

Specifications:

Gene:	hGPR18
Accession:	NP_005283
Insert size:	1009bp
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.

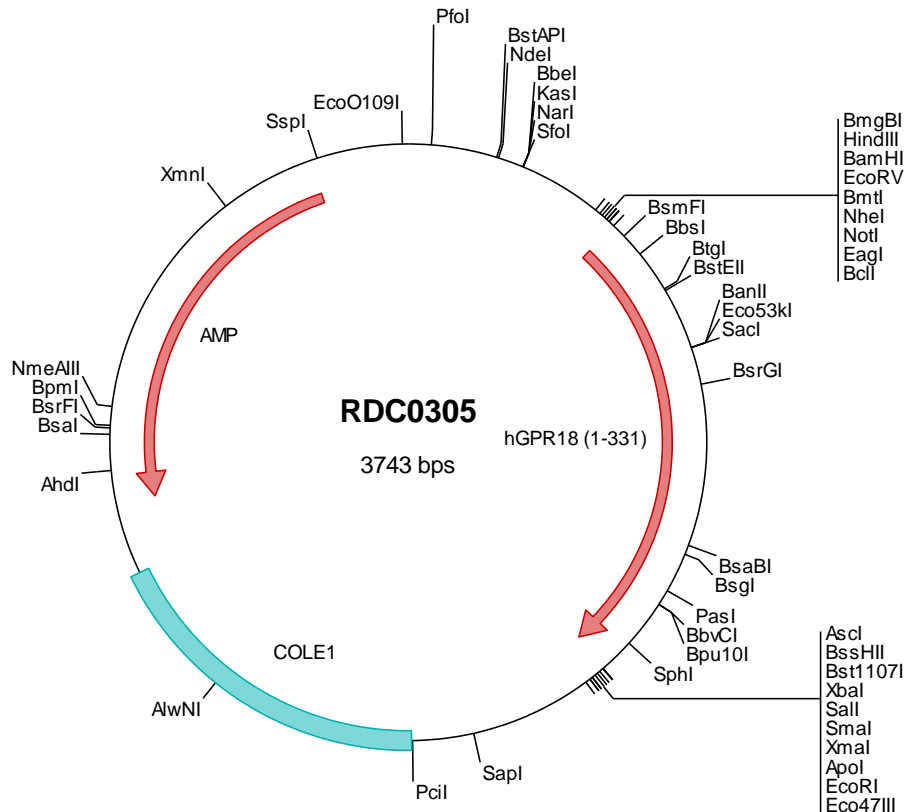
hGPR18 cDNA Plasmid

GPR18 G protein-coupled receptor 18 [*Homo sapiens*]

Also known as: N-arachidonyl glycine receptor; NAGly receptor

Summary:

GPR18, the most abundantly overexpressed orphan GPCR in all melanoma metastases, is constitutively active and inhibits apoptosis, indicating an important role for GPR18 in tumor cell survival. GPR18 is expressed significantly in lymphoid cell lines, but not in non-lymphoid hematopoietic cell lines. N-arachidonylglycine (NAGly) induces an increase in intracellular Ca(2+) concentration in GPR18-transfected cells. GPR18 may be considered a potential novel anticancer target in metastatic melanoma.



FOR RESEARCH USE ONLY

NOT FOR USE IN HUMANS



> RDC0305 Plasmid DNA Sequence

1 tcgcgcggtt cggatgatgac ggtgaaaacc tetgacacat gcaagctccc gagacggtca cagcttgtct gtaagcggat gccgggagca gacaagcccg
101 tcaggggcgc tcagcgggtg ttggcgggtg teggggctgg cttactatg cggcatcaga gcagattgta ctgagagtgc accatattgc ggtgaaata
201 ccgcacagat gcgtaaggag aaaataccgc atcaggcgcc attcgccatt caggctgcgc aactgttggg aaggcgatc ggtcgggccc tcttcgctat
301 taaggcagct ggcgaaaggg ggatgtgctg caaggcgatt aagtgggta acgccagggt ttcccgatc acgacgttg aaaacgacg ccagtgaatt
401 ggagacgtgt taacaagctt ggatccgata tcgctagcgc gggcggcacc atgataccgc tgaacaatca agatcaaacgt gtcctcttta acagctcaaca
501 tccagatgaa tacaaaaattg cagcccttgt cttctatagc tgtaattca taattggatt atttgttaac atcaactgcat tatgggtttt cagttgtacc
601 accaagaaga gaaccacggt aacctatct atgatgaatg tggcattagt ggaactgata ttataatga ctttaccctt tcgaatgttt tattatgcaa
701 aagatgaatg gccatttggg gagtaactct gccagattct tggagctctc acaggttttt acccaagcat tgctttatgg cttcttgcct ttattagtgc
801 tgacagatag atggccattg tacagcggaa gtacgcaaaa gaacttaaaa acacgtgcaa agccgtgctg gcgtgtgtgg gagtotggat aatgaccctg
901 accacgacca cccctctgct actgctctat aaagaccag ataaagactc cactcccgcc acctgctca agatttctga catcatctat ctaaaagctg
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2501 aaagatctc aagaagatcc tttgatcttt tctacggggt ctgacgtca gtggaacgaa aactcacgtt aagggatttt ggtcatgaga ttatcaaaaa
2601 ggatcttccac ctgactcctt ttaaaataaa aatgaagttt taaatcaatc taaagtatat atgagtaaac ttggtctgac agttaaccaat gcttaatcag
2701 tgaggcaact atctcagcga tctgtctatt tctgtcctcc tctgtcctcc atagttgctt gactccccgt cgtgtagata actacgatac gggaggcctt accatctgac
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3501 caaaaaagg aataaggcg acacggaaat gttgaatact catactcttc cttttcaat atttatgaag catttatcag gttattgtc tcatgagcgg
3601 atacatattt gaatgtattt agaaaaataa acaaatagg gttccgcgca catttccccg aaaagtcca cctgacgtct aagaaacat tattatcatg
3701 acattaacct ataaaaatag gcgtatcacg aggcctttc gtc

> RDC0305 Translated Insert Sequence

1 mitlnngdqp vpfnsshpde ykiaalvfys cifiiglfvn italwvfcst tkkrtvtviy mmnvalvqli fimtlpfrmf yyakdewpfg eyfcqilgal
101 tvfypsialw llafisadry maivqpkyak elkntckavl acvgvwimtl ttttplllly kdpdkdstpa tolkisdiy lkavnvlnlt rltffflipl
201 fimigcyulvi ihnllhgrts klkpkkvkeas iriitllvg vlvcfmpfhi cfafmlgtg ensynpwgaf ttflmlnstc ldvilyyivs kqfqravis
301 mlyrnylrsm rksfrsrgsl rslsninsem l