

Publication

A constitutive decay element promotes tumor necrosis factor alpha mRNA degradation via an AU-rich element-independent pathway

JournalArticle (Originalarbeit in einer wissenschaftlichen Zeitschrift)**ID** 153474**Author(s)** Stoecklin, Georg; Lu, Min; Rattenbacher, Bernd; Moroni, Christoph**Author(s) at UniBasel** [Moroni, Christoph](#) ;**Year** 2003**Title** A constitutive decay element promotes tumor necrosis factor alpha mRNA degradation via an AU-rich element-independent pathway**Journal** Molecular and cellular biology**Volume** 23**Number** 10**Pages / Article-Number** 3506-15**Keywords** 1-Phosphatidylinositol 3-Kinase/metabolism; 3' Untranslated Regions; 3T3 Cells; Animals; Base Sequence; Blotting; Northern; Dactinomycin/pharmacology; Gene Deletion; Gene Expression Regulation; Genes; Reporter; Granulocyte-Macrophage Colony-Stimulating Factor/metabolism; Green Fluorescent Proteins; Humans; Luminescent Proteins/metabolism; Macrophages/metabolism; Mice; Mitogen-Activated Protein Kinases/metabolism; Models; Genetic; Molecular Sequence Data; Plasmids/metabolism; Point Mutation; RNA; Messenger/metabolism; Sequence Homology; Nucleic Acid; Time Factors; Transfection; Tumor Cells; Cultured; Tumor Necrosis Factor-alpha/*metabolism; p38 Mitogen-Activated Protein Kinases

Tumor necrosis factor alpha (TNF-alpha) expression is regulated by transcriptional as well as posttranscriptional mechanisms, the latter including the control of mRNA decay through an AU-rich element (ARE) in the 3' untranslated region (UTR). Using two mutant cell lines deficient for ARE-mediated mRNA decay, we provide evidence for a second element, the constitutive decay element (CDE), which is also located in the 3' UTR of TNF-alpha. In stably transfected RAW 264.7 macrophages stimulated with lipopolysaccharide (LPS), the CDE continues to target a reporter transcript for rapid decay, whereas ARE-mediated decay is blocked. Similarly, the activation of p38 kinase and phosphatidylinositol 3-kinase in NIH 3T3 cells inhibits ARE-mediated but not CDE-mediated mRNA decay. The CDE was mapped to an 80-nucleotide (nt) segment downstream of the ARE, and point mutation analysis identified within the CDE a conserved sequence of 15 nt that is required for decay activity. We propose that the CDE represses TNF-alpha expression by maintaining the mRNA short-lived, thereby preventing excessive induction of TNF-alpha after LPS stimulation. Thus, CDE-mediated mRNA decay is likely to be an important mechanism limiting LPS-induced pathologic processes.

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