

Product datasheet for MC202230

Gtf2h5 (NM_181392) Mouse Untagged Clone

Product data:

OriGene Technologies, Inc.

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Product Type:	Expression Plasmids
Product Name:	Gtf2h5 (NM_181392) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Gtf2h5
Synonyms:	2700017P07Rik; 2810432H05Rik; D17Wsu155e
Mammalian Cell Selection:	Neomycin
Vector:	PCMV6-Kan/Neo (PCMV6KN)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>BC029238 sequence for NM_181392 CCGGGCCAGCTCTGGTTTGTTGGAGTTGGAACGCTGCTGCCTCTGCGCGCCTAGAAAAAGAATTCTTGAA GTCATCTGCACCGTCTGCCTGGGAAAAGATGGTCAACGTGCTGAAAGGGGTGCTTATAGAATGTGACCCT GCCATGAAGCAGTTCTTGCTGTACTTGGATGAGGCCAACGCCTTGGGGAAGAAGTTCATCATCAGGACA TTGATGACACGCACGTCTTCGTCATTGCCGAGCTGGTCAACGTCCTCCAGGAGCGAGTAGGGGAACTGAT GGACCAGAATGCCTTTTCTCTTACCCAGAAGTGAGAGAGGCGCTGGTGTTGAGCACTTGGGAATCCCAGCCAC AGACACTTGAGAGAGGCCAGCTCAGTGTCTCGTGGGGTTTATACTTGTTATGAGCATGTCGTGGGAAAGG CTGCAGGGTCATTCATTCAGATGAAAGCCCCTAACTGATTTGGTTGG
Restriction Sites:	Rsrll-Notl
ACCN:	NM_181392
Insert Size:	216 bp
OTI Disclaimer:	Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).
Components:	The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).



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Reconstitution Method:	 Centrifuge at 5,000xg for 5min. Carefully open the tube and add 100ul of sterile water to dissolve the DNA. Close the tube and incubate for 10 minutes at room temperature. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.
Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	<u>BC029238, AAH29238</u>
RefSeq Size:	524 bp
RefSeq ORF:	216 bp
Locus ID:	66467
UniProt ID:	<u>Q8K2X8</u>
Cytogenetics:	17 3.7 cM
Gene Summary:	Component of the general transcription and DNA repair factor IIH (TFIIH) core complex, which is involved in general and transcription-coupled nucleotide excision repair (NER) of damaged DNA and, when complexed to CAK, in RNA transcription by RNA polymerase II. In NER, TFIIH acts by opening DNA around the lesion to allow the excision of the damaged oligonucleotide and its replacement by a new DNA fragment. In transcription, TFIIH has an essential role in transcription initiation. When the pre-initiation complex (PIC) has been established, TFIIH is required for promoter opening and promoter escape. Phosphorylation of the C-terminal tail (CTD) of the largest subunit of RNA polymerase II by the kinase module CAK controls the initiation of transcription. Necessary for the stability of the TFIIH complex and for the

presence of normal levels of TFIIH in the cell.[UniProtKB/Swiss-Prot Function]

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