MK214431

>Seq1 [organism=Hi-Line Brown] GTPase,IMAP family member 5 (GIMAP5) gene, complete cds

ATGCGTCTGCTCCTGGTTGGGAAGACCGGGGGGGGACGAAGTGCCACAGGGAACTCCATTCTGGGGCGATGTGCCTTCGAGTCCAAGCTGGCCACCAAACCAGTGACCCTGAGCTGCCAGAAGGCAGACGGGCTCTGGAATGGCCAAGACATCACAGTGATCGACACGGCCAACATCTTCTATCTGTGGGATGACAATGCTCCGGTGCACAAAGAAATCTTGCACTGCGTCAGGCTGTCCTTCCCGGGCCCCCACGCTCTGCTGTTGGTCACCCAACTGGGCCGCTTCACCCAGGAGGACCAGGAGGCCGTGAAGGGCGTGCAGGACGTCTTCGGATCCAGCGTGCTCAGATACACGATCGTGGTGTTCACCCGTGGAGAAGAGCTGGTGTCAGGGACCCTGGATGACTACGTGACCTACACTGACAACAGAGCTCTGCGTGATGTGATCCAGAGCTGCGGGTACAGGTACTGCAGCATCAACAACCGGGCCACCAGCGCCGAGCGGGACCAGCAGGTCCAGCAGCTGATGGAGAAGGTCGTCCAAATGGTGCAGGAGAACGAGGGCAAGTACTACAGCAACGAGATGTACCTGGATCCCGGCTTAACGGAAGAGAAGGTGATGTATCACGTGAAGATGTACAGAGAAGATAGGAAAAGCAGAGAGCTGCCCTGGTGGAGGAAATACTCGAAGTGCCTCATGGTTGTTGGGGGGGGTGTCATTCTCACGGCTGTGGTTCTCTCATTTTGCCTTACCCGATGGAAGCCGTGA