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                                                  10        20        30        40        50        60        70        80        90
                                                  ATTTCACACAGGAAACAGCTATGACCATGATTACGAATTCGAGCTCGGTACCCGGGGATCCCCACCATGGTGAGCGTGATCACCAGCGAGATG
ZH02-M76150-G0213094-1_M13-77.ab1(1>796) ←        ATTTCACACAGGAAACAGCTATGACCATGATTACGAATTCGAGCTCGGTACCCGGGGATCCCCACCATGGTGAGCGTGATCACCAGCGAGATG
G0213094-1.seq(1>704) →                                                                                GGATCCCCACCATGGTGAGCGTGATCACCAGCGAGATG

                                                 100       110       120       130       140       150       160       170       180
                                                  AAGATCGAGCTGAGGATGGAGGGCGCCGTGAACGGCCACAAGTTCGTGATCACCGGCAAGGGCAGCGGCCAGCCCTTCGAGGGCATCCAGAAC
ZH02-M76150-G0213094-1_M13-77.ab1(1>796) ←        AAGATCGAGCTGAGGATGGAGGGCGCCGTGAACGGCCACAAGTTCGTGATCACCGGCAAGGGCAGCGGCCAGCCCTTCGAGGGCATCCAGAAC
G0213094-1.seq(1>704) →                           AAGATCGAGCTGAGGATGGAGGGCGCCGTGAACGGCCACAAGTTCGTGATCACCGGCAAGGGCAGCGGCCAGCCCTTCGAGGGCATCCAGAAC

                                                 190       200       210       220       230       240       250       260       270
                                                  GTGGACCTGACCGTGATCGAGGGCGGCCCCCTGCCCTTCGCCTTCGACATCCTGACCACCGCCTTCCACTACGGCAACAGGGTGTTCGTGGAG
ZH02-M76150-G0213094-1_M13-77.ab1(1>796) ←        GTGGACCTGACCGTGATCGAGGGCGGCCCCCTGCCCTTCGCCTTCGACATCCTGACCACCGCCTTCCACTACGGCAACAGGGTGTTCGTGGAG
G0213094-1.seq(1>704) →                           GTGGACCTGACCGTGATCGAGGGCGGCCCCCTGCCCTTCGCCTTCGACATCCTGACCACCGCCTTCCACTACGGCAACAGGGTGTTCGTGGAG

                                                 280       290       300       310       320       330       340       350       360       370
                                                  TACCCCGAGGAGATCGTGGACTACTTCAAGCAGAGCTTCCCCGAGGGCTACAGCTGGGAGAGGAGCATGAGCTACGAGGACGGCGGCATCTGC
ZH02-M76150-G0213094-1_M13-77.ab1(1>796) ←        TACCCCGAGGAGATCGTGGACTACTTCAAGCAGAGCTTCCCCGAGGGCTACAGCTGGGAGAGGAGCATGAGCTACGAGGACGGCGGCATCTGC
G0213094-1.seq(1>704) →                           TACCCCGAGGAGATCGTGGACTACTTCAAGCAGAGCTTCCCCGAGGGCTACAGCTGGGAGAGGAGCATGAGCTACGAGGACGGCGGCATCTGC

                                                 380       390       400       410       420       430       440       450       460
                                                  CTGGCCACCAACAACATCACCATGAAGAAGGACGGCAGCAACTGCTTCGTGAACGAGATCAGGTTCGACGGCGTGAACTTCCCCGCCAACGGC
ZH02-M76150-G0213094-1_M13-77.ab1(1>796) ←        CTGGCCACCAACAACATCACCATGAAGAAGGACGGCAGCAACTGCTTCGTGAACGAGATCAGGTTCGACGGCGTGAACTTCCCCGCCAACGGC
G0213094-1.seq(1>704) →                           CTGGCCACCAACAACATCACCATGAAGAAGGACGGCAGCAACTGCTTCGTGAACGAGATCAGGTTCGACGGCGTGAACTTCCCCGCCAACGGC

                                                 470       480       490       500       510       520       530       540       550
                                                  CCCGTGATGCAGAGGAAGACCGTGAAGTGGGAGCCCAGCACCGAGAAGATGTACGTGAGGGACGGCGTGCTGAAGGGCGACGTGAACATGGCC
ZH02-M76150-G0213094-1_M13-77.ab1(1>796) ←        CCCGTGATGCAGAGGAAGACCGTGAAGTGGGAGCCCAGCACCGAGAAGATGTACGTGAGGGACGGCGTGCTGAAGGGCGACGTGAACATGGCC
G0213094-1.seq(1>704) →                           CCCGTGATGCAGAGGAAGACCGTGAAGTGGGAGCCCAGCACCGAGAAGATGTACGTGAGGGACGGCGTGCTGAAGGGCGACGTGAACATGGCC

                                                 560       570       580       590       600       610       620       630       640       650
                                                  CTGCTGCTCCAGGGCGGCGGCCACTACAGGTGCGACTTCAGGACCACCTACAAGGCCAAGAAGGTGGTGCAGCTGCCCGACTACCACTTCGTG
ZH02-M76150-G0213094-1_M13-77.ab1(1>796) ←        CTGCTGCTCCAGGGCGGCGGCCACTACAGGTGCGACTTCAGGACCACCTACAAGGCCAAGAAGGTGGTGCAGCTGCCCGACTACCACTTCGTG
G0213094-1.seq(1>704) →                           CTGCTGCTCCAGGGCGGCGGCCACTACAGGTGCGACTTCAGGACCACCTACAAGGCCAAGAAGGTGGTGCAGCTGCCCGACTACCACTTCGTG

                                                 660       670       680       690       700       710       720       730       740
                                                  GACCACCAGATGGAGATCACCAGCCACGACAAGGACTACAACAAGGTGAAGCTGTACGAGCACGCCAAGGCCCACAGCGGCCTGCCCAGGCTG
ZH02-M76150-G0213094-1_M13-77.ab1(1>796) ←        GACCACCAGATGGAGATCACCAGCCACGACAAGGACTACAACAAGGTGAAGCTGTACGAGCACGCCAAGGCCCACAGCGGCCTGCCCAGGCTG
G0213094-1.seq(1>704) →                           GACCACCAGATGGAGATCACCAGCCACGACAAGGACTACAACAAGGTGAAGCTGTACGAGCACGCCAAGGCCCACAGCGGCCTGCCCAGGCTG

                                                 750       760       770       780       790
                                                  GCCAAGTAACTGCAGGCATGCAAGCTTGGCACTGGCCGTCGTTtTACAACGT
ZH02-M76150-G0213094-1_M13-77.ab1(1>796) ←        GCCAAGTAACTGCAGGCATGCAAGCTTGGCACTGGCCGTCGTTtTACAACGT
G0213094-1.seq(1>704) →                           GCCAAGTAACTGCAG
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