



Activation of NFAT signal by p53-K120R mutant

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ARTICLE INFO

Article history:

Received 3 April 2009

Accepted 27 April 2009

Available online 3 May 2009

Edited by Varda Rotter

Keywords:

Nuclear factor of activated T-cell

p53

Luciferase reporter assay

Calcium signal

ABSTRACT

The tumor suppressor p53 is activated by phosphorylation and/or acetylation. We constructed 14 non-phosphorylated, 11 phospho-mimetic, and 1 non-acetylated point p53 mutations and compared their transactivation ability in U-87 human glioblastoma cells by the luciferase reporter assay. Despite mutations at the phosphorylation sites, only the p53-K120R and p53-S9E mutants had marginally reduced activities. On the other hand, the Nuclear factor of activated T-cells (NFAT)-luciferase reporter was more potently activated by p53-K120R than by wild-type p53 and other mutants in glioblastoma, hepatoma and esophageal carcinoma cells. This suggests that acetylation at Lys-120 of p53 negatively regulates a signaling pathway leading to NFAT activation.

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1. Introduction

The tumor suppressor protein p53 possesses various biological activities such as cell cycle arrest, induction of apoptosis and inhibition of angiogenesis by transactivating p21 [1], Bax [2], PUMA [3], Noxa [4], Bad [5], etc. The transactivation ability and the selectivity of the target genes of p53 are assumed to be regulated mainly by post-translational modifications including phosphorylation, acetylation, ubiquitination, methylation, neddylation and sumoylation [6]. Despite numerous reports on p53, the precise mechanism of selection of the target genes of activated p53 remains to be proven.

Nuclear factor of activated T-cells (NFAT) was first identified as a transcription factor in activated, but not in resting, T cells [7,8]. NFAT is dephosphorylated by calcineurin and translocated into the nucleus [9,10]. p53 also induces proline dehydrogenase 1 (PRODH1, proline oxidase, PIG6) [11,12] and proline dehydrogenase 2 (PRODH2, hydroxyproline oxidase) [13], both of which then activate NFAT pathway through reactive oxygen and Ca²⁺/calcineurin [14]. Although the NFAT pathway was suggested to be involved in induction of apoptosis [15], NFAT has an oncogenic role in pancreatic carcinoma through transcriptional activation of c-myc [16].

In the present study, we show that a non-acetylated mutant, p53-K120R, markedly activated NFAT.

2. Materials and methods

2.1. Plasmids

pCMV-p53WT, pCMV-p53V143A [17], reporter plasmids and pBV-PUMA-Luc [3] were provided by Dr. Bert Vogelstein (Howard Hughes Medical Institute). pGL3-Bax-Luc, pGL3-p21-Luc and pGL3-Bad-Luc [5] were provided by Dr. Mian Wu (University of Science and Technology of China). pGV-B2 Noxa-Luc [4] was provided by Dr. Nobuyuki Tanaka (Nippon Medical School). NFAT-Luc and SV40-Rluc were purchased from Promega (Madison, WI).

2.2. Cell culture

Human glioblastoma U-87, hepatocarcinoma HepG2, esophageal carcinoma YES-4 and bladder carcinoma T-24 cells were cultured in Dulbecco's modified Eagle's minimum essential medium supplemented with 10% fetal bovine serum.

2.3. Construction of p53 point mutants

p53 mutants were constructed from pCMV-p53 using KOD-Plus-Mutagenesis kit (Toyobo, Osaka, Japan). The primers used are listed in Table 1 (Supplementary data). All constructs were confirmed by base sequencing.

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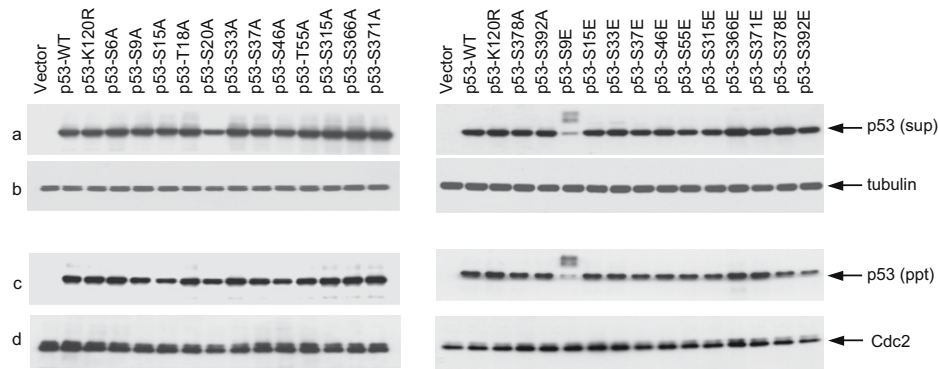


Fig. 1. Expression of p53 in p53-transfected cells. U-87 human glioblastoma cells were transfected with expression plasmids of pCMV-Neo-Bam (Vector), p53-WT or p53 mutants. Expression levels of p53 proteins in the cytoplasmic (A) and nuclear fractions (C) were examined by Western blotting analysis using anti-p53 (DO-1) with loading controls, anti- α/β -tubulin for the cytoplasmic fractions (B) and anti-Cdc2 antibodies for the nuclear fractions (D).

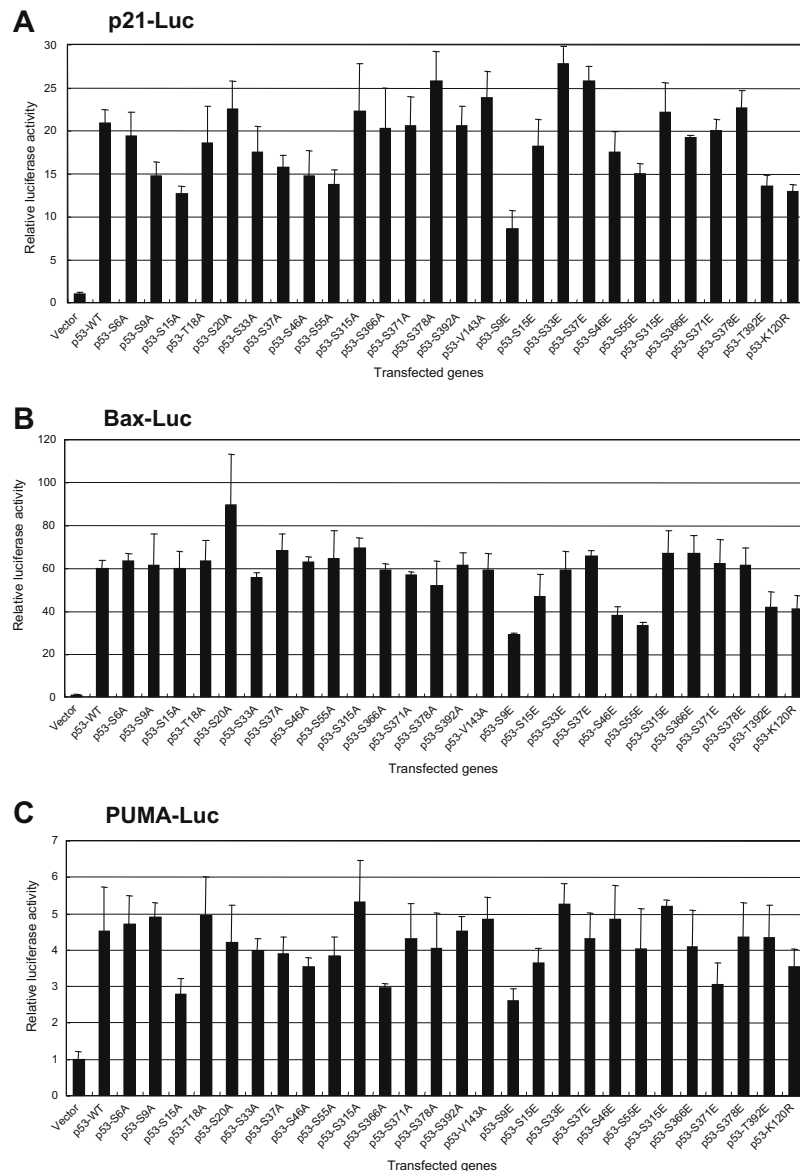


Fig. 2. Activation of p53 reporter plasmids by p53 mutants. U-87 glioblastoma cells (5×10^4 cells) were co-transfected with p53-responsive reporter plasmids (p21-Luc, Bax-Luc, PUMA-Luc, Noxa-Luc, Bad-Luc or NFAT-Luc; 100 ng), transfection standard SV40-Rluc (10 ng), and the expression plasmid (0.5 μ g) of p53-WT or p53 mutants, or control empty vector pCMV-Neo-Bam. Cells were harvested 48 h after transfection and luciferase activities in the cell extracts were measured. The error bars represent S.D. ($n = 3$).

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