*m*GDF-8/Myostatin VersaClone cDNA

Catalog Number: RDC3578

Specifications:

Gene:	*m*Mstn
Accession:	NP_034964.1
Insert size:	1144bp
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

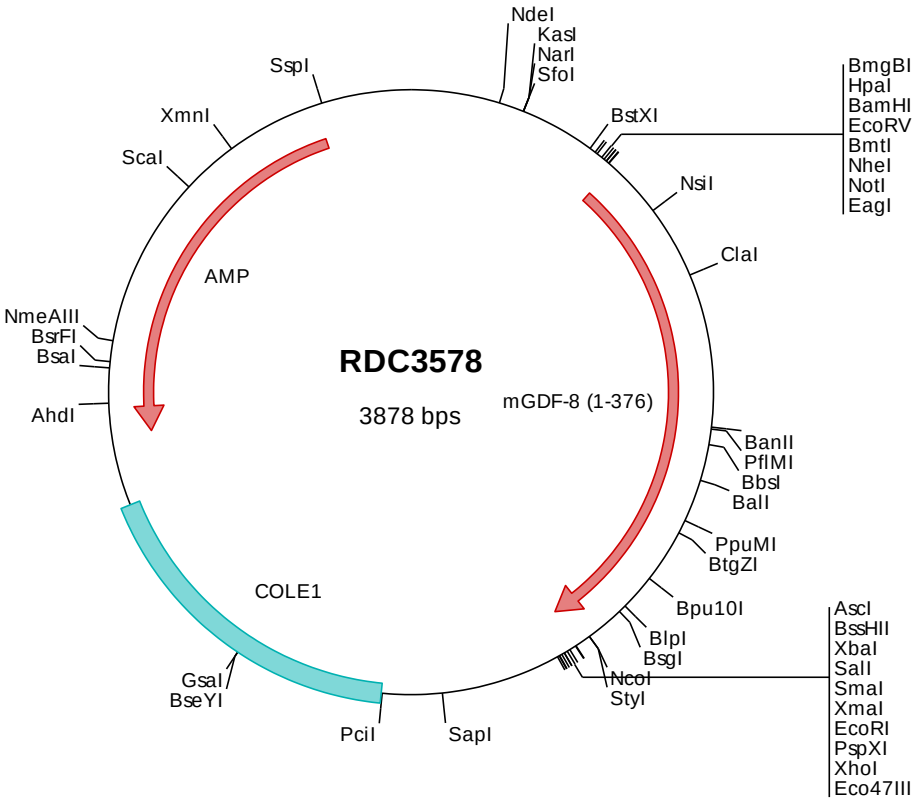
Formulation	cDNA is provided in 10 mM Tris-Cl, pH 8.5
Shipping	Ships at ambient temperature
Stability	1 year from date of receipt when stored at -20°C to -80°C
Storage	Use a manual defrost freezer and avoid repeated freeze-thaw cycles.

*m*GDF-8/Myostatin cDNA Plasmid

Mstn myostatin [*Mus musculus* (house mouse)]

Also known as: Cmpt; Gdf8

Summary:
GDF8 is a secreted ligand of the transforming growth factor-beta (TGF-beta) superfamily of proteins. Ligands of this family bind various TGF-beta receptors leading to recruitment and activation of SMAD family transcription factors that regulate gene expression. GDF8 is proteolytically processed to generate each subunit of the disulfide-linked homodimer. It negatively regulates skeletal muscle cell proliferation and differentiation. Mutations in GDF8 are associated with increased skeletal muscle mass in humans and other mammals.



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NOT FOR USE IN HUMANS

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R&D SYSTEMS

> RDC3578 Plasmid DNA Sequence

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 101  tcagggcgcg tcagcgggtg ttggcgggtg tcggggctgg cttaactatg cggcatcaga gcagattgta ctgagagtgc accatatgcg gtgtgaaata
 201  ccgcacagat gcgtaaggag aaaataccgc atcaggcgcc attcgccatt caggctgcgc aactgttggg aagggcgatc ggtgcgggcc tcttcgctat
 301  tacgccagct ggcgaaaggg ggatgtgctg caaggcgatt aagttgggta acgccagggt tttcccagtc acgacgttgt aaaacgacgg ccagtgaatt
 401  ggagacgtgt taacaagctt ggatccgata tcgctagcgc ggccgccacc atgatgcaaa aactgcaaat gtatgtttat atttacctgt tcatgctgat
 501  tgctgctggc ccagtggatc taaatgaggg cagtgagaga gaagaaaatg tggaaaaaga ggggctgtgt aatgcatgtg cgtggagaca aaacacgagg
 601  tactccagaa tagaagccat aaaaattcaa atcctcagta agctgcgcct ggaaacagct cctaacatca gcaaagatgc tataagacaa cttctgccaa
 701  gagcgcctcc actccgggaa ctgatcgatc agtacgacgt ccagagggat gacagcagtg atggctcttt ggaagatgac gattatcacg ctaccacgga
 801  aacaatcatt accatgccta cagagtctga ctttctaatg caagcggatg gcaagcccaa atgttgcttt tttaaattta gctctaaaat acagtacaac
 901  aaagtagtaa aagcccaact gtggatatat ctcagacccg tcaagactcc tacaacagtg tttgtgcaaa tcctgagact catcaaaccc atgaaagacg
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1501  atgctatatt ttaatggcaa agaacaaata atatatggga aaattccagc catggtagta gaccgctgtg ggtgctcata aaggcgcgcc agtatactct
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1701  cacacaacat acgagccgga agcataaagt gtaaagcctg gggtgcctaa tgagtgagct aactcacatt aattgcgttg cgctcactgc ccgctttcca
1801  gtcgggaaac ctgtcgtgcc agctgcatta atgaatcggc caacgcgcgg ggagaggcgg tttgcgtatt gggcgctctt ccgcttcctc gctcactgac
1901  tcgctgcgct cggtcgttcg gctgcggcga gcggtatcag ctcactcaaa ggcggtaata cggttatcca cagaatcagg ggataacgca ggaaagaaca
2001  tgtgagcaaa aggccagcaa aaggccagga accgtaaaaa ggccgcgttg ctggcgtttt tccataggct ccgccccct gacgagcatc acaaaaatcg
2101  acgctcaagt cagaggtggc gaaacccgac aggactataa agataccagg cgtttccccc tggaagctcc ctcgtgcgct ctcctgttcc gaccctgccg
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2801  taaacttggt ctgacagtta ccaatgctta atcagtgagg cacctatctc agcgatctgt ctatttcgtt catccatagt tgcctgactc cccgtcgtgt
2901  agataactac gatacgggag ggcttaccat ctggccccag tgctgcaatg ataccgcgag acccacgctc accggctcca gatttatcag caataaacca
3001  gccagccgga agggccgagc gcagaagtgg tcctgcaact ttatccgcct ccatccagtc tattaattgt tgccgggaag ctagagtaag tagttcgcca
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3601  ttctgggtga gcaaaaacag gaaggcaaaa tgccgcaaaa aagggaataa gggcgacacg gaaatgttga atactcatac tcttcctttt tcaatattat
3701  tgaagcattt atcagggtta ttgtctcatg agcggataca tatttgaatg tatttagaaa aataaacaaa taggggttcc gcgcacattt ccccgaaaag
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> RDC3578 Translated Insert Sequence

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 101  dssdgsledd dyhattetii tmptesdflm qadgkpkccf fkfsskiqyn kvvkaqlwiy lrpvktpttv fvqilrlikp mkdgtrytgi rslkldmspg
 201  tgiwqsidvk tvlqnwlkqp esnlgieika ldenghdlav tfpgpgedgl npflevkvtd tpkrsrrdfg ldcdehstes rccrypltvd feafgwdwii
 301  apkrykanyc sgecefvflq kyphthlvhq anprgsagpc ctptkmspin mlyfngkeqi iygkipamvv drcgcs
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