

Supplemental Table-A1. Oligonucleotide Sequences of Primers for PCR

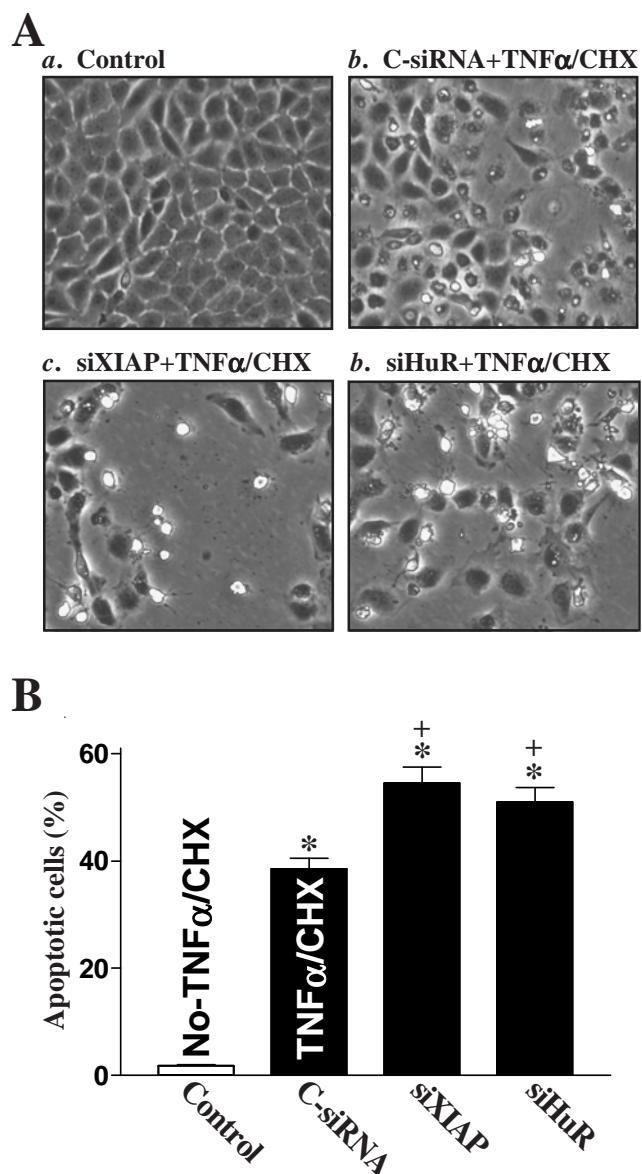
Fragment	Primers for Mapping XIAP Coding Region
Full length	F: (T7)5'-CAAGTTACATGTGAATTCTGCCAGGC-3' R: 5'-CGATCTGTGAAAGAG CAGGCTCTGTACTC-3'
F1	F: (T7)5'-TGACTTTAACAGTTGAAGGATCT-3' R: 5'-TTCCAACAGCTGAGTCTCCAT-3',
F2	F: (T7)5'-GACACAGGAGAATATCCCCAAA-3' R: 5'-ACTCTCTGGGGCTTAAATGG-3',
F3	F: (T7)5'-GCTAGTGGACTCTACTACACG-3' R: 5'-CTGAGTATAGCCATGTTCCAAAAG-3',
F4	F: (T7)5'-TGCAAGAGCTGGATTATG-3' R: 5'-TTCCAAGTGAATGGGTTA-3',
F5	F: (T7)5'-TCTGTGGTAAGAACTGCTGAAAAA-3' R: 5'-TCTTGTAGGCGCCTAGCTG-3',
F6	F: (T7)5'-GCTTGCAAAATCTGTATGGA-3' R: 5'-GCCTACTGT GGTGCTGGATT-3'

Fragment	Primers for Mapping XIAP 3'-UTR
Full length	F: (T7)TCCAGCACC ACAGTAGGCAT R: TGGTAGTCCACCATTAG
F1	F: (T7)TCCAGCACC ACAGTAGGCAT R: AGCTAAGATGGGCATTACAG
F2	F: (T7)CTGAATGCCGATCTTAGCT R: GTCAAGTACTACAAATAT
F3	F: (T7) GTACACTGACTTGT R: TCCCTGCTTATAACAGAACAC
F4	F: (T7) GTGTTCTGTATAAGCAGGGA R: GAATCCTAAAACACAGA
F5	F: (T7) GTGTTTAGGATTCTGTT R: TGGTAGTCCACCATTAG

Supplemental Table -A2. Potential HuR-hits in the XIAP Coding Region

331 atgactttta acagtttga aggatctaga
361 actgttgtac ctgcagacac caataaggat gaagaatttga tagaagagtt
411 Taatagatta aaaacatttgc ctaacttccc aagcagcagt cctgttcag
461 catcaacatt ggccggagcg gggtttctct acactggta aggagacacc
511 gtgcagtgtt tcagttgtca cgcggcagta gatagatggc agtatggaga
561 ctcagctgtt qgaagacaca ggagaatatac cccaaattgc agatttatca
611 atggttttta ttttggaaac ggtgccacac agtctacatc tcctggcatc
661 caaaaatggcc agtacaaaatc tgaaaactgt gtgggaaaca gaaatcatt
711 tgctcttgac aggccgtcg agactcatgc agattatctc ctgagaactg
761 gacaggtgt agatatttca gataccatat acccgaggaa cccggccatg
811 tgttagtgaag aagccagact gaagacgtt cagaactggc cagactatgc
861 ccatttaagc cccagagagt tagctgtgc tggactctac tacacgggga
911 ttgatgatca agtgcataatc tttgttgtg gtggaaact gaaaaattgg
961 gaacctgtg accgtgcctg gtcagagcac aggagacact ttcccaactg
1011 cttcttcgtt ttggggccgga atgttaatgt tcgaagttag tctgggtgtga
1061 gttcagatag gaatttcca aattcaacaa attctccaag aaatccagcc
1111 atggcagaat atgacgcacg gatcgttact tttggaacat ggctataactc
1161 agttaacaag gaggcgttgc caagactgg atttatgtc ttaggtgaag
1211 gtgataaagt gaagtgcttt cactgtggag gagggtcac ggattggaaag
1261 ccaagtgaag acccttggga acagcatgct aagtggatc cagggtgtaa
1311 atatctatttgc gatgagaagg gacaagaata tataaataat attcattttaa
1361 cccatttact tggggaaatct gtggtaagaa ctgctgaaaa aacaccatca
1411 gtaactaaaa aaatcgatga taccatcttgc cagaatcttgc tggcaaga
1461 agctatacga atgggatttca acttcaagga catcaagaaa acaatggaaag
1511 aaaagctcca aacatctgg agcaactatc tatcaacttgc ggttctgatt
1561 gcagatcttgc tgagtgcata gaaagataat tcgcaggatg agtcaagtca
1611 gacttcatttgc cagaaagaca tcagttacttgc agagcagacta aggccctac
1651 aagaggagaa gctttgcataa atctgtatgg atagaaatat tgctatagtt
1711 tttgcttt gtggacatct ggtcacttgtt aaacagtgtg cgaaagcagt
1761 tgacaaatgt cccatgtgttgc acacagtcat tacgttcaag caaaaatttt
1811 ttatgtcttaa

Computationally predicated hits of the HuR motif are indicated by underline in blue color



Supplemental Figure A1. Effects of XIAP silencing and HuR silencing on apoptotic sensitivity in normal IEC-6 cells (without DFMO). Cells were transfected with either siXIAP, siHuR, or C-siRNA for 48 h and then exposed to TNF α /CHX. (A) TNF α /CHX-induced apoptosis in controls and XIAP- or HuR-silenced populations: *a*), control cells; *b*) cells transfected with C-siRNA and then treated with TNF α /CHX; *c*), siXIAP-transfected cells treated with TNF α /CHX; *d*) siHuR transfected cells treated with TNF α /CHX. Apoptosis was measured by morphological analysis 4 h after treatment with TNF α /CHX. Original magnification, $\times 150$. (B) Percentage of apoptotic cells as described in A. Values are means \pm SE of data from six samples. * $P < 0.05$ compared with No-TNF α /CHX. + $P < 0.05$ compared with cells transfected with C-siRNA and then treated with TNF α /CHX for 4 h.