

# IV

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## CELLULAR PROTEINS

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p53 . . . . .	IV-2
Retinoblastoma . . . . .	IV-6
Deleted in Colon Cancer . . . . .	IV-9
E2F1 . . . . .	IV-11
Epidermal Growth Factor Receptor . . . . .	IV-13
ERBB2 . . . . .	IV-16
K-ras . . . . .	IV-19
Monocyte Chemoattractant Protein . . . . .	IV-22
Nuclear Factor 1 . . . . .	IV-24
Oct1 . . . . .	IV-26
Platelet Derived Growth Factor Receptor . . . . .	IV-28
PP2A 72kDa Regulatory Subunit . . . . .	IV-31
PP2A Catalytic Subunit . . . . .	IV-33
PP2A PR55 Regulatory Subunit . . . . .	IV-35
PP2A PR65 Regulatory Subunit . . . . .	IV-37
Transcriptional Enhancer Factor . . . . .	IV-39
Transforming Growth Factor . . . . .	IV-41
Tumor Necrosis Factor . . . . .	IV-42
YY1 . . . . .	IV-45
c-myc . . . . .	IV-47
Cyclin A . . . . .	IV-51
Glucocorticoid Receptor . . . . .	IV-52
junB . . . . .	IV-54
Retinoblastoma-related p107 . . . . .	IV-56

LOCUS p53-php53c1 1317 bp ss-mRNA PRI 22-OCT-1992  
 DEFINITION Human p53 cellular tumor antigen mRNA, complete cds.  
 ACCESSION X02469 M60950  
 KEYWORDS antigen; tumor antigen.  
 SOURCE Human, cDNA to mRNA, clone php53c1 from a cDNA library of SV40 transformed fibroblasts.  
 REFERENCE 1 (bases 1 to 1317)  
 AUTHORS Zakut-Houri,R., Bienz-Tadmor,B., Givol,D. and Oren,M.  
 TITLE Human p53 cellular tumor antigen: cDNA sequence and expression in COS cells  
 JOURNAL EMBO J. 4, 1251-1255 (1985)  
 REFERENCE 2 (bases 1 to 1317)  
 AUTHORS Rideout,W.M.III., Coetzee,G.A., Olumi,A.F. and Jones,P.A.  
 TITLE 5-methylcytosine as an endogenous mutagen in the human ldl receptor and p53 genes  
 JOURNAL Science 249, 1288-1290 (1990)  
 REFERENCE 3 (M14694 and M14695 shown in alignment below)  
 AUTHORS Harris,N., Brill,E., Shohat,O., Prokocimer,M., Wolf,D., Arai,N. and Rotter,V.  
 TITLE Molecular basis for heterogeneity of the human p53 protein  
 JOURNAL Mol. Cell. Biol. 6, 4650-4656 (1986)  
 REFERENCE 4 (K031991 shown in alignment below)  
 AUTHORS Harlow,E., Williamson,N.M., Ralston,R., Helfman,D.M. and Adams,T.E.  
 TITLE Molecular cloning and in vitro expression of a cDNA clone for human cellular tumor antigen p53  
 JOURNAL Mol. Cell. Biol. 5, 1601-1610 (1985)  
 COMMENT Clone php53c1 is the wild-type form of p53, isolated from an SV40 transformed fibroblast cell line. It was first thought that p53, a cell-cycle regulatory protein, was an oncogene. This has since been disproven, for it has been shown that the clones that were first isolated and tested were mutants. In fact, the p53 gene is a tumor suppressor. The mutated forms are thought to be trans-dominant over the wild-type p53. The mutations cause a conformational change in the protein which facilitates its binding to the heat shock protein hsp70. The two forms of p53 and hsp70 become bound together in long-lived unproductive complexes. Since these complexes aggregate in the cytoplasm, the nucleus is deprived of p53. With the negative regulatory effect of p53 reduced or eliminated, the cell is more easily able to proliferate. However, a different mechanism of inactivation is employed by the viral oncoproteins: the large T-antigen of SV40 and the E1B 55kD protein of adenovirus. Instead of the hsp70, mutant p53 complex trapping the wild-type p53 in inactive structures it is the viral oncoproteins that sequester the wild type p53 and inactivate it. Human papillomavirus uses yet another mechanism. The E6 protein of the oncogenic HPV types bind wild-type p53 and stimulate its destruction; although at this point there is no evidence that E6 proteins of noncogenic HPV types, such as HPV-6 or HPV-11, can associate with wild-type p53 in vivo (Werness et al. Science 248, 76-79, Scheffner et al. Cell 63, 1129-1136). It is however possible that the binding assay used to determine this fact was not sensitive enough to detect low affinity binding. The degradation of p53 by HPV E6 is ATP dependant and the ubiquitin-dependant protease system has been suggested to be involved. Since the large T antigen of SV40 also complexes with wild-type p53 and does not target it for degradation, there must be an additional signaling mechanism not yet elucidated involved in the interaction with the HPV E6 protein.  
  
 The p53 wild type php53c1 coding region and four variants are shown in the alignment below. The p53 coding regions which are represented by the accession numbers M14694 and M14695 have both been isolated from the human transformed cell line SV-80. The two forms differ from one another by a single base pair substitution and consequently accounts for the change in electrophoretic mobility of the two proteins. Harris et al. believe that this heterogeneity is due to gene polymorphism.

The coding region represented by accession number K03199 has been isolated from the human vulva carcinoma cell line A431.

BASE COUNT 295 a 408 c 352 g 262 t
ORIGIN 2 bp upstream of XbaI site; chromosome 17p13.
1 gtctagagcc accgtccagg gaggcaggtag ctgctgggct ccggggacac ttgctgttcg
61 ggctgggagc gtgctttcca cgacgggtgac acgcttccct ggattggcag ccagactgcc
121 ttccgggtca ctgccATGga ggagccgcag tcagatccta gctcagacc ccctctgagt
p53 start ->
181 caggaaacat tttcagacct atggaaacta cttcctgaaa acaacgttct gtcccccttg
241 ccgtcccaag caatggatga tttgatgctg tccccggacg atattgaaca atggttcact
301 gaagaccag gtccagatga agctcccaga atgccagagg ctgctcccc cgtGGCCCCCT
->
361 GCACcagcag ctctacacc ggcGGCCCCCT GCACcagccc cctcctggcc cctgtcatct
<- direct rpt -> <- direct rpt
421 tctgtccctt cccagaaaac ctaccagggc agctacggtt tccgtctggg cttcttgcac
481 tctgggacag ccaagtctgt gacttgcacg tactccccctg ccctcaacaa gatgttttgc
541 caactggcca agacctgccc tgtgcagctg tgggttgatt ccacacccc gcccggcacc
601 cgcgtccgcg ccatggccat ctacaagcag tcacagcaca tgacggagggt tgtgaggcgc
661 tgccccacc atgagcgctg ctcatagatg gatggtctgg ccctcctca gcatcttata
721 cgagtggag gaaatttgcg tgtggagtat ttggatgaca gaaacacttt tcgacatagt
781 gtggtggtgc cctatgagcc gcctgagggt ggctctgact gtaccacat ccaactacaac
841 tacatgtgta acagttcctg catggcgggc atgaaccgga ggccatcct caccatcatc
901 acactggaag actccagtgg taatctactg ggacggaaca gctttgagggt gcgtgtttgt
961 gcctgtcctg ggagagaccg gcgcacagag gaagagaatc tccgcaagaa aggggagcct
1021 caccacgagc tgcccccagg gagcactaag cgagcactgc ccaacaacac cagctcctct
1081 cccagccaa agaagaacc actggatgga gaatatttca cccttcagat ccgtgggctg
1141 gagcgcttcg agatgttccg agagctgaat gaggccttgg aactcaagga tgcccaggct
1201 gggaggagc caggggggag cagggtcac tccagccacc tgaagtcaa aaagggtcag
1261 tctacctccc gccataaaaa actcatgttc aagacagaag ggctgactc agactGA
<- p53 end

//
P53.WTPHP53C1 ATGGAGGAGCCGCAGTCAGATCCTAGCGTCGAGCCCCCTCTGAGTCAGGAAACATTTTCAGACCTATGGA 70
p53.X54156 ----- 70
p53.M14695 ----- 70
p53.M14694 ----- 70
p53.K03199 ----- 70
P53.WTPHP53C1 AACTACTTCTGAAAACAACGTTCTGTCCCCCTTGCCGTCCCAAGCAATGGATGATTTGATGCTGTCCCC 140
p53.X54156 ----- 140
p53.M14695 ----- 140
p53.M14694 ----- 140
p53.K03199 ----- 140
P53.WTPHP53C1 GGACGATATTGAACAATGGTTCACTGAAGACCAGGTCCAGATGAAGCTCCAGAAATGCCAGAGGCTGCT 210
p53.X54156 ----- 210
p53.M14695 ----- 210
p53.M14694 ----- 210
p53.K03199 ----- 210
P53.WTPHP53C1 CCCCCGTGGCCCCCTGCACCAGCAGCTCCTACACCGGCGGCCCTGCACCAGCCCCCTCCTGGCCCCCTGT 280
p53.X54156 ----G----- 280
p53.M14695 -----GA----- 280
p53.M14694 ----G-----GA----- 280
p53.K03199 ----- 280
P53.WTPHP53C1 CATCTTCTGTCCCTTCCCAGAAAACCTACCAGGGCAGCTACGGTTTCCGTCTGGGCTTCTTGCATTCTGG 350
p53.X54156 ----- 350
p53.M14695 ----- 350
p53.M14694 ----- 350
p53.K03199 ----- 350
P53.WTPHP53C1 GACAGCCAAGTCTGTGACTTGCACGTACTCCCCTGCCCTCAACAAGATGTTTTGCCAACTGGCCAAGACC 420
p53.X54156 ----- 420
p53.M14695 ----- 420
p53.M14694 ----- 420
p53.K03199 ----- 420

P53 .WTPHP53C1	TGCCCTGTGCAGCTGTGGGTTGATTCCACACCCCCGCCGGCACCCGCGTCCGCGCCATGGCCATCTACA	490
p53 .X54156	-----	490
p53 .M14695	-----	490
p53 .M14694	-----	490
p53 .K03199	-----	490
P53 .WTPHP53C1	AGCAGTCACAGCACATGACGGAGGTTGTGAGGCGCTGCCCCACCATGAGCGCTGCTCAGATAGCGATGG	560
p53 .X54156	-----	560
p53 .M14695	-----	560
p53 .M14694	-----	560
p53 .K03199	-----	560
P53 .WTPHP53C1	TCTGGCCCCCTCCTCAGCATCTTATCCGAGTGAAGGAAATTTGCGTGTGGAGTATTTGGATGACAGAAAC	630
p53 .X54156	-----	630
p53 .M14695	-----	630
p53 .M14694	-----	630
p53 .K03199	-----	630
P53 .WTPHP53C1	ACTTTTCGACATAGTGTGGTGGTGCCTATGAGCCGCTGAGGTTGGCTCTGACTGTACCACCATCCACT	700
p53 .X54156	-----	700
p53 .M14695	-----	700
p53 .M14694	-----	700
p53 .K03199	-----	700
P53 .WTPHP53C1	ACAACCTACATGTGTAACAGTTCCTGCATGGGCGGCATGAACCGGAGGCCATCCTCACCATCATCACACT	770
p53 .X54156	-----	770
p53 .M14695	-----	770
p53 .M14694	-----	770
p53 .K03199	-----	770
P53 .WTPHP53C1	GGAAGACTCCAGTGGTAATCTACTGGGACGGAACAGCTTTGAGGTGCGTGTGTTGTGCCTGTCTGGGAGA	840
p53 .X54156	-----	840
p53 .M14695	-----	840
p53 .M14694	-----	840
p53 .K03199	-----A-----	840
P53 .WTPHP53C1	GACCGGCGCACAGAGGAAGAGAATCTCCGCAAGAAAGGGAGCCTCACCACGAGCTGCCCCAGGGAGCA	910
p53 .X54156	-----	910
p53 .M14695	-----	910
p53 .M14694	-----	910
p53 .K03199	-----	910
P53 .WTPHP53C1	CTAAGCGAGCACTGCCCAACAACACCAGCTCCTCTCCCCAGCCAAAGAAGAAACCACTGGATGGAGAATA	980
p53 .X54156	-----	980
p53 .M14695	-----	980
p53 .M14694	-----	980
p53 .K03199	-----	980
P53 .WTPHP53C1	TTTCACCCCTCAGATCCGTGGGCGTGAGCGCTTCGAGATGTTCCGAGAGCTGAATGAGGCCTTGGAACCTC	1050
p53 .X54156	-----	1050
p53 .M14695	-----	1050
p53 .M14694	-----	1050
p53 .K03199	-----	1050
P53 .WTPHP53C1	AAGGATGCCAGGCTGGGAAGGAGCCAGGGGGAGCAGGGCTCACTCCAGCCACCTGAAGTCCAAAAAGG	1120
p53 .X54156	-----	1120
p53 .M14695	-----	1120
p53 .M14694	-----	1120
p53 .K03199	-----	1120
P53 .WTPHP53C1	GTCAGTCTACCTCCCGCCATAAAAACTCATGTTCAAGACAGAAGGGCCTGACTCAGACTGA	1182
p53 .X54156	-----	1182
p53 .M14695	-----	1182
p53 .M14694	-----	1182
p53 .K03199	-----	1182

LOCUS Rb 4740 bp ss-mRNA PRI 15-JUN-1989  
 DEFINITION Human retinoblastoma susceptibility mRNA, complete cds.  
 ACCESSION M15400  
 KEYWORDS retinoblastoma susceptibility.  
 SOURCE Human fetal retina, cDNA to mRNA, clone RB-[1,5].  
 REFERENCE 1 (bases 243 to 4740)  
 AUTHORS Lee,W.H., Bookstein,R., Hong,F., Young,L.J., Shew,J.Y. and Lee,E.  
 TITLE Human retinoblastoma susceptibility gene: Cloning, identification,  
 and sequence  
 JOURNAL Science 235, 1394-1399 (1987)  
 REFERENCE 2 (bases 1 to 480)  
 AUTHORS Lee,E., Bookstein,R., Young,L.J., Lin,C.-J., Rosenfeld,M.G. and  
 Lee,W.H.  
 TITLE Molecular mechanism of retinoblastoma gene inactivation in  
 retinoblastoma cell line Y79  
 JOURNAL Proc. Natl. Acad. Sci. U.S.A. 85, 6017-6021 (1988)  
 COMMENT Draft entry and computer-readable copy of sequence in [1] kindly  
 provided by R.Bookstein, 27-APR-1987.

The authors [1] identified the retinoblastoma susceptibility (RB) gene encoding a messenger RNA (mRNA) of 4.6 kb on the basis of chromosomal location, homozygous deletion, and tumor-specific alterations in expression. Transcription of the gene was abnormal in 6/6 retinoblastomas while normal in fetal retina and placenta. The sequence presented in this entry is derived from human fetal retina cDNA. The authors [2] also examined cDNA from the retinoblastoma cell line Y79. They found that the mRNA produced was a shortened transcript which reflects a deletion of exons 2-6 and must have occurred by a mechanism other than recombination of homologous sequences.

The Retinoblastoma (Rb) susceptibility gene maps to chromosome 13q14 and encodes a nuclear phosphoprotein of 105kDa. Rb is a tumor suppressor gene; its nuclear location and ability to bind DNA suggests a role in transcriptional regulation. On the side of the protein lies a complementary oncoprotein-binding pocket that is a common target for the three viral proteins: E1A of adenovirus, large T antigen of SV40, and E7 of HPV. Munger et al. (EMBO 8, 4099-4105) determined that the high risk HPV types such as HPV16 and HPV18 bind to pRB with higher affinity than low risk types 6b and 11. The amino acid locations of the pRB binding site of the HPV16 E7 protein have been mapped to a small stretch of amino acids. In addition, pRB has shown to associate with a number of cellular host proteins at the onco-protein binding site. pRB shows differential phosphorylation at different points within the cell cycle. All three of the viral proteins mentioned above interact with the underphosphorylated form of pRB.

BASE COUNT 1508 a 887 c 862 g 1483 t

ORIGIN 1038 bp upstream of EcoRI site; chromosome 13q14.1-q14.2.

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1 ttccggtttt tctcagggga cgttgaaatt atttttgtaa cgggagtcgg gagaggacgg
61 ggcgtgcccc gcgtgcgcgc gcgtcgctct ccccggcgct cctccacagc tcgctggctc
121 ccgcccgcgga aaggcgtcat gccgccaaa acccccgaa aaacggccgc caccgccgcc
181 gctgccgccc cggaaccccc ggcaccgccc ccgcccgcc ctctgagga ggaccacagag
241 caggacagcg gcccgaggga cctgcctctc gtcaggcttg agtttgaaga aacagaagaa
301 cctgatttta ctgcattatg tcagaaatta aagataccag atcatgtcag agagagagct
361 tggttaactt gggagaaagt ttcactctgt gatggagat tgggaggtta tattcaaaa
421 aaaaaggaac tgtggggaat ctgtatcttt attgcacgag ttgacctaga tgagatgtcg
481 ttcactttac tgagctacag aaaaacatac gaaatcagtg tccataaatt ctttaactta
541 ctaaaagaaa ttgataccag taccaaagtt gataatgcta tgtcaagact gttgaagaag
601 tatgatgtat tgtttgcact cttcagcaaa ttggaaagga catgtgaact tatatatattg
661 acacaacca gcagttcgat atctactgaa ataaattctg cattgggtgct aaaagtttct
721 tggatcacat ttttattagc taaaggggaa gtattacaaa tggagatga tctggtgatt
781 tcatttcagt taatgctatg tgtccttgac tattttatta aactctcacc tcccatggtg

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## Retinoblastoma

841 ctcaagaac catataaac agctgttata cccattaatg gttcacctcg aacacccagg  
901 cgaggtcaga acaggagtgc acggatagca aaacaactag aaaatgatac aagaattatt  
961 gaagttctct gtaaagaaca tgaatgtaat atagatgagg tgaaaaatgt ttatttcaaa  
1021 aattttatac cttttatgaa ttctcttggg cttgtaacat ctaatggact tccagagggt  
1081 gaaaaatctt ctaaacgata cgaagaat tttcttaaaa ataaagatct agatcgaaga  
1141 ttatttttgg atcatgataa aactcttcag actgattcta tagacagttt tgaaacacag  
1201 agaacaccac gaaaaagtaa ccttgatgaa gaggtgaata taattcctcc acacactcca  
1261 gtttaggactg ttatgaacac tatccaacaa ttaatgatga ttttaaattc tgcaagtgat  
1321 caaccttcag aaaatctgat ttctatttt aacaactgca cagtgaatcc aaaagaaagt  
1381 atactgaaaa gagtgaagga tataggatac atctttaaag agaaatgtgc taaagctgtg  
1441 ggacaggggt gtgtcgaat tggatcacag cgatacaaac ttggagttcg cttgtattac  
1501 cgagtaatgg aatccatgct taaatcagaa gaagaacgat tatccattca aaattttagc  
1561 aaactctctg atgacaacat ttttcatatg tctttattgg cgtgcgctct tgaggttgta  
1621 atggccacat atagcagaag tacatctcag aatcttgatt ctggaacaga tttgtctttc  
1681 ccatggattc tgaatgtgct taatttaaaa gcctttgatt tttacaaagt gatcgaaggt  
1741 tttatcaaag cagaaggcaa cttgacaaga gaaatgataa aacattttaga acgatgtgaa  
1801 catcgaatca tggaaatcct tgcattggctc tcagattcac ctttatttga tcttattaaa  
1861 caatcaaagg accgagaagg accaactgat caccttgaat ctgcttgcct tcttaattct  
1921 cctctccaga ataactcacac tgcagcagat atgtatcttt ctctgtgtaag atctccaaag  
1981 aaaaaagggt caactacgcy tgtaaatctt actgcaaatg cagagacaca agcaacctca  
2041 gccttccaga cccagaagcc attgaaatct aactctcttt cactgtttta taaaaagtg  
2101 tatcggctag cctatctcgc gctaaatata ctttgtgaac gcctctctgc tgagcacca  
2161 gaattagaac atatcatctg gacccttttc cagcacaccc tgcagaatga gtatgaactc  
2221 atgagagaca ggcatttggg ccaaatatg atgtgttcca tgtatggcat atgcaaagtg  
2281 aagaatctga accttaaat caaaatcatt gtaacagcat acaaggatct tctcatgct  
2341 gttcaggaga cattcaaacg tgttttgatc aaagaagagg agtatgattc tattatagta  
2401 ttctataact cggctttcat gcagagactg aaaacaaata ttttgcagta tgcttccacc  
2461 aggccccccta ccttgtcacc aatacctcac attcctcgaa gcccttacia gtttctagt  
2521 tcacccttac ggattcctgg agggaaacatc tatatttcac ccctgaagag tccatataaa  
2581 atttcagaag gtctgccaac accaacaaaa atgactccaa gatcaagaat cttagtatca  
2641 atttggtgaat cattcgggac ttctgagaag ttccagaaaa taaatcagat ggtatgtaac  
2701 agcagacctg tgctcaaaag aagtgcgtaa ggaagcaacc ctctcaaac actgaaaaaa  
2761 ctacgctttg atattgaagg atcagatgaa gcagatggaa gtaaacatct cccaggagag  
2821 tccaaatttc agcagaaact ggcagaaatg acttctactc gaacacgaat gcaaaagcag  
2881 aaaatgaatg atagcatgga tacctcaaac aaggaagaga aatgaggatc tcaggacctt  
2941 ggtggacact gtgtacacct ctggattcat tgtctctcac agatgtgact gtataacttt  
3001 cccaggttct gtttatggcc acatttaata tcttcagctc tttttgga tataaaatgt  
3061 gcagatgcaa ttgtttgggt gagtccctaa ccactgaaa tgttagtcat tgttattat  
3121 acaagattga aaatcttgyt taaatctgcy catttaaaaa gttgtagcag attgtttcct  
3181 cttccaaagt aaaattgctg tgctttatgg atagtaagaa tggccctaga gtgggagctc  
3241 tgataaccga ggctgtctg actactttgc cttcttttgt agcatatagg tgatgtttgc  
3301 tcttgttttt attaatatt atgtatattt ttttaattta acatgaacac ccttagaaaa  
3361 tgtgtcctat ctatcttcca aatgcaattt gattgactgc ccattcaca aaattatcct  
3421 gaactcttct gcaaaaatgg atattatag aaattagaaa aaaattacta attttaca  
3481 ttagatttta ttttactatt ggaatctgat atactgtgtg cttgttttat aaaattttgc  
3541 ttttaattaa ataaaagctg gaagcaaagt ataaccat atgatactatca tactactgaa  
3601 acagatttca tacctcagaa tgtaaaagaa cttactgatt attttcttca tccaacttat  
3661 gtttttaaat gaggattatt gatagtaactc ttggttttta taccattcag atcactgaat  
3721 ttataaagta cccatctagt acttgaaaaa gtaaagtgtt ctgccagatc ttaggtatag  
3781 aggaccctaa cacagtatat cccaagtgca ctttctaaty tttctgggct ctgaagaatt  
3841 aagatacaaa ttaattttac tccataaaca gactgttaat tataggagcc ttaatttttt  
3901 tttcatagag atttgtctaa ttgcatctca aaattattct gccctcctta atttgggaag  
3961 gtttgtgttt tctctggaat ggtacatgct tccatgtag cttttgaact ggcaattgtc  
4021 tatttatctt ttattttttt aagtcagtat ggtctaacac tggcatgttc aaagccacat  
4081 tatttctagt ccaaaattac aagtaatcaa gggctattat ggggttaggca ttaatgtttc  
4141 tatctgattt tgygcaaaag cttcaaatta aaacagctgc attagaaaa gaggcgcttc  
4201 tccctcccc tacacctaaa ggtgtattta aactatcttg tgtgattaac ttatttagag  
4261 atgctgtaac ttaaaatagg ggatatttaa ggtagcttca gctagctttt aggaaaatca  
4321 ctttgtctaa ctcagaatta tttttaaaaa gaaatctggt cttgttagaa acaaaaattt  
4381 tattttgtgc tcatttaagt tccaaactta ctattttgac agttattttg ataacaatga  
4441 cactagaaaa cttgactcca tttcatcatt gtttctgcat gaatatcata caaatcagtt  
4501 agtttttagg tcaagggctt actatttctg ggtcttttgc tactaagttc acattagaat  
4561 tagtgccaga attttaggaa cttcagagat cgtgtattga gatttcttaa ataatgcttc  
4621 agatattatt gctttattgc ttttttgtat tggttaaaa cgtacattta aaattgctat

4681 gttactatatt tctacaatta atagtttgc tattttaaaa taaattagtt gttaagagtc

## Deleted in Colon Cancer

LOCUS DCC 4608 bp ss-mRNA PRI 04-FEB-1994  
DEFINITION Homo sapiens DCC mRNA.  
ACCESSION X76132  
KEYWORDS cell adhesion molecule; DCC gene; immunoglobulin gene superfamily;  
transmembrane protein; tumor suppressor gene.  
SOURCE Human normal adult and fetal brain, Clontech cDNA libraries  
CAT #HL1065b, HL1003b.  
REFERENCE 1 (bases 1 to 4608)  
AUTHORS Cho, K.  
TITLE ;  
JOURNAL direct submission  
REFERENCE 2 (bases 1 to 4608)  
AUTHORS Hedrick L., Cho K.R., Fearon E.R., Wu T.C., Kinzler K.W.,  
Vogelstein B.  
TITLE The DCC gene product in cellular differentiation and colorectal  
tumorigenesis  
JOURNAL Unpublished  
COMMENT Tumors were induced in nude mice after transplantation of the  
HPV-18 immortalized human keratinocyte cell line (1811) and  
treatment with the carcinogen nitrosomethylurea (NMU). In these  
transformed cells, one allele of the deleted in colon cancer DCC  
tumor-suppressor gene was absent and the other allele was mutated  
(Klingelhutz et al. Oncogene 8: 95-9).  
Related sequence: M32292  
BASE COUNT 1282 a 1179 c 1058 g 1089 t  
ORIGIN  
1 atggagaata gtcttagatg tgtttgggta cccaagctgg cttttgtact cttcggagct  
61 tccttgctca gcgcgcatct tcaagtaacc ggttttcaaa ttaaagcttt cacagcactg  
121 cgcttcctct cagaaccttc tgatgcccgc acaatgccgg gaggaaatgt cctcctcgac  
181 tgctccgcgg agtccgaccg aggagttcca gtgatcaagt ggaagaaaga tggcattcat  
241 ctggccttgg gaatggatga aaggaagcag caactttcaa atgggtctct gctgatataca  
301 aacatacttc attccagaca ccacaagcca gatgagggac tttaccaatg tgaggcatct  
361 ttaggagatt ctggctcaat tattagtcgg acagcaaaag ttgcagtgc aggaccactg  
421 aggttccttt cacagacaga atctgtcaca gccttcattg gagacacagt gctactcaag  
481 tgtgaagtca ttggggagcc catgcccaaca atccactggc agaagaacca acaagacctg  
541 actccaatcc caggtgactc ccgagtggtg gtcttgccct ctggagcatt gcagatcagc  
601 cgactccaac cgggggacat tggaaattac cgatgctcag ctcgaaatcc agccagctca  
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**E2F1**

LOCUS E2F1 2517 bp ss-mRNA PRI 10-AUG-1992  
 DEFINITION Homo sapiens (E2F-1) pRB-binding protein mRNA, complete cds.  
 ACCESSION M96577  
 KEYWORDS DNA-binding protein; pRB-binding protein; transcription factor E2F.  
 SOURCE Homo sapiens fetal brain cDNA to mRNA.  
 REFERENCE 1 (bases 1 to 2517)  
 AUTHORS Helin,K., Lees,J.A., Vidal,M., Dyson,N.J., Harlow,E. and Fattaey,A.  
 TITLE A cDNA encoding a pRB-binding protein with properties of the  
 transcription factor E2F  
 JOURNAL Cell 70, 337-350 (1992)  
 COMMENT E2F, a cellular transcription factor, forms an inactive complex with  
 pRB (retinoblastoma tumor suppressor protein). HPV-16 E7 is one of  
 several DNA virus oncoproteins that dissociates the E2F-pRB complex.  
 This dissociation is coupled with enhanced E2F-dependent transcription.  
 In the S-phase of the cell cycle, E2F complexes with cyclin A, p107,  
 and cdk2 kinase. Arroyo et al. (Mol and Cell Bio 13: 6537-46) report  
 that the E7 protein of HPV-16 associates with the E2F-cyclin A complex.  
 Arroyo et al. (Mol and Cell Bio 13: 6537-46) further note that a high  
 risk HPV has a higher binding efficiency to the E2F-cyclin A complex  
 than the low-risk HPV type. Pagano et al. (Oncogene 7: 1681-6) report  
 that the E2F-cyclin A complex can occur in HPV-18 infected cells  
 independent of pRB.  
 BASE COUNT 454 a 784 c 781 g 498 t  
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## Epidermal Growth Factor Receptor

LOCUS EGFR 5532 bp ss-mRNA PRI 13-JUN-1985  
DEFINITION Human mRNA for precursor of epidermal growth factor receptor  
ACCESSION X00588  
KEYWORDS epidermal growth factor receptor; signal peptide.  
SOURCE  
REFERENCE 1 (bases 1 to 5532)  
AUTHORS Ullrich A., Coussens L., Hayflick J.S., Dull T.J., Gray A.,  
Tam A.W., Lee J., Yarden Y., Libermann T.A., Schlessinger J.,  
Downward J., Mayes E.L., Whittle N., Waterfield M.D., Seeburg P.H.;  
TITLE Human epidermal growth factor receptor cDNA sequence and aberrant  
expression of the amplified gene in A431 epidermoid carcinoma  
cells  
JOURNAL Nature 309:418-425(1984).  
COMMENT Cohen et al. demonstrated that the BPV E5 protein activates the  
EGF receptor through complex formation. They have further  
determined that this activation is specific to the cytoplasmic  
domain of the EGF receptor and is ligand independent (J Virol  
67: 5303-11). Straight et al. (J Virol 67: 4521-32) demonstrated  
that E5 HPV-16 infected keratinocytes had a two to five fold  
increase in EGF receptors.

BASE COUNT 1472 a 1484 c 1337 g 1239 t

### ORIGIN

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## Epidermal Growth Factor Receptor

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**ERBB2**

LOCUS ERBB2 4473 bp ss-mRNA PRI 30-JUN-1987  
 DEFINITION Human c-erb-B-2 mRNA.  
 ACCESSION X03363  
 KEYWORDS c-myc proto-oncogene; cell surface glycoprotein; erbB gene;  
 erbB oncogene; glycoprotein; growth factor; growth factor receptor;  
 kinase; neu oncogene; transmembrane protein; tyrosine kinase.  
 SOURCE Homo sapiens cell line MKN-7  
 REFERENCE 1 (bases 1 to 4473)  
 AUTHORS Yamamoto,T., Ikawa,S., Akiyama,T., Semba,K., Nomura,N.,  
 Miyajima,N., Saito,T. and Toyoshima,K.  
 TITLE Similarity of protein encoded by the human c-erb-B-2 gene to  
 epidermal growth factor receptor  
 JOURNAL Nature 319, 230-234 (1986)  
 COMMENT Milde-Langosch et al. (Verh Dtsch Ges Pathol 75: 363-5)  
 demonstrated an increase in expression of c-erbB2 in  
 4/11 HPV-positive endometrial carcinomas and in 3/19  
 HPV-positive CIN3 lesions.

The c-erb-B-2 protein shows similarity to the epidermal growth factor receptor.

EMBL features not translated to GenBank features:

key	from	to	description
SITE	376	384	pot. glycosylation site
SITE	544	558	pot. glycosylation site
SITE	733	741	pot. glycosylation site
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SITE	1762	1770	pot. glycosylation site
SITE	1885	1893	pot. glycosylation site
SITE	2059	2067	pot. glycosylation site
SITE	2353	3132	aa 727-986, seq. homologous to EGF receptor kinase domain
SITE	2446	2454	pot. glycosylation site
SITE	4455	4460	put. polyA signal
POLYA	4473	4473	polyA site

BASE COUNT 902 a 1383 c 1329 g 859 t

ORIGIN

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# K-ras

LOCUS KRAS 5775 bp ds-DNA PRI 05-JUN-1991  
DEFINITION Human K-ras oncogene, complete cds.  
ACCESSION M54968 M38506  
KEYWORDS K-ras oncogene.  
SOURCE Human tumor DNA.  
REFERENCE 1 (bases 1 to 5775)  
AUTHORS Kahn,S., Yamamoto,F.-I., Almoguera,C., Winter,E., Forrester,K.,  
Jordano,J. and Perucho,M.  
TITLE The c-K-ras gene and human cancer (review)  
JOURNAL Anticancer Res. 7, 639-652 (1987)  
BASE COUNT 1739 a 974 c 1105 g 1957 t  
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## **K-ras**

5761 taaatcatta ccagg

## Monocyte Chemoattractant Protein

LOCUS MCP1 3227 bp ds-DNA PRI 15-JUN-1994  
DEFINITION Human MCP-1 gene for monocyte chemoattractant protein-1.  
ACCESSION D26087  
KEYWORDS monocyte chemoattractant protein-1.  
SOURCE Homo sapiens (library: EMBL3 SP6/T7) placenta DNA.  
REFERENCE 1 (sites)  
AUTHORS Rollins,B.J., Stier,P., Ernst,T. and Wong,G.G.  
TITLE The human homolog of the JE gene encodes a monocyte secretory protein  
JOURNAL Mol. Cell. Biol. 9, 4687-4695 (1989)  
REFERENCE 2 (sites)  
AUTHORS Shyy,Y.-J., Li,Y.-S. and Kolattukudy,P.E.  
TITLE Structure of human monocyte chemotactic protein gene and its regulation by TPA  
JOURNAL Biochem. Biophys. Res. Commun. 169, 346-351 (1990)  
REFERENCE 3 (bases 1 to 3227)  
AUTHORS Ueda,A., Okuda,K., Ohno,S., Shirai,A., Igarashi,T., Matsunaga,K., Fukushima,J., Kawamoto,S., Ishigatsubo,Y. and Okubo,T.  
TITLE NF-kB and Sp1 regulate transcription of human monocyte chemoattractant protein-1 gene  
JOURNAL J. Immunol. (1993) In press  
COMMENT Rosl et al. (J Virol 68: 2142-50) demonstrated that MCP-1 protein is not expressed in HPV-18 positive cervical carcinoma cells, despite the fact that the gene which encodes it, the JE gene, has no mutations. Conversely, in non-malignant cells the tumor necrosis factor alpha induces MCP-1 expression which results in HPV-18 repression.

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3-9 Fukuura, Kanazawa-ku  
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Japan  
Phone: 045-787-2630  
Fax: 045-786-3444.

NCBI gi: 516772

BASE COUNT 850 a 793 c 732 g 852 t  
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## Monocyte Chemoattractant Protein

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LOCUS NF1 3238 bp ss-mRNA PRI 02-NOV-1990  
 DEFINITION Human hepatic nuclear factor 1 (HNF1) mRNA, complete cds, clones HCL10, HCL12, HCL17, and HCL20.  
 ACCESSION M57732 J04771  
 KEYWORDS hepatic nuclear factor 1; transcription factor.  
 SOURCE Homo sapiens liver NF1 cDNA to mRNA.  
 REFERENCE 1 (bases 1 to 3238)  
 AUTHORS Bach, I., Galcheva-Gargova, Z.I., Mattei, M.-G., Simon-Chazottes, D., Guenet, J.-L., Cereghini, S. and Yaniv, M.  
 TITLE Cloning of human hepatic nuclear factor 1 (HNF1) and chromosomal localization of its gene in man and mouse  
 JOURNAL Genomics 8, 155-164 (1990)  
 COMMENT Chong et al. (Nucleic Acids Res 18: 465-70) report that the HPV-16 enhancer is activated by NF1, AP-1 and TEF-2, and that it occurs without cooperation. Taniguchi et al. (Virology 195: 500-10) demonstrated reduced enhancer activity upon mutation of any of the three NF-1 binding sites (GCCAA).

BASE COUNT 651 a 1100 c 955 g 531 t 1 others

ORIGIN

```

1 cgtggccctg tggcagccga gccatgggtt ctaaactgag ccagctgcag acggagctcc
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## Nuclear Factor 1

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LOCUS OCT1 3824 bp ss-mRNA PRI 18-NOV-1993  
 DEFINITION Human octamer binding transcription factor 1 (OTF1) mRNA, complete cds.  
 ACCESSION L20433  
 KEYWORDS octamer binding transcription factor 1.  
 SOURCE Homo sapiens (library: lambda pSH4K) cDNA to mRNA.  
 REFERENCE 1 (bases 1 to 3824)  
 AUTHORS Bhargava,A.K., Zhen,L.I. and Weissman,S.M.  
 TITLE Differential expression of four members of the POU family of proteins in activated and TPA-treated Jurkat T-cells  
 JOURNAL Proc. Natl. Acad. Sci. U.S.A. 93, 10260-10264 (1994)  
 COMMENT Hoppe-Seyler et al. (J Virol 65: 5613-8) demonstrated that Oct-1 can repress transcription of HPV-18. Mutagenesis experiments suggest that this effect is mediated without direct DNA binding. Morris et al. determined that HPV-16 transcription is repressed by Oct-1 only in non-cervical cells. Conversely, in cervical cells Oct-2 is expressed which transactivates HPV-16 transcription.  
 BASE COUNT 896 a 1005 c 1067 g 856 t  
 ORIGIN  
 1 gcgccccctag agctgtcggg gaagcggggac cgcgaggccg gcgcgcggcg ctctgcgcgg  
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 121 cagccgcgcg gaccgcccgc gctgcagcct ccgaagggag gccgggtgag ccggcgctacg  
 181 cactttcccg cggactttcg gagtgtttgt ggatatacat gccaagccgc cacgatgatg  
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 421 gccgtggaca tcgccgtgtc ccagggcaag agccatcctt tcaagccgga cgccacgtac  
 481 cacacgatga acagcgtgcc gtgcacgtcc acttcacagg tgctctggc gcaccaccac  
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## Oct1

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## Platelet Derived Growth Factor Receptor

LOCUS PDGFR 5570 bp ss-mRNA PRI 28-SEP-1992  
 DEFINITION Human platelet-derived growth factor (PDGF) receptor mRNA, complete cds.  
 ACCESSION J03278  
 KEYWORDS cell surface glycoprotein; glycoprotein; kinase; tyrosine kinase.  
 SOURCE Human skin fibroblast cell (SK5), cDNA to mRNA, clone pHPDGFR.  
 REFERENCE 1 (bases 1 to 5570)  
 AUTHORS Gronwald,R.G.K., Grant,F.J., Haldeman,B.A., Hart,C.E., O'Hara,P.J., Hagen,F.S., Ross,R., Bowen-Pope,D.F. and Murray,M.J.  
 TITLE Cloning and expression of a cDNA coding for the human platelet-derived growth factor receptor: evidence for more than one receptor class  
 JOURNAL Proc. Natl. Acad. Sci. U.S.A. 85, 3435-3439 (1988)  
 COMMENT Nilson and DiMaio (Mol Cell Biol 13: 4137-4145, Cohen et al. J Virol 67:5303-11) reported that the BPV-1 E5 protein activates PDGF beta receptors through complex formation resulting in fibroblast transformation. Cohen et al. demonstrated that this activation is specific to the transmembrane domain of the PDGF receptor and that it is ligand independent (Cohen et al. J Virol 67: 5303-11).  
 Draft entry and computer-readable sequence [1] kindly submitted by R.G.K.Gronwald, 06-APR-1988.  
 BASE COUNT 1195 a 1676 c 1530 g 1169 t  
 ORIGIN Chromosome 5q31-q32.  
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 2401 agtactgccc ctacggagac ctgggtggact acctgcaccg caacaaacac accttctctg

## Platelet Derived Growth Factor Receptor

2461 agcaccactc cgacaagcgc cgcccgccca gcgcggagct ctacagcaat gctctgcccg  
2521 ttgggctccc cctgcccagc catgtgtcct tgaccgggga gagcgacggt ggctacatgg  
2581 acatgagcaa ggacgagtcg gtggactatg tgcccattgct ggacatgaaa ggagacgtca  
2641 aatgatgcaga catcgagtc cccaactaca tggcccctta cgataactac gttccctctg  
2701 ccctgagag gacctgccga gcaactttga tcaacgagtc tccagtgcta agctacatgg  
2761 acctcgtggg cttcagctac cagggtggcca atggcatgga gtttctggcc tccaagaact  
2821 gcgtccacag agacctggcg gctaggaacg tgctcatctg tgaaggcaag ctgggtcaaga  
2881 tctgtgactt tggcctggct cgagacatca tgcgggactc gaattacatc tccaaaggca  
2941 gcaccttttt gcctttaaag tggatggctc cggagagcat ctcaacagc ctctacacca  
3001 ccctgagcga cgtgtggctc ttcgggatcc tgctctggga gatcttcacc ttgggtggca  
3061 ccctttacc agagctgccc atgaacgagc agttctacaa tgccatcaaa cggggttacc  
3121 gcatggccca gcctgcccac gcctccgacg agatctatga gatcatgcag aagtgtctggg  
3181 aagagaagtt tgagattcgg ccccccttct cccagctggt gctgcttctc gagagactgt  
3241 tggcggaagg ttacaaaaag aagtaccagc aggtggatga ggagtttctg aggagtgacc  
3301 acccagccat ccttcggtcc caggcccgtc tgcctggggt ccatggcctc cgatctcccc  
3361 tggacaccag ctccgtcctc tatactgccc tgcagcccaa tgaggggtgac aacgactata  
3421 tcattcccct gcctgacccc aaacccgagg ttgctgacga gggcccactg gagggttccc  
3481 ccagcttagc cagctccacc ctgaatgaag tcaacacctc ctcaaccatc tctgtgaca  
3541 gccccctgga gcccaggac gaaccagagc cagagcccca gcttgagctc cagggtggagc  
3601 cggagccaga gctggaacag ttgccggatt cggggtgccc tgcgcctcgg gcggaagcag  
3661 aggatagctt cctgtagggg gctggcccct accctgccct gcctgaagct cccccctgc  
3721 cagcaccag catctcctgg cctggcctga cgggcttcc tgcagccag gctgccctta  
3781 tcagctgtcc ccttctggaa gcttctgtct cctgacgtgt tgtgcccac accctggggc  
3841 tggcttagga ggcaagaaaa ctgcaggggc cgtgaccagc cctctgcctc cagggaggcc  
3901 aactgactct gagccagggg tccccaggg aactcagttt tcccatatgt aagatgggaa  
3961 agttaggctt gatgaccag aatctaggat tctctcctg gctgacaggt ggggagaccg  
4021 aatccctccc tgggaagatt cttggagtta ctgaggtggt aaattaact tttctgttc  
4081 agccagctac cctcaagga atcatagctc tctcctcgca ctttttatcc acccaggagc  
4141 tagggaagag accctagctc cctggctgc tggctgagct agggcctagc cttgagcagt  
4201 gttgcctcat ccagaagaaa gccagctcct tccctatgat gccagctcct gcgttccctg  
4261 gcccagctg gtctggggcc attaggcagc ctaattaatg ctggaggctg agccaagtac  
4321 aggacacccc cagcctgcag cccttgccc gggcacttgg agcacacgca gccatagcaa  
4381 gtgcctgtgt cctgtcctt caggcccctc agtccctggg ctttttctt ataccctca  
4441 gtcttaatcc atccaccaga gtctagaagg ccagacgggc cccgcactct tgatgagaat  
4501 gtaaatgtgc cagtgtggag tggccaagt tgtgtgccag tatatggccc tggctctgca  
4561 ttggacctgc tatgaggctt tggaggaatc cctcaccctc tctgggcctc agtttccct  
4621 tcaaaaaatg aataagtgcg acttattaac tctgagtgc ttgccagcac taacattcta  
4681 gagtattcca ggtggttga catttgtcca gatgaagcaa ggccatatac cctaaacttc  
4741 catcctgggg gtccagctggg ctccctgggag attccagatc acacatcaca cctcggggac  
4801 tcaggaacca tgccccttc ccaggccccc agcaagtctc aagaacacag ctgcacaggc  
4861 cttgacttag agtgacagcc ggtgtcctgg aaagcccaa gcagctgcc cagggacatg  
4921 ggaagaccac gggacctct tcaactacca cgatgacctc cgggggtatc ctgggcaaaa  
4981 gggacaaaaga gggcaaatga gatcacctcc tgcagcccac cactccagca cctgtgccga  
5041 ggtctgcgtc gaagacagaa tggacagtga ggacagttat gtcttgtaaa agacaagaag  
5101 cttcagatgg taccccaaga aggatgtgag aggtggccgc ttggagtgtg ccctcacc  
5161 accagctgcc ccatccctga ggcagcgtc catgggggta tggttttgtc actgcccaga  
5221 cctagcagtg acatctcatt gtcccagcc cagtgggcat tggaggtgcc aggggagtca  
5281 ggggtgtgag caagacgccc ccgcacgggg aggggtggga agggggtgca ggaagctcaa  
5341 cccctctggg caccaaccct gcattgcagg ttggcacctt acttccctgg gatccccaga  
5401 gttgttccaa ggagggagag tgggttctca atacggatc aaagatataa tcacctaggt  
5461 ttacaaaat ttttaggact cacgttaact cacatttata cagcagaaat gctatttgt  
5521 atgctgttaa gttttctat ctgtgtactt tttttaagg gaaagatttt

**PP2A 72kDa Regulatory Subunit**

LOCUS PP2A72 2338 bp ss-mRNA PRI 20-AUG-1993  
 DEFINITION Homo sapiens protein phosphatase 2A 72 kDa regulatory subunit mRNA, complete cds.  
 ACCESSION L12146  
 KEYWORDS phosphoprotein phosphatase 2A; phosphoprotein phosphohydrolase; protein phosphatase 2A 72 kDa regulatory subunit; regulatory subunit.  
 SOURCE Homo sapiens (library: lambda ZAP; Stratagene) female heart muscle cDNA to mRNA.  
 REFERENCE 1 (bases 1 to 2338)  
 AUTHORS Hendrix,P., Mayer-Jaekel,R.E., Cron,P., Goris,J., Hofsteenge,J., Merlevede,W. and Hemmings,B.A.  
 TITLE Structure and expression of a 72-kDa regulatory subunit of protein phosphatase 2A. Evidence for different size forms produced by alternative splicing  
 JOURNAL J. Biol. Chem. 268, 15267-15276 (1993)  
 COMMENT PP2A consists of regulatory subunits and one catalytic subunit. In their review, "Protein Phosphatase 2A and the Regulation of Human Papillomavirus Gene Activity, Schegget and Nooraa (in Human Papillomaviruses, 121-9) report that PP2A is inactivated by high expression of its regulatory subunit, PR55Beta. This inhibition is linked to the activation of the LCR region of HPV-16. Thus, the authors suggest that the regulation of the HPV-16 promoter is correlated to cellular protein phosphorylation.  
 BASE COUNT 694 a 497 c 526 g 621 t  
 ORIGIN

```

1 gaattccgca gctgtgttta taacttgaaa attatgagca ggtattggga gccacagact
61 gtattttggga acagctgcat gtcaggacaa gcatagattg tggctctgtgg aggctggggag
121 agcacaatgc ttgaatttca tttctttggg agataagaag aaaaaatacc ttctactctg
181 caaagattga caacttgcta aatctgatta tcttcagaag tgtgctggac agttgcgaag
241 taaacattta cttttcaaag atgatgatca aggaaacatc tctacgaagg gaccgggatt
301 taaggggaga gctagctttc ctggcaaggg gctgtgattt tgttctccct tcacggttta
361 agaagcggct gaagtcattt cagcagacac agattcaaaa taaccagaa aagaaacctg
421 gaacaccact cccacctcca gccacctctc caagtagtcc ccgacctctc tccccggttc
481 cccatgtgaa taatgttgtg aatgcgccat tgtccataaa cattccacgg ttctactttc
541 ctgaaggact cccagatacc tgtagtaatc atgaacaaac tctaagcaga attgaaactg
601 ctttcatgga tattgaagaa cagaaagcag acattttatga aatggggaaa attgcaaagg
661 tctgtggctg tcctctctat tggaaagccc ccatgttcag ggctgcaggg ggagagaaga
721 caggatttgt gacagcacag tcattcattg ccatgtggag aaagttgctg aataaccatc
781 atgatgatgc ctctaaattc atctgtcttc tagcaaagcc caactgcagc tctctagaac
841 aggaggattt catccctcta cttcaggatg tgggtggatac ccaccctggt ctcacgttcc
901 tgaagatgc tccagaatc cactcccgtc acatcaccac ggttattcag agaatattct
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1021 ttttgcaaac cctagcactt ttggaagaag aggaagatat aaaccaaat acagattact
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1141 acctctacat cagccaggcc gatctgtctc gatacaatga ccaggcttca tcaagcagga
1201 ttattgaaag gatattctct ggtgcagtaa caaggggaaa aacaatacag aaagagggaa
1261 gaatgagcta tgcagatttt gtttggtttt tgatctctga agaagacaaa agaatccta
1321 ccagcattga gtattggttc cgctgcatgg atgtggatgg agacggtgta ctctccatgt
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1561 tctttaatct ggagaaatac ttagaccatg aacagagaga tccctttgcg gtccagaagg
1621 atgttgagaa cgatgggctc gagccctcag actgggaccg gtttgccgct gaggagtatg
1681 agacgcttgg tgcagaggaa tctgccaag cacaattcca ggaaggcttt gaagattatg
1741 aaacagatga acctgcctct ccctctgaat ttggaacaaa aagcaataaa atattaagtg
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1861 ctacaatgaa acgaagatgt gtattttaa tgtttcttcc ttgtgaagag atgttctcgt
1921 ttgcatactg ctttttaaag actttgattt ctccaagtgt gtatcatctg cactaggaac
1981 tttgttttta agcaataggt ctggatacac atttaactta ggaggctcct ccaatttggc
2041 tcaaacctct tacggagctt ctctcagaa gtggtaccat cgccttcaa agtcagcact
2101 ctacactctt gaatgtacca aggatctctt ggcgacagta ccaagcaggg ttctctacac
2161 aggtgactga agttgcctct gtgttggctg gcacccctga gtcccctcgg gctcctatgg
  
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## PP2A 72kDa Regulatory Subunit

```
2221 agcctagaag aaaccttcac ttgcagaaaa cttgagtcag aaaattctgg aactgaaaa  
2281 agtagtaagg gcctccagaa ttgacttagc cctagtaata aaagcactgc cggaattc
```

LOCUS PP2ACAT 2966 bp ds-DNA PRI 04-MAR-1991  
DEFINITION Human protein phosphatase 2A catalytic subunit-alpha gene, complete cds.  
ACCESSION M60483 J05297  
KEYWORDS protein phosphatase-2A catalytic subunit-alpha.  
SOURCE Human placenta leukocyte DNA.  
REFERENCE 1 (bases 1 to 2966)  
AUTHORS Khew-Goodall,Y., Mayer,R.E., Maurer,F., Stone,S.R. and Hemmings,B.A.  
TITLE Structure and transcriptional regulation of protein phosphatase 2A catalytic subunit genes  
JOURNAL Biochemistry 30, 89-97 (1991)  
COMMENT PP2A consists of regulatory subunits and one catalytic subunit. In their review, "Protein Phosphatase 2A and the Regulation of Human Papillomavirus Gene Activity, Schegget and Noordaa (in Human Papillomaviruses, 121-9) report that PP2A is inactivated by high expression of its regulatory subunit, PR55Beta. This inhibition is linked to the activation of the LCR region of HPV-16. Thus, the authors suggest that the regulation of the HPV-16 promoter is correlated to cellular protein phosphorylation.

Although this is a genomic sequence, the introns have been omitted. Intronic sequences have been requested. The data is presented as submitted by the author.

NCBI gi: 190223

BASE COUNT 747 a 728 c 732 g 759 t

ORIGIN

```

1 aaccaccggc gaggagcggg gcgctgga gcgagccggt ccgagagcc caaagaaaag
61 cccaagcctc gcccccgcca tcgcccgcga cgagacacct aggtccgggg acgggtgtgt
121 gccgcggaag tcaggtgcac tgcgcagcac tcccccggtt ggtacacgct cctccacctc
181 cgagtgacct aattacaagg tgccagccgc gccagaggtt ggggggtggtt aatccaagcg
241 gccactcgct gcccgcttct gccccaaaag atgacggaaa cccacacgat tacagagccg
301 cagcacccca gatgagccac ggggtcgcaa ttctcgtttc cgtgatcgga ctgccaggcc
361 ccaggtgagg agctgagttc atcaccagag cggccttccc aggggaacca gttacaggct
421 gccagtgagg ccggcttcca tccggtctgc gcctgcgcgc ggcccaagcc ctgcctctc
481 ctggaatagt gctcagggat tagtccgggt cgcctgctgt ccaactgcga tgctccagct
541 ccatccttcc cttccccac caccocgccc tccgggagcc acgcccacaaa agtcaaggcg
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721 cctccgctc cccgcccctt cctgacgccc ggcgtgacct caccacgccc ggcggcccgc
781 attacagaga gccgagctct ggagcctcag cgagcggagg aggaggcgca gggccgacgg
841 ccgagtactg cggtgagagc cagcgggcca gcccagcctt caacagccgc cagaagtaca
901 cgaggaaccg gcggcggcgt gtgcgtgtag gccctgtgac gggcggcggc gcgggaggag
961 cgcggagcgg cagccggctg gggcgggtgg catcatggag gagaaggtgt tcaccaagga
1021 gctggaccag tggatcgagc agctgaacga gtgcaagcag ctgtccgagt cccagggtcaa
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1141 atgtccagtt actgtctgtg gagatgtgca tgggcaattt catgatctca tggaaactgtt
1201 tagaattggt ggcaaatcac cagatacaaa ttacttgttt atgggagatt atgttgacag
1261 aggatattat tcagttgaaa cagttacact gcttgtagct cttagggttc gttaccgtga
1321 acgcatcacc attcctcgag ggaatcatga gagcagacag atcacacaag tttatggttt
1381 ctatgatgaa tgtttaagaa aatatggaaa tgcaaatgtt tggaaatatt ttacagatct
1441 ttttgactat cttcctctca ctgccttggg ggatgggagc atcttctgtc tacatggtgg
1501 tctctcgcca tctatagata cactggatca tatcagagca cttgatcgcc tacaagaagt
1561 tccccatgag ggtccaatgt gtgacttggc gtggtcagat ccagatgacc gtgggtgggtg
1621 gggatatact cctcgaggag ctggttacac ctttgggcaa gatatttctg agacatttaa
1681 tcatgccaat ggcctcagct tggtgtctag agctcaccag ctagtgtagg agggatataa
1741 ctgggtgcca gaccggaatg tagtaacgat tttcagtgct ccaaactatt gttatcggtg
1801 tgtaaccaa gctgcaatca tggaaactga cgatactcta aaatactctt tcttgagtt
1861 tgaccagaca cctcgtagag gcgagccaca tgttactcgt cgtaccaccag actacttct
1921 gtaataaagt tttaaacttg tacagtattg ccatgaacca tatatcgacc taatggaaat
1981 ggaagagaca acagtaactc caaagtgtca gaaaatagtt aacattcaaa aaacttgttt
2041 tcacatggac caaaagatgt ccaatataaa aatacaaaag cttctgtcat caacagcgt
2101 gaccacttta gaatgaacca gttcattgca tgctgaagcg acattgttgg tcaagaaac

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## PP2A Catalytic Subunit

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2161 agtttctggc atagcgctat ttgtagttac ttttgctttc tctgagagac tgcagataat
2221 aagatgtaaa cattaacacc tcgtgaatac aatttaactt ccatttagct atagctttac
2281 tcagcatgac tgtagataag gatagcagca aacaatcatt ggagcttaat gaacattttt
2341 aaaaaataatt accaaggcct cccttctact tgtgagtttt gaaattgttc tttttatfff
2401 cagggatacc gtttaattta attatatgat ttgtctgcac tcagtfffatt ccctactcaa
2461 atctcagccc catgfffgttc tttgfffattg tcagaacctg gtgagfffgtt ttgaacagaa
2521 ctgffffffcc cccttctgt aagacgatgt gactgcacaa gagcactgca gtgffffffca
2581 taataaactt gtgaactaag aactgagaag gtcaaatttt aattgtatca atgggcaaga
2641 ctggtgctgt ttattaafaaa agfftaaatca attgagtaaa fffftagaatt tgtagacttg
2701 taggtaaaat aaaaatcaag ggcactacat aacctctctg gtaactcctt gacattcttc
2761 agattaactt caggattfat ttgtattfca catattacaa ffffgtcacat tgttgggtgtg
2821 cactfffgtgg gtfcttctctg catattaaact tgtfffgtaaag aaaggaaatc tgtgctgctt
2881 cagtaagact taattgtaaa accatataac ttgagattfa agtctfffggg ttfgtffftta
2941 ataaaacagc atgfffctcag gtagag
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PP2A PR55 Regulatory Subunit

LOCUS PP2APR55 2131 bp ss-mRNA PRI 09-MAY-1991  
 DEFINITION Human protein phosphatase 2A alpha subunit mRNA, complete cds.  
 ACCESSION M64929 J05328  
 KEYWORDS protein phosphatase-2A subunit-alpha; regulatory subunit.  
 SOURCE Human lung fibroblast cell line WI38, cDNA to mRNA.  
 REFERENCE 1 (bases 1 to 2131)  
 AUTHORS Mayer,R.E., Hendrix,P., Cron,P., Matthies,R., Stone,S.R., Goris,J., Merlevede,W., Hofsteenge,J. and Hemmings,B.A.  
 TITLE Structure of the 55 kDa regulatory subunit of protein phosphatase 2A: Evidence for a neuronal specific isoform  
 JOURNAL Biochemistry 30, 3589-3597 (1991)  
 COMMENT PP2A consists of regulatory subunits and one catalytic subunit. In their review, "Protein Phosphatase 2A and and the Regulation of Human Papillomavirus Gene Activity, Schegget and Noordaa (in Human Papillomaviruses, 121-9) report that PP2A is inactivated by high expression of its regulatory subunit, PR55Beta. This inhibition is linked to the activation of the LCR region of HPV-16. Thus, the authors suggest that the regulation of the HPV-16 promoter is correlated to cellular protein phosphorylation.

NCBI gi: 190421

BASE COUNT 665 a 421 c 460 g 585 t

ORIGIN

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1  ccgccgcat  ccgccctctc  tccccccca  tccccagggtg  aggggggtga  gttcaggaag
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121  ggaggggaatg  atattcagtg  gtgtttttct  caggtgaaag  gagcagtaga  tgatgatgta
181  gcagaagcag  atataatttc  tacagtagaa  tttaatcatt  ctggagaatt  actagcaaca
241  ggagataaag  gtggtagagt  tgtcatcttt  caacaggagc  aggagaacaa  aatccagtct
301  catagcagag  gagaataata  tgtttacagc  accttcaga  gccatgaacc  agagtttgac
361  tacttgaaaa  gtttagaaat  agaagaaaag  atcaataaaa  ttaggtggtt  accccagaaa
421  aatgctgctc  agtttttatt  gtctaccaat  gataaaacaa  taaaattatg  gaaaatcagt
481  gaaagggaca  aaagaccaga  agggataaac  ttgaaagagg  aggatggaag  gtagagagat
541  cctactacag  ttactacact  acgagtgcc  gtctttagc  ctatggatct  aatggttgag
601  gccagtccac  gaagaatatt  tgccaatgct  catacatatc  acatcaactc  aatttctatt
661  aatagtgatt  atgaaacata  tttatctgca  gatgatttgc  ggattaatct  ttggcatctg
721  gaaattacag  acaggagttt  taacattgtg  gatatcaagc  ctgccaatat  ggaagagcta
781  acagaggtga  ttacagcagc  agaatttcat  ccaaacagct  gtaacacatt  tgtatacagc
841  agcagtaaag  gaactattcg  gctatgtgac  atgagggcat  ctgocctctg  tgatagacat
901  tctaaattgt  ttgaagaacc  tgaagatccc  agtaaacaggt  catttttttc  cgaaatcctc
961  tcctctatth  cggatgtaaa  attcagccat  agtggctgat  atatgatgac  tagagactat
1021  ttgtcagtca  aaatttggga  cttaaatatg  gaaaacaggc  ctgtggaac  ataccaggtg
1081  catgaatacc  tcagaagtaa  actctgttca  ctgtatgaaa  atgactgcat  atttgacaaa
1141  tttgaatggt  gttggaatgg  atctgacagt  gttgtcatga  ctggatctta  caataatttc
1201  ttcagaatgt  ttgacagaaa  cacaaagcga  gacataacc  tagaagcatc  gcgggaaaac
1261  aataagcctc  gcacagtctc  gaagcctcgc  aaagtctgtg  caagtggcaa  gcgaaagaaa
1321  gatgaaataa  gtgttgacag  cctagacttc  aataagaaaa  tccttcacac  agcctggcac
1381  cccaaggaaa  atatcattgc  cgtagctact  acaacaatc  tgatatatt  tcaagacaaa
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1501  aatctagatc  tcgttcctat  aaaagagaga  ggtccattgt  ggcgcccctt  tccagtgttt
1561  gacagtgctc  cattcgacaa  cacattgtta  tagctacatg  gagaaagctc  tgtggattca
1621  tcaactgggt  gttctccatg  tctgctagcc  atttaggtaa  gggtagggca  cttttaattt
1681  aaatgacttc  ttgcaccatc  ttgcctaata  gactagattg  gactgtatca  acattgattt
1741  actccacttt  ttatgccttc  cattgtgatg  acgtocaaac  cagtgaagc  cttcagtcot
1801  gctatgggat  ttaattgtgt  atcctcatta  ctgtatcatt  tgtgggttac  accccttccc
1861  ccttttttta  aattaaatac  agctcattct  tactgtggct  tgtagcattc  ctccctctct
1921  ggcctcctgg  actgctcccc  ttcctctctt  acccttgccc  cctccaccog  gtcttgggtg
1981  tggtatatta  aaaaaagaaa  gaatgaaagc  acacaaaatg  agtcagtttg  gggtcagttg
2041  tataaagggg  gtatatgttg  caaacaaatg  ttttagtaac  agttggctgt  aatcactcct
2101  cgcggtgtct  ggcactgaaa  ataaggaaaa  g

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## PP2A PR65 Regulatory Subunit

LOCUS PP2APR65 2205 bp ss-mRNA PRI 22-OCT-1992  
DEFINITION Human protein phosphatase 2A regulatory subunit alpha-isotype  
(alpha-PR65) mRNA, complete cds.  
ACCESSION J02902  
KEYWORDS protein phosphatase-2A regulatory alpha-subunit.  
SOURCE Human HeLa cell, cDNA to mRNA, clone lambda-HHPR65-3.  
REFERENCE 1 (bases 1 to 2205)  
AUTHORS Hemmings,B.A., Adams-Pearson,C., Maurer,F., Mueller,P., Goris,J.,  
Merlevede,W., Hofsteenge,J. and Stone,S.R.  
TITLE Alpha and beta-forms of the 65-kDa subunit of protein phosphatase  
2A have a similar 39 amino acid repeating structure  
JOURNAL Biochemistry 29, 3166-3173 (1990)  
COMMENT PP2A consists of regulatory subunits and one catalytic subunit.  
In their review, "Protein Phosphatase 2A and the Regulation  
of Human Papillomavirus Gene Activity, Schegget and Noordaa  
(in Human Papillomaviruses, 121-9) report that PP2A is  
inactivated by high expression of its regulatory subunit,  
PR55Beta. This inhibition is linked to the activation of the LCR  
region of HPV-16. Thus, the authors suggest that the regulation  
of the HPV-16 promoter is correlated to cellular protein  
phosphorylation.

Draft entry and printed sequence [1] kindly submitted by  
B.A.Hemmings, 23-MAR-1990, for release after publication.

NCBI gi: 189427

BASE COUNT 447 a 676 c 639 g 443 t

ORIGIN

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LOCUS TEF 4443 bp ds-DNA PRI 28-JUL-1992  
 DEFINITION Transcriptional enhancer factor DNA, complete CDS.  
 ACCESSION M63896  
 KEYWORDS trans-acting transcriptional activator; transcription enhancer.  
 SOURCE Homo sapiens (library: ZAP-II random primed cDNA) DNA.  
 REFERENCE 1 (bases 1 to 4443)  
 AUTHORS Xiao,J.-H., Davidson,I., Matthes,H., Garnier,J.-M. and Chambon,P.  
 TITLE Cloning, expression, and transcriptional properties of the human  
 enhancer factor TEF-1  
 JOURNAL Cell 65, 551-568 (1991)  
 BASE COUNT 1226 a 1040 c 969 g 1208 t  
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## Transcriptional Enhancer Factor

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4441 ccc

## Transforming Growth Factor

LOCUS TGFA 867 bp ss-mRNA PRI 19-APR-1994  
DEFINITION Human (cell line 1027 F57) transforming growth factor-alpha mRNA,  
complete cds.  
ACCESSION K03222  
KEYWORDS growth factor; transforming gene; transforming growth factor-alpha.  
SOURCE Human renal carcinoma cell line 1027 F57, cDNA to mRNA, clone  
pTGF-C1.  
REFERENCE 1 (bases 1 to 867)  
AUTHORS Derynck,R., Roberts,A.B., Winkler,M.E., Chen,E.Y. and Goeddel,D.V.  
TITLE Human transforming growth factor-alpha: Precursor structure and  
expression in E. coli  
JOURNAL Cell 38, 287-297 (1984)  
COMMENT Pietenpol et al. (Cell 6: 777-785) report that HPV-16 and HPV-18  
infection inhibits growth suppression by TGF-B1 in  
human foreskin keratinocytes. The authors suggest that the inhibitory  
capability of TGF-B1 is linked to the down-regulation of c-myc and  
that HPV-16 E7 expression interferes with this pathway.  
  
[1] also sequenced exons 1 and 2 of this gene from the genomic DNA  
library of Lawn et al. (see separate entries).  
BASE COUNT 175 a 240 c 254 g 198 t  
ORIGIN Chromosome 2p13  
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# Tumor Necrosis Factor

LOCUS TNF 7112 bp ds-DNA PRI 15-JUN-1989  
DEFINITION Human tumor necrosis factor and lymphotoxin genes, complete cds.  
ACCESSION M16441  
KEYWORDS lymphotoxin; tumor necrosis factor.  
SOURCE Human placenta DNA, clone pTNF186.  
REFERENCE 1 (bases 1 to 7112)  
AUTHORS Nedospasov,S.A., Shakhov,A.N., Turetskaya,R.L., Mett,V.A.,  
Azizov,M.M., Georgiev,G.P., Korobko,V.G., Dobrynin,V.N.,  
Filippov,S.A., Bystrov,N.S., Boldyreva,E.F., Chuvpilo,S.A.,  
Chumakov,A.M., Shingarova,L.N. and Ovchinnikov,Y.A.  
TITLE Tandem arrangement of genes coding for tumor necrosis factor  
(TNF-alpha) and lymphotoxin (TNF-beta) in the human genome  
JOURNAL Cold Spring Harb. Symp. Quant. Biol. 51, 611-624 (1986)  
COMMENT Kyo et al. (Virology 200: 130-9) demonstrated that tumor necrosis  
factor (TNF) alpha represses transcription of the early HPV-16  
genes through responsive elements in the LCR. Malejczyk et al.  
(Int J Cancer 56: 593-8) reported this inhibition for weakly  
tumorigenic cell lines, but not for highly tumorigenic cell lines.  
Further Malejczyk et al. demonstrated that TNF-alpha receptor  
expression is reduced in the highly tumorigenic lines and that this  
reduction may inhibit the autocrine TNF-alpha-mediated growth  
limitation.  
  
Draft entry and computer-readable sequence for [1] kindly submitted  
by C.V.Jongeneel, 02-OCT-1988.  
BASE COUNT 1676 a 2005 c 1865 g 1566 t  
ORIGIN 1 bp upstream of EcoRI site; chromosome 6p21.3.  
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 4381 tctccttctc gatcgtggca ggcgccacca cgctcttctg cctgctgca ctttgagtg  
 4441 tcggccccca gaggaagag gtgagtgcct ggcagcctt catccactc cccaccaag  
 4501 gggaaatgga gacgcaagag agggagagag atgggatggg tgaagatgt gcctgatag  
 4561 ggagggatgg agagaaaaaa acgtggagaa agacggggat gcagaaagag atgtggcaag  
 4621 agatggggaa gagagagaga gaaagatgga gagacaggat gtctggcaca tggaggtgc  
 4681 tcaactaagt tgtatggagt gaatgaatga atgaatgaat gaacaagcag atataaat  
 4741 aagatatgga gacagatgtg ggggtgtgaga agagagatgg ggaagaaac aagtgatatg  
 4801 aataaagatg gtgagacaga aagagcggga aatatgacag ctaaggagag agatggggg  
 4861 gataagagga gaagaagata cctgtctctg cacacagaag acactcaggg aaagactgt  
 4921 tgaatgctcg gaaggtgaat acacagatga atggagagag aaaaccagac acctcagggc  
 4981 taagagcgca ggccagacag gcagccagct gttcctcctt taagggtgac tccctcagtg  
 5041 ttaaccatct tccttctccc caacagttcc ccagggacct ctctctaata agccctctgg  
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 5221 atggatggag gtgaaagttag ggggtatttt tctaggaagt ttaagggtct cagctttttc  
 5281 ttttctctct cctcttcagg atcatcttct cgaaccccga gtgacaagcc tgtagcccat  
 5341 gttgtaggtg agagctctga ggatgtgtct tggaaacttg agggctagga tttggggatt  
 5401 gaagccgggc tgatggtagg cagaacttgg agacaatgtg agaggactc gctgagctca  
 5461 agggaaaggg ggaggaacag cacaggcctt agtgggatac tcagaacgct atggccaggt  
 5521 gggatgtggg atgacagaca gagaggacag gaaccggatg tgggggtggg agagctcagag  
 5581 ggcagagatg tggagagtga accgacatgg ccacactgac tctcctctcc ctctctccc  
 5641 cctccagca aacctcaag ctgaggggca gctccagtg ctgaaccgcc gggccaatgc  
 5701 cctcctggcc aatggcgtgg agctgagaga taaccagctg gtggtgcca cagagggcct  
 5761 gtacctcact tactcccagg tctcttcaa gggccaaggg tgcccctca cccatgtgct  
 5821 cctcaccacc accatcagcc gcatcgccgt ctctaccag accaaggtca acctctctc  
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 5941 tgagcccatc tatctgggag gggcttcca gctggagaag ggtgaccgac tcagcgtga  
 6001 gatcaatcgg cccgactatc tcgactttgc cgagtctggg caggtctact ttgggatcat  
 6061 tgccctgtga ggaggaagaa catccaacct tccaaacgc ctcccctgccc

## Tumor Necrosis Factor

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6121 tattaccccc tccttcagac accctcaacc tcttctggct caaaaagaga attgggggct
6181 tagggctcga acccaagctt agaactttaa gcaacaagac caccacttcg aaacctggga
6241 ttcaggaatg tgtggcctgc acagtgaagt gctggcaacc actaagaatt caaactgggg
6301 cctccagaac tcaactggggc ctacagcttt gatccctgac atctggaatc tggagaccag
6361 ggagcctttg gttctggcca gaatgctgca ggacttgaga agacctcacc tagaaattga
6421 cacaagtgga ccttaggcct tcctctctcc agatgtttcc agacttcctt gagacacgga
6481 gccagccct ccccatggag ccagctccct ctatttatgt ttgcacttgt gattatttat
6541 tatttattta ttatttattt atttacagat gaatgtattt atttgggaga ccgggggatc
6601 ctgggggacc caatgtagga gctgccttgg ctcagacatg ttttccgtga aaacggagct
6661 gaacaatagg ctgttcccat gtagccccct ggccctctgtg ccttcttttg attatgtttt
6721 ttaaaatatt tatctgatta agttgtctaa acaatgctga tttggtgacc aactgtcact
6781 cattgctgag cctctgctcc ccaggggagt tgtgtctgta atcgcctac tattcagtgg
6841 cgagaaataa agtttgctta gaaaagaaac atgggtctcct tcttgaatt aattctgcat
6901 ctgcctcttc ttgtgggtgg gaagaagctc cctaagtcct ctctccacag gctttaagat
6961 ccctcggacc cagtcccatc cttagactcc tagggcctg gagaccctac ataaacaaag
7021 cccaacagaa tattcccat ccccaggaa acaagagcct gaacctaatt acctctccct
7081 cagggcatgg gaatttcaa ctctgggaat tc
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LOCUS YY1 2353 bp ss-mRNA PRI 25-MAY-1993  
 DEFINITION Homo sapiens GLI-Krupple related protein (YY1) mRNA, complete cds.  
 ACCESSION M77698  
 KEYWORDS GLI-Krupple related protein.  
 SOURCE Homo sapiens cDNA to mRNA.  
 REFERENCE 1 (bases 1 to 2353)  
 AUTHORS Shi,Y., Seto,E., Chang,L.-S. and Shenk,T.  
 TITLE Transcriptional repression by YY1, a human GLI-Krupple related protein, and relief of repression by adenovirus Ela protein  
 JOURNAL Cell 67, 377-388 (1991)  
 COMMENT YY1 has been shown to be involved in transcriptional repression of both HPV-16 and HPV-18 (May,M., Dong,X.-P., Beyer-Finkler,E., Stubenrauch,F., Fuchs,P., and Pfister,H., EMBO J.:13, 1460-1466 (1994); Bauknecht,T., Angel,P., Royer,H.-D., and zur Hausen,H., EMBO J.:11, 4607-4617 (1992)). It is thought to interfere with transcriptional initiation by binding to sites contained within a transcriptional silencer domain in the LCRs of these HPV types. In HPV-16, this silencer domain is contained within a naturally occurring deletion mutation which was found by May et al. to be present in 40% of the HPV-16 DNA in a lymph node metastasis of a cervical cancer. May et al. contend that this deletion may represent a means by which expression of oncogenic genes is de-regulated in episomal copies of HPV-16 DNA. Since at least one-third of HPV-16-positive cervical cancers contain only extra-chromosomal viral DNA, this deletion may be an important alternative to the destruction of the E2 reading frame, and thereby the E2-dependent negative regulation of E6/E7 expression, that occurs during integration of the viral DNA into the host genome. Mutations of YY-1 binding sites in HPV-18 DNA has been shown to affect transcriptional activity differently in malignant and non-malignant cell lines. In non-malignant cells, YY1 binding-site mutations led to increased transcriptional activity, whereas in certain malignant cell lines, transcriptional activity was apparently reduced by the mutations. This may point to more complex interactions between YY1 and other cellular factors.

BASE COUNT 625 a 598 c 608 g 522 t  
 ORIGIN

```

1 cgccgagacg agcagcggcc gagcagcgc gggcgcgggc gcaccgaggc gagggaggcg
61 ggggaagcccc gccgcgcgccc ccccgccgc cccttcccc gccgcccgcc ccctctcccc
121 ccgcccgcctc gccgccttcc tccctctgcc ttcttcccc acggccgggc gcctcctcgc
181 ccgcccgcctc gccgccttcc tccctctgcc ttcttcccc acggccgggc gcctcctcgc
241 atggcctcgg gcgacaccct ctacatcgcc acggacggct cggagatgcc ggccgagatc
301 gtggagctgc acgagatcga ggtggagacc atcccgggtg agaccatcga gaccacagtg
361 gtggggcgagg aggaggagga ggacgacgac gacgaggacg gcggcggtgg cgaccacgac
421 ggcggggggcg gccacgggca gcgcggccac caccaccacc accatcacca ccaccaccac
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541 caggaggtga tcctggtgca gacgcgcgag gaggtggtgg gcggcgacga ctcggaacggg
601 ctgcgcgccc aggacggcct cgaggatcag attctcatcc cggtgcccgc gccggccggc
661 ggcgacgacg actacattga acaaacgctg gtcaccgtgg cggcggcccg caagagcggc
721 ggcggcggct cgtcgtcgtc gggaggcggc gcgctcaaga agggcggcgg caagaagagc
781 ggcaagaaga gttacctcag cggcggggcc ggcgcggcgg gcggggcggc cgccgacccg
841 ggcaacaaga agtgggagca gaagcaggtg cagatcaaga ccctggaggg cgagttctcg
901 gtcaccatgt ggtcctcaga tgaaaaaaaa gatattgacc atgagacagt ggttgaagaa
961 cagatcattg gagagaactc acctcctgat tattcagaat atatgacagg aaagaaactt
1021 cctcctggag gaatacctgg cattgacctc tcagatccca acaaactggc agaatttctg
1081 agaatgaagc caagaaaaat taaagaagat gatgctccaa gaacaatagc ttgcctcat
1141 aaaggctgca caaagatggt cagggataac tcggccatga gaaaacatct gcacaccacc
1201 ggtcccagag tccacgtctg tgcagaatgt ggcaaagctt ttgttgagag ttcaaaacta
1261 aaacgacacc aactggttca tactggagag aagcctttc agtgacggt cgaaggctgt
1321 gggaaacgct tttcactgga cttcaatttg cgcacacatg tcgcaatcca taccggagac
1381 aggccctatg tgtgcccctt cgatggttgt aataagaagt ttgctcagtc aactaacctg
1441 aaatctcaca tcttaacaca tgctaaggcc aaaacaacc agtgaagaaga agagagaaga
1501 cccttctcga ccacgggaag catcttccag aagtgtgatt ggaataaat atgcctctcc
1561 tttgtatatt atttctagga agaattttaa aaatgaatcc tacacaccta agggacatgt
  
```

# YY1

1621 tttgataaag tagtaaaaat taaaaaaaaa aaactttact aagatgacat tgctaagatg  
1681 ctctatcttg ctctgtaatc tcgtttcaaa aacacagtgt ttttgtaaag tgtggtccca  
1741 acaggaggac aattcatgaa cttcgcatca aaagacaatt cttatacaa cagtgctaaa  
1801 aatgggactt cttttcacat tcttataaat atgaagctca cctgttgctt acaatTTTT  
1861 taatTTTgta ttttccaagt gtgcataatt tacactTTTT tggggatatg cttagtaatg  
1921 ctacgtgtga tttttctgga ggttgataac tttgcttgca gtagattttc tttaaaagaa  
1981 tgggcagtta catgcatact tcaaaagtat tttcctgtaa aaaaaaaaaa agttatatag  
2041 gttttgTTTg ctatcttaat tttggTTgta ttctttgatg ttaacacatt ttgtataatt  
2101 gtatcgtata gctgtattga atcatgtagt atcaaatatt agatgtgatt taatagTgtt  
2161 aatcaattta aaccatttt agtcaacttt tttttccaaa aaaatactgc cagatgctga  
2221 tgTtcagtgt aatttctttg cctgttcagt tacagaaagt ggtgctcagt tgtagaatgt  
2281 attgtacctt ttaacacctg atgtgtacat cccatgtaac agaaagggca acaataaaaat  
2341 agcaatccta aag



LOCUS c-myc 8082 bp ds-DNA PRI 10-AUG-1992  
 DEFINITION Human (Lawn) c-myc proto-oncogene, complete coding sequence and flanks.  
 ACCESSION J00120  
 KEYWORDS Alu repeat; c-myc proto-oncogene; myc oncogene; proto-oncogene; repeat region; transforming gene.  
 SOURCE Human DNA (genomic library of Lawn et al.), clones lambda-M1 [1], and pUC9-myc [2].  
 REFERENCE 1 (bases 3507 to 7559)  
 AUTHORS Colby,W.W., Chen,E.Y., Smith,D.H. and Levinson,A.D.  
 TITLE Identification and nucleotide sequence of a human locus homologous to the v-myc oncogene of avian myelocytomatosis virus MC29  
 JOURNAL Nature 301, 722-725 (1983)  
 REFERENCE 2 (bases 1 to 8082)  
 AUTHORS Gazin,C., Dupont,S., de Dinechin,D., Hampe,A., Masson,J.M., Martin,P., Stehelin,D. and Galibert,F.  
 TITLE Nucleotide sequence of the human c-myc locus: provocative open reading frame within the first exon  
 JOURNAL EMBO J. 3, 383-387 (1984)  
 COMMENT In several cases of HPV-positive cancer, Couturier et al. (J Virol 65: 4534-8) determined that HPV sequences integrate near the myc locus. In several instances, this integration disrupted the integrity of the proto-oncogene sequence or its regulation.

The myc gene is the cellular homologue of the transforming gene carried by the avian myelocytomatosis virus MC29. Unlike the ras proto-oncogenes which obtain transforming potential through mutations within their coding exons (namely mutations within codon 12), the myc gene identified as the cause of Burkitt lymphomas acquires its transforming potential through defects of either transcriptional or translational control. Thus it is not an altered gene product that induces tumors, but a normal product that is present either in the wrong quantity or at the wrong time in the life cycle of the cell.

[2] notes an open reading frame upstream of the c-myc coding exons with an 'atg' start codon at bases 2304-2306 and a 'tag' stop codon at bases 2868-2870. However other researchers have used c-myc and v-myc DNA sequences to probe for mRNA's with homology to c-myc in various human cell lines and none of them have noted any mRNA's beginning upstream of bp 2328 (see other human c-myc entries).

The t(8;14) translocation site in the Burkitt lymphoma cell line BL22 occurs between bp 1316 and 1317 of this sequence.

BASE COUNT 1850 a 2115 c 2135 g 1982 t  
 ORIGIN 198 bp upstream of Sau96A site, on chromosome 8 (q24).  
 1 agcttggttg gccgttttag ggtttgttg aattttttt tcgtctatgt acttgtgaat  
 61 tatttcacgt ttgccattac cggttctcca tagggtagtg ttcattagca gtgggtgatg  
 121 gtaaatattc accatctctt atgcccgttg atagtcacct ctgaaccact ttttctcca  
 181 gtaactcctc tttcttcgga cttctgcag ccaacctgaa agaataaca ggaggtggct  
 241 ggaaacttgt tttaaggaac cgctgtcct tccccgctg gaaacctgc acctcggacg  
 301 ctctgctcc tgccccacc tgacccccgc cctcgttgac atccaggcgc gatgatctct  
 361 gctgccagta gagggcacac ttactttact ttcgaaaacc tgaacgcggg tgetgcccag  
 421 agagggggcg gagggaaaga cgctttgcag caaaatccag catagcgatt ggttgctccc  
 481 cgcgtttgcg gcaaaggcct ggaggcagga gtaatttgca atccttaaag ctgaattgtg  
 541 cagtgcacg gatttgaag ctactatatt cacttaacac ttgaacgctg agctgcaaac  
 601 tcaacgggta ataaccatc ttgaacagcg tacatgctat acacacacc ctttcccccg  
 661 aattgttttc tcttttgagg gtggtggagg gagagaaaag tttacttaaa atgcctttgg  
 721 gtgagggacc aaggatgaga agaatgttt ttgttttca tgccgtggaa taacacaaaa  
 781 taaaaaatcc cgaggaata tacattatat attaaatata gatcattca gggagcaaac  
 841 aaatcatgtg tggggctggg caactagctg agtcgaagcg taaataaaa gtgaatacac  
 901 gtttgcgggt tacatacagt gcactttcac tagtattcag aaaaattgt gagtcagtga  
 961 actaggaat taatgctgg aaggcagcca aattttaatt agctcaagac tcccccccc  
 1021 ccccaaaaa aggcacggaa gtaatactcc tctctcttc tttgatcaga atcgatgat

1081 tttttgtgca tgaccgcatt tccaataata aaaggggaaa gaggacctgg aaaggaatta  
 1141 aacgtccggg ttgtccgggg aggaaagagt taacggtttt tttcacaagg gtctctgctg  
 1201 actcccccg ctcgggtccac aagctctcca cttgcccctt ttaggaagtc cgtcccccg  
 1261 gttcgggtac ccctgcccc tcccatattc tcccgtctag cacctttgat tctcccaaa  
 1321 cccggcagcc cgagactgtt gcaaaccggc gccacagggc gcaaagggga tttgtctctt  
 1381 ctgaaacctg gctgagaat tgggaactcc gtgtgggagg cgtgggggtg ggacgggtgg  
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 1561 gcggggcag aggggcggta tctgctgctt tggcagcaaa ttgggggact cagtctgggt  
 1621 ggaaggtatc caatccagat agctgtgcat acataatgca taatacatga ctcccccaa  
 1681 caaatgcaat gggagtttat tcataacgcy ctctccaagt atacgtggca atgcgttct  
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 1801 tctacactaa catcccagc tctgaacgcy cggccattaa tacccttctt tctccactc  
 1861 tccctgggac tcttgatcaa agcgcggccc tttccccagc cttagcgagg cgcctgagc  
 1921 cctggtacgc gcgtggcgtg gcgggtggcg cgcagtgcgt tctctgtgtg gagggcagct  
 1981 gttccgcctg cgatgattta tactcacagg acaaggatgc ggtttgtcaa acagtactgc  
 2041 tacggaggag cagcagagaa agggagaggg tttgagaggg agcaaaagaa aatggtaggg  
 2101 gcgcgtagtt aatcatgcy gctctcttac tctgtttaca tcctagagct agagtgcctg  
 2161 gctgccccgc tgagtctcct cccacacctc cccacctcc cataagcgc  
 2221 cctcccggt tccaaagca gagggcgtgg gggaaaagaa aaaagatcct ctctcgtaa  
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 2461 ataaaagccg gttttcgggg ctttatctaa ctgcctgtag taattccagc gagaggcaga  
 2521 gggagcgagc gggcgccggg ctagggtgga agagccgggc gagcagagct gcgctgggg  
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 3001 agagtgttta tggtaactgg ggctgggggtg gggggtaatc cagaactgga tcggggtaaa  
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 3421 cgcggcgatt ccaaccgcc ctgatccttt taagaagttg gcatttggct ttttaaaaag  
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 3841 tttatggagg ggtgttaaag cccgcggctg agctcgccac tccagccggc gagagaaaga  
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 4261 cgggggctcg gcgggcacca agccgctggt tcaactaagt cgtctccgag atagcagggg  
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 5761 ggagccagtg aactgcctca agagtgggtg ggctgaggag ctgggatctt ctccagctat  
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 6001 tgaagtgttc ttggtaaagt ccctcaaaaa taggaggtgc ttgggaatgt gctttgcttt  
 6061 ggggtgtgtc aaagcctcat taagtcttag gtaagaattg gcatcaatgt cctatcctgg  
 6121 gaagtgtcac ttttctgtgc catgccataa cccagctgtc ttccccttta tgagactctt  
 6181 accttcatgg tgagaggagt aagggtggct ggctagattg gttctttttt ttttttttcc  
 6241 ctttttttaag acggagtctc actctgtcac taggctggag tgcagtggcg caatcaacct  
 6301 ccaacccccct ggttcaagag attctcctgc ctccagcctc caagtagctg ggactacagg  
 6361 tgcacaccac catgccagcc taatttttgt aatttttagta gagatggggg ttcatcgtgt  
 6421 tggccaggat ggtctctcct gacctcacga tccgcccacc tcggcctccc aaagtctggt  
 6481 gattacaggt gtgagccagg gcaccaggct tagatgtggc tctttgggga gataattttg  
 6541 tccagagacc tttctaacgt attcatgcct tgtatttcta cagcattaat ctggtaattg  
 6601 attattttaa tgtaaccttg ctaaaggagt gatttctatt tcctttctta aagaggagga  
 6661 caaagcaaat gaggaagaaa tcgatgttgt ttctgtggaa aagaggcagg ctccctggca  
 6721 aaggtcagag tctggatcac cttctgctgg agggccacagc aaacctctc acagcccact  
 6781 ggtcctcaag aggtgccacg tctccacaca tcagcacaac tacgagcgc ctcccctcac  
 6841 tcgggaaggac tatcctgctg ccaagagggt caagttggac agtgtcagag tccctgagaca  
 6901 gatcagcaac aaccgaaaat gcaccagccc caggtcctcg gacaccgagg agaattgcaa  
 6961 gaggcgaaca cacaacgtct tggagcgcca gaggaggaac gagctaaaac ggagcttttt  
 7021 tgcccctgctg gaccagatcc cggagtggga aaacaatgaa aaggccccc aaggtagttat  
 7081 ccttaaaaaa gccacagcat acatcctgtc cgtccaagca gaggagcaaa agctcatttc  
 7141 tgaagaggac ttggttcgga aacgacgaga acagttgaaa cacaacttg aacagctacg  
 7201 gaactcttgt gcgtaaggaa aagtaaggaa aacgattcct tctaacagaa atgtcctgag  
 7261 caatcaccta tgaacttggt tcaaatgcat gatcaaatgc aacctcaca ccttggtgta  
 7321 gtcttgagac tgaagattt agccataatg taaactgcct caaattggac tttgggcata  
 7381 aaagaacttt tttatgctta ccatcttttt tttttcttta acagatttg atttaagaat  
 7441 tgttttttaa aaattttaag atttacacaa tgtttctctg taaatattgc cattaatgt  
 7501 aaataacttt aataaacgt ttatagcagt tacacagaat ttcaatccta gtatatagta  
 7561 ctagtatta taggtactat aaacctaat tttttttatt taagtacatt ttgcttttta  
 7621 aagttgattt ttttctattg tttttagaaa aaataaaaata actggcaaat atatcattga  
 7681 gccaaatcct aagttgtgaa tgttttggtt cgtttcttcc cctcccac caccaccatc  
 7741 cctgtttggt ttcatcaatt gcccttcag agggcgtct taagaaaggc aagagttttc  
 7801 ctctgttgaa atgggtctgg gggccttaag gtctttaagt tcttgagggt tctaagatgc  
 7861 ttctgggaga ctatgataac agccagagtt gacagttaga agaatggca gaaggcagg  
 7921 gagaaggatg gaggtaggca aaggagatac aagaggtaa aggtagcagt taagtacaca  
 7981 aagaggcata aggactgggg agttggaggg aaggtgagga agaaactcct gttactttag  
 8041 ttaaccagtg ccagtcacct gctcaactcca aaccaggaa tt

## Cyclin A

LOCUS cyclinA 1181 bp ds-DNA PRI 21-JUN-1993  
DEFINITION Human cyclin A (CCNA) gene sequence.  
ACCESSION M96390  
KEYWORDS cyclin A.  
SOURCE Homo sapiens DNA.  
REFERENCE 1 (bases 1 to 1181)  
AUTHORS Nikaido,T. and Yamamoto,M.  
TITLE p34cdc2 and cyclin A expression may be suppressed by RB and/or p53  
JOURNAL Unpublished (1992)  
COMMENT In the S-phase of the cell cycle, cyclin A complexes with E2F (a cellular transcription factor), p107, and cdk2 kinase. Arroyo et al. (Mol. Cell Biol. 13: 6537-46) report that the E7 protein of HPV-16 associates with the E2F-cyclin A complex. Arroyo et al. (Mol. Cell Biol. 13: 6537-46) further note that the high-risk HPV has a higher binding efficiency to the E2F-cyclin A complex than the low-risk HPV type. Steinmann et al. (Oncogene 9: 387-94) report a 20-fold increase in the level of cyclinA as a result of HPV-16 infection in human keratinocytes. Pagano et al. (Oncogene 7: 1681-6) report that the E2F-cyclin A complex can occur in HPV-18 infected cells independent of pRB.  
BASE COUNT 302 a 279 c 262 g 338 t  
ORIGIN  
1 aagctttgta tattcttata tttatatata aatataaaaa tttgttaaag gcacgtag  
61 ttaagagagt tttattttaa taaggtcata ttgtttttac tatgttttaa aaactttact  
121 ctgaaaggaa cataattata tctaggtcac tagaacgtca ttgtgttttt tgttggttgc  
181 acagcttggg gaaaaataga aaaaaattaa tgactgattt gaatattttg taatgcactg  
241 ctatttatta tatatatcaa cagtagttca aggtgccatc ttaaattaat tgcattctca  
301 ttaggaaaaa taaaagcat aaaacacaaat ttctggttac tatgaataaa cgcctaaatg  
361 ttaagatgac attacagtct tgacacttga gtactgtatt actatgtgag ctccgtgta  
421 aataatttat gcacattatt taatcctaac aaccatata ctgtagtatt tagtcctat  
481 taacacataa gaaaacggag aatcggagat actgaaaaac gtgccccaga ttttagacct  
541 ttggaaaaag tcaacttaagc taactagacg tcccagagct aaaggctggg caacccaat  
601 gatagtcgcc aaagtttaat tccgtttaat tccctaaaag gcttagagtc agcttcggac  
661 agcctcgctc actaggtggc tcagcttaaa ataatcggaa gcgtcggggc ctaaatccta  
721 cctctccccg ccccgcgag cggttttctc cgcgccagc cagtttggtt ctccctctg  
781 ccccgccccg gtcagtttc ctttggttta cccttcactc gccctgacct tgtcgcttg  
841 aatgacgtca aggcgcgag cgctttcatt ggtccatttc aatagtcgag ggatacttga  
901 actgcaagaa cagccgccgc tccagcgggc tgctcgtgac atctctgggc gtctttggct  
961 cgccacgctg ggcagtgctt gctcgcctc ttcgcaacct cctcggccct gcgtggctc  
1021 gagctgggtg agcagcggg cgggctggta ggctggcctg ggctcgcacc ggcggctacg  
1081 actattcttt ggccgggtcg gtgcgagtgg tcggctgggc agagtgcacg ctgcttggcg  
1141 ccgaggctg atcccgcct cactcccgg gagcagtgat g

# Glucocorticoid Receptor

LOCUS GCORTICOIDR 2518 bp ds-DNA PRI 15-JUN-1990  
DEFINITION Human glucocorticoid receptor gene, partial.  
ACCESSION M32284  
KEYWORDS glucocorticoid receptor.  
SOURCE Human blood leukocyte DNA.  
REFERENCE 1 (bases 1 to 2518)  
AUTHORS Zong,J., Ashraf,J. and Thompson,E.B.  
TITLE The promoter and first, untranslated exon of the human glucocorticoid receptor gene are GC rich but lack consensus glucocorticoid receptor element sites  
JOURNAL Mol. Cell. Biol. 10, 5580-5585 (1990)  
COMMENT Authorin copy of sequence [Unpublished (1990) Univ. of TX Med. Branch, Galveston TX 77550] kindly submitted by B.E.Thompson, 22-FEB-1990.  
Mittal et al. (J Virol 67: 5656-9) identified three glucocorticoid response elements in the LCR of HPV-16 at position 7640, 7385, and 7474. This element, which binds the glucocorticoid receptor, is a partial palindrome and consists of the consensus TGTACANNNTGTCAT (Chan et al. J Virol 63: 3261-9). In the presence of glucocorticoids, up-regulation of the P\$<sub>97</sub> promoter is observed.

BASE COUNT 403 a 802 c 882 g 431 t  
ORIGIN

```
1 aggattaggt ggagctgagg cagcctcccg cccgtgtcag gagctggcaa gcgatgtcta
61 cctgtgtgtg cgcaaaagtt acctccccaa accctaaaacc cacacagcac aacctttccc
121 agatcacaaa aatcataatc tgtgtctgca caaggtagga ggctcggctc cggcatcgtc
181 gaagccttcc cgacgcggcg agctggggaa gggagctggg gcgggggctt cccgcacggg
241 caccctcgcg cccacgcccct ctccctttctc aggacggacc acgagttccc ttccccttgg
301 actgagggggg aagctcctaa caggaacatc tgtagggagt tgaacgctgg cattttaaag
361 ctgcctgtat tttgttttat ttgtaggggc aggggtccta tgaacgtgat aggggtgagca
421 acgcacagag tcgagggcag caaatgtcaa gattcggggg tggggcctgc accgggaact
481 tggacgcggg ccctggccgg ggtggaagaa gaggtcagga gttcggaaag gggctatatt
541 tcgccagcaa ctactatatt cgcctgcaac ttgcttttaa gctgcgcgcc cctgtctttc
601 cttaatacata ataataaaaa aaaagtgcaa agaaatccag ctcgctggag gttttgcatt
661 tggcgtgcaa cttccttctg gtgtgagcac attgggcggg aggggtgggg gttgaacttg
721 gcaggcggcg cctccttctg ccgcccgcgc gcctcgcag actcggggaa gaggggtggg
781 gacggtcggg gcgcccggga ggggtgggtc tgctttgcaa ctctctccc agtgcgagag
841 cgcggcggcg gcagctgaag acccggccgc ccagatgatg cgggtggggg ggacctgccg
901 gcacgcagca tccccccggg cccaagtac gtabgcgcgc acccccgcta tcccgctcct
961 tcctgaagc ctcccagag ggcgtgtcag gccgcccggc cccgagcgcg gccgagacgc
1021 tgcggcaccg tttccgtgca accccgtagc ccttttgaa gtgacacact tcacgcaact
1081 cggcccggcg gcggcggcgc gggcactca cgcagctcag ccgcccggag cgcgccggct
1141 cttgtggccc gcccgctgtc accgcagggg cactcggcgc ttgccgcaa ggggcagagc
1201 gagctcccga gtgggtctgg agccgcggag ctgggggggg gcgggaagga ggtagcagga
1261 aaagaaactg gagaaaactg gtggccctct taacgcgcgc ccagagagac caggctggcc
1321 ccgcccctgc cgcgcgccacc ctttttctg gggagttggg gcgggggggc gaagcgcggc
1381 gcaccgggcg gggcggccac gccaggggac gcggcgtgc aggcgcgtc gggccggggt
1441 ggcggggccc gcgaggagg cgtgggggca gggaccgcgc gcccccctgc agttgccaag
1501 cgtcaccaac aggttgcatc gttccccgcg ccgcccgcgc gccctcggg cggggagcgg
1561 ccgggggtgg agtgggagcg cgtgtgtgcg agtgtgtgcc gccgtggcgc gcctccacg
1621 ccgctccccg ctcggtcccg ctcgctcgcc caggccgggc tgcccttctg tgtccgcgct
1681 ctcttccctc cgcgcgcgcc tcctccattt tgcgagctc tgtctgtgac gggagcccga
1741 gtcaccgctt gccgtcgggg acggattctg tgggtggaag gagaccgca ccggagcggc
1801 cgaagcagct gggaccggga cgggcacgcg cgcgggaag ccccgaccgc cggagcccgg
1861 cgcggggcgg agggctggct tgtcagctgg gcaatgggag actttcttaa atagggctct
1921 cccccacc atggagaaag gggcggctgt ttacttctt tttttagaa aaaaaaatat
1981 atttccctcc tgctccttct gcgttcacaa gtaagttgt ttatctcggc tgcggcggga
2041 actgaggacg gtggcgggag agcggctcct ctgccagagg taagaagcga ggcgggaggg
2101 ggcgggggcg cgctcgtcc cccgaggtgc cgctgggacc ggagacaact cgggggcccg
2161 cgcgggagcc tacaactttt tattagcctc ggggagtggt ggtggggggc tggcaagggc
2221 cgggcgacgg tgacgaaagg gtacgcgcgc ggtgacagcg ctggcctctt cctctcctc
2281 cgccggcgtc cctggccggg ccgaggggga ggaacctgac ctcgacggc gagcggagcc
2341 ctgtcgaact ccggggcttc gagcctctca ttctcgggg aatcctggcc tcttttctc
2401 ccctagtgtc ccttttctc caagggggtc gcccgacacc cgttttctg gtgagactaa
2461 gccgctctg aattttact gcccaatat ttcaaccacc ccgcccagc gcgagccc
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junB

LOCUS junB 2136 bp ds-DNA PRI 30-MAR-1994  
 DEFINITION Human transactivator (jun-B) gene, complete cds.  
 ACCESSION M29039  
 KEYWORDS c-myc proto-oncogene; transactivator.  
 SOURCE Homo sapiens DNA.  
 REFERENCE 1 (bases 1 to 2136)  
 AUTHORS Schuette,J., Viallet,J., Nau,M., Segal,S., Fedorko,J. and Minna,J.  
 TITLE jun-B inhibits the transforming and transactivating activities of c-jun  
 JOURNAL Cell 59, 987-997 (1989)  
 COMMENT Authorin copy of sequence [1] kindly submitted by J.D.Minna 10-OCT-1989.

Thierry et al. ( J Virol 66: 3740-8 ) demonstrated that JunB binds AP-1 target sites within the LCR of the HPV-18 P\$<sub>105</sub> promoter during human keratinocyte infection. Mutation of these two target sites destroyed P\$<sub>105</sub> promoter activity in these cells. Bossy-Wetzel et al. (Genes Dev 6: 2340-51) reported that junB or c-jun are overexpressed during the aggressive fibromatoses or fibrosarcoma stage of BPV infection. They also correlated this overexpression to changes in fibroblast cell shape and anchorage dependence.

BASE COUNT 387 a 716 c 671 g 362 t  
 ORIGIN

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1 cgcgagccgc ctccctccc tccccacgct cgaggagggg ggcgcggggg cccggctccg
61 gcgacggcca atcgagcgc acttccgtgg ctgactagcg cggataaag gcgtgtggct
121 caggctgagc ggctgggacc ttgagagcgg ccaggccagc ctccggagcca gcagggagct
181 gggagctggg ggaacgcagc ccaggaaagc tatcgcgcca gagagggcga cgggggctcg
241 ggaagcctga cagggctttt gcgcacagct gccggctggc tgctaccgcg ccgcgccagc
301 ccccgagaac gcgcgaccag gcaccagtc cggtcaccgc agcggagagc tcgcccgtcg
361 ctgcagcgag gcccgagcgc gcccgcagg gacctcccc agaccgcctg gcccgcccg
421 atgtgacta aaatggaaca gcccttctac cacgacgact catacacagc tacgggatac
481 ggcggggccc ctggtggcct ctctctacac gactacaaac tcctgaaacc gagcctggcg
541 gtcaacctgg ccgacccta ccggagtctc aaagcgcctg gggctcgcg acccggccca
601 gagggcgggc gtggcgggag ctacttttct ggtcagggct cggacaccgg cgcgtctctc
661 aagctcgctt ctccggagct ggaacgcctg attgtcccca acagcaacgg cgtgatcacg
721 acgacgccta cccccggg acagtacttt taccgccgcg ggggtggcag cgttggaggt
781 gcagggggcg cagggggcg cgtcaccgag gagcaggagg gcttcgccga cggctttgtc
841 aaagccctgg acgatctgca caagatgaac cacgtgacac ccccaacgt gtcctggggc
901 gctaccgggg ggcggggggc tgggcccggg ggcgtctacg ccggcccggg gccacctccc
961 gtttacacca acctcagcag ctactcccca gcctctgcgt cctcgggagg cgcgggggct
1021 gccgtcggga ccgggagctc gtaccgcagc accaccatca gctacctccc acacgcgccg
1081 cccttcgcag gtggccaccg gcgcgagctg ggcctggggc gcggcgccct cacttcaag
1141 gaggaaccgc agaccgtgcc ggaggcgcgc agccgggagc ccacgcgccg ggtgtcccc
1201 atcaacatgg aagaccaaga gcgcatcaaa gtggagcgca agcggctgcg gaaccggctg
1261 gcggccacca agtgccggaa gcggaagctg gagcgcatcg cgcgcctgga ggacaagggtg
1321 aagacgctca agcccgagaa cgcggggctg tcgagtaccg ccggcctcct cggggagcag
1381 gtggcccagc tcaaacagaa ggtcatgacc cacgtcagca acggctgtca gctgctgctt
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1501 gacggctggg cacacgcctc cactgggggt ccaggagca ggcgggtggc acccacctg
1561 ggacctaggg gcgccgaaa ccacactgga ctccggccct cctaccctgc gccagctct
1621 tccacctega cgtttacaag ccccccttc cactttttt tgtagtgtt tttctgctg
1681 gaaacagact cgattcatat tgaatataat atatttgtgt atttaacagg gagggaaga
1741 gggggcgatc gcggcgagc tggcccgcgc gcttggtact caagcccgcg gggacattgg
1801 gaaggggacc ccgcccctt gccctcccct ctctgcaccg tactgtggaa aagaaacag
1861 cacttagtct ctaaagagtt tattttaaga cgtgtttgtg tttgtgtgtg tttgtcttt
1921 ttattgaatc tatttaagta aaaaaaaaaa tggttcttta ttaatttctg ttgtctttt
1981 ttccaagctg ggagggcggg gggaaaaaaa aagcactggg ttgccccag ctcagtgtg
2041 ttggtggctc ggtcctgtat gtgtcccctt cgtcggttcg gcgcaggcat cttgtgttcc
2101 cagcccagga gtcccacct tcccgcgtcc ccagat

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LOCUS p107Rb 3960 bp ss-mRNA PRI 06-MAY-1993  
 DEFINITION Human retinoblastoma related protein (p107) mRNA, complete cds.  
 ACCESSION L14812  
 KEYWORDS cell cycle regulation protein; retinoblastoma protein;  
 tumor suppressor.  
 SOURCE Homo sapiens cDNA to mRNA.  
 REFERENCE 1 (sites)  
 AUTHORS Ewen,M.E., Xing,Y., Bentley-Lawrence,J. and Livingston,D.M.  
 TITLE Molecular cloning, chromosome mapping and expression of the cDNA  
 for p107 a retinoblastoma gene product-related protein  
 JOURNAL Cell 66, 1155-1164 (1991)  
 REFERENCE 2 (bases 1 to 3960)  
 AUTHORS Zhu,L., van den Heuvel,S., Helin,K., Fattaey,A., Ewen,M.,  
 Livingston,D., Dyson,N. and Harlow,E.  
 TITLE Inhibition of cell proliferation by p107, a relative of the  
 retinoblastoma protein  
 JOURNAL Genes Dev. (1993) In press  
 COMMENT p107, an Rb related protein, has been shown to complex with  
 Rb (Carloti et al. J Gen Virol 74: 2479-86 ). Davies et al.  
 (J Virol 67: 2521-8) demonstrated that the HPV-16 E7 protein  
 binds p107 and that this binding involves the Rb target site.  
 B-myb, a gene involved in cell cycle progression, is  
 inappropriately transcribed during G1 in HPV-16 E7 transactivated  
 cells. Lam et al (EMBO J 13: 871-8) demonstrated that B-myb  
 transcription is regulated through interactions with p107.  
 BASE COUNT 1257 a 771 c 869 g 1063 t  
 ORIGIN  
 1 cgggtagcgc gcctgggagg gagaagaag tcgggggccc tggcgcgcag cccgcggggc  
 61 ctgaagggat gttcgaggac aagccccacg ctgagggggc gccggtggtc gccgcagccg  
 121 gggaggcgcct acaggccctg tgccaggagc tgaacctgga cgaggggagc gcggccgaag  
 181 ccctggacga ctttactgcc atccagggca actacagcct agaggggagaa gttacacact  
 241 ggttggcatg ttcattatat gttgcatgcc gcaaaagcat tattcccacg gttggaagg  
 301 gtatcatgga aggcaactgt gtttacttta ccagaatact acgttcagct aaattaagtt  
 361 taatacaatt ttttagtaaa atgaagaaat ggatggacat gtcaaatcta ccacaagaat  
 421 ttcgtgaacg tatagaagg ctagagagaa attttgaggt gtctactgta atattcaaaa  
 481 aatgatgacc aatTTTTTTA gatataattc aaaatccata tgaagaacca ccaaagtac  
 541 cacgaagccg gaagcagagg aggattcctt gcagtgttaa ggatctgttt aatttctgtt  
 601 ggacactttt tgtttatact aagggtaat ttcggatgat tggggatgac ttagtaaac  
 661 ctatcatctt acttctatgc tgcttggatc tgatttttgc caatgcgatt atgtgccc  
 721 atagacaaga cttgctaaat ccatcattta aaggtttacc atctgatttt catactgctg  
 781 actttacggc ttctgaagag ccaccctgca tcattgctgt actgtgtgaa ctgcatgatg  
 841 gacttctcgt agaagcaaaa ggaataaagg agcactactt taagccatat atttcaaac  
 901 tctttgacag gaagatatta aaaggagaat gcctcctgga ctttcaagt tttactgata  
 961 atagcaaaag agtgaataag gagtatgaag agtatgttct aactgttggg gattttgatg  
 1021 agaggatcct tttgggagca gacgcagaag aggaatttgg aacacctcga aagttcactc  
 1081 gtgacacccc attagggaaa ctgacagcac aggctaagt ggagtataac cttcaacagc  
 1141 actttgaaaa aaaaaggta tttgcacctt ctacccact gacgggacgg agatatttac  
 1201 gagaaaaaga agcagtcatt actcctgttg catcagccac ccaaagtgtg agccggttac  
 1261 agagtattgt ggctggtctg aaaaatgcac caagtgacca acttataaat atttttgaat  
 1321 cttgtgtgcy taatcctggt gaaaacatta tgaataactt aaaaggaata ggagagactt  
 1381 tctgtcaaca ctatactcaa tcaacagatg aacagccagg atctcacata gactttgctg  
 1441 taacagact aaagctggca gaaatTTTgt attataaaat actagagact gtaatggttc  
 1501 aggaaacacg aagacttcat ggaatggaca tgtcagttct ttagagcaa gatataattc  
 1561 atcgttcctt gatggcttgt tggttggaaa ttgtgctctt tgctatagc tcacctcgta  
 1621 cttttccttg gattattgaa gttctcaact tgcaaccatt ttacttttat aagggttattg  
 1681 aggtggtgat ccgctcagaa gaggggctct caagggacat ggtgaaacac ctaaacagca  
 1741 ttgaagaaca gattttggag agtttagcat ggagtcaaga ttctgcaact tggggaggctc  
 1801 tccaggtttc tgcaaaaaa gttcctacct gtgaagaagt tatattcca aataactttg  
 1861 aaacaggaaa tggaggaaat gtgcaggac atcttcccct gatgccaatg tctcctctaa  
 1921 tgacccaag agtcaaggaa gttcgaactg acagtgggag tcttcgaaga gatatgcaac  
 1981 cattgtctcc aatttctgtc catgaacgct acagtctctc taccgcaggg agtgctaaga  
 2041 gaagactctt tggagaggac cccccaaagg aaatgcttat ggacaagatc ataacagaag  
 2101 gaacaaaatt gaaaatcgct ccttcttcaa gcattactgc tgaaaatgta tcaattttac  
 2161 ctgggtcaaac tcttctaaca atggccacag cccagtaac aggaacaaca ggacataaag

## Retinoblastoma-related p107

2221 ttacaattcc attacatggt gtcgcaaatg atgctggaga gatcacactg atacctcttt  
2281 ccatgaatac aaatcaggag tccaaagtca agagtcctgt atcacttact gctcattcat  
2341 taattgggtc ttctccaaaa cagaccaatc tgactaaagc acaagaggta cattcaactg  
2401 gaataaacag gccaaagaga actgggtcct tagcactatt ttacagaaag gtctatcatt  
2461 tggcaagtgt acgcttacgt gatctatgtc taaaactgga tgtttcaaat gagttacgaa  
2521 ggaagatatg gacgtgtttt gaattcactt tagttcactg tcctgatcta atgaaagaca  
2581 ggcatttggg tcagctcctc ctttgtgctt tttatatcat ggcaaaggta acaaaagaag  
2641 aaagaacttt tcaagaaatt atgaaaagtt ataggaatca gccccaagct aatagtcacg  
2701 tatatagaag tgttctgctg aaaagtattc caagagaagt tgtggcatat aataaaaaata  
2761 taaatgatga ctttgaaatg atagattgtg acttagaaga tgctacaaaa acacctgact  
2821 gttccagtgg accagtgaaa gaggaaagaa gtgatcttat aaaattttac aatacaatat  
2881 atgtaggaag agtgaagtca tttgcaactg aatacgactt ggccaatcag gaccatatga  
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