A variety of macrolide activities as biological response modifiers have already been reported on (1, 3, 6, 7, 9, 11, 14). We (13) also recently reported finding that in a rat tumor model, clarithromycin (CAM) has an activity which induces a beneficial therapeutic outcome, possibly by inhibiting the production of tumor-derived factors which induce cachexia (2, 5) or immunosuppression (15). In this study, we examined in more detail the effects of CAM on the production of cytokines by a tumor to define a characteristic property of CAM.

The tumor we used was the 13762NF (subclone MTLn3) mammary adenocarcinoma (12) originating from an F-344 rat, and it was maintained in vitro in RPMI 1640 medium containing 10% fetal calf serum (FCS). CAM (Abbott Co. Ltd.) was dissolved with 100% methanol (1 mg/ml) and then further diluted in RPMI culture medium. Gentamicin sulfate (GM) and cefotiam dehydrochloride (CTM) were also diluted in RPMI medium. Expression of genes was measured by the reverse transcription-PCR method as described previously (13). The primers used for the amplification of genes were as follows: matrix metalloproteinase-9 (MMP-9) gene, 5'-GGTCCCTCCGACCTGCTGCGCCTCTA CGGCC-3' and 5'-GTCTCCAGGGCACTGAAAGATGTC ATAGGT-3'; transforming growth factor β (TGF-β) gene, 5'-GGCTCGGACACCTATTGC-3' and 5'-GCTGCACCTTGACAGGCACGAC-3'; tumor necrosis factor alpha (TNF-α) gene, 5'-CAAGGAGGAGAAGTTCCCAA-3' and 5'-CGGACTTCCGATGTCGAG-3'; inhibitor of metalloproteinase (TIMP-2) gene (24 h), and no significant effect was observed for these genes (data not shown), and further dilution in RPMI culture medium to reach final concentrations. Gentamicin sulfate (GM) and cefotiam dehydrochloride (CTM) were also diluted in RPMI medium. Expression of genes was measured by the reverse transcription-PCR method as described previously (13). The primers used for the amplification of genes were as follows: matrix metalloproteinase-9 (MMP-9) gene, 5'-GGTCCCTCCGACCTGCTGCGCCTCTA CGGCC-3' and 5'-GTCTCCAGGGCACTGAAAGATGTC ATAGGT-3'; transforming growth factor β (TGF-β) gene, 5'-GGCTCGGACACCTATTGC-3' and 5'-GCTGCACCTTGACAGGCACGAC-3'; tumor necrosis factor alpha (TNF-α) gene, 5'-CAAGGAGGAGAAGTTCCCAA-3' and 5'-CGGACTTCCGATGTCGAG-3'; inhibitor of metalloproteinase (TIMP-2) gene (24 h), and no significant effect was observed for these genes (data not shown), and an increase was observed for the IL-6 gene (Fig. 1E).

The gelatinolytic activity in the culture medium was shown to be inhibited by treatment of tumor cells with CAM (data not shown). We also examined the effects of two other antimicrobial agents, CTM and GM, on the expression of the MMP-9 gene (24 h), and no significant effect was observed for these agents (Fig. 2). We further examined the effect of CAM on the expression of the TIMP-1 or TIMP-2 gene (8) in 13762NF tumor cells. In the tumor cells, the TIMP-2 gene was shown to be expressed highly but the TIMP-1 gene was not (data not shown). As shown in Fig. 3, no significant change due to CAM (5 μg/ml) treatment was observed.

As a consequence, three patterns were observed after CAM treatment: suppression (MPP-9, TGF-β, and TNF-α), transient enhancement (IL-6), and no change (TIMP-2). It is difficult to explain why these different patterns appeared following CAM treatment. Such a modulatory effect was not observed for CTM or GM. We suppose that the effect observed in this study is specific to the macrolides, probably specific to the 14-membered ring macrolides. For example, Yatsunami et al. (Kyushu University School of Medicine) showed, in the B16 tumor-C57BL6 mouse model, that CAM and roxithromycin (14-membered ring) could inhibit angiogenesis but josamycin (16-membered ring) and azithromycin (15-membered ring) could not (personal communication). We have no explanation for the difference in action among the 14-, 15-, and 16-membered ring macrolides.

It is interesting that macrolide antibiotics cause a beneficial therapeutic outcome in hosts bearing cancer (3, 10, 13). We
FIG. 1. Effects of CAM (5 μg/ml) treatment time on expression of the MMP-9 (A), TGF-β (B), TNF-α (C), and IL-6 (D) genes. 13762NF tumor cells were treated with CAM at 5 μg/ml for different lengths of time, and then total RNAs were extracted for analysis. (E) Effect of CAM concentration on expression of the IL-6 gene. 13762NF tumor cells were treated with different concentrations of CAM for 24 h, and then total RNAs were extracted for analysis. GAPDH, glyceraldehyde-3-phosphate dehydrogenase.
hope that CAM may become a beneficial tool for the treatment of certain cancers.

REFERENCES


