

Specifications:

Gene:	hEEA1
Accession:	NP_003557
Insert size:	4249bp
Concentration:	10µg at 0.2µg/µL

hEEA1 cDNA Plasmid

EEA1 early endosome antigen 1 [*Homo sapiens* (human)]

Also known as: MST105; ZFYVE2; MSTP105

Summary:

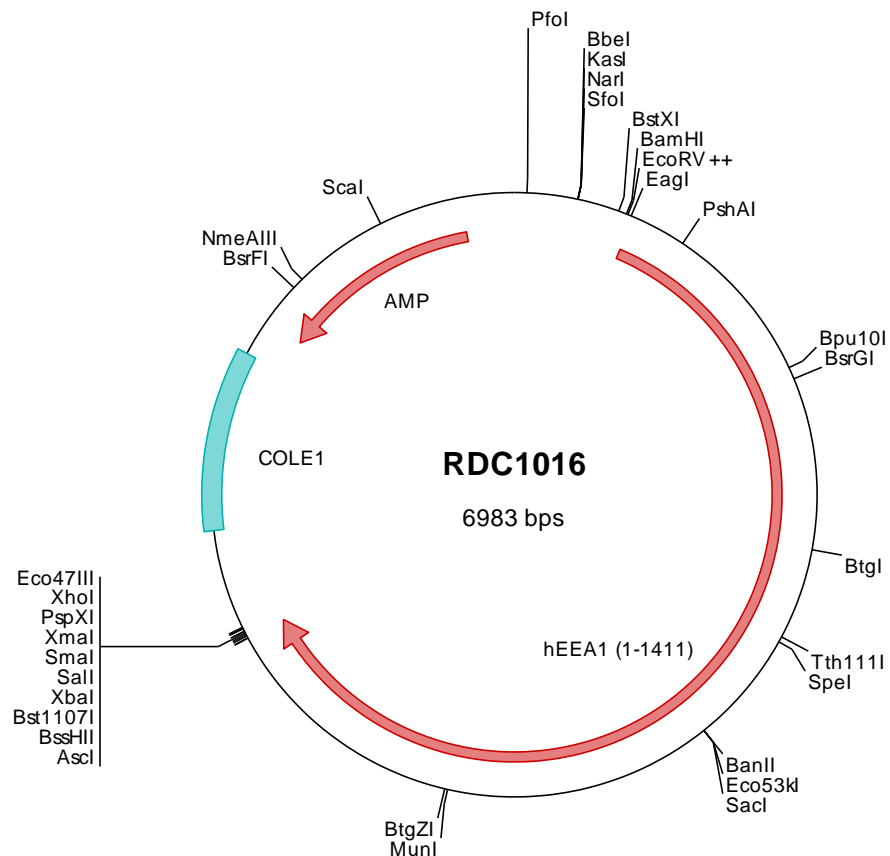
EEA1 serves as an identifying marker for early endosomes. It is ubiquitously expressed, and found in both the cytosol and on cellular membranes. Its activity has been described as that of a tethering factor which links endosomes to endocytic vesicles, allowing for their fusion via a SNARE complex. Normally, EEA1 exists as a homodimer in the cytoplasm and appears to make transient contacts with endosome membrane phosphatidylinositol.

Description

This shuttle vector contains the complete ORF for the gene of interest, along with a Kozak consensus sequence for optimal translation initiation. It is inserted NotI to AscI. The gene insert is flanked with convenient multiple cloning sites which can be used to easily cut and transfer the gene cassette into your desired expression vector.

Preparation and Storage

Formulation	cDNA is provided in 10 mM Tris-Cl, pH 8.5
Shipping	Ships at ambient temperature
Stability	1 year from date of receipt when stored at -20°C to -80°C
Storage	Use a manual defrost freezer and avoid repeated freeze-thaw cycles.



FOR RESEARCH USE ONLY

NOT FOR USE IN HUMANS



> RDC1016 Translated Insert Sequence

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1 mlrrilqrtp grvsgqgsdl dssatpintv dvnnessseg ficpqcmsl gsadelfkhy eavhdagns ghggesnlal krddvtllrq evqdlqasl  
101 eekwyseeek kelekyqglq qgeakpdglv tdssaelqsl eggleeaqte nfniqmkdl feqkaaqlat eiadikskyd eerslreaae qkvtrlteel  
201 nkeatviqdl ktellrpgi edvavlkkel vqvqtlmdnm tlerererek lkdeckklqs qyasseatis qlrselakgp qevavyvqel qklkssvnel  
301 tqknqtlten llkkeqdytk leekhneesv skkniqatlh qkdldcqqllq srlsasetsl hrihvelsek geatqklkee lsevetkyqh lkaefkqlqq  
401 qreekeghl qlqseinqlh sklleterql geahgrlkeq rqlssekimd keqqvadlql klsrleeqlk ekvtnstelq hqldktkqgh qeqqalqgst  
501 taklreaqnd leqvlrqigd kdqkiqnlea llgkskenis llekeredly akiqageget avlnqlqekn htlqeqvtql teklnqses hkqaqenlhd  
601 qvqeqkahlr aaqdrvlsl tsvnelnsql neskekvsq diqikaktel llsaeaakta qradlqnhd taqnalqdkq qelnkittql dqvtaklqdk  
701 qehcsqlesh lkeykekyls leqkteeleg qikkleadsl evkaskeqal qdlqqqrqln tdlelratel skqlemekei vsstrldlqk ksealesikq  
801 kltkqeeekq ilkqdfetls qetkiqheel nnriqttvte lgkvkmekea lmtelstvk d klskvsdsl nsksefeken qkgkaaidl ektckelkhq  
901 lqvqmentlk eqkelkksle kekeashqk lelnsmqeql iqaqntlqn ekeeqqlqgn inelkqsseq kkkqiealqg elkiavlqkt elenklqqq  
1001 tqaaqelaae kekisvlqnn yeksqetfkq lqsdfygres ellatrqdk sveeklsiaq edlismnqi gnqnkliqel ktakatleqd sakkeqqlq  
1101 rckalqdiqk ekslkekely neksklaeie eikcrqekei tklneelksh klesikeitn lkdaklliq qklelqgkad slkaaveqek rnqqilkdqv  
1201 kkeeeelkke fiekeaklhs eikekevgmk kheeneaklt mqitalnenl gtvkkewqss qrrvselekq tddlrgeiav leatvqnnqd errallercl  
1301 kgegeiekq kvlvelqrkl dnttaavqel grenqslqik htqalnrrwa ednevqncma cgkgfsvtvr rhhcrqcgni fcaecsakna ltpsskkpvr  
1401 vcdacfnldl q
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