

NCBI Multiple Sequence Alignment Viewer, Version 1.25.3						
Sequence ID		Start	Alignment	End	Organism	
			<div><div></div><div>110120130140150160167</div></div>			
Query_2285929	(+)	1	MHWGTL CGFLWLWPYLFYVQA VPIQKVQDDTKTLIKTIVTRINDISHTQSVSSKQKVTGLDFIPGLHPILTL SKMDQTLAVYQQILTSMP SRNVIQISNDLENLRDLLHVLA FSK SCHLPWASGLETLD SLGGVLEASGYSTE VVALSR LQGS LQDMLWQ LDLS PGC	167		
P41159.1	(+)	1	MHWGTL CGFLWLWPYLFYVQA VPIQKVQDDTKTLIKTIVTRINDISHTQSVSSKQKVTGLDFIPGLHPILTL SKMDQTLAVYQQILTSMP SRNVIQISNDLENLRDLLHVLA FSK SCHLPWASGLETLD SLGGVLEASGYSTE VVALSR LQGS LQDMLWQ LDLS PGC	167	Homo sapiens	
Q28504.1	(+)	1	MYWRTLW GFLWLWPYLFYIQA VPIQKVQSDTKTLIKTIVTRINDISHTQSVSSKQ RVTGLDFIPGLHPVLTLSQMDQTLAIYQQILINLPS RNVIQISNDLENLRDLLHLLA FSK SCHLP LA SGLETLE SLGDVLEASLYSTE VVALSR LQGS LQDMLWQ LDLS PGC	167	Macaca mulatta	
O02750.1	(+)	1	VPIQKVQDDTKTLIKTIVTRINDISHTQSVSSKQKVTGLDFIPGLHPILTL SKMDQTLAVYQQILTSMP SRNM IQISNDLENLRDLLHVLA FSK SCHLPWASGLETLD SLGGVLEASGYSTE VVALSR LQGS LQDMLWQ LDLS PGC	146	Pan troglodytes	
Q95189.1	(+)	1	VPIQKVQDDTKTLIKTIVTRISDISHTQSVSSKQKVTGLDFIPGLHPILTL SKMDQTLAVYQQILTSMP SRNM IQISNDLENLRDLLHVLA FSK SCHLPWASGLETLD SLGGVLEASGYSTE VVALSR LQGS LQDMLWQ LDLS PGC	146	Gorilla gorilla gorilla	
Q9N2C1.1	(+)	1	M LCGP LCR FLWLWPYLSYVEA VPIRK VQDDTKTLIKTIVTRINDISHTQSVSSKQ R VAGLDFIPGLHPVLSLSKMDQTLAIYQQILTGLPS RNVVQISNDLENLRDLLHLLA SSKNCPLPR AGLETLE SLGGALEASLYSTE VVALSR LQAS LQDMLW R LDLSPGC	167	Felis catus	
Q95234.1	(+)	1	VPIQKVQDDTKTLIKTVITRINDISHTQSVSSKQKVTGLDFIPGLHPILTL SKMDQTLAVYQQILTSMP SRNV IQISNDLENLRDLLHVLA FSK SCHLPWASGLETLD RLGGVLEASGYSTE VVALSR LQRS LQDMLWQ LDLS PGC	146	Pongo pygmaeus	
Q29406.1	(+)	1	MRCGP LCRFLWLWPYLSYVEA VPIW RVQDDTKTLIKTIVTRISDISHMQSVSSKQ R VTGLDFIPGLHPVLSLSKMDQTLAIYQQILTS LPSRNVIQISNDLENLRDLLHLLA SSKSCPLPQARA LETLE SLGGVLEASLYSTE VVALSR LQGA LQDML RQ LDLSPGC	167	Sus scrofa	
P50595.1	(+)	1	MRCGP LYRFLWLWPYLSYVEA VPIRK VQDDTKTLIKTIVTRINDISHTQSVSSKQ R VTGLDFIPGLHP LLSLSKMDQTLAIYQQILTS LPS RNVVQISNDLENLRDLLHLLA ASKSCPLPQVRA LESLE SLGVVLEASLYSTE VVALSR LQGS LQDML RQ LDLSPGC	167	Bos taurus	
Q5J732.1	(+)	1	MRCGP LYQFLWLWPYLSYVEA VPIRK VQDDTKTLIKTIVTRINDISHTQSVSSKQ R VTGLDFIPGLHP LLSLSKMDQTLAIYQQILTS LPS RNVVQISNDLENLRDLLHLLA ASKSCPLPQVRA LESLE SLGVVLEASLYSTE VVALSR LQGS LQDML RQ LDLSPGC	167	Bubalus bubalis	