

Group G Sequences

HPV1 HPV4
HPV41 HPV63
HPV65

INTRODUCTION

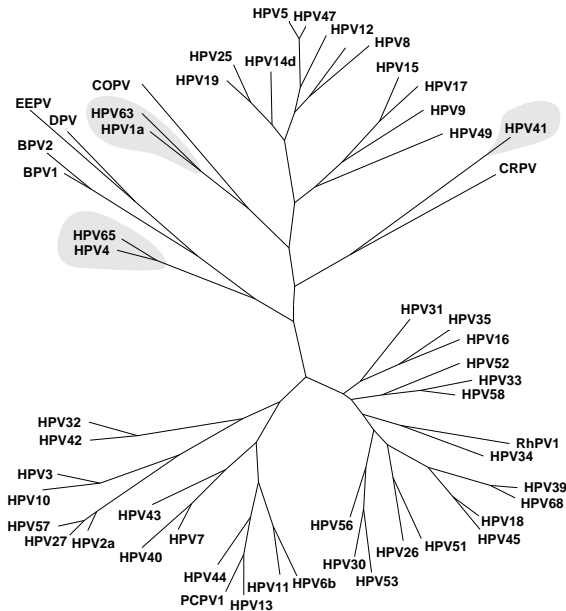
Group G consists of the human papillomaviruses HPV-1, HPV-4, HPV-41, HPV-63 and HPV-65, a group primarily associated with the benign cutaneous lesions, commonly seen in the general population.

HPV-1 and HPV-4, in conjunction with HPV-2, a group F virus, are the major etiologic agents of benign cutaneous papillomas in the general population. HPV-1 is primarily associated with deep palmo-plantar warts, while HPV-4 has been correlated with common warts and keratotic flat lesions on the hands and feet [1,2]. HPV-4 is also frequently present in hand warts of meat handlers [3]. HPV-41, like HPV-4, has been linked to flat warts, which are mainly found on the face and feet. Unlike the others in this group, HPV-41 has been detected in skin squamous cell carcinomas and its precursor lesions [4]. Also unique to HPV-41 is the absence of typical E2 binding sites in the LCR. However, modified E2 sites, as reported for BPV-1, have been located near the E6 gene [5]. HPV-65 has been linked to pigmented common warts and keratotic flat lesions, while HPV-63 is associated with multiple punctate keratotic lesions of the foot [2].

The viruses HPV-1, HPV-4, HPV-63 and HPV-65 produce intracytoplasmic inclusion bodies in most infected epidermal cells. The inclusion bodies primarily contain E4 proteins that can be used to histologically identify these viruses. HPV-4 like HPV-65, is associated with a homogeneous type of ICB. HPV-63 is associated with a filamentous type of ICB (FI-ICB) and HPV-1 presents a granular type (Gr-ICB) [2].

The primary target tissue of the group G viruses is the epithelium, however rare mucosal infection has been reported for HPV-1, HPV-4 and HPV-41. Both HPV-1 and HPV-41 have been identified in benign anogenital warts [4, 6, 7]. Conversely, HPV-4 has been identified in isolated cases of both normal and malignant oral lesions [8].

The members of Group G were placed together primarily on the basis of their clinical associations. Within the group are three subsets, each of which is at least as distant from the other subsets of Group G as any intergroup comparison of Groups A–F. For the purposes of analysis “by group” we chose to create two distinct subgroups, one consisting of HPV-1a and HPV-63, the other of HPV-4 and HPV-65. The latter pair of sequences are nearly close enough in terms of non synonymous base changes to be considered “close types”. HPV-41 is closer to the former subgroup than the latter, but sufficiently distant that it merits separation when analysis is performed. The members of Group G are generally closer to non primate animal papillomaviruses than to other human papillomaviruses.



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- [1] Danos,O., Katinka,M., and Yaniv,M. Human papillomavirus 1a complete DNA sequence: a novel type of genome organization among Papovaviridae. *EMBO* **1**: 231–236 (1982)
- [2] Egawa, K., Delius,H., Matsukura,T., Kawashima,M., and de Villiers,E.M. Two novel types of human papillomavirus, HPV 63 and HPV 65: comparisons of their clinical and histological features and DNA sequences to other HPV types. *Virology* **194**: 789–99 (1993)
- [3] Melchers,W., de Mare,S., Kuitert,E., Galama,J., Walboomers,J., van den Brule,A.J. Human papillomavirus and cutaneous warts in meat handlers. *J Clin Microbiol* **31**: 2547–9 (1993)
- [4] Grimmel,M., de Villers, E.M., Neumann,C., Pawlita,M., and zur Hausen, H. Characterization of a new human papillomavirus (HPV 41) from disseminated warts and detection of its DNA in some skin carcinomas. *Int. J. Cancer* **41**: 5–9 (1988)
- [5] Hirt,L., Hirsch-Behnam,A., de Villiers,E.M. Nucleotide sequence of human papillomavirus (HPV) type 41: an unusual HPV type without a typical E2 binding site consensus sequence. *Virus Res* **18**: 179–89 (1991)
- [6] Krzyzek,R.A., Watts,S.L., Anderson,D.L., Faras,A.J., and Pass,F. Anogenital warts contain several distinct species of human papillomavirus. *J Virol* **36**: 236–44 (1980)
- [7] Gissmann,L., deVilliers,E.M., and zur Hausen,H. Analysis of human genital warts (condylomata acuminata) and other genital tumors for human papillomavirus type 6 DNA. *Int J Cancer* **29**: 143–6 (1982)
- [8] Yeudall,W.A., and Campo,M.S. Human papillomavirus DNA in biopsies of oral tissues. *J Gen Virol* **72**: 173–6 (1991)

LOCUS HPV1a 7815 bp ds-DNA Circular VRL 06-JUL-1989
 DEFINITION Human papillomavirus 1a (HPV-1a), complete genome.
 ACCESSION V01116 X03321
 KEYWORDS circular; genome; origin of replication.
 SOURCE Human papillomavirus type 1a DNA.
 REFERENCE 1 (bases 1 to 7811)
 AUTHORS Danos,O., Katinka,M. and Yaniv,M.
 TITLE Human papillomavirus 1a complete DNA sequence: a novel type of genome organization among papovaviridae
 JOURNAL EMBO J. 1, 231-236 (1982)
 REFERENCE 2 (bases 1 to 7815)
 AUTHORS Danos,O.
 TITLE Direct Submission
 JOURNAL Submitted (23-JAN-1985) Danos O.
 REFERENCE 3 (bases 7574, 7691, 7692 and sites; revision)
 AUTHORS Palermo-dilts,D., Broker,T., and Chow,L.
 TITLE Human papillomavirus type 1 produces redundant as well as polycistronic mRNAs in plantar warts
 JOURNAL J. Virology 64, 3144-3149 (1990)
 COMMENT HPV-1a has a strong preferential association with deep plantar warts, characterized by a highly thickened corneal layer (hyperkeratosis). Generally, they are present in a single location. Skin warts are transmitted by direct contact with infected tissue or with contaminated objects. A majority of warts regress, spontaneously within two years. This is thought to be the result of a cell-mediated immune response. HPV-1 DNA has also been detected in anogenital tissue, however rarely.

The 7815 bp genome of HPV-1a is the embl corrected version of a previously published sequence. In addition to the embl corrections this sequence has been updated as noted in reference 3: a C has been changed to a T at nt 7574 and a CC has been changed to a GG at nts 7691-7692.

Palermo-dilts et al. (J. Virology 64, 3144-3149) report that the HPV1a mRNAs are highly analagous to those produced by HPV11 and HPV6 in splice site usage and major promoter locations. The exception to this similarity is the existence of a URR promoter in HPV1a, not found in either HPV11 or HPV6. This promoter, located at nt 7490, is active in plantar warts but not in primary keratinocytes. Thus, the HPV1a genome contains three putative promoters: one within the E7 ORF, one preceeding the E6 ORF, and one in the URR. The splice locations and their coding potentials are shown in the table below. All splice junctions shown annotated in the sequence are experimentally determined [3].

mRNA species	Splice donor/acceptor pair	Coding potentials
a	827/3200	E1 [^] E4, E5a
b	1231/3200	E1M, E2C, E5a
c	827/2545	E2, E5a
e	827/3200	E1 [^] E4, L1
	3592/5431	
f	7710/3200	L1
	3592/5431	
g	7710/5431	L1
h	7710/3200	E5a, E5, L2

HPV1a

```
BASE COUNT      2396 a   1482 c   1666 g   2271 t
ORIGIN
   1 gttaactacc atcattcatt attctagtta caacaagaac cTAGgagtta tatgccagaa
                                     E8 orf start ->
   61 gtaagccTAT AAAAtacaca ggTAAgactc tgcacaggac cagATGgcga caccaatccg
      signal ->      E6 orf start ->      E6 cds ->
  121 gaccgtcaga cagctttccg aaagcctctg tatcccatat attgATGttt tattgccttg
                                     E8 cds ->
  181 taatTTTTgt aattatTTTT tgtctaagtc tgagaagctg ctttttgatc attttgatt
  241 gcaccttTgc tggagagaca atttggtgTt tggatgctgt caaggggtgTg ctagaactgt
  301 tagcctattg gagtttTgtt tatattatca ggagTcttat gaggtaccgg aaaTAGaaga
                                     <- E8 end
  361 aatTTTggac agacctttat tgcaaattga actccgtTgt gttacatgca taaaaaaact
  421 gagTgtTgct gaaaaattgg aggtTgtgTc aaacggagaa agagTgcata gagttagaaa
  481 cagacttaaa gcaaagTgTA GttTgtgTcg cttgtatgTc ATATAAcaAT Ggtgggcgaa
      E7 orf start ->      E7 cds ->
                                     <- E6 end
                                     signal ->
  541 atgccagcac taaaggacct gtttcttcaa cttgaaccaa gcgtcctaga tttagatctt
  601 tattgttTac aggaggTgcc tcttgatgac atagaggagg agttagtgtc gcctcagcaa
  661 ccttatgctg tcgttgcttc ctgtgcctat tgcgagaaac tggttcgatt gaccgtcctc
  721 gcgcatcaca gcgccattag acagctggag gaactccttc tgcgatcttT GAacatcgtg
                                     E1 orf start ->
  781 tgcccactgt gcaccctaca gcgacagTAA aATGgcagat AATAAAgGTa ctgaaaacga
      E1 cds ->      <- E7 end      -> signal
      <- E7 end      -> signal
  841 ttggtTTTTg gtggaggcga cagattgtga ggaaacgtta gaggaaACCT CACTTGGTga
                                     -> E2 bind
  901 cctagataat gtttcttTgt ttagcgactt atctgattta ttagacgagg cgccgcaaag
  961 ccaggggaat tccctggaat tgttccaca gcaagaatcg ctggaaagcg aacaggaact
 1021 taatgcttta aaacgaaagt tactttacag tctcaggcg agaagcgcg acgaaacaga
 1081 cattgctagc attagtccta gattagaaac tatttctatt acaaagcaag acaaaaaaag
 1141 gtatcgaaag caactgtttt ctcaggatga tagtggttta gagctatcgc tgcttcagga
 1201 tgaactgaa aatattgatg aatcgacaca gTTagatcaa cagcagaaag aacatactgg
      5' sj /\
 1261 ggaagtTggg gccgctgggg tgaacatTtt gaaagctagt aatatccgcg ccgcattatt
 1321 aagcagattt aaagatacgg ctggcgtcag ttttacagAC CTGACGCGGT cgtacaagag
      -> E2 bind
 1381 caacaaaacc tgtTgtggag attgggtttt ggcagttTgg ggtgtccgtg aaaatTtaat
 1441 tgacagtTga aaagaattat tgcaaacca ttgtgtgtat attcaattgg aacatgcagt
 1501 aactgaaaaa aatagatttt tatttttatt ggtacgattt aaagcccaga aaagtagaga
 1561 gactgtgata aaacttataa ccacaattct tccagttgat gctagctata tttTgtctga
 1621 gcctccaaaa tcaagaagtg tggctgctgc attattttgg TATAAAagat ctatgtcttc
      signal ->
 1681 aactgttttt acatggggta caactttgga tgggattgca cagcaaacc ttattaatca
 1741 tcagttagat tccgaaagtc cctttgagct ttgtaaaat gttcagTggg cctatgataa
 1801 tggacataca gaagagtTga aaattgcata ttattatgct gttttagcag atgaggatga
 1861 aaatgcaagg gcatttctaa gctctaattc acaggcaaaa tatgtgaaag actgtgcaca
 1921 aatggtaaga cactattTac gtgctgagat ggcacaaatg tctatgtcag agtggatttt
 1981 tagaaaacta gataatgtag aaggttctgg taattggaaa gaaattgtaa gatttttaag
 2041 atttcaagaa gttgaattta taagctttat gattgcattt aaagattTgt tatgtgTtaa
 2101 gccaaagaaa aactgtTtTg taatattTgg acctccaaat acaggaaaat caatgtTttg
 2161 tacaagTtta ttaaagTtTg taggagggaa agtgattTca tactgtaaca gtaaaagTca
 2221 gttttggTtg cagcctctgg ctgatgctaa gatagggcta ttagatgatg caacaaagcc
 2281 atgtTgggat tatatggaca tttatatgag aaatgcattg gatggtaaca ctattTgtat
 2341 tgattTaaaa catagagctc ctcaacaaat taaatgocca ccttactta ttactagtaa
 2401 tattgatgtt aaatcagata cctgtTggat gtattTgcat agtagaatat cagctttTaa
 2461 attTgctcat gagtttccat ttaaagacaa tggTgatcca ggattTtct taacagacga
 2521 aaattggaaa tctttctTtg aaAGgtTttg gcaacagTta gaatTAAGtg accaagaaga
      /\ 3' sj      E2 orf start ->
 2581 cgagggaaac gATGgaaaac ctcagcagtc gcttagactt actgcaagag cagctaatga
      E2 cds ->
 2641 acctataTGA acaggacagt aaattgatag aagatcaaat taagcagTgg aatcctaatta
      <- E1 end
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2701 gacaagaaca agttcttttc catttcgcca gaaaaaatgg ggtaatgaga attggattgc
2761 aggcagttcc atcttttagcg tcctcacagg agaaggcaaa gacagctatt gaaatgggtg
2821 tacatttaga gtcttttaaag gactcacctt atggcacaga ggattggtca cttcaagaca
2881 ctagcagaga gctgtttttg gcacccccag ctggcacctt caagaagagt ggcagcacac
2941 ttgaggttac ctatgacaat aacctgata atcagacaag gcacacaatt tggatcatg
3001 tgtattatca aaatggggac gatgtatgga gaaaagtac cagtgggtgtt gatgctgtag
3061 gagtgtacta tttagaacac gatggcTATA AAaattatta tgtgttattt gctgaggagg
          signal ->
3121 cctctaagta cagcacaaca ggacaatatg ctgTAAatta caggggtaaa aggtttcaa
          E4 orf start ->
3181 ATGttatgtc ttccactAGc tcccccaagg ctgctggggc tcctgcagta cactccgact
E4 cds ->          /\ 3' sj
3241 acccaacct atccgagagT GAcaccgcc agcaatcgac gtccatcgac tacaccgaac
          E3 orf start ->
3301 tcccaggaca gggggagacc tcgcaggctc gacaaagaca gcagaaaaca cctgtacgca
3361 gacggcctta cggacggcga agatccagaa gtcccagagg tggaggacga agagaaggag
3421 aatcaacgcc ctctAGgaca cccggatctg tccttctgc gcgagacgtt ggaagtatac
          <- E3 orf end
3481 acacaacgcc tcaaaagga cattcttcaa gacttagacg acttctgcag gaagcttggg
3541 atccaccctg ggtctgtgTA Aaaggggtg ccaatcagct taagtgtctc agGTacagac
          <- E4 end          5' sj /\
3601 ttaaagcatc tactcaagtt gactttgaca gcataagcac cacatggcat tggacagata
3661 gaaaaaacac cgagaggata ggtagtgcta gaatgttagt aaagtttatt gatgaggctc
3721 aacgagagaa gtttcttgag agagttgctt tgcccagatc agtgtctgtg tttttgggac
3781 agtttaatgg gtctTAAaat taatggaagt tgattttgct tggacgtgtg tacatagctc
          <- E2 end
3841 ctgTATATAT tccccctcta cccccacata cctTGAagct tgcaacattg TAAcaaATGt
          signal ->          E5 orf start ->          L2 orf start ->
          L2 cds ->
3901 atcgcctacg tagaaaaacgc gctgccccca aagaTATATA cccctcatgc aaaatatcaa
          signal ->
3961 acacctgccc acctgacatt caaAATAAAA ttgagcatac aacaattgct gataaaatat
          signal ->
4021 tgcaatatgg cagtctggga gtttttttgg gaggtttggg cattggaaca gccagaggct
4081 ctggaggaag aattggttat actcccctcg gtgagggtgg tggggttaga gttgctactc
4141 gtccaactcc agTAaggcct acaatacctg tggaaacagt aggccccagt gaaattttcc
          <- E5 orf end
4201 ccatagatgt tgtagatcct acaggccctg ctgttattcc cctacaagat ttaggtagag
4261 acttcccaat accaactgty caggttattg cagaaattca ccctatttct gacataccaa
4321 acattggtgc atcttcaaca aatgaaggag aatctgccat attagatgtg ttacgaggga
4381 atgcaacat acgcactggt tcaagaacac aatacaataa cccctcttcc actggtgcat
4441 ctacatctaa tataagtgct ggagaagcat caacatcaga tattgtattt gttagcaatg
4501 gttcagggtga caggggtggt ggcgaggata tccccttggg agaattaaac ttaggccttg

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HPV1a

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4561 aaacagacac atcttctgtt gtacaagaaa cagcattttc cagcagcaca CCAATtgctg
                                signal ->
4621 aaagaccctc ttttaggccc tcaagattcT ATAATAGgcg tctatatgaa caggtgcaag
                                signal ->
4681 tacaagacc  taggttcggt gagcagccac agtcaatggt cacttttgat aatccagcat
4741 ttgagccaga gcttgatgag gtgtctatta ttttccaaag agacttagat gctcttgctc
4801 agacaccagt gcctgaattt agagatgtag tttatctgag caagcccaca ttttcgctgg
4861 aaccaggggg acggttaagg gttagccgcc ttggcaaaag ttcaactatt cgtacacgcc
4921 tgggcacagc aatggcgcc  agaaccoact ttttctatga ttttaagttct attgctccag
4981 aagactcaat tgaattattg ctttaggtg agcatagtca aacaacagtc attagttcca
5041 acttaggtga cacagcattt atacaagggt agacagcaga ggatgactta gaagttatct
5101 ctttagaacc accacaatta tattcagaag aagagctttt agacacaaac gaaagtgtgg
5161 gcgaaaattt gcaacttact attactaact cagaggggtga ggtttctata ctagatttaa
5221 cacaaagcag agtcaggcca ctttttgcca ctgaagatac tagcttgcat gtatattacc
5281 caaattcttc taaagggact ccaataatta atcctgaaga atcatttaca ctttggtta
5341 ttaTAGctct taacaactca acaggggatt ttgagttaca tcctagtctt agaaagcgtc

L1 orf start ->
5401 gtaaaagagc ttATGtaTAA tgtttttcAG atggctgtct ggttaccagc gcagaataag
      L1 cds ->   <- L2 end   /\ 3' sj
4561 ttctatcttc ctccccagcc catcactaga atcctgtcca ctgatgaata tghtaaccaga
5521 accaatctct tctaccatgc aacatctgaa cgtctactgc tggctggaca tcctttgttt
5581 gagatctcca gtaatcaaac tgtaactata ccaaaagtgt caccaaatgc atttagagtt
5641 tttaggggtc gttttgctga tccaaataga tttgcatttg gggataaggc aatttttaat
5701 ccagaaacag aaagatttag ttggggccta agagggatag agataggtag aggccagcct
5761 tttagtatag gaataacggg ccaccctctt ttaaataagt tagatgatgc agaaaatcca
5821 acaaatata  ttaactacta tgcaaatgga gattctagac aaaactactgc ttttgatgca
5881 aaacagacac aaatgttcct cgtcggctgt actcctgctt caggtgaaca ctggacaagt
5941 agtcgttgcc caggggaaca agtgaaactt ggggactgcc ccagggtgca aatgatagag
6001 tctgtcatag aagatggtga catgatggat attggttttg gggctatgga ttttgctgct
6061 ttacagcaag acaagtctga tgtcccctta gatgttgctt aagcaacatg caaatatcct
6121 tattatatca gaatgaacca tgaagcctat ggcaactcta tgtttttttt tgcacgtcgc
6181 gagcaaatgt ataccaggca cttttttact cgcggggggt cgggtgggtga taaggaggca
6241 gtcccacaaa gcctgtattt aacagcagat gctgaaccaa gaacaacttt agcaacaaca
6301 aattatgtag gcacaccaag tggtctatg gtttcatctg atgtccaatt gttaataga
6361 tcttactgyc ttcagcgatg tcaaggccag aataatggca tttgctggag aaaccagtta
6421 tttattacag ttggagataa taccagagga acaagtttat ctatcagtat gaaaaacaat
6481 gcaagtacta catattccaa tgctaatttt aatgattttc taagacatac tgaagaattt
6541 gatctttctt ttatagttca gctttgtaaa gtaaagttaa ctcccgaaaa tctagcctac
6601 attcatacaa tggaccctaa tatttttagag gattggcaac tatctgtatc tcaaccacct
6661 accaatcctc tagaagatca atataggttt ttagggctct ccttggcagc aaaatgtcca
6721 gaacagggcg ctctgagcc  ccagactgat ccttatagtc aaTATAAATt ctgggaagtc
                                signal ->
6781 gatctcacag aaaggatgtc cgaacaatta gaccaatttc cactaggaag gaaatttcta
6841 tatcaaatg  gcatgacaca acgtactgct actagtcca ccacaaagcg caaacagtg
6901 cgtgtatcta cgtcagccaa gcgcagcgct aaggctTAGT ATATAtTATA TATAactata
                                <- L1 end
                                signal ->
6961 tttattagta gattatttat TATATAtttt tatattttta tactttttat acttgtttag
                                signal ->
7021 ttctaataag acatgtaaga tttacattag tataagtagg catgtattta cataaaatag
7081 tcttggaaac cttttattag tgaacatca tttacaatag tgacatcata gttcatctgc
7141 aattgctatt ccatcgttct tcacatattc tacagtagtg ttctctagat tgtattgcta
7201 ttttctgtt aggcacacaa caacatctgt acatggacca aacaaccac tttcatttta
7261 ttgtgctgca tatattccag attggtgagg atttatttgt ttagactccg gtgcattata
7321 cacaagtgtg cttttttgt  gttctctgat tgattgtgtg ttattttcct gcaatatgcA
                                signal ->
7381 ATAAAagtg  gctgtccttt ctttttgta atccctocct actccAATAA Aaaatcccta
                                signal ->
7441 ccctaaaat ctgtttgctg tggttttatt aataattgcg ctctttTATA TAATAagtac
                                signal ->
                                |-> mRNA start site
                                from P(7490)
                                promoter
7501 tattaacACC GCACCCGTTg tggctaacc cttatggat ttaaagact acacctacag

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```
      -> E2 bind
7561 gatgtattgt ctttattggt tatggtttac cgcgctCCAA AGACGGTTG CCCAAAGACG
      -> repeat region start
7621 GTTTGCcaac cgcggttagg acttgtttca atttgctgcc aaacttatct ggtcgtgctc
      <- repeat region end
7681 caacggggtt ggtgccaagc acctaaaacg GTaggtgtgt actcttttca agaattaaca
      5' sj /\
      -> E2 bind
7741 aaggagattt ctcccgcaa attagtttctg agcgACCGAA TTCGGTcgta aaaatctaaa
      -> E2 bind
7801 gtgatgattg ttggt
```

HPV4

LOCUS HPV4 7353 bp ds-DNA VRL 04-OCT-1993
DEFINITION Human papillomavirus type 4 (HPV-4), complete genome.
ACCESSION X70827
SOURCE Human papillomavirus type 4 DNA.
REFERENCE 1 (bases 1 to 7353)
AUTHORS Egawa,K., Delius,H., Matsukura,T., Kawashima,M. and
De Villiers,E.M.
TITLE Two novel types of human papillomavirus, HPV63 and HPV65
comparisons of their distinct clinical and histological features
and their DNA sequences to other HPV types
JOURNAL Virology 194, 789-799 (1993)
COMMENT Submitted (27-JAN-1993) on tape to the EMBL Data Library by: H.
Delius, Deutsches Krebsforschungszentrum, Abt ATV, Im Neuenheimer
Feld 506, W -6900 Heidelberg, FRG.

HPV-4 has a strong preferential association with common warts,
characterized by a dome shape with multiple conical projections
(papillomatosis). Most often they are multiple and are found on
the hands. Skin warts are transmitted by direct contact with
infected tissue or with contaminated objects. A majority of warts
regress spontaneously within two years. This is thought to be the
result of a cell-mediated immune response.

Infection from HPV-4 produces intracytoplasmic inclusion bodies
(ICB) in most cells of the epidermis. These inclusions have been
previously shown to contain mainly E4 proteins for HPV-1. To
date, three morphologically different types of ICBs have been
distinguished, and certain types of HPV are preferentially found
in one type of ICB or another. HPV-4, like HPV-65, is associated
with a homogenous type of ICB (Hg-ICB). The author notes that
histologically, these Hg-ICBs were usually present as a single
inclusion in each cell, surrounding an extremely vacuolated
nucleus. HPV-4 and HPV-65 have both been detected in common
warts and keratotic flat lesions on the hands and feet.

BASE COUNT 2363 a 1290 c 1543 g 2157 t
ORIGIN

```
1 gtctgtaatg atagttggca acaatcatta cttatagcta tataTAAccg gaagagatac
                                     E6 orf start ->
61 atataaaaag ggacagtgca tttctactaa atcctgtcca gATGgcagat ggcagacctg
                                     E6 cds ->
121 caaccttggc cgacttctgc agacgattcg acatttccct ttttgatttg cgccttactt
181 gtattttttg ttctcatact gtcgatcttg cggatcttgc tttattctat ctaagaaac
241 ttagttagt attagagga aattgttatt atgcatgttg ttctgaatgc ttaagattaa
301 gtgcaactgt tgaacaagag aattattttc aatgttctat taaagctgta catttggagg
361 aaattgctca gaaaaagatt aaggaaattt gcattagatg catttgctgc cttagattac
421 ttgatattgt tgagaaatta gatttattat actctgacga gacttgctat ttaataaggg
481 gtttggggag gggctattgc agaaattgta tTAGgaaaca ATGAgaggag cagcgccac
                                     E7 cds -><- E6 end
                                     E7 orf start ->
541 ggttcagat cttaatttag aactaaatga cttagtgtta ccagcaaacc tgctgagtga
601 ggaggtcttg caatcttcag atgatgagta tgagattaca gaggaggagt cggtggttcc
661 attagaata gacacctgtt gctatagatg tgaagttgct gtaagaatta cattgtagc
721 tgctgagctc ggactacgga ccttgaaca acttcttgta gaaggaaagc TGAcgttttg
                                     E1 orf start ->
781 ctgaccgct tgtgcaagaa gtcttaacag aaATGgcaga TAAaggtaga gacaattttg
                                     E1 cds -> <- E7 end
841 acttagaagg gaataattgg tatattgtcc atgaagcaga atgcaactgac agtatagata
901 cgttggatga tttatgagc gaaagtaatg acgatcaca catttctaac ttaattgatg
961 acgatgtcgt tgatcagggg aattcccttg cgctgtacaa tgcacaaata aatgaggatt
1021 gtgacaatgc actagcacac ctaaaacgaa agtataacaa aagtccagag caggcagtcg
1081 ctgaattgag tccgcagttg caggctgtga aaataactcc tgaagacac agcaaaagga
1141 gattatttca ggacagtggg attttcgaag atgaagctga aaattctctt acacaggtag
1201 aatccgagag ccaggctgga ccttctagcc aagatggcgg cggagatatt aattgtgtg
1261 tgttacaaag tagtaacagg agggcaaca tgctagcaaa gtttaagaa tggtatgggg
1321 tctatacaaa tgaataaaca agaatttata aaagtata atctttagt gataattggg
```



```

1381 taatagttat ttttagagct gctggtgaag tattagaaag ttcaaagatt gttttaaagc
1441 agcattgtac atatatcaa gttaagatct ttggatttcc agctttatat ttagtacagt
1501 ttaaaagtgc gaaaagtaga gaaactgtac aaaagttgat gtgttctata ttaaatatcc
1561 aagaatatca aatgttatgt gatcctccaa aattacgaag tgtaccaca gcattatact
1621 tttataagca tgctatgta acagagagtt ctgttttgg acaaacaccg gattggatcg
1681 caaaacaac tctcgtaat catcaagcag caactactgc agagactttt gagttatcta
1741 gaatggttca gtggcatac gataataatt atgtggatga atgtgacatt gcttatcact
1801 atgcaatgta cgcagaggag gatgcaaatg ctgctgctta tttaaaaagt aataatcaag
1861 taaagcatgt acgagattgt agtacaatgg tcaggatgta taaaagatat gaaatgagag
1921 atatgtcaat gtcagaatgg atttataaat gttgtgatga atgttctgaa gaaggagatt
1981 ggaagccaat ctcacagttt ttaaaatata aagggtgtaa tatattatcc tttcttatag
2041 tgcttaaatc atttttaaaa ggtattccaa aaaaaactg tatagttatt catggccac
2101 cagatacagg aaaatcatta ttttgttatt cttttataaa atttttaaaa ggaaaagtag
2161 tttcatatgt aaatagaagt agccattttt ggttgcagcc tctgatggat tgcaaggtag
2221 gatttatgga tgatgctacc tatgtgtgct ggacatata agatcaaaa ttaaggatg
2281 cattagatgg taatccaatg tgtattgacg ctaaacacag agcaccaca caattaaaat
2341 taccaccaat gctaataacg tcaaatattg atattaaaca ggaacaatct ttaatgtatt
2401 tacacagtag aatacagtggt ttaattttc ctaaaaaat gcctatttta gatgatgga
2461 gtccatgta tacatttact gacggtactt ggaatcttt ttccaaaag cttggcagac
2521 aattagaatT AAcagatcct gaagaggaaa acaATGgagt ccctagtcgc acgtttcgat
E2 orf start -> E2 cds ->
2581 gcacttcaag aagcaattct gactcatatT GAGtcacag agagcacttt ggaatcccaa
<- E1 end
2641 atccaatatt gggaaaatat cagaaaagaa aatgctataa tgcattatgc tcgaaaacaa
2701 ggccataACCA AATTAGGTct acaaccactt cctacactag cagtaactga atacaatgca
-> E2 bind
2761 aagcaagcta ttcagataca ttttaactttg caatcattgt taaaatctcc ctttgcattc
2821 gaacgggtgga cattgacaga tgttagtgca gaactgataa atacctctcc acaaaaactgt
2881 ttaaaaaagg gaggttatga tgttgctgtg tggtttgata atgatagaca gaatgcaatg
2941 ctgtacacaa attgggactt tttatattat caagataTGA ATGaacagtg gcacaaagtt
E4 orf and cds start ->
3001 aaagggtgaag tggattatga tggcttatac tttacagacc atacgggaga aagagcttat
3061 tttacattat ttagctctga tgctcaaaga tttagcagaa ctggactgtg gactgtgcat
3121 tttaaaacc aagttatttc ctcccctatt gttagctcta catactctc ctccctcgac
3181 actgaggaac aacagttacc cgggcccctcc accagctact ccgaagtac cgagcaggcg
3241 agccctactc gaaggaggaa accgaggaaa tccgacgcga cctccaccac gtcccctgaa
3301 accgagggag tacgactacg acgaagacga cgagaaggaa aatcagggcc cgggtcagga
3361 gaaacccccc gcaaaagaag aagaggagga ggaagaggag gaggagagac cgaattggga
3421 tctgcaccat ctccctgcaga agtggggagc agacatcgac aagttgaaag acaaggctcg
3481 tcgcgacttg gactcttaca agcagaagct agggatccgc ctaTGAatatt gttaaagggc
<- E4 end
3541 acagcaaat ctttgaatg ttggagatat agaaaagtta actcaaatg ctgcaacttc
3601 ttattcatga gtactgtttg gaactgggtt ggagattgct cacataatca tagtcgcatg
3661 cttattgcat ttgatagcac tgaccaaaga gacgcttttg taaaacacaa ccttttctc
3721 aaactgtgta catataccta cggctcatTG AatagtttaT AAaATGcaaa gcttgagtag
<- E2 end
L2 orf start -> L2 cds ->
3781 aaggaaaaga gattcagttc caaatcttta tgcaaatgt caactgtctg gcaattgcct
3841 acctgatgta aaaAATAAAg tagaagctga tactcttgct gatcgtttgc tgagatggtt
-> signal
3901 gggagagtga atatacctag gaggtttggg tattggtact gggagagga gtggggggtc
3961 aactgggtat aatccaattg gagctccaag tagagtcaca cctagtggta ctttagtaag
4021 gctacagtg cctgtggaaa gtttgggacc ctcaaaaata atccaatag atgcaataga
4081 cccaacaaca tcttctgttg tgccattaga ggatctgacc atcccagatg tcacagtaga
4141 tagtggagat acaagaggaa tagggggagac tactcttcag cctgcacaag tagatatttc
4201 aacatcacat gaccctatat cagatgtcac tgggtctagc agccacccta caatcatatc
4261 tggcagggat aacgccattg cagtgtttaga tgtgtcccct atagaacctc ccacaaaacg
4321 gatagcattg gcaactaggg gagcctcagc aactccacat gtaagtgtca tatctggcac
4381 aACCGAATTC GGTcagtcac ctgatctgaa tgtattttgt aatgccacat tttcaggoga
-> E2 bind
4441 ttccattgggt tatacagaag aaattccatt agaaccgttg aaccctttc aagaattoga
4501 aatagaaaagc cctccaaaaa ctagtacacc acgtgacgtt ttaaactcgt caataggaag
4561 agcacgggat ttatataata gaagggttca gcaaacacct actaggaacc cagctttact
4621 gacacagcct tcccgcgcaa tagtatttgg atttgaat cccgccttg atgctgacat

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HPV4

```
4681 cactcaaaca tttgagcggg atttagaaca ggttgacgca gctccagatg ctgactttgc
4741 agacatagtc actatagggc gtccaaggtt ttcagagaca gatgctggtc aaattagagt
4801 tagcaggcctt ggacgccgag gcacaataaa aactagaagt ggtgtgcaaa ttgggcaggg
4861 ggttcattttt tattacgacc taagtacaat agatactgct gatgctattg aattatctac
4921 tttagggtcaa cattcaggag aacaaagcat tgttgatgct atgatagaaa gcagcttaat
4981 agatcctttt gaaatgcccg atcctacttt tacagaagaa caacagcttt tagatccact
5041 tacagaagat tttagtcagt cacacttggg gcttactagt agcagacgtg ggacatcatt
5101 tactatacct acaataccac ctggattagg tcttagaatt tatgtagatg atgtaggttc
5161 tgattttattt gtttcctatc cagaatctag agtaatacct gctggagggt taccaactga
5221 gccatttgtt cctctagaAC CAGCTTTGTT atctgatata tttagtagcg attttgtata
                                -> E2 bind
5281 tcgtcctagt ttatatcgca agaaacggaa acgattagaa atgtttTAA tgttttgcag
                                <- L2 end
                                L1 orf start ->
5341 gaacATGtcg agttggttat ctacaacggg taaagtctac ttacctccag ctcaacctgt
L1 cds ->
5401 ggcaagagtt ttggaaactg acgaatatac cactggaaca tctctgtatt tccacgctgg
5461 tacagaaagg cttttaactg taggccatcc ttattttcca gtgaaagatg tacaggaacc
5521 tcacaaagta ttagttccta aggtttcagg aagtcaattt agagtgttta gattcaattt
5581 gccagaccce aacagatttg ctttaattga taatggcttt tatgattctg atcatgaacg
5641 cctagtatgg aaactgaggg gaatagaaat aggaagagga ggaccgcttg gtagaggtac
5701 tacagggtcat cctttatata ataagtttgg agacacagaa aatcctaata gctacaaaaa
5761 gcaatcagat gataatagac aggatgtctc tttagaocca aaacaaacac agatgtttat
5821 tataggttgc actcctgcaa taggtgaaca ttgggataaa gctgaacctt gtcccagccc
5881 tgctccgcaa caggagatt gccaccaat agagcttcta aattcataca ttcaagatgg
5941 agatagtgtg gacattggat ttggggcttt caattttaa gctttgcagg ctgataaatc
6001 tagtgctcct ttggatgtca ttgccacagt ttgtaaatgg ccagattttt taaaaatggg
6061 gaaagatate tatggagata gcttgttttt ctttgggaaga agagaacaac tatatgccag
6121 acatttcttt gtcagagcag gcaccatggg agatgctcta ccagaacctt ttgaagctac
6181 ctcagattat tttattggtg ctcaaaaacca acaagatcag tacactttag gacctcatat
6241 ttatgtaggg acccctagt gctctttagt atccagtga tcccagttgt ttaatcgACC
                                E2 bind ->
6301 GTATTGGTTa aacagagctc aggggtacaaa taatggaatt tgttgggata atcagttgtt
6361 tgttactctt gtagataaca ctcataatac aaactttaca atttctgtga agtcagatgg
6421 tgctaagtac aattatcagt ataaagctag tgattttaa cagtaacctc gacatataga
6481 ggagtttgaa atggaattta ttttcaact ttgtaaagtt cctctaactg cagatgttat
6541 ggctcattta aatgtaatga atcctaatac tttggataat tggcagttaa attttgttcc
6601 accacctccc tctggaattg aggatcaata tagatttttg caatctagag ctacaagatg
6661 ccctacacag acccctgcaa ctgaaaaaga agatccatat aaagatttgt ctttttgggt
6721 tgttgattta agtgaagaat tttccagtga attgagccaa ttttccttag gcaggcggtt
6781 tttatatcaa agtggtttaa ttaatggttc tctaaaacgt aaaagaataa taagttcttc
6841 tcatgcacaa actaatacca aacgttctgc caaacgaaaa cggctctctga aaTAAcaatg
                                <- L1 end
6901 tgaactcttc tggaaatggtt tattctgcca ggaaaacctt caactgagcc aaattattat
6961 ataactgctc ttaatctcaa aattgagcta attatataag atttgcaaac gtgtatgtat
7021 ctgtttttgt gaactatagt gaaataaact gccacatact tgccagtgct cagtctctct
7081 gagtcatttg gtcaacatgc gtccgcaccc caataattat ttgcatacac agatcagtag
7141 gagaggcgcc aagacggaca tatcctcttc aaatttccctt aaaattattg aatttaacaa
7201 ctgtaagcta caaaagACCG TTATCGTTtc ctctaactt gggaaaaagg tgagtgaag
                                -> E2 bind
7261 ttttattgca ccttttgtga gtcaatttgt ctggcggcgc tgaacgaatt tggctgtcag
7321 cctttgcACC GGGAGTGGTg gaaaatagtt tct
                                -> E2 bind
```

LOCUS HPV41 7614 bp ds-DNA VRL 04-JAN-1993
 DEFINITION Human papillomavirus type 41 (HPV-41), complete genome.
 ACCESSION X56147
 KEYWORDS papillomavirus.
 SOURCE Human papillomavirus type 41 DNA.
 REFERENCE 1 (bases 1 to 7614)
 AUTHORS De Villiers,E.M., Hirsch-Behnam,A. and Hirt,L.
 TITLE Nucleotide sequence of human papillomavirus (HPV) type 41: an unusual HPV type without a typical E2 binding site consensus sequence
 JOURNAL Virus Res. 18, 179-190 (1990)
 COMMENT *source: tissue=facial wart; *source: is_proviral=N; From EMBL entry PAP41CG; dated 14-JUN-1991.

 HPV-41 was originally isolated from biopsies taken from a 15-year-old female patient with facial, peri-anal and foot warts (Grimmel et al. Int. J. Cancer 41, 5-9). The patient had a history of dermatitis atopica since early childhood, but no clinical or histological symptoms of epidermodysplasia verruciformis or any type of immunodeficiency at the time of treatment. This suggests that HPV-41 caused the lesions.

 Subsequent screenings of tissues taken from 106 biopsies from benign and malignant skin lesions as well as 71 malignant tumors from non-cutaneous tissues indicated the presence of HPV-41 DNA in two out of ten squamous-cell carcinomas and one out of three arsenic keratoses (the latter being regarded as a precursor lesion to the former). HPV-41 DNA was not detected in any non-cutaneous tissues.

 Sequence analysis of HPV-41 reveals it to be highly divergent from all other known types. Overall nucleotide similarity to other sequenced HPV types is less than 50%, and the highly conserved E2 protein binding motif (ACCN6GGT) is found only in a modified form in the upstream regulatory region of the genome. (These modified sequences have been demonstrated to bind to the E2 protein in the BPV-1 genome.)

 The author observes the presence of three short ORFs (X, Y, and Z) in the HPV41 genome. At the time of publication, they were no observed analogs in other sequenced papillomavirus genomes. Because of this, these ORFs are probably not functional. The author notes that these ORFs have the following characteristics: X ORF 922-1212 with first methionine at 979, Y ORF 2811-3050 with first methionine at 2832, and Z ORF 5631-5882 with first methionine at 5652.

 BASE COUNT 2101 a 1665 c 1908 g 1940 t
 ORIGIN
 1 aCAATCATaa tcatcgccct ttcgtgttat ttcttgtAAC GAATTCGTTa caaaacacac
 signal -> -> E2-bind (modified)
 61 acacagTATA TAAgatagag gaacggattg gtacaccaca gATGgcatca acaagcggtg
 signal -> E6 cds ->
 121 tgggatccgt cgggcctgca agctgttgcg agacgcagaa gccacatacc atacgggagt
 181 tgtgtttggc gcagcagata acttatccat gcatacagct ctgctgcat tattgctata
 241 agatccttag cgtattggat atttacgctt tgcaccagag ctgtctgtac ttatcctggg
 301 gagaaggggg gccaacgggt atttgttctc agtgacttag agtgcttgca aggctggagt
 361 tcaactgcac gcacgaagtg tcttgtgcag ccagccgtct gccgcacttt ataggacaga
 421 gcctcagcga ccttgagggt aggtgtgtga ggtgcctagc tcttctacaa tctgtggaaa
 481 aggattacat attgcgggaa gacttgtctg tgcaTAGaat tggcgggatc tggaggggaa
 E7 orf start ->
 541 cttgtgttcg ATGtatggta ggactgtatT AGctgtgaga ctaatatact gtttgcgtga
 E7 cds -> <- E6 end
 601 ttgtattgct gtaatcgtgc gtaaattgct ataccctgta ataatgagag ggaatagtgt
 661 tgacctgcaa gaaattgtgc ttgttcagca gggggaggta cctgagaatg ctgcagtgca
 721 ttcaggggag cattctgatg atgaggggtga gagcgaggag gaggagcggg aacaggtgca
 781 gcaagtcccc acaccagga gaacattata cctggtagag agtcagtgtc cattttgcca
 841 ggctatcata cgatttgtat gcgTAGcaag caacactggg atacggaatc tacaggcact

HPV41

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                                E1 orf start ->
901 cctgggtcaac agtcacctTG Acctcgcttg tcacgcctgt gtcgagcaga ATGgcggtcca
                                "X" ORF start ->                                E1 cds ->
961 ggggtctcaga caccgggcaat ggcaaTGAAa acaaagagaa tgaagGTaca Gtggcatctg
                                <- E7 end      5' sj /\
                                (putative E1,E4 fusion site)
1021 atcattctga ggcgcggtgt agctatatat tatttgaggc tgaatgtagc gatggcgggg
1081 acgatgagga aagtatggag gatagcttgg tggaagacct tgtggatgat gcttctgtgc
1141 atcagggaaa ttcttctgctg ctgtttcatg cccaaactgt cgaggaatac gagggagaga
1201 tccagagcctT AAaacgaaa gttatctctga gtccttgca tagggatgtg gcagaactaa
                                <- "X" ORF end
1261 gcccgcgctct ggcggggtgtt tccctggaag aaaaccgtgg gaaaaaggct cgcaaattctc
1321 tgttccacga tgacagtggc atagacagca ggcagtgga agtctcccag ctatctagta
1381 cgccatcagc tccagggcca gacatccggc tgcctaaacc ctcagatata gatctagagc
1441 cactgttcca aagccgccag cgctgtacgc atatgtatag caaatttaa gctgtgtacg
1501 gggtagctt tacagatata accaggccat tcaaaagcga caaaacaaca tcacagcatt
1561 ggggtgtggc cgcctactat ttagcttttg atagtगत atagtगत gaggttttgc
1621 tgcgacaaca atgccaattt ttatacattg acaacaatga tggcattata ctgttcttcc
1681 tggaatacaa cgtgcagaaa tctaggacta cagtgtacaa ttggttcaca gccaaatttcc
1741 attataatga aatagaatg ctagctaatc cgccaaggac acgaaacatg cctgctgctt
1801 tattcttcta tcatagattt atgggtacag ggggtataaa acatggcgca atgccagaaa
1861 taattgtaaa ccagtgcgtg gtgtctaatc agcagacaga cacctttgaa ttatcacgta
1921 tggtacagtg ggcactggac aacgatctgc aagatgaaca tatgttagct ttagagtatg
1981 ctttgcttgc tgaagtgat ggcaatgcgc gggctttttt aaagcagaat aatcagccaa
2041 tgatagttaa gaattgtagc ataatggta gacactacaa gacagcgctg gtcgcaaaaa
2101 tgtctatttc acagtatgtg aataagcggg gtctggacca tggggaagct gatgaaaaca
2161 gctggcgggg aattgtgcat tttctgaggt atcaaggcca ggaattcctg cccttcatgt
2221 gtaaaatgca caatttctca caccatagac caaagaaatc aacacttcta ttatgtggac
2281 cgtcggacac aggcaaatca ttttttgcca atggctctaa caaatttttg gatggacacg
2341 tgctgagctt tgtcagcaat gggtcacatt tttggttatc accattacgt ggggcacggg
2401 gctgtctaat agacgatgcg accctcacgt tttggaggta cgcggaccaa aacatgaggg
2461 cactgctaga tggatatgag atttccattg atgcaaaaca cagaaccca atgcaacta
2521 gagcaccacc attaataata accacaaatg aggacattat gcgattagat gaattcaaat
2581 atctgcaaac cagaacaatg tatgtgtact ttaacaagcc atttctctt aaaggaaatg
2641 ggcaccggtt atattacatt gatgggtata catggaactc ttttttagg aaattttggc
2701 gtcacctaaa tcTAAaagac cctgaggATG agtcagatgg agagactcct ggaacgatta
                                E2 orf start ->                                E2 cds ->
2761 gactatatac aagagcagat actgacacta taTGaгааag atagtgtTGA cctagaggat
                                <- E1 end
                                "Y" orf start ->
2821 catataaggc tATGgaatct gctaaggagg gaaaatgcaa tctggtatgt actcagacag
2881 gaaggacacg caagggtcgg cggcagagcg gtgccggcaa tgacggtatc ggaagccaat
2941 gccaaattcg caatagaat gcagataaag ctagaatcac taaaggccag tcctatgcg

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3001 gccgagggct ggtcattgca agaaaccacc aaggaacggt acttggcTGA accgtctcgg
                                     <- "Y" end
3061 acatttaaga aattaggcca gccagttacc ctaatgtttg acaatgatcc cgaaaaacctt
3121 acagaagttg tatttgaggaa atgggtttat tatattacac caacagatga atgggtataaa
3181 gctagagggtg gcattgatga cactgggtata tactacattg accacgagtc tgtaaanaatg
3241 tactatgtga gatttgacat ggaagcggag aactttagcg agacaggcac tgtcacctac
3301 cggctaggca gcgccctggt aaatgtacct gaacctgTAA ctgttaccga cagctcctcc
                                     E4 orf start ->
                                     NH2 terminus unknown
3361 acgAGggaga gaaccccaaa ggtactacga ccgaggggt cgagacgacg cagaaaacgag
                                     /\ 3' sj
(putative E1,E4 fusion product)
3421 gaaacggggg agccggtcgc cccagcccct aagcgaagac gaggagctta cggacgcaga
3481 tcctccccga aggcccaacg caggaccgcg gcgtcgctg tttctagagg aaacggagga
3541 tcgtctgact tcacttctgg agagtctgac gaaggacatc gagtcagaca tagagcactt
3601 cgaaagaaaa ctgcgggtgt tgctccagca gaaggacact atcTAGttgg cgccaaaggt
                                     <- E4 end
3661 ccagtgaata gcctgcgggtg cttaaggtac aaatggaaaa acaagtatag cggtgacaTA
3721 Atgtatctgg ggactacttt cacatggacg gagtctgacg ggacagaacg gtgtgggtcg
E5 orf ->
start
3781 gggcgctttt tttgtgcttt ctctaATGaa acaaaaagag aaaagttcct caaatctgtc
                                     E5 cds ->
3841 aagattccta aaaacattgg gctgtttcgc gcacacgcag aaaagctgTG Acctgtgtat
                                     <- E2 end
3901 catTAAacaA TGcttctag gcaaaggggt aaacgcgcta atcctgaaca actgtataag
                                     L2 cds ->
L2 orf start ->
3961 acatgcaaag caacgggggg cgattgtcca cccgatgtTA TTAAAcgcta tgagcaaact
                                     signal ->
4021 acacctgctg atagtatATT AAAgtatggg agtgtagggg ttttctttgg cggctctgggc
                                     signal -> <- E5 end
4081 attggcacag gacgtggtgg cgggtggcaca gtgcttgggg ctggggcagt tgggggacgc
4141 ccgtccatat ccagtgtgac aattggtccc cgggatattt tgccaattga atcagggggg
4201 ccttactctg cagagaaaat acctctgctt cccatggcac cccgtgtgcc aaggcctaca
4261 gatccctttc ggccgtcagt gctggaagag ccttttatta taaggcctcc tgaacgcccc
4321 aacatttttg atgagcagcg tttccctaca gacgctgcac cattedgaca tggcaacaca
4381 gaaatcacia ccattcctag ccaatatgat gttagtgggg gaggggttga cattcagata
4441 attgaactcc ctagtgtgaa tgaccccggg cctcgggtt ttaccgcac acaatacaac
4501 aatccaacgt ttgaggtgga ggtgtccact gacattagtg gagaaacctc atcaacggac
4561 aacattattg taggagctga aagcgggtgg acatccctag gtgacaatgc tgaactgata
4621 cctttgctag atatatcccg gggggacaca attgacacaa caatacttgc cctggegag
4681 gaggagactg cctttgtgac cagcactcct gaacgtgtgc ctatacagga gcgattaact
4741 attagccct atggcagaca gtatcagcaa gtgcgagta ccgaccctga atttttagac
4801 agcgtctcag tacttctctc tttagagaat ccagtgtttg atgcagacat tactctcagc
4861 tttgaggatg atctgcagca ggcactacgt agtgacacag acctgcggga cgtgcgtcgc
4921 ctcagtagac cttattacca gaggcgcact actggccttc gtgttagtgc cctggggcaa
4981 cgtcggggta ctatatccac gcgctctggt gttcaggtag gctccgctgc tcattttttc
5041 caggacatta gtccaatcgg ccaggctatt gagccaattg atgcaattga actagatgta
5101 ctgggtgagc aatccggtga gggactatt gtgagaggag accctacgcc ttctattgag
5161 caagacatag gactaacccg tttgggggac aacattgaaa atgaattgca ggaaatagat
5221 ttattaactg cggatggtga agaagaccag gagggcagag acctgcagtt ggtattttcc
5281 actggcaatg atgaggtggt tgatattaTG Actataccta tacgtgcagg cggggATGac
                                     L1 orf start ->                                     L1 cds ->
5341 aggccttcag tatttatttt tagcgtatgat ggcactcaca ttgtctatcc tactagcaca
5401 acagccacca cccactcgt gcctgcacag cccagcgatg tgccctacat tgttgttgac
5461 ttgtatagtg gaagtatgga ttatgatata catcctagcc tgttgcgcag gaaacgtaaa
5521 aaacgcaaac gtgtttattt ttcagatggc cgtgtggctt ccaggcccaa aTAGatttta
                                     <- L2 end
5581 cttaccccct caacctatac aacggacatt gaacacagag gaatacgTGA gacgcaccag
                                     "Z" orf start ->
5641 tactttcttc cATGctgcca ctgaccgttt gcttactgtt ggacatccat ttacaatat
5701 tactaatgcy gatggcaaag aggtggtccc taaagtttcc tctaatacgt tcagggcctt
5761 ccgtgtccgt ttcccaaatc ccaatacctt tgcattttgt gataagtccc tttttaacc

```

HPV41

```

5821 tgacaaggag cgtctggtct ggggtattcg tgggattgag gtttctaggg gacagcctT
5881 AGgtattggt gtaacagggg accctttttt taataagttt gatgatgctg aaaatcccta
    <- "Z" end
5941 caatgggata aacaaaaata acattactga ccaaggttca gactcaaggt tgagcattgc
6001 atttgaccct aagcaaacac agctgctgat agtaggtgct aaacctgcaa aggggtgagta
6061 ctgggacggt gctgcaacat gtgaaaaccc tccactgacc aaagcagatg acaaatgtcc
6121 tgctctagag cttaagtctt catacattga ggatgcagac atgagtgaca taggcctggg
6181 aaacttgaat ttttctacac tgcagagaaa caaatccgat gccccattag atattgtgga
6241 ttctatctgc aaatatcctg actacctgca aatgatagaa gaactatatg gagaccacat
6301 gtttttctat gtgcggcgctg aagctctgta tgctaggcat ataatgcaac acgcggggcaa
6361 gatggatgct gagcaatttc ccacttctct gtacatagac tcctctgtag aagggtgagaa
6421 attaaattcc ttgcagcgca ctgataggta tttcatgaca cccagcggct cctgggtagc
6481 tactgagcag cagctgttta acaggccctt ttggctgcag agatcccagg gccataacaa
6541 tggcatactg tggcacaacg aggcctttgt aacattggtt gacactacca ggggaactaa
6601 ctttaccatc agtgttcctg agggggatgc ttcttcatat aacaattcta agtttttga
6661 gtttttaagg cacaccgagg agtttcagct tgcctttatt ctacagctgt gtaaggtaga
6721 ccttaccctt gagaatttgg cttacataca cacaatggat ccatccatta ttgaagactg
6781 gcatttagct gtcacttcac ctoccaaattc tgtactggag gatcattata ggtacatact
6841 gtccattgca actaaatgtc cctctaagga tgcagatgat acctccactg acccatacaa
6901 agatcttaag ttttgggagg ttgatctacg ggatcgtatg acagagcaat tggaccagac
6961 tccccttggc aggaagtttt tgtttcaaac tggatcact cagtcacat caaataagcg
7021 ggtgtccacg cagtctactg cccttactac ctacagggcg cctactaagc gccgcgggaa
7081 ggcTAAAcg aatgtctggt atgtggtgct ggtgtcctcg acgggccatg tgtcatctta
    <- L1 end
7141 taatcacttg gtcagtccag ggtacaccac tccattatct atttacttcg catgtatttc
7201 tctgttatgt tcctgtatgg gttatgaatg tgttAATAAA atatgttggg aacgctgtgc
    signal ->
7261 acggggtttg tcacgttcat gtctcatgat TTGGCAcccc tgtattcccg cgcgcccccg
    NF-1 bind ->
7321 ggggatcgca gatataatcc ccaaACCCAA AGCGTTccaa caTTGGCAaa cgtctctggc
    -> E2-bind -> NF-1 bind
    (modified)
7381 cccgatacAA CTGAAACGGT ctgtctTGCC AAtagcccca tctggcgggg attcAACTGA
    -> E2-bind -> NF-1 bind -> E2-bind
    (modified)
7441 AACGGTgtgt acTGCCAAgt aacatttttg ttattggAAC GCCTCCGGTg ctggcggaag
    -> NF-1 bind -> E2-bind
    (modified)
7501 cgcaaggatt taggcgcgaa gacagtttta tTGCCAAaac cttttggttg cTGCCAAtag
    NF-1 bind -> NF-1 bind ->
7561 caggcgtggt ctcAACGAAT TCGTTgcggc aaTAGgtatg taccatgggt atga
    -> E2-bind E6 orf ->
    (modified) start

```

LOCUS HPV63 7348 bp ds-DNA VRL 04-OCT-1993
 DEFINITION Human papillomavirus type 63 (HPV-63), complete genome.
 ACCESSION X70828
 SOURCE Human papillomavirus type 63 DNA recovered from punctate keratotic lesions of the foot.
 REFERENCE 1 (bases 1 to 7348)
 AUTHORS Egawa,K., Delius,H., Matsukura,T., Kawashima,M. and De Villiers,E.M.
 TITLE Two novel types of human papillomavirus, HPV63 and HPV65 comparisons of their distinct clinical and histological features and their DNA sequences to other HPV types
 JOURNAL Virology 194, 789-799 (1993)
 COMMENT Submitted (27-JAN-1993) on tape to the EMBL Data Library by: H. Delius, Deutsches Krebsforschungszentrum, Abt ATV, Im Neuenheimer Feld 506, W -6900 Heidelberg, FRG.

The 7348 bp genome of HPV-63 was isolated from punctate keratotic lesions of the foot. Infection from HPV-63 produces intracytoplasmic inclusion bodies (ICB) in most cells of the epidermis. These inclusions have been previously shown to contain mainly E4 proteins for HPV-1. Up to date three morphologically different types of ICBs have been distinguished, and certain types of HPV are preferentially found in one type of ICB or another. HPV-63 is associated with a filamentous type of ICB (FI-ICB). The author notes that histologically, these FI-ICBs were seen as heavily stained keratohyalin-like substances with filamentous brush-like structures within vacuolated cells, seen either as single cells or clumped in nests.

BASE COUNT 2239 a 1359 c 1612 g 2138 t

ORIGIN

```

1 gtTAACAact atcaggcgat tctctagttc taacacgaac gtttacggtc gttgccagct
E6 orf start ->
61 ttttccttat aaaactctgg tgggaatttc tcttgggaca gATGgacctg acatctgtac
E6 cds ->
121 attcggttcg ggatctgagt tctgctctcc gtatoccatt tattgatttg gttgttcctt
181 gcaatTTTTg cttgaaatTT cttacaaaatg ctgaaaaatt gctgtttgat tattttgact
241 tgcacTcttat ctggcgagat aatttcgtgt ttgcttgttg tcagtgctgt gctaggcatg
301 ttagtctgct tgagtttatg ctttattatc aggagtcttt cgaggTatct gaagtagaag
361 aattacttaa tcaacctctt gtaaatattg gtttaaggTg tgttacatgc acaaaaaaac
421 tgactgtttc agaaaagTta gctgttgttt ctgctggaga aagagttcat aaagtaagga
481 acaaaTtcaa agcaaaTgTc agtttTgTca gactctacat tataTAGttt gtgcagactc
<- E6 end
541 tataTAAtta acaATGgtTg gagagcagcc aaatataggt gatttTgtga gtcaagaaga
E7 orf start -> -> E7 cds
601 accaagcgtc ctagatctaa attgttatga ggatatacct gctgaggagg aggagtctga
661 atatccatat gcaattgtgc ttccttTgtg tttgtgcat cagctgttaa ggctgacctg
721 cgtttctgac ctgtctactc ttacgcgtct ggaggagctg ctgttaggct cactGAGgat
E1 orf start ->
781 cgtgtgtccc ctgtgtgcca ttcgacacca acgacacTAA gATGACCGAC AGAGGTacaa
E1 cds -> -> E2 bind
<- E7 end
841 ataatgatga ttggtatatt gtggatgagg cagaatgtcg ggatgatgat gagagcgaat
901 tggaggattt ggaggacacc tataattcat tgtttaatag atctgaaagt gacatatcag
961 atctattaga cgatacgag caaagtcag gaaattccct ggaactgttc cacttacagg
1021 agcacttgca gaacgagcag gacctaaaata ccctaaaaacg aaagtactta aacagtcctc
1081 cgcaggcaag tgccacagag actgcctgca atagcctcag tcccagattg gaatctataa
1141 caatttcgca gagggaaaaa aaggcaagaa agcaactatt tacacaaaat gacagtggca
1201 tagagttatc gctatgccag gatgaagTtg acaatattaa cgaagcgtt caggagcagg
1261 tagacatcgt acagtctctg ggaggtgggg tgcgtgactg tataggagtg gacattttga
1321 aatgcagtaa tacaagatct gctctacttg ccaaatTtaa agacacagta ggtgtcagtt
1381 ttactgacct caccagagca tacaAAAaca acaagacatg ctgtagtTac tgggtcatag
1441 cagtgtgggg agtaacatct acgtctgtgg acgttTgtgaa aactgtattc caagttcagT
1501 gtaattatat gcatgtagaa cattgtTtaa ctgaaaaaaa taagtttcta attgtattag
1561 ctggctTtaa agctcaaaaa agtagagaaa cagtgtTtaa tctcgtactc agcagttTga

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HPV63

```
1621 atgtgcaaag taattacata atggctgaac caccaaaaaa tagaagtatg gcggcagcgt
1681 tatattggta taggagatct atgtctccag ctgtatatac ctggggagaa atgccagatt
1741 ggatggcgca gcagacattg ttgaatcatc aattagcatc agaaaagcat tttgaattgt
1801 cacaaatggt acaatgggct tatgataatg gctatacaga tgaaagtgat attgcatact
1861 attatgctat tttagcagaa gaagatgaaa atgcaaaagc attcttggct tctaatgcac
1921 aagcaaaaata tgtaaggac tgtgctagaa tggttagtca ttacaaaagg gcagaaatga
1981 gtagtatgtc tatgtcagca tggatttata aaagactgga ggaagttgaa aatgggtggtg
2041 actggaaaca tattgtaaag ttcttgaggt ttcaagaagt agaatttata agtttcatga
2101 tagcatttaa ggaattgtta agtggtaaac caaagaaaaa ttgtcttgta atatatggtc
2161 caccaaatac tggtaaatct atgttttgta tgagtttgtt gagagtatta aaaggaaaaag
2221 taatatctta tgtaaatagc aaaagtcaat tttggttga accactagct agcactaaaa
2281 tagcattatt agatgatgca acaaaaaccag catgggatta tattgattta tttttgagaa
2341 atgctttaga tgggaatcct atttgtgtag atctgaaaca taaggcacca caacaaataa
2401 aatgtcctcc acttatgata acttctaata taaatgtaa ggctgatgta tgttggatgt
2461 atttacatag taggataaca tgttttgaat ttaaacaacc ttttccattt gatgaaaatg
2521 gtcaaccggc attttcctta acagacatca attggaaatc tttttttgaa aggttttggg
2581 gccagttaga ctTAAGtgac caagaagacg aggagagtGA TGgaaagcct caacaaccgc
      E2 orf start ->                E2 cds ->
2641 ttagactggc tacaagagca gcttctaact ctataTGAg aagacagtaa agatattgaa
      <- E1 end
2701 gatcagataa tgcagtggaa tctacttaga caggaacaag tgttattcca ctatgcccca
2761 aaaaagggaa taatgcgact tggcctgcaa gttgtgcctt cccttgcagc ttcccaggat
2821 aaagcaaaaa cagctataga aatgactctt tatcttagtg gcctcagaga ctcacaatat
2881 ggttctgaac agtggctctt acaagatact agcagagaaa tcttttttagc accaccagat
2941 catacattca aaaagggagg gcaacaatt gaggtaatct atgatgagga tcccaataat
3001 agcaccagac atactgtatg gcgcatata tattatcaaa acggtgataa cagatggaga
3061 aaagcagcta gtgatgtaga tgttcatggt gtgttttatt TAGaatATGa tgggtgcaaa
      E4 cds ->
      E4 orf start ->
3121 aactactatg ttgactttca agaagaggcc aatcgatata gcaaaacagg tcgatatact
3181 gttcaatag agggtaaaag gttcacaat gttatgtctc ctgtcaatag ctccccacta
3241 cggacttctg ggtctcctac agacaccaac ccagccacc aaggacaatc cacccaaact
3301 gccagaaaag cagagacgaa ggggtcgaga caccaccgca aatcgccggc tgttcgcaag
3361 cgacggccct acggacgaa ggggtccaga agtcccagag ataccaccct cagacgagga
3421 gaaggagaat cggccagagc ctctgcgggt agtggagaac ggggtggcatt catttctccg
3481 ggagacgttg gaacatcaac taggtcgctt ccaaagggag gtcaatcaag acttcgaaga
3541 cttatacagg aggctcggga tccaccaTA Atttgtctga agggggggccc taatcaactt
      <- E4 end
3601 aagtgcctaa ggtatagat taaagcttca aatcatctg actttgaaag tatcagtagt
3661 acatggcatt ggttacataa taaatgcaca gatagagtag gtcatgcacg tatgctgggtg
3721 cgttttatat caacagaaca acgtgaccga tttttagata aggtgggtgg gcctaaatct
3781 gtttctgtta ttttaggggc attTGAcgggt tccTAAGgggt ggggtgtggg gtatattttg
      L2 orf start ->                <- E2 end
```



```

3841 taatcATGtt aagagtacgt aaacgacgag ctgctccaca agatatttat cctgcttgta
L2 cds ->
3901 aggttgcaaa caattgcccc cctgatatac aaAATAAAat tgaacaaaca acagttgctg
signal ->
3961 acaagatttt acaatatggg agtttgggaa tattcctggg aggtttgggt attggtactg
4021 gcaaggggtg ggggtggccg tatggttata cacctctagg ggacagtggg gcggtgcgag
4081 ttgggtggcag aagtacacct gtaagaccaa cagtacctgt ggagactgta ggaccaaggg
4141 atataattacc tatagattca ttggatcctt tagggcctc agtcattgaa ctagaagata
4201 ttccagccac aacagtggaa gtagtggctg aagtgcaccc catatctgat actccacaaa
4261 taccggcacc tactactgat gaatctagtt cagctgttct tcatattcca caagaaagtc
4321 ctgctgcacg tacaatcaca cgttcccaat acaataatcc tttattcagg atcacagcta
4381 gtgcagacat agcatcaggt gaagcttcag catctgataa ttttttata gatgtagata
4441 cgccgggtca aatagtagga caagaaatac cactagttaa ttttgatatg ggacctatat
4501 ctactgaagg tgagcttgaa actgagttca caactagtac accaagaacc acacaagtac
4561 aggaaaggcc tacacgtttc tataatagac gctattatga acaagtgcc gttactgcac
4621 ctgaatttat cacaaggcct gcttccttag ttacttttga gaatcctgca tttgaaagga
4681 gtgtttcttt gatttttgaa caagatttag aagatatttt aaatgctcct gatcaggatt
4741 tttagagacat tgtttattta agcagaccaa catacagtcg tgcccctgat ggccgcacgc
4801 gcctaagccg cctgggacgc agagccacta taagtACCAG AAGTGGTgtt actataggtg
-> E2 bind
4861 ctcaatcaca cttttatag gatattagct ctatctctc aaatgatggc attgagttac
4921 aaacactggg tgaagcttct ggcgagactg tggtgcaaag ttctcttct gcatcggatc
4981 ctattgaagc agaacattca ttcattgaac cagcaccatc tatagatagt tatgatattg
5041 tttcacttca gtctgagact tattcagatg aacatttgtt agatatgtat gaacctgtag
5101 gttcttctct gcaattacaa atatcagacg tcagaggtcg gccaaactgtt attgatattc
5161 ctttttagacc ccgcaggcct ccattaggtc ctataaatgc tgggtgtgat atctatagtc
5221 caactgctag tgttggatca cctactataa atcctactga tcttgacatt ccattaatta
5281 ttatacattt agataattca acaggggatt atgatttaca tccaagtttg cgtaaacgtc
5341 gcaaatTAGt tcatattTGA tattttacag ATGgctgttt ggcttctgc ccagaataag
L1 orf start -> <- L2 end -> L1 cds
5401 ttttaccctc ctaccaacc gatcaccaag attctaagca gcatgatta tgtgtctcgc
5461 accaacatct tctatcacgc taccagtgat cgactgctca ttgtgggaca cccgctctat
5521 gaggttacc gtgcaaatga taacactatg actgtgcta aagtttctcc aaatcagtat
5581 agagtcttcc gtggttagatt tccagatcct aaccgatttg cttttggaga taaggatatt
5641 tttgaccag aaactgagag actagtttgg ggtcttagag gcatagaaat cggtaggggt
5701 caaccattag gtgtgggtat ttcaggcaat ccattattaa ataggtttga tgatgctgaa
5761 aatcctagca gatataataa tacacatgca actggtgata ataggcaaaa tgttgctttt
5821 gatgcaaaac aaaccctaat gtttctaatt ggctgtacac cagccactgg ggaacactgg
5881 tcaatagctc gacgctgtgc aggaacacag tttcagcttg gagattgtcc tccatagaa
5941 ttagttaaca cagttattga ggatggtgat atgtttgaca taggtctagg tgctatggac
6001 tttggttctt tgcaagcaaa caaagcagat gctccttgg atattgcagg cactgtctgc
6061 aaatatccag attatattaa aatgggacag gaagtacatg gtaattctct gtttttcttt
6121 gctgcgagag aacaaatgta ttttaaggcat gtatttacac atgctgggat tgttagtgaa
6181 aaagaaagag tccctaccag tgcatatatt gctgctaaag ccgagcaacc ccaaaact
6241 attgctacag ataattattt tgtagctccc agtggatctt tagtgcctc tgatgtgcaa
6301 atttttaata ggccctattg gttacaacgt tctcaaggac agaacaatgg tatctgttgg
6361 agaaatgagt tatttgtaac tgtagctgat aataccagag gaaccacgat gaatataaat
6421 gttcttaaca aagcaacccc tgagacttat gatagcgag attataatga gtatactgc
6481 catgtggagg aatatgagtt atcctttata gttcagcttt gtaaggtaaa actaacacct
6541 gaaaatttag catttttgca taatatggat ccaacaatta tcgattcctg gcagtttaaca
6601 gtttctcaac ctctgcaaa tgctatagag gacaagtata gattttattga atcattagca
6661 acaaaatgct ctgataacgt gccccaccc actcctactg atccttaca agatttacgt
6721 ttttgggatg tagacctcag tgagcgaatg tcggagcagc ttgatcaatt tcttttaggc
6781 cgcaaatttt tgtatcaaa tggtcttgca cagcgttctg ttccaaaaac tgtgaatttc
6841 agaaaacgta gatcctccaa tactactgtg gccaaaacgga ggcgacgggc cTGAatatac
<- L1 end
6901 atgtgaatgt tgaatTATAT AAtgtgaatt gtgaattctt gactttggca cttgcacttt
signal ->
6961 attcttggca tactgatact tgaacttgt tcaatgcttg aaggttacac acctgtacag
7021 tattgttaat aaacgtttat gctgctgtca tttacctgtc ttcgagtcac tattgcctag
7081 tcatatagcc tcatgacttg gcatgcaatt ggtatgtggc agatacttca aacaggatac
7141 tggtatcctt tttggcgcgc gcgcgaattt tgaagttacc actgttcca cttgttctga
7201 gacgtctgga tctgatcccg ACCGTGTCG TTactgcaaa agacgaaagt ggtaggcgcg
-> E2 bind

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HPV63

```
7261 aACCGTTTGT GGTttccctg gggctagcag aaactcttta ggttgcgACC GTTTTCGGTc  
    -> E2 bind                               -> E2 bind  
7321 gggccaataa tctctttcga tcgttgtt
```

LOCUS HPV65 7308 bp ds-DNA VRL 04-OCT-1993
 DEFINITION Human papillomavirus type 65 (HPV-65), complete genome.
 ACCESSION X70829
 SOURCE Human papillomavirus type 65 DNA recovered from a black verrucous lesion.
 REFERENCE 1 (bases 1 to 7308)
 AUTHORS Egawa,K., Delius,H., Matsukura,T., Kawashima,M. and De Villiers,E.M.
 TITLE Two novel types of human papillomavirus, HPV63 and HPV65 comparisons of their distinct clinical and histological features and their DNA sequences to other HPV types
 JOURNAL Virology 194, 789-799 (1993)
 COMMENT Submitted (27-JAN-1993) on tape to the EMBL Data Library by: H. Delius, Deutsches Krebsforschungszentrum, Abt ATV, Im Neuenheimer Feld 506, W -6900 Heidelberg, FRG

The 7308 bp genome of HPV-65 was isolated from a black verrucous lesion. Lesions of this type are mainly distributed on the palmoplantar or lateral surfaces of the hands and feet. Infection from HPV-65 produces intracytoplasmic inclusion bodies (ICB) in most cells of the epidermis. These inclusions have been previously shown to contain mainly E4 proteins for HPV-1. Up to date three morphologically different types of ICBs have been distinguished, and certain types of HPV are preferentially found in one type of ICB or another. HPV-65, like HPV-4, is associated with a homogenous type of ICB (Hg-ICB). Egawa et al. note that histologically, these Hg-ICBs were usually present as a single inclusion in each cell, surrounding an extremely vacuolated nucleus. HPV-4 and HPV-65 have both been detected in common warts and keratotic flat lesions on the hands and feet, as well as in two biopsies from cutaneous horns. All of the lesions in this study were pigmented.

BASE COUNT 2280 a 1306 c 1575 g 2147 t
 ORIGIN

```

1 attattaatg atagttggca acaaccatca tctaaagtaa tataTAACcg gaagagatac
                                     E6 orf start ->
61 atataaaaag acacattgta ttctgataa atcccatcca gATGgcagat ggcagacctg
                                     E6 cds ->
121 cagctttgga cgacttctgc agacgatttg atatttcttt ttttgatttg catcttactt
181 gtattttttg ttctcactact gttgatctgc aagatcttgc ttcgttttat cttagaagc
241 ttagttagt tttcaggggg ggttgctact atgcatgctg ttctgaatgc ttacgtttaa
301 gtgcaagatt tgagcaagag aattattttc agtgttctat aaaagctggt aatttggag
361 aagtagctca aaggaaaatt aaggagattt gcatcagatg tatatgttgt ttgagactac
421 tagacattgt tgagaaaatta gacttgctgt actctgacca ggcctgctac ttaataaggg
481 gtttgtggag gggctattgc agaaattgca tTAGGaaaca ATGAgaggag cagcaccag
                                     E7 orf start -> -> E7 cds
                                     <- E6 end
541 ggttgcatg cttaatgtag aattaaatga cttggtgttg cctataaac tgctgagtga
601 ggaggtcttg caaccttcag atgatgagtc tgaggctcca gaggaggagc tttttccttt
661 tagaatagac acctgttgct atagatgtga agttaatgta aggattactc tgtttgctgt
721 ggaatttggc cttcgagcgt tggaacaact catagtggac ggaagctga cgttttgctg
781 tactacttgt gcaagaacct TAAGaaATGg cagaTAAagg tacagaaaat ttgacttag
                                     E1 orf start -> -> E1 cds <- E7 end
841 aagggagtag ttggtatatt gtgcatgaag cagaatgcac agacagtata gacactttgg
901 aggatttgtg cgacgaaagc gattcaaatg tttccaattt aattgatgac gatgtggttg
961 atcaggggaa ttccctggca ctgtacaatg caaaaataac tgatgattgt gataatgcaa
1021 tagcacacct aaaacgaaag tataacaaaa gtccagagca ggcagttgct gaattgagcc
1081 ctcagttgca ggctgtgaaa attactcctg aaagaaaacag caaaaggaga ttatttcagg
1141 aggacagtgg gatatttgaa gatgaagctg aaaattctct tacacaggta gaatctaaca
1201 gccagactgg tggcaacagc caagatggcg gaggagatat taacttactg ttgttataaa
1261 ctagtaatag gctgtgtaca atgcttgcaa aatttaaaga ttggatggg gtttcatata
1321 atgaaataac tagagtttat aaaagtgata agtctttagt tgataattgg gttatagtta
1381 tctttcgagc tgctgttgaa gttttggaaa gctctaaaat tgttttgcaa cagcactgta
1441 catatattca agttaaataa tttgattttt cagctttata ttattacaa ttcaaaagtg
1501 cgaaaagcag agaaacggta caaaagttaa tgtgttctat gttgaatatt caagagtttc

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HPV65

```

1561 aaatattgac tgaccctcct aagttacgaa gtgtgcctac agctttatac ttctataagc
1621 aagccatggt aacagagagc tttgttttcg gacagacacc agattggatt gcaagcaaaa
1681 ccttagtaag ccatcaagca gcaactactg cagaaacatt cgaattatca aaaatggttc
1741 agtgggctta tgataataat ctgctggaag aatgtgatat tgcgtatcat tatgctatgt
1801 atgctgatga agatgctaag gctgcagcct atttaaagag taataatcaa gtaaaacatg
1861 ttagagattg cagtacaatg gttagaatgt ataaaagata tgaatgaggg gatatgtcaa
1921 tgtcagaatg gatttataaa tgttgtgatg aatgcACCGA AGAAGGTgat tggaaaccta
                                -> E2 bind
1981 tttctcaatt tttaaaatat caaggtgtca atattctttc ttttcttata gttctaaagt
2041 cttttttaa aggtattcct aagaaaaatt gcatagtcat tcatggccca ccagatactg
2101 gaaaatctct gttttgttat tctctagtga aatttcttaa aggtaaagta gtatcctatg
2161 tgaatcgaag tagccacttc tggttgcagc ctttaagtga ttgtaaggta ggatttatgg
2221 atgatgcaac ctatgtatgc tggacataca tagatcagaa tttgaggaat gcattagatg
2281 gaaatccaat gtgtatagat gcaaaacata gagctccaca gcaattaaaa ttACCTCCTA
                                -> E2 bind
2341 TGTTaataac atcaaatatt gatgtgaaac aggaacaatc attaatgtag ttgcatagta
2401 ggggccagtg ttttagtttt cctaataaaa tgcctttttt agatgatggg tctcccatgt
2461 atacatttac tgacgcaacg tggaaatcct ttttccaaaa gcttggcaga caattagagc
2521 TAAcagatcc tgaagaggaa agcaATGgag tccctagtcg cgcgtttcga tgcacttcaa
E2 orf start ->                                E2 cds ->
2581 gaagcaatc tgactcatat TGAgtctcag gacgatactt tggaaatcca aattcgatat
                                <- E1 end
2641 tgggaaaata tcagaaaaga aatgcaata atgcattttg cacgaaaaca aggcctaaca
2701 aaattaggtc tgcaaccact tcccacatta gcagtaactg aatataatgc aaaacaagca
2761 atacaatac atttgacttt cgagtcatta ttaaagtctc cttatggatc agagcgttgg
2821 actctgccag aagttagtgc agaactgatt aatactgctc cacagaactg tTAAgaaa
                                E4 orf start ->
2881 ggaggttATG atgtgtctgt atggtttgat aatgatagat ataatgctat ggtgtataca
                                E4 cds ->
2941 aattgggatt atctatatta tcaagatgtc aatgaaatat ggcataaagt taaaggtgaa
3001 gtggattatg atggcttata ctttacagac catacaggag aacgtgcata ctttactctg
3061 tttagcacag atgctcacag atttagcaga actggactat ggactgtgca ttttaaaaca
3121 caagttatth cctcctctgt tgtcagctca acaaacaccc cctccttcga ctttgaggaa
3181 caacaactac cggggcctc aacaccacc tacaccgagc ttaccaggc gagcccttgt
3241 ggtaggggga aatcgaggga atctcaaccg acctccaaa cgtccccga aacctcgggg
3301 ctacgagtac gacgaggacg acgacaaaga aaatcagggc cggggccagg agagacccc
3361 agcaaaagaa gaagaggagg agggagagga ggaggagaga ccagattgga gtctgcgcca
3421 tctcctgggg aagtgggaat cagacataga acagttgaaa gacaaggctc gtcgagactt
3481 ggacaactac aagctgaagc tagggatcca cccaTGATat tgttaaaggg cacagcaaat
                                <- E4 end
3541 tcggtgaaat gttggagata taggaaacaa aattccagta actgtgggtt tctctttatg
3601 agcactgttt ggaactgggt tgggtgatgtg tcagaaaatc acagtgcgat gtaattgtct
3661 tttaaaagtc ctggtcagag agactcattt gttaaacaca atctattccc aaaactctgt

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3721 acatatacct atgggtcttT AAatagctta TAAaATGcaa gcctcacgca gaacaaaaag
      L2 orf start ->      L2 cds ->
                                <- E2 end
3781 agactcaata ccaaacctct atgcaaaatg tcaattatct ggcaactgtc ttccccgatgt
3841 aaaaAATAAAA gtagaagctg atacccttgc tgatcgtttg ctcagatggt tgggaagtgt
      signal ->
3901 catatactta ggaggttttg gaattggggac ggggaagaggc agtggagggt cttctgggta
3961 taatccactg ggagccccta gcagagtaac acctagtgga actgttataa gaccaacagt
4021 ccctgtagaa ggtctggggc ctagtgaat catccctgta gatgttgtaa ACCCTGGCAG
                                -> E2 bind
4081 TTcttctgtg gttcctttag aggatttaac agttccagaa gtcaccatag atagcgggga
4141 ggtgggagga ggtggctctc atccttctga gatagatggt gtcacttctc cagacctat
4201 ttcagatgtc actggtacaa gtagccaccc tacaataata tctggcgaag ataacgcat
4261 tgcagtctca gatgtttctc caacagagcc cctaccaaaa cgcatagctt taggaactag
4321 gggcgcaaca tcaacaccac atataagtgt aatatcaggc acgACCGAAT TCGGTcagtc
                                -> E2 bind
4381 ttcagaccta aatgtgtttg ttaatgccac attttctgga gattctattg gatatacaga
4441 agaaattcct ttagaagagt tgaatactat ccagcaattt gaaatagaaa cccctccaaa
4501 aactagcaca ccacgtgaga ctattggacg tgctttggaa agagcgcgag atctttataa
4561 cagaagagtg cagcagatag ccactagaaa tccagcaatg ctaggacagc cttccccgcg
4621 aatagttttt ggatttgaaa atcccgcctt tgacgctgac atcactcaag tatttgagcg
4681 ggacttgtaa caggttgtag cagctccaga tgctgatttt gctgatatag tcagaatagg
4741 gcgtccaaga ttttctcaga cagacactgg gcaaataga attagcagac ttgggcgccc
4801 aggaactatt aaaactagaa gtggtttgca aattggtcag gctgtgcatt tttattatga
4861 cttgagtaca atagacactg ctgatgcaat tgagctatct acacttggtc agcattcagg
4921 agaacaaagc attgtagatg caatgataga gagcagtttc gtagatccct ttgaaactcc
4981 tgatcctaca tacacagaag agcaacagct tctagatccc cttacagagg attttagcaa
5041 ctctcacttg gttttaacta gcagtagacg tggatcttca ttcagtatcc cactatttcc
5101 tcttggttta ggcttgagaa tttatgttga tgatgtaggc tccgatttat ttgtatctta
5161 cccagaaaca agagttatac ctgcccggagg cttacctact gagcctttta ccccgtaga
5221 gcctccattt ttttcagagt tttacagttc tgactttgta tatcgtccTA Gttgtatcg
                                L1 orf start ->
5281 caagaaacga aaacgatcag atatatTTA Attttttgca ggaacATGgc gagttgggta
                                <- L2 end -> L1 cds
5341 tctgcaaagg gtaaagtgta cctacctccc gctcaacctg tggcaagagt tttggaaact
5401 gacgaatata tcaactggaac atctctttat ttccacgctg gaacagaaa gcttttaact
5461 gtaggccatc cttattttcc agtgaagat gtgcaggatc aacacaagg tttagtctc
5521 aaagtttctg gaagtcagta tagagttttc agattctatt taccagatcc taataggttt
5581 gcattgatg ataagtgttt ctatgattct gaccatgaac gtcttggttg gagactcaga
5641 ggaatagaaa taggtagggg tggctcccctt gggattggaa caacaggcca tcctttatat
5701 aataagtttg gagattcaga aaatcctaag ggatacagga aacaatcaga tgacaacaga
5761 caggatgttt ctttagatcc aaaacagaca caaatgttta tcatagggtg cactcctgct
5821 attggggaac attgggataa ggctgagcct tgtgcaagtc ctgttctca accaggagat
5881 tgtcctccta tagagcttgt gaacacatat attgaggatg gggatatgty tgatatggt
5941 tttggagctt ttaattttaa agctttgcag caggataaat ctagcgtccc attggatgta
6001 gttgctacaa tgtgcaaat gccagatttt ctaaaaatga gtaaagatgt ctatggagat
6061 agcttattct tttatgtag gcgagaacag ttatatgcca ggcaactttt tggtagagca
6121 ggagcaatgg gagatgcact accagaACCT TTCGAGGTa aaactgatta ctggattcct
                                -> E2 bind
6181 gctcaagaag gtcaggatca gaatacttta ggtccacata tatatatagg cactcctagt
6241 ggatctttag tttcaagtga gtcccagtta tttaatcgAC CGTATTGGTT aaacagagct
                                -> E2 bind
6301 cagggtagcaa ataatggaat atgctgggat aatcaattat ttgtgacact tgttgacaat
6361 actcataata ctaattttac tatttctgta aaaacagaaag cagctgatga atcctataaa
6421 tataaagcag gtgattttaa acagtattta agacacattg aggagtttga aatggaattt
6481 atttttcagc tttgcacagt tcctcttaca gccgatgtca tggctcactt aaatgtaatg
6541 aatcctaata tcttagacaa ttggcaatta aattttgtgc cccctcctcc ttcaggaatt
6601 gaggatcaat atagatttat acaatccaga gctacgagat gtcctactca gtctcccagt
6661 actgaaaagg aagaccctta taaagattta tcattttggg gtgtagattt aacagaaga
6721 ttttccagtg aacttagcca attttcttca gggagacgct ttttatatca aagtggttta
6781 attaatggaa cacttaaaag aaaacgtact attaatctc aagctccaac tagtattaaa
6841 cgttctgcca aacgaaaacg gtctatataa cagTAGatta atcctataa gaatgcattt
                                <- L1 end
6901 gttctgcaaa tagaaactta aactgagcca attatTATAT AAtctttgga aacgtgtatg

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HPV65

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                                signal ->  <-
6961 tatttgcttt ttctgtgaga aacttgagta aAATAAActg ccacattctt gccactgtcc
                                signal ->
7021 atattgtctg actcattggt caacatgtgt gtccgcaccc cagtaattac ctggatcgcc
7081 tccacatccg tcggatgcgc gccaaaagag gaacgtcccc ttctatTTTT ctaaaaaatt
7141 accgTTTctg cagctgcaaa attgtgtaag ACCGTATCG TTccctgtaa ccttggcaca
                                -> E2 bind
7201 aaagtgagt gaaagTTTTa tagtaacgtt tatgagtcaa tttgtctggc ggcgctgaac
7261 gaattggct gtcagccttt gcACCGGAG TGGTagaaaa tagtttct
                                -> E2 bind
```