CLUSTAL muttiple sequence	acignment by hosele (5:0)
ENSMUST00000184554.7 ENST00000556895.T0	CACCCGAGCTGTCAGGCGTTCCGCCGCGGCCGGCGGGGGGGG
ENSMUST00000184554.7 ENST00000556895.T0	CGCCCGGACGCCAGCAGGCCCGCGGGAGTGGGGCTGCCGCGGCTGAGGCGAGGCGGGCC
ENSMUST00000184554.7 ENST00000556895.T0	-CGCGCG-CGATGTGGGAGCTCGCGGCCGGAGCGCCCGGGGAGGCCGGGCCCACGACGCC GCGCGCGTCGGCGTCACAGCCCGGGCAGAGGCGCCCAGGGCGGCCCACGACGCC ****** ** ** *** *** **************
ENSMUST00000184554.7 ENST00000556895.T0	CGTGGCTCCGCTGCGGCAGCGGCGGTGCTGGTGTCGCGCGCG
ENSMUST00000184554.7 ENST00000556895.T0	GCCGCGGGCGGAGGCTGCGGCGGGCGACCCGTCCTGGACACGCGAGGAAGAGCGAGC
ENSMUST00000184554.7 ENST00000556895.T0	ATGGCGGCAGGGCCGCGCTTCGACCCGGTAACTTAGAAGATGATAATTAAT
ENSMUST00000184554.7 ENST00000556895.T0	TGATAATTCTGAATAAATACAGCTTTTATCCCAGGTGTGCCATTTTGAAGACTGAG TGTTGGTTCTGAACAAATAATGAGTTCTTTTATTTGAGGTATGCCATTTTGAAGACTGAG ** * ****** ****** * ***************
ENSMUST00000184554.7 ENST00000556895.T0	ACCATAGAGTTCTAAGAATAAAGGAAAGAGCCCTTGGGAAATTATTATAT ACGTTGGAGTTTTATCCTAGAGGATAAAGGAAATCTTTGGGAAAGTCAGTATTTTAT ** * ***** ** ** ******** * ******* * ****
ENSMUST00000184554.7 ENST00000556895.T0	ATAGCAAAAATGTGAATCCTCAGATGGAATGAAAGGCCTGCACCATAGACATAGCAAAAATATGAA-CCTCAAACTGAATCCTCTAAAGGCCCCATACAATGAAGTAGAC**********
ENSMUST00000184554.7 ENST00000556895.T0	ATCGAAGCATTTAAATTTTTTCTTCTAATTTTTTATGAAGCACCCCGCTTGAAGA ATTGGTGGAAACATTTACACCCAGTCAGCTTGAAGA ** *** *******
ENSMUST00000184554.7 ENST00000556895.T0	GTTTGAAATGGACTTTACCACTGAGAAATCAAGATGGCAGCCCATTATGGGGAATTGAGG GTTTCAAATGGACCCTACCACTGAGAAATCAAGATGGCAGTCTACTATGGGGAATTGAGG **** ******* **********************
ENSMUST00000184554.7 ENST00000556895.T0	-AAAATGGATTAATGCAAGAATGCTGTAATATTATACAACCAAC
ENSMUST00000184554.7 ENST00000556895.T0	GTGGATTCCATGAAATGAATGA—————TTCTTACCCAACACAAATGGACAGTGGA ATGGATTCCATGCAGTGAATGAAGACATAGGTTTCTTACCCAACACAAATGGACAGTGGA *********************************
ENSMUST00000184554.7 ENST00000556895.T0	ATTTACTTCCTAAAGACTTGTTACATGTCATGTACATTTTTGACATCTGGAGAAGACTCT TTTGACTTTCTAAAGACTTTTTTTGTGGACA-TTTTGACATTTGGAGAAG ** *** ********** * * * **********
ENSMUST00000184554.7 ENST00000556895.T0	ACAATTCTACAAATGGTAGTTTGTATTCCTGGAATTTCTTGCAGTTTGATCTGAAGTGAC
ENSMUST00000184554.7 ENST00000556895.T0	CTTATGGAATGTTAACTTTAATAAAATCTCTAAAAACTTAAAAA AAAG

CLUSTAL multiple sequence alignment by MUSCLE (3.8)
