





> [ab1075698.1|KSU75698](#) **ED** Human herpesvirus 8 type M, partial genome  
 Length=137508

Score = 549 bits (297), Expect = 2e-157  
 Identities = 297/297 (100%), Gaps = 0/297 (0%)  
 Strand=Plus/Plus

Query	1	AGTGGGTTTGCTGGCCCTTGGTTGCCATGGATGCTACCTAGGCAAGTCCAGATATAATAA	60
Sbjct	127583		127642
Query	61	CCGGGGTGAGAAACAGAAACGGCCAAATAACCCATGTTTTTCGAAACCACACACACCT	120
Sbjct	127643		127702
Query	121	TAAACAAAATCATGTACACCTGGTATTACTATTTCCACACATCTTATAGCATTTCAAAG	180
Sbjct	127703		127762
Query	181	ATAAGGTGCCCTTACGGGCCCGCCGAAACAAGTGGGGCGGCTACTCAGTGTATAAAG	240
Sbjct	127763		127822
Query	241	TCAGCCGGACCAAGCTGCTGCTCTTGGGGACGTGACTGCTTCGTGGCGCAGTGCCT	297
Sbjct	127823		127879

Promoter Region Blast