



Human CTAGE5 cDNA Clone

Catalog Number: CSB-CL006131HU1

Synonyms	CTAGE-5A, cTAGE-5B, cTAGE-5C, cTAGE-5D, MEA6, MGEA, MGEA11, MGEA6
Accession Number	BC064355
cDNA Size	2430
Definition	Homo sapiens acyl-Coenzyme A binding domain containing 4, mRNA (cDNA clone MGC:35411 IMAGE:5188183), complete cds.
Insert site	rrnB_T1, rrnB_T2
GeneID	4253
Vector	pENTR223.1 (spectinomycin)
Sequence	<p>ATGGAGGAGCCCGGGCTACCCCTCAACCGATTGGGGCTGCTCCTGGAGGAGCTACGCAGGGTTGTGGCAGCA CTGCCTGAAGGTATGAGACCAGATTCTAATCTTTATGGTTTTCCATGGGAATTGGTGATATGTGCAGCTGTTGTTGGATT TTTTGCTGTTCTCTTTTTTTGTGGAGAAGTTTTAGATCGGTTAGGAGTCGGCTTTATGTGGGACGAGAGAAAAAGCTT GCTCTAATGCTTTCTGGACTAATTGAAGAAAAAGTAACTACTTGAAAAATTTAGCCTTGTTCAAAAAGAGTATGAAGG CTATGAAGTAGAGTCATCTTTAAAGGATGCCAGCTTTGAGAAGGAGGCAACAGAAGCACAAAGTTTGGAGGTAGAAAA TCAAATGGCAACCTGTGAAAAGCTGAACAGGTCCAATTCTGAACTTGAGGATGAAATACTCTGTCTAGAAAAAGAGTTA AAAGAAGAGAAATCCAAACATTCTGAACAAGATGAATTGATGGCGGATATTTCAAAAAGGATACAGTCTCTAGAAGATG AGTCAAAATCCCTCAAATCACAAGTAGCTGAAGCCAAATGACCTTCAAGATATTTCAAATGAATGAAGAACGACTGAA CATAGCAATAAAAGATGCTTTGAATGAAAATTCTCAACTTCAGGAAAGCCAGAAACAGCTTTTGAAGAAGCTGAAGTA TGGAAAGAACAAGTGAGTGAACCTAATAAACAGAAAGTAACATTTGAAGACTCCAAAGTACATGCAGAACAAGTTCTAA ATGATAAAGAAAGTCACATCAAGACTCTGACTGAACGCTTGTTAAAGATGAAAGATTGGGCTGCTATGCTTGGAGAAGA CATAACGGATGATGATAACTTGAATTAGAAATGAACAGTGAATCGGAAAATGGTGCTTACTTAGATAATCCTCCAAAAG GAGCTTTGAAGAACTGATTCATGCTGCTAAGTTAAATGCTTCTTTAAAACCTTAGAAGGAGAAAGAAACCAAATTTAT ATTCAGTTGTCTGAAGTTGATAAAACAAAGGAAGAGCTTACAGAGCATATTTAAAATCTTCAGACTGAACAAGCATCTTT GCAGTCAGAAAACACACATTTTAAAATGAGAATCAGAAGCTTCAACAGAACTTAAAGTAATGACTGAATTATATCAAG AAAATGAAATGAACTCCACAGGAAATTAACAGTAGAGGAAAATTATCGGTTAGAGAAAGAAGAGAACTTTCTAAAGTA GATGAAAAGATCAGCCATGCCACTGAAGAGCTGGAGACCTATAGAAAGCGAGCCAAAGATCTTGAAGAAGAATTGGAG AGAACTATTCTTATCAAGGGCAGATTATTTCCCATGAGAAAAAAGCACATGATAATTGGTTGGCAGCTCGGAATGC TGAAAGAAACCTCAATGATTTAAGGAAAGAAAATGCTCACAACAGACAAAATTAAGTAAACAGAGCTTAAATTTGAAC TTTTAGAAAAAGATCCTTATGCACTCGATGTTCAAATACAGCATTGGCAGAGAGCATTCCCATATGGTCCCTCACCA TTGGGTTGGCCTTCATCTGAAACAAGAGCTTTTCTCTCTCCTCCAACCTTTGTTGGAGGGTCCACTCAGACTCTCACCT TTGCTTCCAGGGGGAGGAGGAAGAGGCTCACGAGGCCAGGGAATCCTCTGGACCATCAGATTACCAATGAAAGAG GAGAATCAAGCTGTGATAGGTTAACCGATCCTCATAGGGCTCCCTCTGACACTGGGTCTCTGTACCTCCATGGGACC AGGACCGTAGGATGATGTTTCTCCGCCAGGACAATCATATCCTGATTAGCCCTTCTCCACAAAGGCAAGACAGAT TTTGTTCTAATTCTGGTAGACTGTCTGGACCAGCAGAACTCAGAAGTTTTAATATGCCTTCTTTGGATAAAATGGATGGG TCAATGCCTTCAGAAATGGAATCCAGTAGAAATGATACCAAAGATGATCTTGGTAATTTAATGTGCCTGATTATCTCTC CCTGCTGAAAATGAAGCCACTGGCCCTGGCTTTGTTCTCCACCTCTTGCTCCAGTCAGAGGTCCATTGTTCCAGTG GATGCAAGAGGCCCATCTTGAGAAGAGGACCTCCTTTCCCCCACCTCCTCCAGGAGCCATGTTTGGAGCTTCTCG AGATTATTTCCACCAGGGGATTTCCAGGTCCACCACCTGCTCCATTTGCAATGAGAAATGTCTATCCACCAGGGGG TTTTCTCCTTACCTTCCCCAAGACCTGGATTTTTCCCCCACCTTCTGAAGGTAGAAGTGAGTTCCCTCA GGTTTGATTCCACCTTCAAATGAGCCTGCTACTGAACATCCAGAACCACAGCAAGAAACCTGA</p>