

## Product datasheet for PH303695

### C6orf211 (ARMT1) (NM\_024573) Human Mass Spec Standard

#### Product data:

Product Type:	Mass Spec Standards
Description:	C6orf211 MS Standard C13 and N15-labeled recombinant protein (NP_078849)
Species:	Human
Expression Host:	HEK293
Expression cDNA Clone or AA Sequence:	RC203695
Predicted MW:	51 kDa
Protein Sequence:	>RC203695 representing NM_024573 <span style="color: red;">Red</span> =Cloning site <span style="color: green;">Green</span> =Tags(s)

MAVVPASLSGQDVGSFAYLTIKDRIPQILTKVIDTLHRHKSEFFEKGEEGVAEKKAISLLSKLRNELQ  
 TDKPFIPLVKFDVTDIWNQYLEYQQSLLNESDGKSRWFYSPWLLVECYMYRRIHEAIIQSPPIDYFDVF  
 KESKEQNFYGSQESIIALCTHLQQLIRTIEDLDENQLKDEFFKLLQISLWGNKCDLSLGGESSSQNTNV  
 LNSLEDLKPFILLNDMEHLWSLLSNCKKTREKASATRVYIVLDNSGFELVTDLILADFLSSELATEVHF  
 YGKTIPWFVSDTTIHDFNLIEQVKHSNHKWSKCGADWEEYIKMGKWVYHNHIFWTLPEHYCAMPQVAP  
 DLYAELQKAHLILFKGDLNRYKLTGDRKWEFSVPFHQALNGFHPAPLCTIRTLKAEIQVGLQPGQGEQLL  
 ASEPSWWTGKYGIFQYDGPL

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

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- <sup>13</sup> C <sub>6</sub> , <sup>15</sup> N <sub>4</sub> ]-L-Arginine and [U- <sup>13</sup> C <sub>6</sub> , <sup>15</sup> N <sub>2</sub> ]-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><a href="#">NP_078849</a></u>
RefSeq Size:	2572
RefSeq ORF:	1323
Synonyms:	C6orf211


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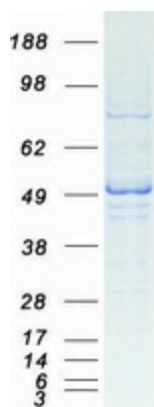
Locus ID: 79624

UniProt ID: [Q9H993](#)

Cytogenetics: 6q25.1

**Summary:** Metal-dependent phosphatase that shows phosphatase activity against several substrates, including fructose-1-phosphate and fructose-6-phosphate (By similarity). Its preference for fructose-1-phosphate, a strong glycating agent that causes DNA damage rather than a canonical yeast metabolite, suggests a damage-control function in hexose phosphate metabolism (By similarity). Has also been shown to have O-methyltransferase activity that methylates glutamate residues of target proteins to form gamma-glutamyl methyl ester residues (PubMed:25732820). Possibly methylates PCNA, suggesting it is involved in the DNA damage response (PubMed:25732820).[UniProtKB/Swiss-Prot Function]

## Product images:



Coomassie blue staining of purified ARMT1 protein (Cat# [TP303695]). The protein was produced from HEK293T cells transfected with ARMT1 cDNA clone (Cat# [RC203695]) using MegaTran 2.0 (Cat# [TT210002]).