

LymTBH

cDNA ATGGATGGACACGCTTGGTCGATCTCGGCTGTCGCTGGCTCGTGGTTCTCTCGGCTGACCACTCGGTTCAAGCGTATCA 80
TSA 80

GTATCAAGTGACCCTAGACCCCGACAGCAGGTACCTGTTCCAGTGGTCAGTGGACTACAAGGAAAGTCTTATCAATGTCC 160
..... 160

AGCTCACGTGCAAGGTCACGCCAGAGAGCTGGCTTGCTTTTGGATTCTCGGACTACGGTGACGTCACCTTCCGCTGACCTA 240
..... 240

ATCTTGTTTTGGACAGACGGTGATGGCAAGCACCATTTCTTCGATGGTCACACGACACCGGATGGGATATTTCTACCTGA 320
..... 320

CCGGCAACAGGACTACCATCTGACATCAGTCGCCGATGACCGTGGGTGAGTTGTGCTGGATTTCTACAGGCACTTCAACA 400
..... 400

CATGCGACCCAGAGGACTACGCCCTGGACAACGGCACCACGCATCTCGTGTACGTAGAGTCGGCGCAGCCCGAGGGCCCG 480
..... 480

CCCCTCGCCCGTGACGTCACGCGACTACGTCACGGGGTCCAACGCCCTGCAGCTGTTAAAACCCGAGATTTCCGCCCCAGT 560
..... 560

GTTCGCCGAGGACACGTGGTCGTTTTGAGGTCAGGGCGCCAGAGGTGTTGGTGCCAGCCGAGGAGACGACCTACTGTTGGC 640
..... 640

ACACTACCATCTTACCGGACATGCCATCACCCACCACATCATAACAGTACGAGGGGATCGTGCCGAGGGAAGCGGTGAC 720
..... 720

CTCGTCCACCACATGGAAGTCTTCCACTGTCAGGTCCAAAAGGTCACGGCGTCCCGTACTACAATGGGCCCGGCATAGC 800
..... 800

TGAAGGGAAGCCCGAAGGCCCTGGAGGTGTGCAGGAAGGTCATCGGCCGTTGGGCCATGGGGCGGAGGCGATGATATACC 880
..... 880

CCGAGGAGGCAGGGGTCCCCGTGGGAGGCCAGGGGTTCTCGCGCTTTGCCTTGCTTGAGGTCCACTACAACAACCCACAG 960
..... 960

AAGAAATCTGGTCGAATGGACTCGTCCGGGATCCGCTTCCACGTGACGTCACAACCTCCGGAAGTATGACGCGGGGATCAT 1040
..... 1040

GGAGCTCGGACTGGAGTACGTCAACAAGATGGCCGTGCCGCCCGACAGCGGGACTTCAAGCTGAGCGGTTACTGCGTAC 1120
..... 1120

ACAAATGTACACAAATGAGCTTACCACCTGCCGGCATCCACGTGTTCCGCTCCAGCTCCACACTCACCTGACGGGGCGC 1200
..... 1200

CGCGTGTACACCAAGCACGCGCGGGATGGCGCGGAGCTTCCGGAAGTCAACCGGGACAACCACTACAGCCCCACTTCCA 1280
..... 1280

GGAGATCAGGAGGCTGCCGACGCCGACCACGTCTCCCGGGCAGCTGCTGATAACCACGTCGAGTACGACACAACGA 1360
..... 1360

AGAGGTCAAAGGCTACAGTTGGCGGCTTCTCCATAACAGATGAGATGTGTCTCAACTACGTCCACTACTACCCAGGTCT 1440
..... 1440

GACCTTGAAGTCTGCAAGAGCTCAGTCAGGACTGATTGCTTCCACATTTTTCTTTTTGCTCAACAGATTGAGAATTC 1520
..... 1520

AAAGGTGCTCCAGAGCGGGTGACAGGGCAAACATGAAAGCATTGAGTGGTCTCCCCTGAATGTCCGGCTCCTGGAGG 1600
..... 1600

ATCTGTACAGCACCTCGCCGCTCTCCATGCAGTGCAACAGGTCAGACGGCACCCGGTCCCGGGCAGTGGGAGCACGTT 1680
..... 1680

CGAGTCCCAGACATTGTCCGACCGTTGGTTGTGGACACGTCAGAAGTTTGTCTCGGGCCACGTCACAGCGGAATAG 1755
..... 1755