

Specifications:

Gene:	hPRKAA1
Accession:	NP_006242
Insert size:	1693bp
Concentration:	10µg at 0.2µg/µL

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.

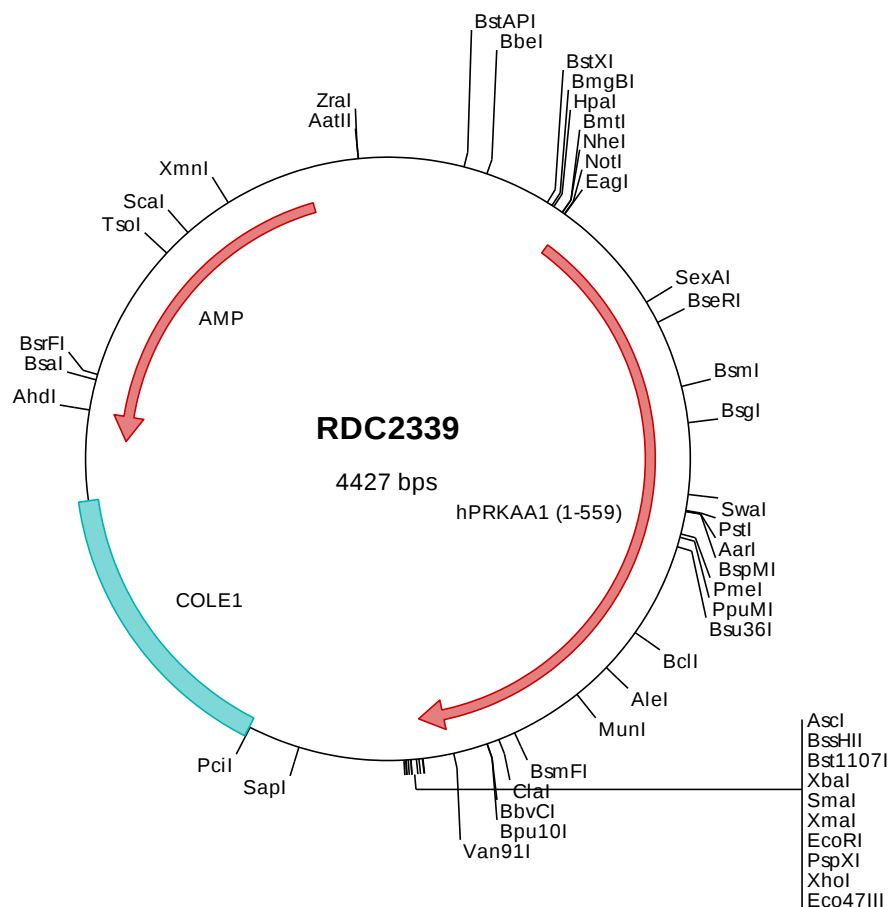
hAMPK α1 cDNA Plasmid

PRKAA1 protein kinase AMP-activated catalytic subunit alpha 1 [*Homo sapiens* (human)]

Also known as: AMPK; AMPKα1

Summary:

PRKAA1 belongs to the ser/thr protein kinase family. It is the catalytic subunit of the 5'-prime-AMP-activated protein kinase (AMPK). AMPK is a cellular energy sensor conserved in all eukaryotic cells. The kinase activity of AMPK is activated by the stimuli that increase the cellular AMP/ATP ratio. AMPK regulates the activities of many key metabolic enzymes through phosphorylation. It protects cells from stresses that cause ATP depletion by switching off ATP-consuming biosynthetic pathways. Alternatively spliced transcripts encoding different proteins have been described.



> RDC2339 Plasmid DNA Sequence

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201  ccgcacacgat gcgtaaggag aaaataccgc atcaggcgcc attcgccatt caggctgcgc aactgttggg aagggcgatc ggtgcgggcc tcttcgctat
301  tacgcccagct ggcgaaaagg ggatgtgctg caaggcgatt aagttgggta agcccgagggt tttcccagtc acgacgttgt aaaacgacgg ccagtgtaatt
401  ggagacgtgt taacaagctt ggatccgata tcgctagcgc ggccgcccacc atgcgcagac tcagttcctg gagaagatg gcgacagccg agaagcagaa
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701  atcctcatat aattaaactg taaccaggtc tcagtacacc atctgatatt ttcatgttga ttggaatatg ctacggagga gagctatttg attatatctg
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> RDC2339 Translated Insert Sequence

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201  yagpevdiws sgvilvallyc gtlpfdddhv ptlfkkicdg ifytpqylnp svslslkxhl qvdpmkrtati kdirehewfk qdlpkylfpe dpsysstmid
301  dealkevcek fecseeevls clynrnhdqp lavayhliid nrrimneakd fylatsppds flddhhltrp hpervpflva etprarthld elnpqskshq
401  gvrkakwhlg irsqsrpndi maevcraikg ldyewkvvnp yyllrvrrkn vtstyskmsl qlyqvdsrty lldfrsidde iteaksgtat pqrsgsvsny
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