Corrigendum

A new member of the polyomavirus family: the hamster papovavirus. Complete nucleotide sequence and transformation properties

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On page 1280, in Figure 2 a line of the sequence corresponding to nucleotides 3361 to 3420 had been omitted. The corrected sequence is:

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   1 CCCCTTGCCTCCTTAGCTCTCAAGTAGAAAAGGAAGAGAGGCTTTTGGGGCTTTTTGGCT
  61 TTAAGCCTCATTTTATGAGCAGGAGGAGCTTGTTGCAACTTGAGAGGCGTTTTGAGGCTT
 121 CCAGGCAGAGAATACTCACAGACCCCACACAGTCTAGACGCTCAGAAGCATCTCTAGCTG
 181 CAACAAGCAAGATGGATAGAATTCTTACTAAAGAAGAAAAGCAAGCCTTAATAAGTTTAC
 241 TAGATTTGGAGCCACAATATTGGGGAGACTATGGACGAATGCAGAAATGCTACAAGAAAA
 301 AGTGTCTTCAACTGCATCCTGATAAAGGTGGCAATGAAGAGCTCATGCAACAGCTTAATA
 361 CCCTGTGGACCAAACTAAAAGATGGTCTTTACAGAGTTAGGCTGTTACTTGGGCCTAGTC
 421 AGGTAAGAAGACTTGGAAAAGATCAGTGGAATTTATCTTTACAGCAAACATTTTCTGGTA
 481 CCTACTTTAGGAGGCTCTGCAGACTCCCCATTACCTGCCTAAGAAACAAGGGAATTAGTA
 541 CCTGCAATTGCATACTTTGTTTGCTCAGAAAACAGCATTTTCTGCTAAAGAAGTCCTGGA
 601 GAGTACCTTGCCTGGTGTTAGGAGAATGCTACTGCATAGACTGCTTTGCCTTATGGTTTG
 661 GCCTGCCAGTTACCAATATGCTGGTTCCATTATATGCACAATTTCTTGCTCCAATACCTG
 721 TGGATTGGCTTGATCTGAATGTTCATGAGGTCTACAAATCCGGCCTCAGGTATGTATGAAT
 781 ATGGGGGGCTTATAGTTGTAACTGTACAAGTTTAAAATGTGCTTTTTTCAGGACCCTAAT
 841 GCTTCCACCTCCACCAGCAGACCCGGAGAGTTCTACAATCCTGACACAGGAGGATACTGG
 901 TCCTACTCTTATGGGTCAGCAGGATACTCTGACCAGCAGAAGAAATACTGGGAAGAGTTT
 961 TTCTCTAAGTGGGATGTTAATGAGGACCTCACCTGCCAAGAAGAGTTATCATCATCAGAA
1021 GATGAATTCACCCCCTGGCATCCCAATCCCCCCCCCTCCCCTGTTTCTATTTCCAGTGAC
1081 AGCTCCAGTTCCTCCTGTGACGAGGAATACCCAAGAAACTCAAGCAGAAAGAGAAAACGA
1141 GTACATGCCAATGGCTCCCCAAATACACCTATACAGCCAAATAAGAGAGCCCACACACCA
1201 GGAGGAGGAAGAACCACAATACGAGGAGATACCGATATACCTAGAACTCCTGCCAGAGAA
1261 TCCCAATCAACATTTGGCTCTTACTTCAACAGCACGGAGGAGCTTGAGGAGGAAATATCA
1321 CAAACACAACAGTCACATCATAACACAACGCCAAAGAAACCGCCTCCGACGGTTAGTCCT
1381 GATGATTTTCCTACTATCCTTAGGGGGTTTCTTTCTCACGCTATTTTTTCTAATAAAACG
1441 CAAAATGCATTTATAATCTACAGTACTAAGGAAAAATGTGAAGTACTTTATGAACAAATA
1501 GACAAATATAATCCAGACTATAAAGGTATCTTCATTATGAAACAAACAGAAGCATTTGTA
1561 ATGTTTATGACTCCTGGAAAACATAGAGTAGCTGCAGTTAAAAGTTACTGTTGTAAATTT
1621 TGTACCGTTAGCTTCCTGCTATGCAAAGCTGTTACAAAACCGTTAGAGTTGTATAACTGT
1681 GTGGCTAAATGTGATGACTTTCAAATTTTAAAAGAAAATAAGCCTGGTCTATATCATTTT
1741 GAATTCTGTGATGAAAAAAAAGAGGTGAAGCAAATAGACTGGAATTTCCTAACATCTTTT
1801 GCAGTTGAAAATGAGTTAGATGATCCTCTTGTAATTATGGGACATTATCTAGAATTTAGT
1861 CAGTGTGAAAGCTCTTGCAAAAAGTGTGCAGAAGCTTTACCAAGGATGAAAGTCCACTGG
1921 GCTAACCACAGTCAGCACTTAGAGAATGCTGAGCTTTTCTTACACTGCAAACAACAGAAA
1981 AGTATCTGTCAGCAAGCAGCAGATAATGTTCTGGCAAGGAGAAGATTAAAGGTCCTTGAA
2041 TCAACAAGACAAGAATTGTTGGCAGAGAGACTGAACAAACTGTTAGACCAATTAAAAGAT
2101 TTATCTCCTGTAGATAAGCATTTATATCTTGCTGGAGTAGCCTGGTACCAATGTATGTTT
2161 CCTGATTTTGAGATGATGTTATTAGATATTTTAAAATTGTTTACTGAAAATGTTCCAAAA
2221 AAAAGAAATGTACTTTTTAGAGGTCCTGTAAATTCAGGGAAAACTAGCCTTGCTGCAGCT
2281 ATCATGAATCTTGTAGGAGGAGTTGCCCTCAATGTTAATTGTCCTGCAGATAAGCTCAAC
2341 TTTGAACTTGGTGTTGCTATAGATAAATTTGCAGTAGTCTTTGAAGATGTCAAAGGACAA
2401 ACCGGAGATAAGAGACACCTACAGTCTGGACTTGGAATTAATAACCTTGATAACCTGAGA
2461 GATTACCTTGATGGAAGTGTAAAGGTTAATTTAGAAAAGAAGCATGTAAATAAGAGGTCC
2521 CAGATATTTCCTCCTTGTATTGTTACTGCTAATGAATATTTTTTCCTCAAACACTCTAT
2581 GCCAGATTCCATAAAGTTTATAACTTTGAAGTGAAGGATTTTCTTGCCAAGAGCCTTGAG
2641 GAAAACAGTTACATGGGGAGACATAGAGTCTGTCAAAGTCCACTTACAATGCTGATAGCA
2701 TTGCTTTGGAATGTACCCACTGAAAATTTTGATAAGTCTCTCAAAGAGAAGGTGGAAACA
2761 GAAAAGAAGGTTTTGTCTGATATGTGTAACTTTACTACATTTGCAGAAATGTGTCTCAAT
2821 ATTCAGAGGGGTGCTGATCCCCTTGAGGCATTGTAATTGAGGAGGAAACAATAATTGATG
2881 AATAAAGCATTTATTAGAAGCTCTGTGTACAGTCATTTTTCAAGCATTAGTTTGCTGGTT
2941 TTGCAGGGGGTTTAGTATGCTGTTGGCCATACTTGTCAATGAACCTATTCACATCTGGGT
3001 CACCAGGAACAGCCTCTGTACCCTCATAAATCCTGACTTCTTCTACCTGAGCAGCTTCTC
3061 CTTCCATGGGCTGGCCTTCAATTGTTGGAAGCATATTGTTGTACAAAGAAGCTAGCAAGC
3121 TTGTAACTGGGTAAGGATTTTTCACCCATCTTTTTCTCAAGGTCACATTAAAATATCTAG
3181 GCAGCCCCCTCCAATGCCAGCCTGCACTGTTGTATTCTATGTACCAGCCCATAACATCTG
3241 CTGCACTGAGATAAAGCCCATCTCCTTTGCAAAGAGGCCCAACCCCATTTTCATCCAGAA
3301 GCACAGTAGTCAAGGTATTAGTAAACTGCATCACTGGTGGAGTACCAGTACCACCTGTGA
3361 AATTCCCATAGTATCTGCTGTTTTCATTTTTAGAAGGATCAGGACCCCATATTTCTATGG
3421 GGTACCTACCATCCTTGTCCAATTTTGCTTTTGCAGTAGGGTCCAGCACCTGGTTTGTGG
3481 AAGTCATTGCTTTGCCAGTAACAGTTTTGATACTAACAATAGCTGCCTCATAATTTGCAT
3541 TATAGTTCTGCACTAGGCCTTGCAAATCTAATGGTTCTCCTCCCACTGCAAACATGTGGT
3601 AAGTTGTACCCTCAACTGGTTTGGAAATTCCAATATCCTTTGTCTCACTTCTGGAGCCAT
3661 ATCCATGCACATTTAGAAGGGATCCCACTCCAACAACTTCAGTTTTTACAGATACAGCCT
3721 CCCACATTTGAAGGGTATCACAGGTCAAATCTTCATTCAGTGTTGGAAGCTGTATTTTAG
3781 CCATACTGTAATATGGCAGTTGATTAGCCTTCACTTCATCAGCAGTAAGGGAGCTATTTA
3841 CTTTAATACTCTGGGAGAACCCATAATACTGGCCATCAGTTCCTGTGCCAGGCTTGTTCT
3901 GACCCATTCTAGGATTAAGGTAGGCCTCAATTTGTGTGATACTGTCTTCTCCTGTTACAA
3961 GATCAAGCACACCCACACCACCCCGCATAATAAGCTTGGGAACATTAGCAGGCTTTGGAC
4021 AGGGTTTCCACAGGGGTTTGCACATCTACTGGAAGCGCCGCTTTTTCTTTTTGGGGCCAT
4081 ACTCAACCTCATCAATGTATGTCTGCCAAGTAGGACTAATGTCTCCGTACAATCCTAGAA
4141 TTAAAGGAAGCATCCAATCAGGTGTCACTCTTTGGTGGGCTCCACCAGGAGCAAAATACC
4201 TCATGATATTTGCCCCTGATTCAAACCAACTAGAAGAGTCTTCCTGCTGTTGACTTCTTC
4261 TCTGGACATCAGGTCTCCCTAGTTCAGCTTCTAATACTTGTCTGCTATTGGCATCTTCAA
4321 TAGAAGGTCTATTACTGTATTCTAAAGCTCTTTCTATTTGTCTTCTTTGAGCTGGATTAA
4381 TTCCTGGAAGTTCTGCATAGTAGTTTTGTAGGCCACCATATATTCTACTATAGGCCTCTC
4441 TAGGTAAATTAGTAACAACCCATCTACTATTTTCCATCATTCTGGCAATGGCATCTAAAA
4501 ATTGATGGGTGGTCTGCAAACTTAAGTCCCTCACAGCAGATTCTACAGCTCCTTGAGTTT
4561 CCCTTCTCAAAGTATCCCAGATATACTCTCCTACAGATTGAAATAATGAGTGGCCCCAGC
4621 CATGAATAACATCCAAGGCATGGGTAAATGACTGTACACCAGGGAATAATATATCATAGT
4681 AGTCAGCTGGTCTCCTGGGAATAAGTGCCATGTTTCTATTCACAATCGGTACTTCGTGAG
4741 CAAGATAGCCGTGTAGACTTCCCAAAGAGAAGGCTGCTGAACCTGCAACAGTTTGAAAAA
4801 TAAATGCTGTCTGGACTGATTCTCTCACAAACTCAGTCATTACTGTTGATGTTAACTCAG
4861 GGGCAGCTTGCATAAATATAAACATGTCTTCACTTAGGCCAATTGAAGACAAAGCTGTCT
4921 CAGCTCCTAAAAACCCCTCCATAGTTATTAATGAAGTAACTTGGGCATCTATCGCGGCAA
4981 AGGCTTCTCCACTAAGTATGGCCTCTACTGAAATTCCAGTAACTGATGAAATTTCGGAGA
5041 GGTAGCTGATCATCTCAATAATCACTGAAATGGCAGATCCCATGTTGACTTACTTGAACA
5101 GTTTGAAAATCTTCTGAACTGTTTCAGGCAGGTTTTTAGGCCGAATTCTAAAGAAACAGA
5161 AAGCAAACACTCAGCGCCGAAGAGCAGGAAATGGCTGACCACTGCACTTGGGCGACACGA
5221 CACGCCTAGCGATAAGGAAGTCACCATGGCAACATAACCGCAGCACTGCTGTTGTCACAG
5281 TTGCCTAGCAAATGACAGACTCAGCAACCACAGGAGAGGAAATGATAGGGCTAGCATTTT
5341 TTCAAATGTAAACCAGAGGCTAGGGG
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Fig. 2. Nucleotide sequence of HapV DNA. The sequence is 5366 nucleotides long and only the early strand is shown from 5′ to 3′. We have arbitrarily designated as the beginning of the sequence a C on one side of a hairpin structure which may be a possible originof replication.