

Multiple sequence alignment of the E6 protein sequence from various strains of HPV16

CLUSTAL O (1.2.4) multiple sequence alignment

A0X22811.1	MHQKRTAMFQDPQERPRKLPQLCTELQTTIHDIILKCVYCKQQLLRREVYDFAFRDLCIV	60
AVN72846.1	MHQKRTAMFQDPQERPRKLPQLCTELQTTIHDIILECVYCKQQLLRREVYDFAFRDLCIV	60
AVN75252.1	MHQKRTAMFQDPQERPRKLPQLCTELTTIHDIILECVYCKQQLLRREVYDFAFRDLCIV	60
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AVN80458.1	MHQKRTAMFQNPQERPRKLPQLCTELQTTIHDIILECVYCKQQLLRREVYDFAFRDLCIV	60
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AVN72949.1	MHQKRTAMFQDPQERPRKLPQLCTELQTTIHDIILGCVYCKQQLLRREVYDFAFRDLCIV	60
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AJP00121.1	YRDGNPYAVCDKCKVFYISKISEYRHYCYSLYGTTLEQQYNKPLCDLLIRCINCQKPLCPE	120
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XCY57405.1	YRDGNPYAVCDKCLKFYISKISEYRHYCYSLYGTTLEQQYNKPLCDLLIRCINCQKPLCPE	120
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AAV91676.1	EKQRHLDKKQRFHNIIRGRWTGRCMSCCRSSRTTRETQL	158
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AVN81959.1	EKQRHLDKKQRFHNIIRGRWTGRCMSCCRSSRTTRETQL	158
ALU65727.1	EKQRHLDKKQRFHNIIRGRWTGRCMSCCRSSRTTRETQL	158
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AVN72876.1	EKQRHLDKKQRFHNIIRGRWTGRCMSCRSSRTTRETQL	158
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AVN76710.1	EKQRHLDKKQRFHNIIRGRWTGRCMSCCRSSRTTRETQL	158
AVN84665.1	EKQRHLDKKQRFHNIIRGRWTGRCMSCCRSSXTTRETQL	158
ALU65774.1	EKQRHLDKKQRFHNIIRGRWTGRCMSCCRSSTTTRETQL	158
AVN74104.1	EKQRHLDKKQRFHNIIRGRWTGRCMSCCTSSRTTRETQL	158
AVN79003.1	EKQRHLDKKQRFHNIIRGRWTGRCMSCCRSSRTTRETQL	158
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AFS64227.1	EKQRHLDKKQRFHNIIRGRWTGRCMSCCRSSRTTRETQL	158
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