

## Product datasheet for **KN202659**

### MRPS18C Human Gene Knockout Kit (CRISPR)

#### Product data:

**Product Type:** Knockout Kits (CRISPR)  
**Format:** 2 gRNA vectors, 1 GFP-puro donor, 1 scramble control  
**Donor DNA:** GFP-puro  
**Symbol:** MRPS18C  
**Locus ID:** 51023  
**Components:** **KN202659G1**, MRPS18C gRNA vector 1 in pCas-Guide CRISPR vector (GE100002), Target Sequence: ACACATCCCGGGACTCACAC  
**KN202659G2**, MRPS18C gRNA vector 2 in pCas-Guide CRISPR vector (GE100002), Target Sequence: GAAGTTGACACACTTGGTAA  
**KN202659D**, donor DNA containing left and right homologous arms and GFP-puro functional cassette.

Homologous arm and GFP-puro sequences:

pUC vector backbone in gray; **Left arm sequence in blue**; **GFP-puro in green**; **Right arm in violet**

```
AAGGCGAGTT ACATGATCCC CCATGTTGTG CAAAAAAGCG GTTAGCTCCT TCGGTCCTCC GATCGTTGTC
AGAAGTAAGT TGGCCGAGT GTTATCACTC ATGGTTATGG CAGCACTGCA TAATTCTCTT ACTGTCATGC
CATCCGTAAG ATGCTTTTCT GTGACTGGTG AGTACTCAAC CAAGTCATTC TGAGAATAGT GTATGCCGGC
ACCGAGTTGC TCTTGCCCGG CGTCAATACG GGATAATACC GCGCCACATA GCAGAATTTT AAAAGTGCTC
ATCATTTGAA AACGTTCTTC GGGGCGAAAA CTCTCAAGGA TCTTACCCTG GTTGAGATCC AGTTTCGATGT
AACCCACTCG TGCACCCAAC TGATCTTCAG CATCTTTTAC TTTACCAGC GTTTCTGGGT GAGCAAAAAC
AGGAAGGCAA AATGCCGCAA AAAAGGGAAT AAGGGCGACA CGGAAATGTT GAATACTCAT ACTCTTCCTT
TTTCAATATT ATTGAAGCAT TTATCAGGT TATTGTCTCA TGAGCGGATA CATATTTGAA TGTATTTAGA
AAAATAACA AATAGGGGTT CCGCGCAT TCCCCGAAA AGTGCCACCT GACGTCTAAG AAACCATTAT
TATCATGACA TTAACCTATA AAAATAGGCG TATCACGAGG CCCTTTCGGG TCGCGGTTT CGGTGATGAC
GGTAAAACC TCTGACACAT GCAGCTCCCG TTGACGGTCA CAGCTTGCT GTAAGCGGAT GCCGGGAGCA
GACAAGCCCG TCAGGGCGCG TCAGCGGGTG TTGGCGGGTG TCGGGGCTGG CTTAACTATG CGGCATCAGA
GCAGATTGTA CTGAGAGTGC ACCATAAAAT TGTAACGTT AATATTTTGT TAAAATTCGC GTTAAATTTT
TGTTAAATCA GCTCATTTTT TAACCAATAG GCCGAAATCG GCAAAATCCC TTATAATCA AAAGAATAGC
CCGAGATAGG GTTGAGTGTT GTTCCAGTTT GGAACAAGAG TCCACTATTA AAGAACGTGG ACTCCAACGT
CAAAGGGCGA AAAACCGTCT ATCAGGGCGA TGGCCCACTA CGTGAACCAT CACCAAATC AAGTTTTTTG
GGTTCGAGGT GCCGTAAAGC ACTAAATCGG AACCCCTAAG GGAGCCCCCG ATTTAGAGCT TGACGGGGAA
AGCCGGCGAA CGTGCGGAGA AAGGAAGGGA AGAAAGCGAA AGGAGCGGGC GCTAGGGCGC TGGCAAGTGT
AGCGGTACAG CTGCGCGTAA CCACCACACC CGCCGCGCTT AATGCGCCGC TACAGGGCGC GACTATGGT
TGTTTTGACG TATGCGGTGT GAAATACCGC ACAGATCGCT AAGGAGAAAA TACCGCATCA GGCGCCATTC
GCCATTCAGG CTGCGCAACT GTTGGGAAGG GCGATCGGTG CGGGCCTCTT CGTATTACG CCAGCTGGCG
AAAGGGGAT GTGCTGCAAG GCGATTAAGT TGGGTAACGC CAGGGTTTTT CCAGTACGA CGTTGTAATA
CGACGGCCAG TGAATTGGAG GCTACAGTCA GTGGAGAGGA CTTTCACAG CTGTCGCCGT GCTCATTTGA
```



[View online »](#)

TAACTGCCCG TTTTCATGCG ACACCGGGGA ATCTGAGAGT AGAAGGGGCT GTACCTCAAC CGGCAGTACG  
 CCCCGGTTT TCCGCTCCT GTTCTCAGCC ACCATTTCTT CCTCTCTTT CCCCTCATCT CCGGGCACGA  
 GCTCGGCCG GGTGGGAGCG CCAAAAATAC ACCCCAAGT TGGACGGTTC CTTTTGGGGA GAGACACCCG  
 CCGGCGGATG CGGAACCAC ATTCATCCAT GGCAAGGACC CAGGGCCCTA TTCAGACGTC GTTCTCAGTG  
 ACCCAGACGC TAAGCCATA TGAAGGGAG AGTGGGACGC CGGAGCCCGC TTCTACATCC ACCTTGGGAA  
 AAGACCCCAA GTTAGCTCTC AGGGCTCGCG GACCGGAAGC ACGCATAAAC TTCTACCCTG TCCAATCATA  
 AGCCTCACGT GACCTGCCGC GTAGGGAGGG CACGAACAGA AAGGTGCTGC GCATCGGTAT TGAAAAACCT  
 CACCTACTCT CGCGGGTCTT CAGCGTTCTC CTGCGGAACC TTTGAACGGG GTACTCGAGC CCACAGGGGA  
 AGAGCAGCGG AAGGGGCCTT TCGGAACGAT TTGGAACGAA AGGAAGTGGA AGAAAACGCGG AACCACTAGC  
 ATGGAGAGCG ACGAGAGCGG CCTGCCCGCC ATGGAGATCG AGTGCCGCAT CACCGGCACC CTGAACGGCG  
 TGGAGTTCGA GCTGGTGGGC GCGGAGAGG GCACCCCGA GCAGGGCCGC ATGACCAACA AGATGAAGAG  
 CACCAAAGGC GCCCTGACCT TCAGCCCTA CCTGCTGAGC CACGTGATGG GCTACGGCTT CTACCACTTC  
 GGCACCTACC CCAGCGGCTA CGAGAACCC TTCCTGCACG CCATCAACA CGGCGGCTAC ACCAACACCC  
 GCATCGAGAA GTACGAGGAC GCGGCGTGC TGCACGTGAG CTTCAGCTAC CGCTACGAGG CCGGCCGCGT  
 GATCGGCGAC TTCAAGGTGA TGGGCACCGG CTCCCCGAG GACAGCGTGA TCTTCACCGA CAAGATCATC  
 CGCAGCAACG CCACCGTGA GCACCTGCAC CCCATGGGCG ATAACGATCT GGATGGCAGC TTCACCGCA  
 CCTTCAGCCT GCGCGACGGC GGCTACTACA GCTCCGTGGT GGACAGCCAC ATGCACTTCA AGAGCGCCAT  
 CCACCCAGC ATCCTGCAGA ACGGGGGCCC CATGTTCCGC TTCCGCGCG TGGAGGAGGA TCACAGCAAC  
 ACCGAGCTGG GCATCGTGA GTACCAGCAC GCCTTCAAGA CCCCAGATGC AGATGCCGTG GAAGAAAGAG  
 TTTAAGAATT CCGATCATAT TCAATAACCC TTAATATAAC TTCGTATAAT GTATGTATA CGAAGTTATT  
 AGGTCTGAAG AGGAGTTTAC GTCCAGCCAA GCTTAGGATC TCGACCTCGA AATTCTACCG GGTAGGGGAG  
 GCGCTTTTCC CAAGGCAGTC TGGAGCATGC GCTTTAGCAG CCCCCTGGG CACTTGGCG TACACAAGTG  
 GCCTCTGGCC TCGCACACAT TCCACATCCA CCGTAGGGC CCAACCGACT CCGTTCTTTG GTGGCCCTT  
 CGCGCACCT TCTACTCCTC CCCTAGTCAG GAAGTTCGCC CCGCCCGC AGCTCGGTC GTGACGAGC  
 TGACAAATGG AAGTAGCACG TCTCACTAGT CTCGTGCAGA TGGACAGCAC CGCTGAGCAA TGAAGCGGG  
 TAGGCCTTTG GGCAGCGGC CAATAGCAGC TTTGCTCCTT CGCTTTCTGG GCTCAGAGC TGGGAAGGG  
 TGGTCCGGG GCGGGCTCA GGGGCGGCT CAGGGCGGG GCGGGCGCC GAAGGTCTC CGGAGGCCG  
 GCATTCTGCA CGTTCAAAA GCGCACGTCT GCCGCGTGT TCTCCTCTC CTCATCTCCG GGCCTTTGCA  
 CCTGCATCCA TCTAGATCTC GAGCAGCTGA AGCTTACCAT GACCGAGTAC AAGCCCACGG TGCCTTCGC  
 CACCCGCGAC GACGTCCCA GGGCCGTACG CACCCTCGC GCCGCTTCG CCGACTACC CGCCACGCG  
 CACACCGTCG ATCCGGACCG CCACATCGAG CGGGTCACCG AGCTGCAAGA ACTCTTCTC ACGCGCTCG  
 GGCTCGACAT CGCAAGGTG TGGTCCGGG ACGACGCGC CCGGTGGCG GTCTGGACCA CGCCGGAGAG  
 CGTCGAAGCG GGGGCGGTG TCGCCGAGAT CGGCCGCGC ATGGCCGAGT TGAGCGGTTT CCGGCTGGCC  
 GCGCAGAAC AGATGGAAGG CCTCTGGCG CCGCACCGC CCAAGGAGCC CGCGTGGTTT CTGGCCACC  
 TCGGCGTCTC GCCGACCAC CAGGGCAAGG GTCTGGGAG CCGCGTCTG CTCCCGGAG TGGAGGCGG  
 CGAGCGCGC GGGGTGCCG CTTTCTGGA GACCTCCGCG CCCCACAACC TCCCCTTCTA CGAGCGGCTC  
 GGCTTACCG TCACCGCCGA CGTCGAGGTG CCCGAAGGAC CCGCACCTG GTGCATGACC CGCAAGCCG  
 GTGCCTGACG CCCGCCAC GACCCGCAGC GCCGACCGA AAGGAGCGCA CGACCCATG CATCGATGAT  
 ATCAGATCCC CGGGATGCAG AAATTGATGA TCTATTAAC AATAAAGATG TCCACTAAA TGAAGTTTT  
 TCCTGTATA CTTTGTAAAG AAGGGTGAGA ACAGAGTACC TACATTTTGA ATGGAAGGAT TGGAGCTACG  
 GGGGTGGGG TGGGTGGGA TTAGATAAAT GCCTGCTCTT TACTGAAGGC TCTTACTAT TGCTTTATGA  
 TAATGTTTCA TAGTTGGATA TCATAATTA AACAAGCAA ACCAAATTAA GGGCCAGCTC ATTCCTCCA  
 CTCATGATCT ATAGATCTAT AGATCTCTG TGGGATCATT GTTTTTCTT TGATTCCCAC TTTGTGGTTC  
 TAAGTACTGT GGTTCAAA TGTGTCAGTT TCATAGCTG AAGAACGAGA TCAGCAGCTT CTGTTCCACA  
 TACTTTCAT TCTCAGTATT GTTTTGCCAA GTTCTAATTC CATCAGAAGC TGGTCGAGAT CCGGAACCT  
 TAATATAACT TCGTATAATG TATGCTATAC GAAGTTATTA GGTCCCTCGA AGAGGTTTAC TAGGCGCGC  
 ACCTTAGTCC TAATGGTTAG AGTCGTCCCT GTTAGCAAAA CGACTCTGGT TTCTAGGCGA TTAAGTTTCT  
 CCAAGATGGG TGGGAATCTG TAATTGCCTT TACTGTTTT AGTGTCGCTT AATCCACTCC TTTAAAGTAT  
 TTAACGATG GAGGAAGTTG GAGTGGTCT TAAAAAGCAG AGACTTTCCT AAGATCTCTG GATAATCTTA  
 GTGAGGAAGG GACATTTCCA GAGTCGCCA GCAGCAAATT CCAGATGTCT AAGGTCCCA AACAGAACA  
 AATTGCATA TTTTGTCAAT TTTATCTTCT CTGTGCTTCT CTATAAATTC TTTCTGGGAC GGGTCTGTTA  
 CTTTCTTTT GGTATGACAG TGCCTTCTG TTGCTTCTC TATAAGAATT TGCATTAAGA ATTTTCTCTA

ACCATGGAAT GAAGGAAAA TAAATGTCAG AAAGAGCAGG TATGGGCGGG AAAATATGTT CTTTTCCCCC  
 TAAGCCCTGG AAAGCACACC CATGTTGAGT CTTACTCTGT CCCAAATATA AAAGTTTTTA ACCATTCTGA  
 ATATAATACA AAACCTTTCT TTAATATCGT GTCCTTAAT TCACTCTCGC CGGTTGGACT TTAGATCAGA  
 AGGGATCTTG CTGCCGCCCG AAAGAGGAAG GGCTGGAAGA GGAAGGAGCT TGGCGTAATC ATGGTCATAG  
 CTGTTTCTCG TGTGAAATTG TTATCCGCTC ACAATCCAC ACAACATACG AGCCGGAAGC ATAAAGTGTA  
 AAGCCTGGGG TGCCTAATGA GTGAGCTAAC TCACATTAAT TGC GTTGGC TCACTGCCCG CTTTCCAGTC  
 GGGAAACCTG TCGTGCCAGC TGCATTAATG AATCGGCCAA CGCGCGGGGA GAGCGGTGTT GCGTATTGGG  
 CGCTCTTCCG CTTCTCGCT CACTGACTCG CTGCGCTCGG TCGTTCGGCT GCGGCGAGCG GTATCAGCTC  
 ACTCAAAGGC GGTAATACGG TTATCCACAG AATCAGGGGA TAACGCAGGA AAGAACATGT GAGCAAAGG  
 CCAGCAAAG GCCAGGAACC GTAAAAAGGC CGCGTTGCTG GCGTTTTTCC ATAGGCTCCG CCCCCCTGAC  
 GAGCATCACA AAAATCGACG CTCAAGTCAG AGGTGGCGAA ACCCGACAGG ACTATAAAGA TACCAGGCGT  
 TTCCCCTGG AAGCTCCCTC GTGCGCTCTC CTGTTCCGAC CCTGCCGCTT ACCGGATACC TGTCCGCCTT  
 TCTCCCTCG GGAAGCGTGG CGCTTTCTCA TAGCTCACGC TGTAGGTATC TCAGTTCGGT GTAGGTCGTT  
 CGCTCCAAGC TGGGCTGTGT GCACGAACCC CCCGTTACG CCGACCGTG CGCCTTATCC GGTAACTATC  
 GTCTTGAGTC CAACCCGTA AGACACGACT TATCGCCACT GGCAGCAGCC ACTGGTAACA GGATTAGCAG  
 AGCGAGGTAT GTAGGCGGTG CTACAGAGTT CTGAAGTGG TGGCCTAACT ACGGCTACAC TAGAAGAAACA  
 GTATTTGGTA TCTGCGCTCT GCTGAAGCCA GTTACCTTCG GAAAAAGAGT TGGTAGCTCT TGATCCGGCA  
 AACAAACCAC CGCTGGTAGC GGTGGTTTTT TTGTTTGCAA GCAGCAGATT ACGCGCAGAA AAAAAGGATC  
 TCAAGAAGAT CCTTTGATCT TTTCTACGGG GTCTGACGCT CAGTGGAAAC AAAACTCACG TTAAGGGATT  
 TTGGTCATGA GATTATCAA AAGGATCTTC ACCTAGATCC TTTTAAATTA AAAATGAAGT TTTAAATCAA  
 TCTAAAGTAT ATATGAGTAA ACTTGGTCTG ACAGTTACCA ATGCTTAATC AGTGAGGCAC CTATCTCAGC  
 GATCTGTCTA TTTGTTTCAT CCATAGTTGC CTGACTCCCC GTCGTGTAGA TAACTCCAGT ACGGGAGGGC  
 TTACCATCTG GCCCCAGTGC TGCAATGATA CCGCAGAAC CACGCTACC GGCTCCAGAT TTATCAGCAA  
 TAAACCAGCC AGCCGGAAGG GCCGAGCGCA GAAGTGGTCC TGCAACTTTA TCCGCCCTCCA TCCAGTCTAT  
 TAATTGTTGC CGGGAAGCTA GAGTAAGTAG TTCGCCAGTT AATAGTTTGC GCAACGTTGT TGCCATTGCT  
 ACAGGCATCG TGGTGTACG CTCGTCGTTT GGTATGGCTT CATTACGCTC CGTTTCCAA CGATC

**GE100003**, scramble sequence in pCas-Guide vector

**Disclaimer:**

These products are manufactured and supplied by OriGene under license from ERS. The kit is designed based on the best knowledge of CRISPR technology. The system has been functionally validated for knocking-in the cassette downstream the native promoter. The efficiency of the knock-out varies due to the nature of the biology and the complexity of the experimental process.

**RefSeq:**

[NM\\_001297767](#), [NM\\_001297769](#), [NM\\_001297770](#), [NM\\_016067](#)

**UniProt ID:**

[Q9Y3D5](#)

**Synonyms:**

CGI-134; MRP-S18-1; MRP-S18-c; MRPS18-1; mrps18-c; S18mt-c

**Summary:**

Mammalian mitochondrial ribosomal proteins are encoded by nuclear genes and help in protein synthesis within the mitochondrion. Mitochondrial ribosomes (mitoribosomes) consist of a small 28S subunit and a large 39S subunit. They have an estimated 75% protein to rRNA composition compared to prokaryotic ribosomes, where this ratio is reversed. Another difference between mammalian mitoribosomes and prokaryotic ribosomes is that the latter contain a 5S rRNA. Among different species, the proteins comprising the mitoribosome differ greatly in sequence, and sometimes in biochemical properties, which prevents easy recognition by sequence homology. This gene encodes a 28S subunit protein that belongs to the ribosomal protein S18P family. The encoded protein is one of three that has significant sequence similarity to bacterial S18 proteins. The primary sequences of the three human mitochondrial S18 proteins are no more closely related to each other than they are to the prokaryotic S18 proteins. Pseudogenes corresponding to this gene are found on chromosomes 8p, 12p, 15q, and 22q. [provided by RefSeq, Jul 2008]

**Product images:**
