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Product datasheet for SC206909

C1orf172 (KDF1) (NM_152365) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	C1orf172 (KDF1) (NM_152365) Human 3' UTR Clone
Symbol:	C1orf172
Synonyms:	C1orf172; ECTD12
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_152365
Insert Size:	537 bp
Insert Sequence:	>SC206909 3'UTR clone of NM_152365 The sequence shown below is from the reference sequence of NM_152365. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC GGGGCACCCTTGCTCCAGGTGTACTGCTAACCCCTGCCAGGCCCAGCTGCCACACCCTTTCTGGGAGAA GCATGGCCTACAGAATGAAGAGGGGGACCAGGAACCCCTGTGGGAGAGGCTTAGACCTGAAGCAGTGCC CACTCTGGCTCCTCCTGCCTTGGCTGACTGGGTTCCTGGACCATGTGCATTTCACTGGGCCATGGGATC TACATCTCCTTGCATCCCCAGCTGGTCTGATCCCTGCCAGGGCCCCTTCCTTCCTGCTCATGGTCTTCA GGTGGCCTGATCATGGAAAGTAAGGAGTTAGGCATTACCTTCTGGGAGTGAACCCTGACTCCATCCCCC TATTGCCACCCTAACCAATCATGCAAACTTCTCCCTCCCTGGGGTAATTCAACAGTTAAAAGAAGCTTA TCTTAAATGTATTGTATTGGGGGGTGGGCAGGGCCCACTCTATGTTATGTTAAGGAGTTGGTTCTGGTT CTTGGCTGATGTTCTGTATCTTAACATGACCACAGTTTGTAAGTACAAAGGTAA ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG
Restriction Sites:	SgfI-MluI
OTI Disclaimer:	Our molecular clone sequence data has been matched to the sequence identifier above as a point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences , e.g., single nucleotide polymorphisms (SNPs).

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Components:	The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The package also includes 100 pmols of both the corresponding 5' and 3' vector primers in separate vials.
RefSeq:	NM_152365.3
Summary:	Plays a role in the regulation of the epidermis formation during early development. Required both as an inhibitor of basal cell proliferation and a promoter of differentiation of basal progenitor cell progeny (By similarity).[UniProtKB/Swiss-Prot Function]
Locus ID:	126695
MW:	18.9