our results showed, interestingly, that UVB radiation has no or little effects on the promoter activity, although IL-6 clearly induced the promoter activity of the MT gene in our study. This discrepancy implies non-transcriptional mechanisms such as an increase in mRNA stability, suggesting complex molecular regulation of the MT gene.

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