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Product datasheet for SC204619

DLX5 (NM_005221) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	DLX5 (NM_005221) Human 3' UTR Clone
Symbol:	DLX5
Synonyms:	SHFM1; SHFM1D
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_005221
Insert Size:	370 bp
Insert Sequence:	>SC204619 3'UTR clone of NM_005221 The sequence shown below is from the reference sequence of NM_005221. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC CTGGCGCTGGCCTCCGGGACACTCTATTAGATGGGCTGCTCTCTCTTACTCTCTTTTTTGGGACTACTG TGTTTTGCTGTTCTAGAAAATCATAAAGAAAGGAATTCATATGGGGAAGTTCGGAAAACTGAAAAAGAT TCATGTGTAAAGCTTTTTTTTTGCATGTAAGTTATTGCATTTCAAAAGACCCCCCCTTTTTTTACAGAGG ACTTTTTTTGCGCAACTGTGGACACTTTCAATGGTGCCTTGAAATCTATGACCTCAACTTTTCAAAAGA CTTTTTTCAATGTTATTTTAGCCATGTAAATAAGTGTAGATAGAGGAATTAAACTGTATATTCTGGATA AATAAAATTATTTCGACCATGAAAA 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).
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.

View online »

RefSeq:	NM_005221.6
Summary:	This gene encodes a member of a homeobox transcription factor gene family similiar to the Drosophila distal-less gene. The encoded protein may play a role in bone development and fracture healing. Mutation in this gene, which is located in a tail-to-tail configuration with another member of the family on the long arm of chromosome 7, may be associated with split-hand/split-foot malformation. [provided by RefSeq, Jul 2008]
Locus ID:	1749
MW:	14.3