

## Product datasheet for **SC215556**

### **FPGT (NM\_003838) Human 3' UTR Clone**

#### **Product data:**

Product Type:	3' UTR Clones
Product Name:	FPGT (NM_003838) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	FPGT
Synonyms:	GFPP
ACCN:	NM_003838
Insert Size:	2000 bp



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**Insert Sequence:** >SC215556 3'UTR clone of NM\_003838  
 The sequence shown below is from the reference sequence of NM\_003838. The complete sequence of this clone may contain minor differences, such as SNPs.  
 Blue=Stop Codon Red=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
GAAATCAGTTTTAAAAAGCAGTTTGTAGTAGAGATATTTTAAATATTGTACTTTGCCTTTTTGAGTAA
CATTCCAGAGATAGGTATTTTTGTAGGCTGTTTCACTGAACTCAGTTAATGAAAAGTATTAAACATA
ATTGTTGTAGCATAATATTAATAGTGCAAAAGTACATATAAGTCATTTTGTATGAAAAATATTCCAAGAC
TAAGTTGAGAAAAGAGATACTATTTTGGATGTGTATCAGTATTTTGTTTTTAATAATGATTGATTTGT
GGAGCATTGTTTTTACATAATTAGTTTTAAAGGTAATTTTCTAAGCATACCTTTGGAATTTTCCAT
CTTTTTGAGGCTTTTGGTCCAGTGAAGTTCTAAGTATTCAGTGGCACTTCTCTCCTCACTGTAATTC
TATTTTTAATAAAAAATGGCATACTGTAGGCTTTCAGAGTAGTGTAGGAATACTGTAGAAATACTT
TTTCAGAAAACGAATCCATAGCTGACAAAATTCAGTCCCAATATATTGTGATTATTTTCGTTGATA
AAGAACTAGATACAAAGACCTCTGAAATTTGATGATAAAATTTGTATCTCATTATTTTATCAAAATGAA
ACTAAAAGTACATATGTATTATACACTTGAACATGTGTTTGTATATCTTTAAAATTTTCTTTATGTTC
CATTCCATAGGAAAACACACATATGCACACAAAAGTATCTGTGGGAAAAGTGGTAGTAATAAATT
GAGATTCATCAATAATCATATAATTTCACTATAGACATTACTTGCATTATCTCCTATCAGTCCTTCAA
CAAACTTCAATTACCAGATGAAGTACTTTAGTTTTTCTTATTTTTTGTCCACATGCTGGTGTATGC
TGAGAAAACAATAATTGGCCCAATGCAGACATCAAGAAGCAGGCAGGAAAATGAAAAAGTGAATAAT
ACACAGGTGATAAACAGTTTCGAGAAAAATAAAATTTGGCTAGAATCTCAAGTTATGTGTTTCTATATTG
GTATAAGCATATAAGTGAAGTCTTATAAGTGAATAGAGAAAAGATTGTCAGTGTGTTTTTTTTAAA
TGAAATAACTAGTCTGTGCTACTTTATGTCAATATAAAAAATTTGGTAAACTAGAAGTAACTGTCCACAA
CCCTCAGTTATGATACTTATGTGCGTGTTTTTTTTTCCAAAGTTTTTCCAAAGGAGAAGTACAAATAT
ATGGTAGCTATTGTTTTAAAAATGATTGATTTACTTGCAGATTTTTTCCAGAGGATGTATGTCTTTGTCAT
TCATTAATGTTCAAAATTTAGTTTTTACCAAATCTGAAGTGCCATCAGTTATAAGAATCACCATTATT
TTATGTTCCATTAAGAAAACACAATAAAACATGACACTGCTTTTTTTGTTACTTGAAATTTTCTATTT
TATTGGAGAGCTCTTTTAGATGTTTTTAAATACAGATTTTTTATTTAACCTCTTGCATATACATATAAA
ATAACAAAAAAATAAAATATTGAGATATCATCAAGAAATTTCTACAGCTTCACTTCAAAGTCTTCCAGC
ACTTTCCAGAGCTATTGATATGCATATGTCCACCTAGTACCATCTTCTCTGCTTTTCCAGAAATTTGGTAA
GGCTGCATTTTCCAGTGTCTACCACACTTATGTCTGGATTTTTTTTACCAGCTACAGACCCACACT
CCTGCAAGATTTGATACCTTCTATTTGGCAAAAGTCAAGTTTCACTTTTCTTTCAGTTTAAATCACAGGT
CCTTAGAAGATCGGACCTGTGATTCTATTCAAAGTCCAGAAAGAGATTTTCCACAAATTTGACGTCATATT
AACATGTCAATATGAAAATTAATTGTTTGGTAAACCAGAAAGTACCTGTCTGAAGCCCTCAATTGTGGT
ATTTGTGTGTTTGTTTTTTCTTTTTCAAGATCTTTTTTAAAGAAAATATATGGTAGCTGTTGTTTAAAC
ACGCGTAAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCACCGCCGCTTCTATGAAAGG
  
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**Restriction Sites:** Sgfl-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.

**RefSeq:** [NM\\_003838.5](#)

**Summary:**

L-fucose is a key sugar in glycoproteins and other complex carbohydrates since it may be involved in many of the functional roles of these macromolecules, such as in cell-cell recognition. The fucosyl donor for these fucosylated oligosaccharides is GDP-beta-L-fucose. There are two alternate pathways for the biosynthesis of GDP-fucose; the major pathway converts GDP-alpha-D-mannose to GDP-beta-L-fucose. The protein encoded by this gene participates in an alternate pathway that is present in certain mammalian tissues, such as liver and kidney, and appears to function as a salvage pathway to reutilize L-fucose arising from the turnover of glycoproteins and glycolipids. This pathway involves the phosphorylation of L-fucose to form beta-L-fucose-1-phosphate, and then condensation of the beta-L-fucose-1-phosphate with GTP by fucose-1-phosphate guanylyltransferase to form GDP-beta-L-fucose. Alternative splicing results in multiple transcript variants. Read-through transcription also exists between this gene and the neighboring downstream TNNI3 interacting kinase (TNNI3K) gene. [provided by RefSeq, Dec 2010]

**Locus ID:**

8790

**MW:**

78.8