

Specifications:

Gene:	mGPR18
Accession:	NP_877958
Insert size:	1009bp
Concentration:	10µg at 0.2µg/µL

mGPR18 cDNA Plasmid

Gpr18 G protein-coupled receptor 18 [*Mus musculus* (house mouse)]

Summary:

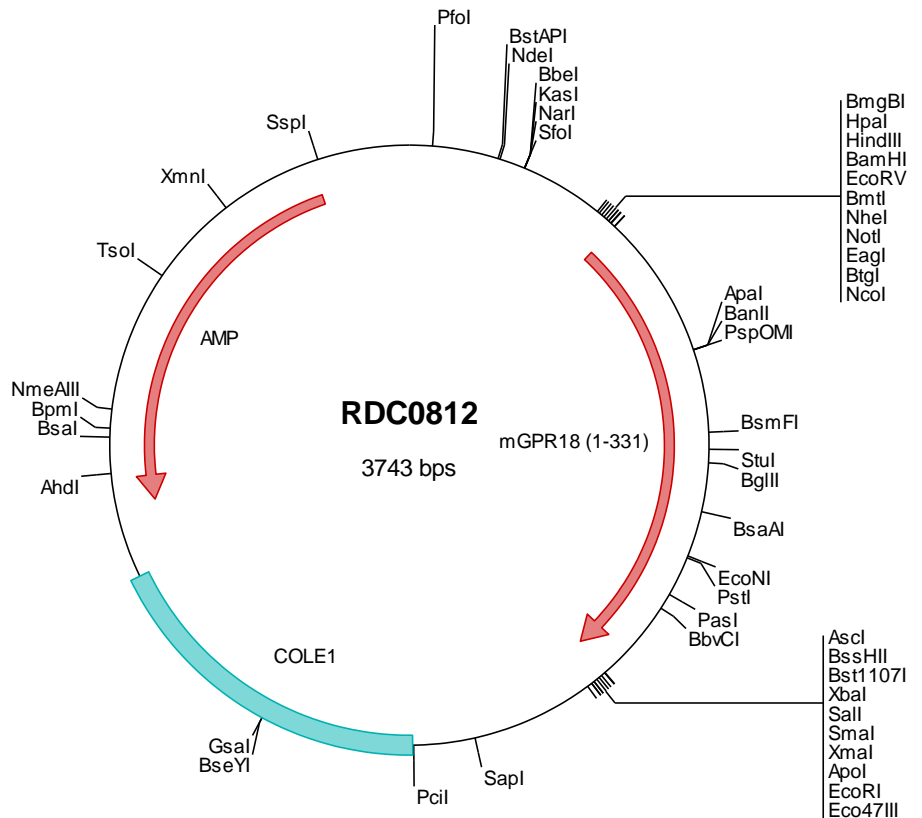
GPR18 is the receptor for N-arachidonyl glycine and belongs to the G protein-coupled receptor 1 family. The activity of GPR18 is regulated by G proteins which inhibit adenylyl cyclase and may contribute to the regulation of the immune system.

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



> RDC0812 Plasmid DNA Sequence

1 tcgcgcggtt cggatgatgac ggtgaaaacc tetgacacat gcaagctccc gagacggta cagcttgtct gtaagcggat gccgggagca gacaagcccg
101 tcaggggcgc tcagcgggtg ttggcgggtg tetggggctgg cttactatg cggcatcaga gcagattgta ctgagagtgc accatattgc gtgtgaaata
201 ccgcacagat gcgtaaggag aaaataccgc atcaggcgcc attcgccatt caggctgcgc aactgttggg aaggcgatc ggtcggggcc tcttcgctat
301 taaggcagct ggcgaaaggg ggatgtgctg caaggcgatt aagtgggta acgccagggt ttcccgatc acgacgtgtg aaaacgacgg ccagtgaatt
401 ggagacgtgt taacaagctt ggatccgata tetgtagcgc gggcggcaacc atggccacc tgagcaatca caaccagett gatctttota atggctcaaca
501 cccagaggaa tacaaaaatg cagccctagt cttctacagc tgcattctoc tgattgggtc gtttgttaat gtcactgcgt tgggggtttt cagctgtaag
601 accaagaaaa gaaccacagt gaccatctac atgatgaacg ttgcaact ggaacctgta ttatactca gtctgccctt tcggatgttt tactatgcaa
701 aaggcgagtg gccatttggg gagtaattct gccacattct tggggccctg gtgtgtttt acccaagcct cgctctgtgg cttctgtctt toattagtgc
801 tgacagatag atggccatcg tacagccaaa atatgccaag gagctgaaga acaccggcaa gcccgtgctt gcgtgtgggg gggctgggtt aatgaccctg
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1501 ctctagcttg gcgtaatcat ggtcatagct gtttctgtg tgaaattgtt atccgctcac aattccacac aacatagcag ccggaagcat aaagtgtaaa
1601 gcctggggtg cctaatgagt gagctaactc acattaattg cgttgccctc actgcccctt ttccagctcg gaaacctgtc gtgccagctg cattaatgaa
1701 tcggccaacg cgcggggaga ggcggtttgc gtattggcgc ctctccctc ctctcctca ctgactcgtc gcgctcggtc gttcggctgc ggcgagcgtg
1801 atcagctcac tcaaaagcgg taatacgggt atccacagaa tcaggggata acgcaggaaa gaactgtgta gcaaaagccc agcaaaagcc caggaaccgt
1901 aaaaagggcg cgttgcctgg gtttttccat aggtcccgcc cccctgacga gcatcaciaa aatcgacgtc caagtcaag gtggcgaaac ccgacaggac
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2201 gaccgctcgc cttatcccg taactatcgt cttgagcca acccggttaag acacgacta tcgccactgg cagcagccac tggtaacagg attagcagag
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2401 taccttcgga aaaagagttg gtatgctctg atccggcaaa caaacaccgc ctggtagcgg tggttttttt gtttcaagc agcagattac gcgcagaaaa
2501 aaaggatctc aagaagatcc tttgatcttt tctacgggtg ctgacgtca gtggaacgaa aactcacgtt aagggatttt ggtcatgaga ttatcaaaaa
2601 ggatcttccac ctatgctctt ttaaaataaa aatgaagttt taaatcaatc taaagtatat atgagtaaac ttggtctgac agttaaccaat gcttaatcag
2701 tgaggcaact atctcagcga tctgtctatt tcgttcaccc atagttgctt gactccccct cgtgtagata actacgatac gggaggcctt accatctgac
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3001 aggcacgtg gtgtcagcct cgtcgtttgg tatgcttca ttcagctccc gttcccaacg atcaaggcga gttacatgat cccccatggt gtgcaaaaaa
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3201 tgccatccgt aagatgcttt tctgtgactc gtgagtactc aaccaagtca tcttgagaat agtgtatgcg gcgaccgagt tgccttggcc cggcgtcaat
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3501 caaaaaaggg aataaggcgc acacggaaat gttgaatact catactctc cttttcaat attattgaag catttatcag gttattgtc tcgatgagcg
3601 atacatattt gaatgtattt agaaaaataa acaaataggg gttccgcgca cttttccccg aaaagtcca cctgacgtct aagaaacatc tattatcatg
3701 acattaacct ataaaaatag gcgtatcacg aggcctttc gtc

> RDC0812 Translated Insert Sequence

1 matlsnhnql dlsngshpee ykiaalvfys cifliglfvn vtalwvfcst tkkrtvtviy mmnvalldlv filslpfrmf yyakgewpfg eyfchilgal
101 vvfypslalw llafisadry maivqpkyak elkntgkavl acggvwmvl tttvplllly edpdkasspa tclksidith lkavnlnft rliffflipl
201 fimigyvvi ihslrrgts klkpkvkeks iriimtlilq vlvcvfpfhi cfavlmllqg ensyspwgaf tflmlnstc ldvlyyivis kqfqrarvis
301 mlyrnylrsv rrksvrsgsl rslsnmsem 1