

Product datasheet for **RG216154**

Collagen VI (COL6A3) (NM_057165) Human Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Collagen VI (COL6A3) (NM_057165) Human Tagged ORF Clone
Tag:	TurboGFP
Symbol:	COL6A3
Synonyms:	BTHLM1; DYT27; UCMD1
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-AC-GFP (PS100010)
E. coli Selection:	Ampicillin (100 ug/mL)
ORF Nucleotide Sequence:	>RG216154 representing NM_057165 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGATCGCC**

ATGAGGAAACATCGGCACTTGCCTTAGTGCCGCTCTTTGCCTCTTCTCTCAGGCTTTCCTACAACCTC
ATGCCCAGCAGCAGCAAGCAGCACAAGACTCTGCTGACATTATTTTCCTTATTGATGGATCAAACAACAC
CGGAAGTGTCAATTCGCAGTCATTCTCGACTTCTTGTAAATCTCCTTGAGAACTCCCAATTGGAACCT
CAGCAGATCCGAGTGGGGTGGTCCAGTTAGCGATGAGCCAGAACCATGTTCTCCTTGGACACCTACT
CCACCAAGGCCAGGTTCTGGGTGCAGTAAAGCCCTCGGTTTGTGGTGGGGAGTTGGCCAATATCGG
CCTCGCCCTTGATTTCTGTTGGGAGAACCACTTACCCGGGCAGGGGGCAGCCCGCTGGAGGAAGGGTT
CCCCAGGTGCTGGTCTCATAAGTGCCGGGCTTCTAGTGACGAGATTGCTACGGGGTGGTAGCACTGA
AGCAGGCTAGCGTGTCTCATTGCGCCTTGAGGCCAGGCCGCTCCAGGGCAGAGCTTCAGCACATAGC
TACCGATGACAACCTTGGTGTACTGTCCCGGAATTCGTAGCTTTGGGGACCTCCAGGAGAAAATTACTG
CCGTACATTGTTGGCGTGGCCAAAGGCACATTGTCTTGAACCAGCAACATTGTCACACAAGTCATTG
AAGTCAACAAGAGAGACATAGTCTTCTGGTGGATGGCTCATCTGCACTGGGACTGGCAACTTCAATGC
CATCCGAGACTTCATTGCTAAAGTCATCCAGAGGCTGAAATCGGACAGGATCTTATCCAGTGGCAGTG
GCCAGTATGCAGACACTGTGAGGCCTGAATTTATTTCAATCCCAACAAAAGGGAGTGCATAA
CCGCTGTGCGGAAAATGAAGCCCTGGACGGCTCGGCCCTGTACACGGGCTCTGCTCTAGACTTTGTTCG
TAACAACCTATTACGAGTTCAGCCGGCTACCGGGCTGCCGAGGGGATTCTAAGCTTTTGGTGTGATC
ACAGGTGGTAAGTCCCTAGATGAAATCAGCCAGCCTGCCAGGAGCTGAAGAGAAGCAGCATAATGGCCT
TTGCCATTGGGAACAAGGGTGGCGATCAGGCTGAGCTGGAAGAGATCGCTTTCGACTCCTCCCTGGTGT
CATCCAGCTGAGTTCGAGCCGCCATTGCAAGGCATGCTGCCTGGCTTGTGGCACCTCTCAGGACC
CTCTCTGGAACCCCTGAAGTTCCTCAACAAAAGGGATATCATCTTTCTTTGGATGGATCAGCCAACG
TTGAAAAACCAATTTCCCTTATGTGCGGACTTTGTAATGAACCTAGTTAACAGCCTTGATATTGAAA
TGACAATATTCGTGTTGGTTTAGTGCAATTTAGTGACACTCTGTAACGGAGTTCTTTAAACACATAC



[View online »](#)

CAGACCAAGTCAGATATCCTTGGTCATCTGAGGCAGCTGCAGCTCCAGGGAGGTTTCGGCCCTGAACACAG
GCTCAGCCCTAAGCTATGTCTATGCCAACCACTTCACGGAAGCTGGCGGCAGCAGGATCCGTGAACACGT
GCCGCAGCTCCTGCTTCTGCTCAGAGCTGGGCAGTCTGAGGACTCCTATTTGCAAGCTGCCAACGCCTTG
ACACGCGCGGGCATCCTGACTTTTTGTGTGGGAGCTAGCCAGGCGAATAAGGCAGAGCTTGAGCAGATTG
CTTTTAACCCAAGCCTGGTGTATCTCATGGATGATTTAGCTCCCTGCCAGCTTTGCCTCAGCAGCTGAT
TCAGCCCTAACACATATGTTAGTGGAGGTGTGGAGGAAGTACCACTCGCTCAGCCAGAGAGCAAGCGA
GACATTCGTTCCTCTTTGACGGCTCAGCCAATCTTGTGGGCCAGTTCCTGTTGTCCGTGACTTTCTCT
ACAAGATTATCGATGAGCTCAATGTGAAGCCAGAGGGGACCCGAATTGCGGTGGCTCAGTACAGCGATGA
TGTC AAGGTGGAGTCCCCTTTTGTATGAGCACCAGAGTAAGCCTGAGATCCTGAATCTTGTGAAGAGAATG
AAGATCAAGACGGGCAAAGCCCTCAACCTGGGCTACGCGCTGGACTATGCACAGAGGTACATTTTTGTGA
AGTCTGCTGGCAGCCGGATCGAGGATGGAGTGCTTCAAGTTCCTGGTGTGCTGGTGCAGGAAGGTCATC
TGACCGTGTGGATGGGCCAGCAAGTAACCTGAAGCAGAGTGGGTTGTGCCTTTCATCTTCAAGCCAAG
AACGCAGACCCTGCTGAGTTAGAGCAGATCGTGTGTCTCCAGCGTTTATCCTGGCTGCAGAGTCGCTTC
CCAAGATTGGAGATCTTCATCCACAGATAGTGAATCTCTTAAAAATCAGTGCACAACGGAGCACCAGCACC
AGTTTCAGGTGAAAAGGACGTGGTGTCTGCTTGTGATGGCTCTGAGGGCGTCAGGAGCGGCTTCCCTCTG
TTGAAAGAGTTTTGTCAGAGAGTGGTGGAAAGCCTGGATGTGGGCCAGGACCGGGTCCGCGTGGCCGTGG
TGCACTACAGCGACCCGACCGAGCCGAGTTCACCTGAATTCATACATGAACAAGCAGGACGTCGTC A A
CGCTGTCCGCCAGCTGACCCTGCTGGGAGGGCCGACCCCAACACCGGGGCCGCCCTGGAGTTTGTCCCTG
AGGAACATCCTGGTCAGCTCTGCGGGAAGCAGGATAACAGAAGGTGTGCCCCAGCTGCTGATCGTCTCA
CGGCCGACAGGCTCGGGGATGATGTGCGGAACCCCTCCGTGGTGTGAAGAGGGGTGGGGCTGTGCCAT
TGGCATTGGCATCGGGAACGCTGACATCACAGAGATGCAGACCATCTCCTTCATCCCGGACTTTGCCGTG
GCCATTCACCTTTCCGACGCTGGGGACCGTCCAACAGGTCACTCTGAGAGGGTGACCCAGCTCACCC
GCGAGGAGCTGAGCAGGCTGCAGCCGGTGTGACGCCCTTACCGAGCCAGGTGTTGGTGGCAAGAGGGA
CGTGGTCTTTCTCATCGATGGGTCCCAAAGTGCCGGGCCCTGAGTTCCAGTACGTTCCGACCCCTCATAGAG
AGGCTGGTTGACTACCTGGACGTGGGCTTTGACACCACCCGGGTGGCTGTATCCAGTTCAGCGATGACC
CCAAGGTGGAGTTCCTGCTGAACGCCATTCCAGCAAGGATGAAGTGCAGAACGCGGTGCAGCGGCTGAG
GCCAAGGGAGGGCGGCAGATCAACGTGGGCAATGCCCTGGAGTACGTGTCCAGGAACATCTTCAAGAGG
CCCCTGGGAGCCGATTGAAGAGGGCGTCCCGCAGTTCCTGGTCTCATCTCGTCTGGAAGTCTGACG
ATGAGGTGGACGACCCGGCGGTGGAGCTCAAGCAGTTTGGCGTGGCCCTTTCACGATCGCCAGGAACGC
AGACCAGGAGGAGCTGGTGAAGATCTCGCTGAGCCCCGAATATGTGTTCTCGGTGAGCACCTTCCGGGAG
CTGCCAGCCTGGAGCAGAACTGCTGACGCCATCACGACCCTGACCTCAGAGCAGATCCAGAAGCTCT
TAGCCAGCACTCGCTATCCACCTCCAGGTGAGATGGGGGCGTCGGAGTTCTCCTTGAGCATTTTCCAT
A

ACGCGTACGCGGCCGCTCGAG – GFP Tag – GTTTAA

Protein Sequence: >RG216154 representing NM_057165
 Red=Cloning site Green=Tags(s)

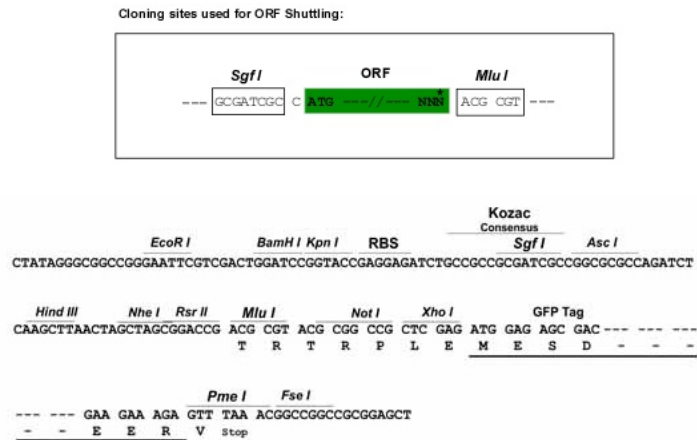
MRKHRHLPLVAVFCLFLSGFPTTHAQQQQAQDSADIIFLIDGSNNTGSVNFVAVILDFLVNLEKLPIGT
 QQIRVGVVQFSDEPRTMFLSDTYSTKAQVLGAVKALGFAGGELANIGLALDFVVENHFTRAGGSRVEEGV
 PQVLVLISAGPSSDEIRYGVVALKQASVFSFGLGAQAASRAELQHIATDDNLVFTVPEFRSFGDLQEKLL
 PYIVGVAQRHIVLKPPTIVTQVIEVNRDIVFLVDGSSALGLANFNAIRDFAKVIQRLEIGQDLIQVAV
 AQYADTVRPEFYFNHTPKREVITAVRKMPLDGSALYTGSSALDFVRNMLFTSSAGYRAAEGIPKLLVLI
 TGGKSLDEISQPAQELKRSSIMAFIAGNKGADQAELEEIAFDSSLVFIPAEFRAAPLQGMLPGLLAPLRT
 LSGTPEVHSNKRDIIFLLDGSANVGKTNFPYVRDFVMNLVNSLDIGNDNIRVGLVQFSDTPVTEFSLNTY
 QTKSDILGHLRQLQLQGGSGLNTGSALSYVYANHFTEAGGSRIREHVPQLLLLLTAGQSEDSYLAQANAL
 TRAGILTCV GASQANKAELEQIAFNPSLVYLMDDFSSLPALPQQLIQPLTTYVSGGVVEEPLAQPESKR
 DILFLFDGSANLVGQFPVVRDFLYKIIDELNVKPEGTRIAVAQYSDDVKVESRFDEHQSKPEILNLVKRM
 KIKTGKALNLGYALDYAQRVIFVKSAGSRIEDGVLQFLVLLVAGRSSDRVDGPASNLKQSGVVPFIFQAK
 NADPAELEQIVLSPAFILAAESLPKIGDLHPQIVNLLKSVHNGAPAPVSGEKDVFLLDGSEGVRS GFPL
 LKEFVQRVVESLDVGDQDRVRVAVVQYSDRTRPEFYLN SYMNKQDVVNAVRLTLLGGPTNTGAALFVFL
 RNILVSSAGSRITGVPLLIVLTADRSGDDVRNPSVVVKRGGAVPIGIGIGNADITEMQTSIFIPDFAV
 AIPTFRQLGTVQQVISERVTLTREELSRLQPVLPSPGVGGKRDDVFLIDGSQSAGPEFYVVRTLIE
 RLVDYLDVGFDTTRVAVIQFSDDPKVEFLLNAHSSKDEVQNAVQRLRPKGGRRQINVGNALEYVSRNIFKR
 PLGSRIEEGVPQFLVLISGKSDDEVDDPAVELKQFGVAPFTIARNADQEELVKISLSPEYVFSVSTFRE
 LPSLEQKLLTPITTLTSEQIQKLLASTRYPPPGE MGASEVLLGAFSI

TRTRPLE - GFP Tag - V

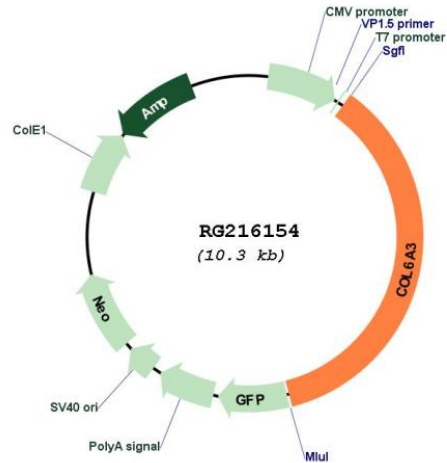
Restriction Sites:

SgfI-MluI

Cloning Scheme:



Plasmid Map:



ACCN: NM_057165

ORF Size: 3711 bp

OTI Disclaimer: The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. [More info](#)

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method:

1. Centrifuge at 5,000xg for 5min.
2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
3. Close the tube and incubate for 10 minutes at room temperature.
4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

RefSeq: [NM_057165.5](#)

RefSeq Size: 4088 bp

RefSeq ORF: 3714 bp

Locus ID: 1293

UniProt ID: [P12111](#)

Cytogenetics:	2q37.3
Protein Families:	Druggable Genome
Protein Pathways:	ECM-receptor interaction, Focal adhesion
Gene Summary:	<p>This gene encodes the alpha-3 chain, one of the three alpha chains of type VI collagen, a beaded filament collagen found in most connective tissues. The alpha-3 chain of type VI collagen is much larger than the alpha-1 and -2 chains. This difference in size is largely due to an increase in the number of subdomains, similar to von Willebrand Factor type A domains, that are found in the amino terminal globular domain of all the alpha chains. These domains have been shown to bind extracellular matrix proteins, an interaction that explains the importance of this collagen in organizing matrix components. Mutations in the type VI collagen genes are associated with Bethlem myopathy, a rare autosomal dominant proximal myopathy with early childhood onset. Mutations in this gene are also a cause of Ullrich congenital muscular dystrophy, also referred to as Ullrich scleroatonic muscular dystrophy, an autosomal recessive congenital myopathy that is more severe than Bethlem myopathy. Multiple transcript variants have been identified, but the full-length nature of only some of these variants has been described. [provided by RefSeq, Jun 2009]</p>