

## Specifications:

Gene:	<i>hFZD6</i>
Accession:	AAD41637
Insert size:	2134bp
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.

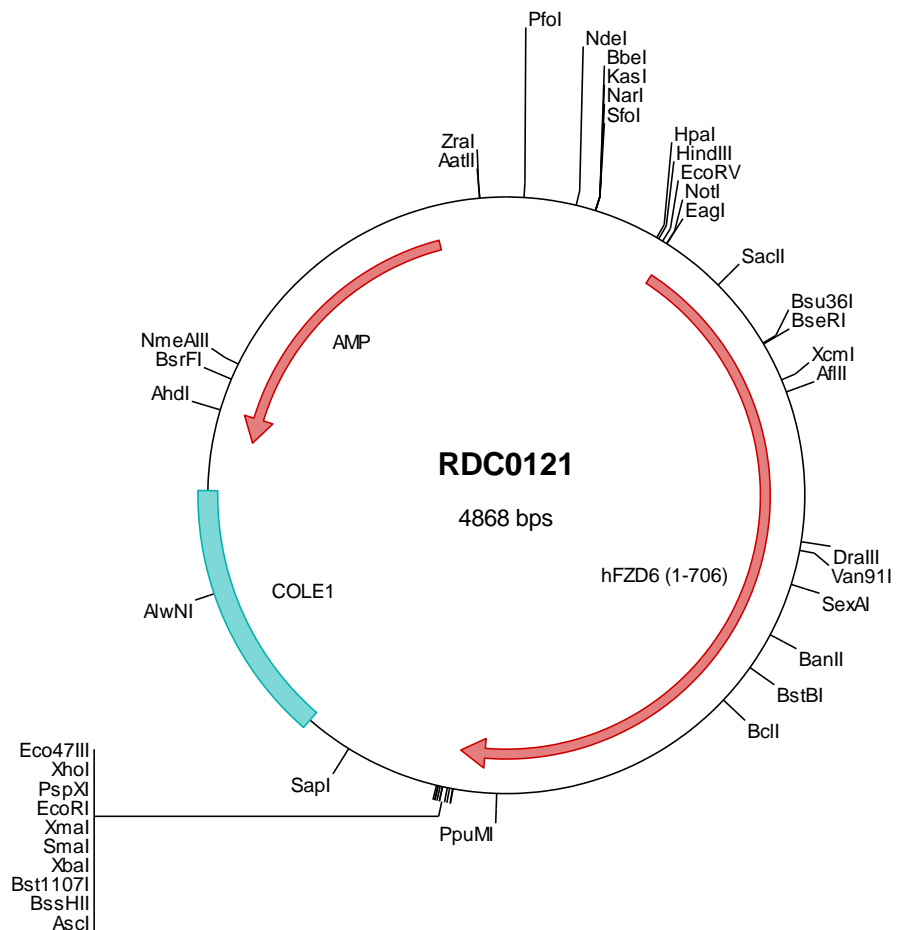
## hFrizzled-6 cDNA Plasmid

**FZD6 frizzled family receptor 6 [ *Homo sapiens* ]**

**Also known as:** FZ6; FZ-6; HFZ6; NDNC10

### Summary:

FZD6 is a 7-transmembrane glycoprotein of the Frizzled family within the G protein-coupled receptor superfamily. FZD6, unlike other family members, does not contain a C-terminal PDZ domain-binding motif. It functions as a negative regulator of the canonical Wnt/beta-catenin signaling cascade, thereby inhibiting the processes that trigger oncogenic transformation, cell proliferation, and inhibition of apoptosis. Activation of Wnt4/FZD6 signaling through a "beta-catenin-independent" pathway plays a role in proliferation and survival of the pituitary adenoma cells.



FOR RESEARCH USE ONLY

NOT FOR USE IN HUMANS



> RDC0121 Plasmid DNA Sequence

1 tcgcgcggtt cggatgatgac ggtgaaaacc tetgacacat gcaagctccc gagacggtea cagcttgtct gtaagcggat gccgggagca gacaagcccg
101 tcaggggcgc tcagcgggtg ttggcgggtg tccgggctgg cttactatg cggcatcaga gcagattgta ctgagagtgc accatatgcg gtgtgaaata
201 ccgcacagat gcgtaaggag aaaataccgc atcaggcgcc attcgccatt caggctgcgc aactgttggg aaggcgatc ggtcggggcc tcttcgctat
301 taacggcagct ggcgaaaagg ggatgtgctg caaggcgatt aagtgggta acggcagggt ttcccgatc acgacgtgtg aaaacgacgg ccagtgaatt
401 ggagacgtgt taacaagcct ggatccgata tccgtagcgc gggcggcaac atggagagt ttacattttt gttgacgtgt atttttctac ccctcctaag
501 agggcacagt ctctcaacct gtgaaccaat taactgtccc agatgtatga aatggccta caacatgacg tttttcccta atctgatggg tcattatgac
601 cagagtattg ccgcggtgga aatggagcat tttctctccc togoaaatct ggaatgtca ccaaacattg aaactttcct ctgcaaaaga tttgtaccaa
701 cctgcataga acaaatcatg gtggttccac cttgtcgtaa actttgtgag aaagatatt ctgattgcaa aaaattaatt gacacttttg ggtaccgatg
801 gcoctgaggag cttgaaatgtg acagattaca atactgtgat gagactgttc ctgtaacttt tgatccacac acagaatttc ttggtctcca gaagaaaaca
901 gaacaagtcc aaagagacat tggattttgg tgtccaaggc atcttaagac tctcggggga caaggatata agtttctggg aattgaacag tgtgcgctc
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1101 ccttaactttt ttaattgtac ttagaagatt cagatacca gaagagccaa ttatatatta ctctgtctgt tacagcattg tatctcttat gtaacttoatt
1201 ggatttttgc tgggocgatg cacagcctgc aataaggcag atgagaagct agaacttggg gacactgttg tctaggctc tcaaaaataag gcttgcaacc
1301 ttttgttcat gcttttgtat tttttcacia ttgctggcac tgtgtgtgg gtgattctta ccattacttg gttcttagct gcaggaagaa aatggagttg
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2001 tgaagctcga agatgactac aggaatcact tgagtttttc ttaagcaaca attcctaagt taacacaaa aagaagcact ataaaccaag ttaacacaag
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2601 cccggggaat tccctogagcg ctctctctca gcttggcgta atcatggtca tagctgtttc ctgtgtgaaa ttgttatccg ctcacaattc cacacaacat
2701 acgagccgga agcataaagt gtaaacgctg ggtgctctaa tgatgtagct aactcacatt aattgcgttg cgctcaactgc ccgctttcca gtcgggaaac
2801 ctgtcgtgcc agctgcatta atgaatcggc caacgcggcg ggagagcggg ttgtgctatt gggcgtctt ccgcttctc gctcaactgac tctgtcgctt
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3801 ctgacagtta ccaatgctta atcagtgagg cacctatctc agcgatctgt ctatttctgt catccatagt tgcctgactc cccgtctgt agataactc
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4001 agggccgagc gcagaagtgg tctgcaact ttatccgctt ccatccagtc tattaattgt tgccgggaa ctagagtag tagttcgcca gttaatagtt
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4801 cgtctaaaga accattatta tcatgacatt aacctataaa aataggcgta tcacagggcc ctttctgct

> RDC0121 Translated Insert Sequence

1 memftlltc iflpllrghs lftcepitvp rcmkmaynmt ffpnlmghyd qsiaavemeh flplanlecs pnietflcka fvptcieqih vppcrklce
101 kvysdckkli dtfgirwpee lecdrlgycd etvpvtfdfh teflpgqkkt eqvqrdigfw cprhltksgg qykflgidq cappcpnyf ksdelefaks
201 figtvsifcl catlftfltf lidvrrfryp erpiiyysvc ysivslmyfi gflldgstac nkadeklelg dtvvlgsqnk actvlfmlly fftmagtvww
301 viltitwfla agrkwsceai eqkavwfhav awgtpgfltv mllalnkvveg dnisgvcfvg lyldasryf vllplclcvf vglslllagi islnhvrqvi
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501 tewagffkn rkrdpiserv rvlqseceff lkhnskvkhh kkhykpsshk lkvisksmgt stgatanhgt savaitshdy lggetlleiq tspetsmrev
601 kadgastprl reqdcgepas paasisrlsg eqvdgkqqag svsesarseg rispkdtdt tglqsnllq vpssepsl1 kgstslslvhp vsqvrkeqgg
701 gchsd