miRNA: mir-130

Human predicted target: ENSG00000184371

MACROPHAGE COLONY STIMULATING FACTOR-1 PRECURSOR (CSF-1)

(MCSF)

pIS36w

ggtagatctgcgatctgcatctcaattagtcagcaaccatagtcccgccctaactccgcccatactccgcccagt teegeceatteteegeceeategetgaetaattttttttatttatgeagaggeegaggeegeeteggeetetgagetatteeagaagt agtgaggaggcttttttggaggcctaggcttttgcaaaaagcttggcattccggtactgttggtaaagccaccatggaagacgcca aaaacataaagaaaggcccggcgccattctatccgctggaagatggaaccgctggagagcaactgcataaggctatgaagag atacgccctggttcctggaacaattgcttttacagatgcacatatcgaggtggacatcacttacgctgagtacttcgaaatgtccgtt ccggtgttgggcgcgttatttatcggagttgcagttgcgcccgcgaacgacatttataatgaacgtgaattgctcaacagtatggg a a attattat catggatt ctaa a acggattac catgggatt t cattagt catgtac acgt to get catcate the categories of the content of the categories ogtgtcgctctgcctcatagaactgcctgcgtgagattctcgcatgccagagatcctatttttggcaatcaaatcattccggatactgc gatttta agtgttgttccattccatcacggttttggaatgtttactacactcggatatttgatatgtggatttcgagtcgtcttaatgtatagatttgaagaagagetgtttetgaggageetteaggattacaagatteaaagtgegetgetggtgeeaaccetatteteettettegee aaaagcactctgattgacaaatacgatttatctaatttacacgaaattgcttctggtggcgctcccctctctaaggaagtcggggaa geggttgccaagaggttccatctgccaggtatcaggcaaggatatgggctcactgagactacatcagctattctgattacacccgagggggatgataaaccgggcgcggtcggtaaagttgttccattttttgaagcgaaggttgtggatctggataccgggaaaacgct gggcgttaat caa agaggcgaactgtgtgtgagaggtcctatgattatgtccggttatgtaaacaatccggaagcgaccaacgccgtctctgattaagtacaaaggctatcaggtggctcccgctgaattggaatccatcttgctccaacaccccaacatcttcgacgcaggtgtcgcaggtcttcccgacgatgacgccggtgaacttcccgccgctgttgttgttttggagcacggaaagacgatgacggaaa aaggtettaeeggaaaactegaegeaagaaaateagagagateeteataaaggeeaagaaggeggaaagategeegtgta $attetaggagctc GAGGCTCCCCTCATGAAGGAAGCCA \\ \textbf{TTGCACT} GTGAACACTGTA$ CCTCCACTCTCCAGCCTCTCCCAGCCTCCTGCACTGAGCTGGCCTCACCAGT CGACTGAGGGAGCCCCTCAGCCCTGACCTTCTCCTGACCTGGCCTTTGACTCC CTTTAGGCTGTGTTCGCCCAGGTTTCTGCATCTTGCACTTTGACtctagcgttct agagteggggeggeeggetegageagacatgataagatacattgatgagtttggacaaaccacaactagaatgcagtga aaaaaatgetttatttgtgaaatttgtgatgetattgetttatttgtaaccattataagetgeaataaacaagttaacaacaactagea tteattttatgttteaggtteaggggggggtgtgggaggttttttaaageaagtaaaacetetacaaatgtggtaaaategataagga totgaacgatggagcggagatgggcggaattgggcggagttagggcggagttagggcggagtatggttget gacta att gag at geat get tt geat act tet geet geg gag act tt gag act tt ceae acc t g g tt get gacta att gag at the calculation of the contraction of the contrgeatgetttgeataettetgeetgetgggageetggggaettteeaeaecetaaetgaeaeaeatteeaeageggateegtegaecgatgcccttgagagccttcaacccagtcagctccttccggtgggcgcggggcatgactatcgtcgccgcacttatgactgtcttc tttat cat g caact c g tag g a cag g t g c g g cag c g ct ctt c c g ct cact g a ct c g c t c g g t c g t c g ct c g ct cact g a ct c g ct c gggcgagcggtatcagctcactcaaaggcggtaatacggttatccacagaatcaggggataacgcaggaaagaacatgtgagcaggaaagaacatgaggaaagaacatgagaagaacatgagcaggaaagaacatgagcaggaaagaacatgagaagaacatgagcaggaaagaacatgagaagaacatgagaagaacatgagaagaacatgagaagaacatgagaagaacatgagaagaacatgagaagaacatgagaagaacatgagaagaacatgagaagaacatgagaaacatgagaagaacatgagaagaacatgagaacatgagaacatgaagaacatgaacatgagaacatgagaacatgagaacatgagaagaacatgaacatgagaacatgaacatgagaacatgagaacatgagaacatgagaacatgaacatgaacatgagaacatgaaa aaggccag caa aaggccaggaaccgtaa aa aaggccgcgttgctggcgtttttccataggctccgccccctgacgagcatcacaaaaatcgacgctcaagtcagaggtggcgaaacccgacaggactataaagataccaggcgtttccccctggaagctccctc gtgcgctctcctgttccgaccctgccgcttaccggatacctgtccgcctttctcccttcgggaagcgtggcgctttctcatagctcacgctgtaggtatctcagttcggtgtaggtcgttcgctccaagctgggctgtgtgcacgaacccccgttcagcccgaccgctgcg

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