

Product datasheet for PH312345

FUT8 (NM_178154) Human Mass Spec Standard

Product data:

OriGene Technologies, Inc.

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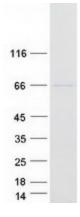
Product Type:	Mass Spec Standards
Description:	FUT8 MS Standard C13 and N15-labeled recombinant protein (NP_835367)
Species:	Human
Expression Host:	HEK293
Expression cDNA Clone or AA Sequence:	RC212345
Predicted MW:	66.3 kDa
Protein Sequence:	>RC212345 representing NM_178154 Red=Cloning site Green=Tags(s)
	MRPWTGSWRWIMLILFAWGTLLFYIGGHLVRDNDHPDHSSRELSKILAKLERLKQQNEDLRRMAESLRIP EGPIDQGPAIGRVRVLEEQLVKAKEQIENYKKQTRNGLGKDHEILRRRIENGAKELWFFLQSELKKLKNL EGNELQRHADEFLLDLGHHERSIMTDLYYLSQTDGAGDWREKEAKDLTELVQRRITYLQNPKDCSKAKKL VCNINKGCGYGCQLHHVVYCFMIAYGTQRTLILESQNWRYATGGWETVFRPVSETCTDRSGISTGHWSGE VKDKNVQVVELPIVDSLHPRPPYLPLAVPEDLADRLVRVHGDPAVWWVSQFVKYLIRPQPWLEKEIEEAT KKLGFKHPVIGVHVRRTDKVGTEAAFHPIEEYMVHVEEHFQLLARRMQVDKKRVYLATDDPSLLKEAKTK YPNYEFISDNSISWSAGLHNRYTENSLRGVILDIHFLSQADFLVCTFSSQVCRVAYEIMQTLHPDASANF HSLDDIYYFGGQNAHNQIAIYAHQPRTADEIPMEPGDIIGVAGNHWDGYSKGVNRKLGRTGLYPSYKVRE KIETVKYPTYPEAEK
	SGPTRTRRLEQKLISEEDLAANDILDYKDDDDKV
Tag:	C-Myc/DDK
Purity:	> 80% as determined by SDS-PAGE and Coomassie blue staining
Concentration:	>0.05 μg/μL as determined by microplate BCA method
Labeling Method:	Labeled with [U- 13C6, 15N4]-L-Arginine and [U- 13C6, 15N2]-L-Lysine
Buffer:	25 mM Tris-HCl, 100 mM glycine, pH 7.3
Storage:	Store at -80°C. Avoid repeated freeze-thaw cycles.
Stability:	Stable for 3 months from receipt of products under proper storage and handling conditions.
RefSeq:	<u>NP 835367</u>
RefSeq Size:	3291
RefSeq ORF:	1725



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	FUT8 (NM_178154) Human Mass Spec Standard – PH312345
Synonyms:	MGC26465
Locus ID:	2530
UniProt ID:	<u>Q9BYC5</u>
Cytogenetics:	14q23.3
Summary:	This gene encodes an enzyme belonging to the family of fucosyltransferases. The product of this gene catalyzes the transfer of fucose from GDP-fucose to N-linked type complex glycopeptides. This enzyme is distinct from other fucosyltransferases which catalyze alpha1-2, alpha1-3, and alpha1-4 fucose addition. The expression of this gene may contribute to the malignancy of cancer cells and to their invasive and metastatic capabilities. Alternative splicing results in multiple transcript variants. [provided by RefSeq, May 2011]
Protein Families:	Transmembrane
Protein Pathways	: Keratan sulfate biosynthesis, Metabolic pathways, N-Glycan biosynthesis
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Product images:



Coomassie blue staining of purified FUT8 protein (Cat# [TP312345]). The protein was produced from HEK293T cells transfected with FUT8 cDNA clone (Cat# [RC212345]) using MegaTran 2.0 (Cat# [TT210002]).

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