

Specifications:

Gene:	hGLR
Accession:	NP_000151
Insert size:	1447bp
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.

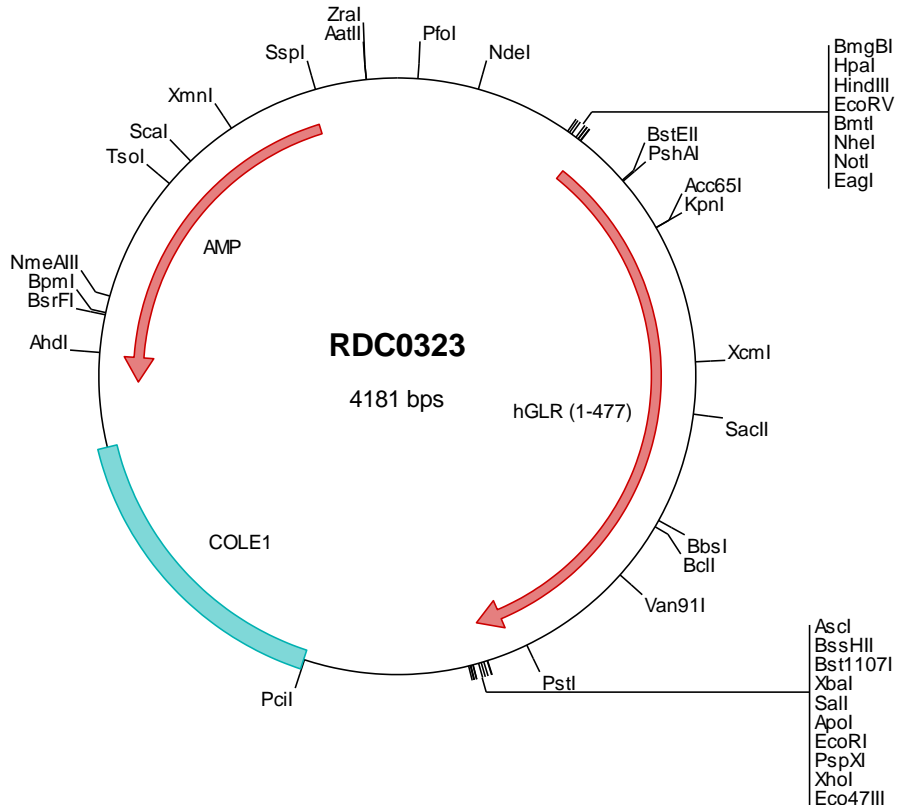
hGLR cDNA Plasmid

GCGR glucagon receptor [*Homo sapiens*]

Also known as: GGR

Summary:

GLR is a receptor for glucagon which plays a central role in regulating the level of blood glucose by controlling the rate of hepatic glucose production and insulin secretion. The activity of GLR is mediated by G proteins which activate adenylyl cyclase and also a phosphatidylinositol-calcium second messenger system. Defects in GLR are a cause of non-insulin-dependent diabetes mellitus.





> RDC0323 Plasmid DNA Sequence

1 tcgcgcggtt cggatgatgac ggtgaaaacc tetgacacat gcagctccc gagacggtca cagcttgtct gtaagcggat gccgggagca gacaagcccg
101 tcaggggcgc tcagcgggtg ttggcgggtg tetggggctg cttactatg cggcatcaga gcagattgta ctgagagtgc accatattgc gtgtgaaata
201 ccgcacagat gcgtaaggag aaaataccgc atcaggcgcc attcgccatt caggctgcgc aactgttggg aaggcgatc ggtgcgggcc tcttcgctat
301 taacgacgct ggcgaaaggg ggatgtgctg caaggcgatt aagtgggta acgcccagggt ttcccgatc acgacgtgtg aaaacgacgg ccagtgaatt
401 ggagacgtgt taacaagctt ggatccgata tcgctagcgc ggccgccacc atgccccct gccagccaca gcgacccctg ctgctgttgc tgcctgtgct
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2101 ccagtcggga aacctgtcgt gccagctgca ttaatgaatc ggccaacgcg cggggagagg cggtttgcgt atggggcct cttccgcttc ctgcctcact
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> RDC0323 Translated Insert Sequence

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101 gpdgqwrvgp rgqpwrdasq cqmdgeeiev qkevakyss fqvmytvvys lslgalllal ailgglslkh ctrnaihanl fasfvlkass vlvidgllrt
201 rysqkigddl svstwlsdga vagcrvaavf mqygivanyc wllvegilyh nllglatlpe rsffsllylgi gwgapmlfvv pwavvkclfe nvqcwtsndn
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401 cflinkevqse lrrrwhrwl gkvlweernt snhrasssp hpppskelqf grgggsqds aetplagglp rlaesp