

## Product datasheet for **RG235279**

### DRIP130 (MED23) (NM\_001270521) Human Tagged ORF Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	DRIP130 (MED23) (NM_001270521) Human Tagged ORF Clone
Tag:	TurboGFP
Symbol:	MED23
Synonyms:	ARC130; CRSP3; CRSP130; CRSP133; DRIP130; MRT18; SUR-2; SUR2
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-AC-GFP (PS100010)
E. coli Selection:	Ampicillin (100 ug/mL)
ORF Nucleotide Sequence:	>RG235279 representing NM_001270521 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCC**CGATCGCC**

ATGGAGACGCAACTGCAGAGCATTTTCGAAGAGGTGGTAAAACGGAAGTTATAGAAGAGGCTTTTCCTG  
GCATGTTTATGGATACTCCTGAAGATGAGAAAACAAAATAATTAGCTGTTTGGGGCCCTCAGACAGTT  
TTGGGGTGGTCTTTCTCAGGAGTCTCATGAACAGTGTATCCAGTGGATTGTTAAGTTTATCATGGTCAG  
CATAGTCTAAAAGAATTTCTTTCTTTATGACTGCTTAGCAATGGCAGTTGAGACTGGTCTCCTCCAC  
CCAGGCTGGTTTGTGAATCCCTGATAAACTCTGACTCTTGAGTGGAAAGAACACAGCTTTGGGCCTT  
AACATTTAACTGGTTCGAAAATAATTGGGGGAGTGGATTACAAGGGTGTTCGAGATCTCTAAAAGTG  
ATTTTGGAGAAGATTTTGACAATTCCTAATACAGTGAGCTCTGCTGTTGTACAGCAGCTTCTGGCAGCAA  
GAGAGGTTATAGCATATATCTTGAAAGAAATGCCTGCTTATTACCAGCCTATTTTGCAGTCACTGAGAT  
CAGGAACTGTATCCTGAAGGCAAACCTCCACTGGTACTTGGAAACCTAGTATCAGACTTTGTGGAT  
ACCTTCAGGCCACAGCAAGGATAAACTCCATTTGTGGTCGCTGTAGTCTTCTGCCAGTTGTAATAATT  
CGGGTCCATTTGTAATTCATGGAACTGGATCCTGCTACTCTTCGTTTTCTTTGAAAGCCTTTTGCC  
ATATGATAAGGATCTGTTTGAACCACAGACTGCTTTGTTGAGATATGTATTGGAGCAGCCTTATCCAGG  
GATATGGTCTGCAATATGCTAGGTTTAAATAAGCAGCACAAGCAGCGCTGCCCTGTGCTGGAGGACCAGT  
TGGTGGATCTGGTTGTTTATGCCATGGAGCGATCTGAGACCGAGGAGAAGTTTGACGATGGGGAAACAAG  
CCAACCTCTGTGGCAGCATCTCTCAAGTCAAGTCAATTTCTTTGTGCTTTTCCAGTTTGAAGTTTCCA  
CATATGGTCTTTCTCTTCATCAGAAGTTAGCAGGGCAGGACTGATTAAGGCAGAGATCATCTTATGT  
GGGTTCTCCTGCAATTCATTTCTGGAAGTATTCAGAAAAATGCACTAGCTGATTTTCTCCCTGTGATGAA  
GCTCTTCGACTTGCTATACCCAGAAAAAGAATATATCCAGTTCTGATTAACAAACCCAGTCAACC  
CATGCCTTGAATGACCTGATTTGGATTCATCTCAATAGAAAAGCTCAAAATGACAACTCCAAGCTAC  
AGATTCGAATACCTCATTCCCTAAGACTTCACCATGAGTTCCTGCAGCAGAGTCTAAGAAATAAAGTTT  
ACAGATGAATGACTATAAGATTGCTCTATTGTGAATGCATACTCAAAATTCAGAATGTTTTACATTA



[View online »](#)

CCCATGGGAGCTCTGGTAGAAACTATTTATGGAATGGAATTATGAGGATACCTCTCCCTGGAACAACT  
GTATGGCTTCAGGATCTATTACCCCTTACCTATGAACCTCCTGGATTCACTGACAGTTCATGCCAAAT  
GAGCCTTATTCACAGCATTGCAACCAGGGTGATAAACTTGCTCATGCAAAGTCCAGTGTGGCCTTGGCT  
CCAGCCCTAGTGGAACTTACAGTCGTTTATTGGTCTATATGGAATAGAGTCTTTGGGCATCAAAGGAT  
TTATCAGTCAGCTTTTGCCAACGTGTTTCAAATCATATGCATGGGGGATCTTACACACACTCCTTGAGAT  
GTTTAGCTACCGGATGCATCATATTCAGCCTCATTACAGAGTTCAGCTCCTGAGTCATCTTCATACTTTG  
GCTGCAGTTGCACAAACAAACCAGAACCAGCTCCATCTTTGTGTGAGAGCACTGCTCAGGCTTATAA  
CAGCATTAGGTAGCTCAGAGGTACAACCCAGTTTACACGCTTCCTTAGTGATCCCAAAACAGTGCCTC  
AGCAGAATCTGAAGAAGTGAACCGAGCCTTGATATTGACCTTGGCTAGAGCAACTCATGTAACAGATTTT  
TTTACAGGCTCTGATTCAATTCAGGGAACCTTGGTGAAAGACATACTTCAGACCATCATGAGTTTCACTC  
CTCATAATTGGGCTTACACACCCTGAGCTGTTTTCCAGGCCACTACAGGCATTCTTCAAACAAAATAA  
TGTGCCTCAGGAAAGCCGTTTTAATCTGAAAAAAATGTGGAGGAGGATATAGGAAGTGGAAAGTCAATG  
AGCAACGAAAACGACATTATTACCCACTTCTATGCAGGGCTCCCTCCTCTCTTTCTTGTCTTCTCT  
GGAAAATGCTCTTGAAACAGATCATATTAATCAGATTGGCTATAGAGTATTAGAGAGAATTGGAGCCAG  
GGCCTTGGTAGCCCATGTGAGGACATTTGCAGATTTCTGGTATATGAGTTTTCTACATCAGCAGGGGT  
CAGCAACTCAATAAATGCATTGAAATCTTAATGACATGGTATGGAAGTATAACATTGTTACACTGGACA  
GATTAATTCTCTGCCTGGCCATGCGTAGTCACGAAGGAAATGAAGCCAGGTTTGTATTTCATAATTCA  
GTTGCTGTTACTCAAACCAAACGATTTTAGAAATCGAGTAAGTGACTTTGTGAAGGAAAATCCCCAGAG  
CACTGGTTACAGAATGACTGGCACACCAAGCACATGAATTATCACAAGAAATATCCAGAGAAGTTGATT  
TTGAGGGCCTCGCGAACAGGTGGATCCTCCTGTACAGATCCAGTCTCCCTATCTGCCATCTATTTTGG  
GAATGTGTGCTTCGATTCCCTCCAGTATTTGATATAGTAATCCACAGATTTTTAGAGTTGCTCCGGTA  
TCCAAATCACTGGAGACTCTACTGGATCATCTAGGAGGCTTATAAAATTCATGATCGTCCAGTGACTT  
ATCTGTATAACTCTGCATCTATTAGAAATGCACCTGAGAGACCGGCATTTCTCAAACGAAAATCGT  
CCATGCGATCATTGGCTCTCTGAAGGATAATCGACCGCAGGGCTGGTGTCTAAGTGACACTTACCTGAAA  
TGCCTATGAATGCACGAGAGGAAAATCCTTGGGTTCCAGATGACACCTACTATTGCAGATTGATTGGCA  
GACTAGTCGATACGATGGCTGGCAAATCTCCTGGTCCCTTTCAAACGTGACTGGAGATTCAATGAGTT  
TCCCAACCCAGCTGCCCATGCTCCTCATGTTACTTGTGTGGAGCTCATGGCCTTGGCAGTTTCAGGCAA  
GAAGTTGGGAATGCCCTTCTAAATGTTGCCTAAAAAGTCAGCCTTTAGTGCCAAGAGAGAACATTACAG  
CATGGATGAATGCAATTGGTTTGTATCATCACTGCCCTACCAGAGCCATATTGGATTGTTCTTCATGATCG  
AATTGTGAGTGCATCAGCAGCCCAGCTTACGCTCTGAAACAGAGTGGGTTGGCTATCCATTCCGCCTC  
TTTGATTTCACTGCCTGTATCAGTCTACTCTGAGATGAGTTGTAGCTATACGTTAGCTCTTGACATG  
CTGTGTGGCACCATTCTAGCATCGGACAACCTTTCTCTATTCAAAGTTTCTTACTGAAGTACTTCTCC  
TATAGTGAAGACCGAATTCAGTTGCTTTATGTATACCATCTTGTGGACCAATTTTACAAGATTTTCAG  
CAAGAGAGAACTCGTTGTATGATAGAGATTGGTGTGGCGTTTTATGACATGCTGTGAATGTTGACCAGT  
GTAGCACCCATTTAAATTACATGGATCCCATCTGTGACTTCTCTATCACATGAAGTATATGTTTACTGG  
TGACAGCGTGAAGAGCAAGTAGAGAAGATTATCTGTAACCTTAAAACAGCTTTAAAACCTCGTCTTCGA  
TTCATCACACACATTAGCAAGATGGAGCCAGCTGCAGTGCCTCCACAAGCCATGAACAGTGGGTCTCCAG  
CACCTCAGTCTAATCAG

ACGCGTACGCGGCCGCTCGAG - GFP Tag - GTTTAA

**Protein Sequence:** >RG235279 representing NM\_001270521  
 Red=Cloning site Green=Tags(s)

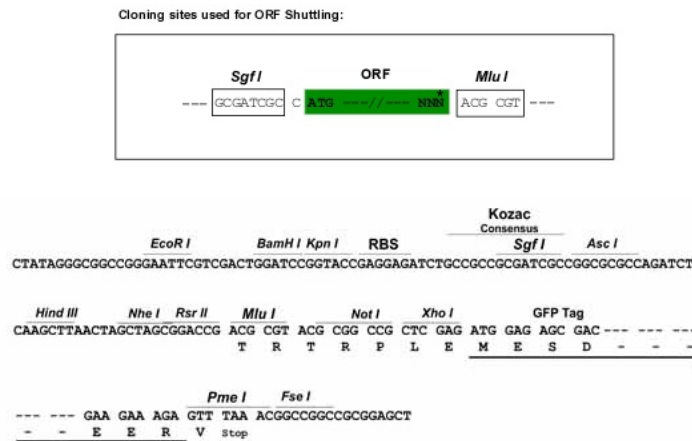
METQLQSIFFEEVVKTEVIEEAFPGMFMDTPEDEKTKLISCLGAFRQFWGGLSQESHEQCIQWIVKFIHGQ  
 HSPKRISFLYDCLAMAVETGLLPPRLVCELSINSDTLEWERTQLWALTFKLVRKIIGGVVDYKGVDRDLLKV  
 ILEKILTIPNTVSSAVVQQLLAAREVIAYILERNACLLPAYFAVTEIRKLYPEGKLPHWLLGNLVSDFVD  
 TFRPTARINSICGRCSLLPVVNSGAICNSWKLDPATLRFPLKGLLPYDKDLFEPQTALLRYVLEQPYSR  
 DMVCNMLGLNKQHKQRCPVLEDQLVDLVVYAMERSETEEKFDDGGTSQLLWQHLSSQLIFFVLFQFASFP  
 HMVLSLHQKLAGRGLIKGRDHLMWVLLQFISGSIQKNALADFLPVMKLFDLLYPEKEYIPVPDINKPQST  
 HAFAMTCIWIHLNRKAQNDNSKLQIPIPHSLRLHHEFLQQSLRNKSLQMNDYKIALLCNAYSTNSECFTL  
 PMGALVETIYNGIMRIPLPGTNCMASGSITPLPMNLLDSLTVHAKMSLIHSAITRVIKLAHAKSSVALA  
 PALVETYSRLLVYMEIESLGIKGFISQLLPTVFKSHAWGILHTLLEMFSYRMHHIQPHYRVQLLSHLHTL  
 AAVAQTNQNLHLCEVESTALRLITALGSSEVQPQFTRFLSDPKTVLSAESEELNRALILTLARATHVDF  
 FTGSDSIQGTWCKDILQTIMSFTPHNWSHTLSCFPGPLQAFFKQNNVQESRFNLKKNVEEYRKKWSM  
 SNENDIITHFSMQGSPPLFLCLLWKMLLETDHINQIGYRVLERIGARALVAHVRTFADFLVYEFSTAGG  
 QQLNKCIEILNDMVWYKIVITLDRILILCLAMRSHEGNEAQVCYFIIQLLLKPNDFRNRVSDFVKENSPE  
 HWLQNDWHTKHMNYHKKYPEKLYFEGLAEQVDPVQIQSPYLPYIFGNVCLRFLPVFDIVIHRFLELLPV  
 SKSLETLLDHLGGLYKFHDRPVTYLYNTLHYYEMHLRDR AFLKRKLVHAIIGSLKDNRPQGWCLSDTYLK  
 CAMNAREENPWPDDTYCRLIGRLVDTMAGKSPGPFPCDWRFNFPNPAHALHVTCVELMALAVSGK  
 EVGNALLNVVLSQPLVRENITAWMNAIGLIITALPEPYWIVLHDIRIVSVISSPLTSETEWVGYPFRL  
 FDF TACHQSYSEMSCSYTLALAHAVWHSSIGQLSLIPKFLTEVLLPIVKTEFQLLYVYHLVGPFLQRFQ  
 QERTRCMIIEIGVAFYDMLLNVDQCSTHLNYMDPICDFLYHMKYMTGDSVKEQVEKIIICNLKPKALKRLR  
 FITHISKMEPAAVPPQAMNSGSPAPQSNQ

TRTRPLE - GFP Tag - V

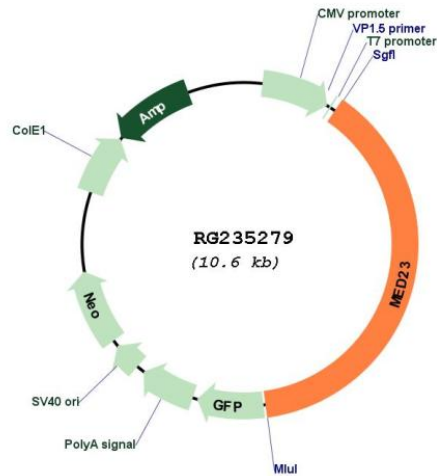
**Restriction Sites:**

Sgfl-MluI

**Cloning Scheme:**



## Plasmid Map:



ACCN: NM\_001270521

ORF Size: 4077 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\\_001270521.2](#)

RefSeq Size: 4612 bp

RefSeq ORF: 4080 bp

Locus ID: 9439

UniProt ID: [Q9ULK4](#)

Cytogenetics: 6q23.2

**Protein Families:** Druggable Genome, Transcription Factors

**Gene Summary:** The activation of gene transcription is a multistep process that is triggered by factors that recognize transcriptional enhancer sites in DNA. These factors work with co-activators to direct transcriptional initiation by the RNA polymerase II apparatus. The protein encoded by this gene is a subunit of the CRSP (cofactor required for SP1 activation) complex, which, along with TFIID, is required for efficient activation by SP1. This protein is also a component of other multisubunit complexes e.g. thyroid hormone receptor-(TR-) associated proteins which interact with TR and facilitate TR function on DNA templates in conjunction with initiation factors and cofactors. This protein also acts as a metastasis suppressor. Several alternatively spliced transcript variants encoding different isoforms have been described for this gene. [provided by RefSeq, Jul 2012]