

Multiple sequence alignment of the E7 protein sequence from various strains of HPV18

CLUSTAL O (1.2.4) multiple sequence alignment

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WWB01568.1      MHGPKATLQDIVLHLEPQNEIPVDLLCHEQLSDSEEENDEIDGVNHQHLPARRAEPQRHT      60
WWB01584.1      MHGPKATLQDIVLHLEPQNEIPVDLLCHEKLSDEEENDEIDGVNHQHLPARRAEPQRHT      60
ATL15240.1      MHGPKATVQDIVLHLEPQNEIPVDLLCHEQLSDSEEENDEIDGVNHQHLPARRAEPQRHT      60
AGU90384.1      MYGPKATLQDIVLHLEPQNDIPVDLLCHEQLSDSEEENDEIDGVNHQHLPARRAEPQRHT      60
CAB53099.1      MHGPKATLQDIVLHLEPQNEIPVGLLCHEQLSDSEEENDEIDGVNHQHLPARRAEPQRHT      60
CAB53098.1      MHGPKATLQDIVLHLEPQNEIPVDLLCHEQLSDSEEENDEIDGVNHQHLPARRAEPQRHT      60
ABP99785.1      MYGPKATLQDIVLHLEPQNEIPVDLLCHEQLSDSEEENDEIDGVNHQHLPARRAEPQRHT      60
BCD58635.1      MHGPKATLQDIVLHLEPQNEIPVYLLCHEQLSDSEEENDEIDGVNHQHLPARRAEPQRHT      60
ADC35661.1      MHGPKATLQDIVLHLEPQNEIPVDLLCHEQLSDSEEENDEIDGVNHQHLPARRAEPQRHT      60
WWB01592.1      MHGPKATLQDIVLHLEPQNEIPVDLLCHEQLSDSEEENDEIDGVNHQHLPARRAEPQRHT      60
CAB53097.1      MHGPKATLQDIVLHLEPQNEIPVDLLCHEQLSDSEEENDEIDGVNHQHLPARRAEPQRHT      60
BCD58525.1      MHGPKATLQDIVLHLEPQNEIPVDLLCHEQLSDSEEENDEIDGVNHQHLPARQAEPQRHT      60
BBP06117.1      MHGPKATLQDIVLHLEPQNEIPVDLLCHEQLSDSEENNEIDGVNHQHLPARRAEPQRHT      60
AGU90416.1      MYGPKATLQDIVLHLEPQNEIPVDLLCHEQLSDSEEENDEIDGVNHQHLPARRAEPQRHT      60
ABP99745.1      MHGPKATLQDIVLHLEPQNEIPVDLLCHEQLSDSKEENDEIDGVNHQHLPARRAEPQRHT      60
P06788.2        MHGPKATLQDIVLHLEPQNEIPVDLLCHEQLSDSEEENDEIDGVNHQHLPARRAEPQRHT      60
NP_040311.1      MHGPKATLQDIVLHLEPQNEIPVDLLCHEQLSDSEEENDEIDGVNHQHLPARRAEPQRHT      60
ARS43460.1      MHGPKATLQDIVLHLEPQNEIPVDLLCHEQLSDSEEENDEIDGVNHQHPARRAEPQRHT      60
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WWB01568.1      MLCMCKCEAXXXVVESSADDLRAFQQLFLNTLSFVCPWCASQQ      105
WWB01584.1      MLCMCKCEARIELVVESSADDLRAFQQLFLNTLSFVCPWCASQQ      105
ATL15240.1      LLCMCKCEARIELVVESSADDLRAFQQLFLNTLSFVCPWCASQQ      105
AGU90384.1      MLCMCKCEARIELVVESSADDLRAFQQLFLSTLSFVCPWCASQQ      105
CAB53099.1      MLCMCKCEARIELVVESSADDLRAFQQLFLKTLFVCPWCASQQ      105
CAB53098.1      MLCMCKCEARIELVVESSADDLRAFQQLFLKTLFVCPWCASQQ      105
ABP99785.1      MLCMCKCEARIELVVESSADDLRAFQQLFLSTLSFVCPWCASQQ      105
BCD58635.1      MLCMCKCEARIELVVESSADDLRAFQQLFLNTLSFVCPWCASQQ      105
ADC35661.1      MLCMCKCEARIELVVESSADDLGAFQQLFLNTLSFVCPWCASQQ      105
WWB01592.1      MLCMCKCEARIELVVESSADDLRAFQQLFRNTLSFVCPWCASQQ      105
CAB53097.1      MLCMCKCEARIELVVESSADDLRAFQQLFLKTLFVCPWCASQQ      105
BCD58525.1      MLCMCKCEARIELVVESSADDLRAFQQLFLNTLSFVCPWCASQQ      105
BBP06117.1      MLCMCKCEARIELVVESSADDLRAFQQLFLNTLSFVCPWCASQQ      105
AGU90416.1      MLCMCKCEARIELVVESSADDLRAFQQLFLNTLSFVCPWCASQQ      105
ABP99745.1      MLCMCKCEARIELVVESSADDLRAFQQLFLNTLSFVCPWCASQQ      105
P06788.2        MLCMCKCEARIKLVVESSADDLRAFQQLFLNTLSFVCPWCASQQ      105
NP_040311.1      MLCMCKCEARIELVVESSADDLRAFQQLFLNTLSFVCPWCASQQ      105
ARS43460.1      MLCMCKCEARIELVVESSADDLRAFQQLFLNTLSFVCPWCASQQ      105
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