

Product datasheet for SC125512

GAA (NM_000152) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	GAA (NM_000152) Human Untagged Clone
Tag:	Tag Free
Symbol:	GAA
Synonyms:	LYAG
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL6</u>
E. coli Selection:	Ampicillin (100 ug/mL)
Fully Sequenced ORF:	>OriGene ORF within SC125512 sequence for NM_000152 edited (data generated by NextGen Sequencing)

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ATGGGAGTGAGGCACCCGCCCTGCTCCACCGGCTCCTGGCCGTCTGCGCCCTCGTGTCC
TTGGCAACCGCTGCACTCCTGGGGCACATCCTACTCCATGATTTCTGCTGGTTCCTCCGA
GAGCTGAGTGGCTCCTCCCACTGCTGGAGGAGACTCACCCAGCTCACCAGCAGGGAGCC
AGCAGACCAGGGCCCGGGATGCCAGGCACACCCGGCCGTCCAGAGCAGTGGCCACA
CAGTGGCAGCTCCCCCAACAGCCGTTTCGATTGCGCCCTGACAAGGCCATCACCCAG
GAACAGTGGCAGGCGCCGGCTGTTGCTACATCCCTGCAAAGCAGGGGCTGCAGGGAGCC
CAGATGGGGCAGCCCTGGTGTCTTTCCACCCAGCTACCCAGCTACAAGCTGGAGAAC
CTGAGCTCCTCTGAAATGGGCTACACGGCCACCCTGACCCGTACCACCCACCTTCTTC
CCCAAGGACATCCTGACCCTGCGGCTGGACGTGATGATGGAGACTGAGAACCCTCCAC
TTCACGATCAAAGATCCAGCTAACAGGGCCTACGAGGTGCCCTTGAGACCCCGCATGTC
CACAGCCGGGCACCGTCCCACTCTACAGCGTGGAGTTCTCCGAGGAGCCCTTCGGGGTG
ATCGTGGCCCGGAGCTGGACGGCCGCTGCTGCTGAACACGACGGTGGCGCCCTGTTC
TTTGGGACCACTTCCCTCAGCTGTCCACCTCGCTGCCCTCGCAGTATATCACAGGCCTC
GCCGAGCACCTCAGTCCCTGATGCTCAGCACCAGCTGGACCAGGATCACCTGTGGAAC
CGGGACCTTGGCCACGCCCCGGTGCAGAACCTCTACGGGTCTCACCTTTCTACCTGGCG
CTGGAGGACGGCGGGTGGCACACGGGGTGTTCCTGCTAAACAGCAATGCCATGGATGTG
GTCTTCCAGCCGAGCCCTGCCCTTAGCTGGAGGTCGACAGTGGGATCCTGGATGTCTAC
ATCTTCCGGGGCCAGGCCAAGAGCGTGGTGCAGCAGTACCTGGACGTTGTGGGATAC
CCGTTTCATGCCGCATACTGGGGCTGGCTTCCACCTGTGCCGCTGGGGCTACTCCTCC
ACCGCTATCACCCGCCAGGTGGTGGAGAACATGACCAGGGCCACTTCCCTGGACGTC
CAGTGGAAACGACCTGGACTACATGGACTCCCGAGGGACTTACGTTCAACAAGGATGGC
TTCCGGGACTTCCCGCCATGGTGCAGGAGCTGCACCAGGGCGGGCGGCTACATGATG
ATCGTGGATCCTGCCATCAGCAGCTCGGGCCCTGCCGGGAGCTACAGGCCCTACGACGAG
GGTCTGGCGAGGGGGTTTTTCATACCAACGAGACCGGCCAGCCGCTGATTGGGAAGGTA
TGCCCCGGGTCCACTGCCTTCCCGACTTACCAACCCACAGCCCTGGCCTGGTGGGAG

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GACATGGTGGCTGAGTTCCATGACCAGGTGCCCTTCGACGGCATGTGGATTGACATGAAC
 GAGCCTTCCAACCTCATCAGGGGCTCTGAGGACGGCTGCCCAACAATGAGCTGGAGAAC
 CCACCCTACGTGCCTGGGGTGGTTGGGGGACCCTCCAGGCGGCCACCATCTGTGCCTCC
 AGCCACCAGTTTCTCTCCACACACTACAACCTGCACAACCTCTACGGCCTGACCGAAGCC
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 AGGAATAACACGATCGTGAATGAGCTGGTACGTGTGACCAGTGGGGAGCTGGCCTGCAG
 CTGCAGAAGGTGACTGTCTGGCGTGGCCACGGCGCCACGAGGTCTCTCCAACGGT
 GTCCTGTCTCCAACCTCACCTACAGCCCGACACCAAGGTCTGGACATCTGTGTCTCG
 CTGTTGATGGGAGAGCAGTTTCTCGTCAGCTGGTGTAG

Clone variation with respect to NM_000152.3

5' Read Nucleotide Sequence:

>OriGene 5' read for NM_000152 unedited
 TTTCCCGCCCGTTGNCGCATTGGGCGGTAGGCGTGTACGGTGGGAGGTCTATATAAGCA
 GAGCTCATTTAGGTGACACTATAGAATAACAAGCTACTTGTCTTTTGCAGCGGCCGCGA
 ATTCGGCACGAGGGTGGCCGGGCGGGGCTGCGGGCTTCCCTGAGCGGGCCGGGTC
 GGTGGGGCGGTGCGCTGCCCGCGGGCCTCTCAGTTGGGAAAGCTGAGGTTGTGCGCGGG
 GCCGCGGGTGGAGGTGCGGGATGAGGCAGCAGGTAGGACAGTACCTCGGTGACGCGAAG
 GACCCCGCCACCTCTAGGTTCTCCTCGTCCGCCGTTGTTACGCGAGGGAGGCTCTGCG
 CGTGCCGAGCTGACGGGAAACTGAGGCACGAGCGGGCCTGTAGGAGCTGTCCAGGCC
 ATCTCAAACCATGGGAGTGAGGCACCCGCCCTGCTCCACCGGCTCCTGGCCGTCTGCGC
 CCTCGTGTCTTGGCAACCGCTGCACTCCTGGGGCACATCCTACTCCATGATTTCTGCT
 GGTCCCCGAGAGCTGAGTGGCTCCTCCCAGTCTGGAGGAGACTCACCCAGCTACCA
 GCAGGGAGCCAGCAGACCAGGGCCNCGGGATGCCAGGCACACCCGGGCGTCCCGA
 GCAGTGCCACACAGTGCAGCTCCCCCCACAGCCGCTTCGATTGCGCCCTGACANGG
 CCATCACCCAGNAACAGTGCGAGGCCCGGGTGGTGCTACATCCCTGNCAAGCAGGGG
 CTGCAGGGAGCCANATGGGGCAGCCTGGTGTCTTTTCCACCAGTACCCAGCTACAGCT
 GGAGAAGTGAAGTCTGAAATGGGCTACAGGCACCCTGACCGTACACCCCACTTCTTC
 CCAGGACANCTGACCTGCGNNTGACGTGA

3' Read Nucleotide Sequence:	>OriGene 3' read for NM_000152 unedited TCNCCCCAAGATCGTGGNACGCGGNCCCAGCATTTTANGANCGNGTTTTTTTTTTTTTTTTTTTT TTCCAAAGCCCCCTTTTATTAATAAATA ATTGCACTTAAAAATTTAAATCTCCAAGTCTGGGAATTTCTGAGTCCCAGGCTTG CTCGGGGACCGGCAGGCATCCAGCCCTTGGGCAGCCGGGCAGCGGTTCGGTTGGGGCAA AGCAACAGGCCCGAAGCAGGAGCTCAGGTGCATACCCACACCTCCACCTGAGCACCCCC TTTTCCGGTGCTGGGAACAAATGGAGGGGGTGGCTAATACTACCCGCATGCCGGCAACA GGTGAGGAGGGCAAACCTTACAATCTTATTAACACAGAGCAGCCCTCCAGGGCCCCGGCC CACAGTGCATCTAGGGAAAAAGCTCTCCTAAACACGTTGGGGGCCAAAGCCCCGGCCCA GGAGGGGGAAACAAGCGATGCGGCGCTTGAATGGTTAGTGCCAGCTCCAGGTGACAC ATGCAACCCCCAGGCCCGCACACAGTTGCTGCCCGCACACAGGCTCTGCTTCCCTGGGGA ACCAGCTCCCTCTGGAGAGACTAACACACTCCGCCCGGCTAACACCAGCTGACGAGAAA CTGCTCTCCATCAACAGCGAGACACAGATGTTCCAGGACCTTGGTGTGGGGCTGTAGG TGAAGTTGGAGACAGGGACCCGTTTGGAAAGGACCTGCTGGGGGCCCGGCCACCCC CAAGAAGTCACCTTCTGGAGGTGCAGGCCAACTCCCTTACTGGTCAAACGTACCAACTTA TTCAGAATCGGGTTATTCCTGGCCGA
Restriction Sites:	NotI-NotI
ACCN:	NM_000152
Insert Size:	3700 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).
Reconstitution Method:	<ol style="list-style-type: none"> 1. Centrifuge at 5,000xg for 5min. 2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA. 3. Close the tube and incubate for 10 minutes at room temperature. 4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom. 5. 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.
RefSeq:	NM_000152.2 , NP_000143.1
RefSeq Size:	3846 bp
RefSeq ORF:	2859 bp
Locus ID:	2548
UniProt ID:	P10253
Cytogenetics:	17q25.3
Domains:	Glyco_hydro_31, PD
Protein Families:	Druggable Genome, Transmembrane

Protein Pathways:	Galactose metabolism, Lysosome, Metabolic pathways, Starch and sucrose metabolism
Gene Summary:	<p>This gene encodes lysosomal alpha-glucosidase, which is essential for the degradation of glycogen to glucose in lysosomes. The encoded preproprotein is proteolytically processed to generate multiple intermediate forms and the mature form of the enzyme. Defects in this gene are the cause of glycogen storage disease II, also known as Pompe's disease, which is an autosomal recessive disorder with a broad clinical spectrum. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jan 2016]</p> <p>Transcript Variant: This variant (1) represents the longest transcript. Variants 1, 2, and 3 all encode the same protein.</p>