

Product datasheet for **KN317757RB**

Sting1 Mouse Gene Knockout Kit (CRISPR)

Product data:

Product Type:	Knockout Kits (CRISPR)
Format:	2 gRNA vectors, 1 RFP-BSD donor, 1 scramble control
Donor DNA:	RFP-BSD
Symbol:	Sting1
Locus ID:	72512
Components:	<p>KN317757G1, Sting1 gRNA vector 1 in pCas-Guide CRISPR vector (GE100002), Target Sequence: AGCGGTGACCTCTGGGCCGT</p> <p>KN317757G2, Sting1 gRNA vector 2 in pCas-Guide CRISPR vector (GE100002), Target Sequence: TGGCCAGCCTGATGATCCTT</p> <p>KN317757RBD, donor DNA containing left and right homologous arms and RFP-BSD functional cassette.</p>

Homologous arm and RFP-BSD sequences:

pUC vector backbone in gray; **Left arm sequence in blue**; **RFP-BSD in green**; **Right arm in violet**

```

AAGGCGAGTT ACATGATCCC CCATGTTGTG CAAAAAAGCG GTTAGCTCCT TCGGTCCTCC GATCGTTGTC
AGAAGTAAGT TGGCCGCAGT GTTATCACTC ATGGTTATGG CAGCACTGCA TAATTCTCTT ACTGTCATGC
CATCCGTAAG ATGCTTTTCT GTGACTGGTG AGTACTCAAC CAAGTCATTC TGAGAATAGT GTATGCGGGC
ACCGAGTTGC TCTTGCCCGG CGTCAATACG GGATAATACC GCGCCACATA GCAGAATTTT AAAAGTGCTC
ATCATTGGAA AACGTTCTTC GGGGCGAAAA CTCTCAAGGA TCTTACCCTG GTTGAGATCC AGTTCGATGT
AACCCACTCG TGCACCCAAC TGATCTTCAG CATCTTTTAC TTTACCAGC GTTTCTGGGT GAGCAAAAAC
AGGAAGGCAA AATGCCGCAA AAAAGGGAAT AAGGGCGACA CGGAAATGTT GAATACTCAT ACTCTTCTT
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 AACCCATAAG GGAGCCCCCG ATTTAGAGCT TGACGGGGAA
AGCCGGCGAA CGTGCGGAGA AAGGAAGGGA AGAAAGCGAA AGGAGCGGGC GCTAGGGCGC TGGCAAGTGT
AGCGGTACG CTGCGGTAA CCACCACACC CGCCGCGCTT AATGCGCCGC TACAGGGCGC GACTATGGT
TGCTTTGACG TATGCGGTGT GAAATACCGC ACAGATCGCT AAGGAGAAAA TACCGCATCA GGCGCCATTC
GCCATTCAGG CTGCGCAACT GTTGGGAAGG GCGATCGGTG CGGGCCTCTT CGTATTACG CCAGCTGGCG
AAAGGGGAT GTGCTGCAAG GCGATTAAGT TGGGTAACGC CAGGGTTTTT CCAGTCACGA CGTTGTAATA
CGACGGCCAG TGAATTGGAG GCTACAGTCA GTGGAGAGGA CTTTCACAG CTGTCGCCGT GCTCATTTGA

```



[View online »](#)

TAACTGCCCG TTATTCATGC GACACGGGGG GGGGTGCTTT GGCAGGAAAC ACCAAAAAGA ATAAAGAGCT
 GGCCAGATGG CTAGCTTCCCT CCCAGAGCAC TGAAAACGTG GCTCTGGCAA GGGCCAGGGC TGTGGTTTTA
 CTGAGGAAGC GAGCTGAGTT GCAGTTGTTT TTAACATTTT GTTTTCAGTC TCCCCATTCA GAAGCCACTT
 GCTAGTAGCT ACTGAAAGGC TCTTCATTGT CTCTTCTGCT CCAGGAACAC CGGTCTAGGA AGCAGAAGGT
 AGGATTAGAA ATGGGGCAGT ATTCTCCTGG TACTTCGTGG TGGGAGAAAC CCTGGGCTGG CAGAACACTC
 TAAGGTCTAG GTCTTGGATC AGTTTCTTCC TTCTTTCTCA CACCAGTGAG CTCAGTGGGG CTCACATGTA
 CAGGCTCTGT TTAATATGAA CCTCTCCTAG ACAGGTGCTG TAGGATGCTA TGTGCCCAGG GCGTCTCCTT
 GAGGTGTATC CAAGAGTAGC CCATGGGACT AGCTGGGCTG GGAGACCCAA GGAGATGGAT GACTAGTCAG
 GACCTTAGGC CCTGAAGGAC AGGAGAACC CAGAAGCATA GCTGTGGATT TCTTGATGTC TACAGCGCAC
 GAACCTGGAC TACTGTTGAA AACCTCTGC TGTCTGGCTG AAGAGCTGTG CCATGTCCAG TCCAGGTAAC
 CCTCTGTGGT CTCCACGATG ACTTGATACC CAGCAAGTAT CACACCCTTG ACTCCTAAC TTTGGCCTTCC
 CTGAACATCA TCAGCTGAGT TGGGCTGGGA TCTGAGGCTA GGCTGTCCCC TGCAGGTACC AGGGCAGCTA
 CTGGAAGGCT GTGCGCGCCT GCCTGGGATG CCCCATCCAC TGTATGGCTA TGATTCTACT ATCGTCTTAT
 TTCTATTTCC TCCAAAACAC TGCTGACATA TACCTCAGTT GGATGTTTGG CTTTCTGGTC CTCTATAAGT
 CCCTAAGCAT GCTCCTGGC CTTCAGGTAT GAAGCAGAAG AGGTGGGTGG TGTGGGAAGA GGGGACTATG
 GCTAGTGCTG GTGTGCCTGA GCTCAGTCT GAGACTCAGA CTAATTTAAA GGTTGGAGAC CTGGGTGTGG
 AGCTATGAAG GCTTGGGATG ATGGGTTTAA TAGCAGTCT GAGAGCAAAG TGGCAGCAGG TTGGGAAAGT
 TTTCTGCAAG AGAAGGGCTT TGGACATCCC CTTGTCTACT CTCGCCGTT GGACTTTAGA TCAGAAGGGA
 TCTTGCTGCC GCCCGAAAGA GGAAGGGCTG GAAGAGGAAG GAGCTTGGCG TAATCATGGT CATAGCTGTT
 TCCTGTGTGA AATTGTTATC CGCTCACAA TCCACACAAC ATACGAGCCG GAAGCATAAA GTGTAAAGCC
 TGGGGTGCCCT AATGAGTGAG CTAACTCACA TTAATTGCGT TGCCTACT GCCCCTTTT CAGTCGGGAA
 ACCTGTCTGT CCAGCTGCAT TAATGAATCG GCCAACGCGC GGGGAGAGGC GGTTCGCTA TTGGGCGCTC
 TTCCGCTTCC TCGCTCACTG ACTCGTCCG CTCGGTCTGT CGGCTGCGGC GAGCGTATC AGCTCACTCA
 AAGGCGGTAA TACGGTTATC CACAGAATCA GGGGATAACG CAGGAAAGAA CATGTGAGCA AAAGGCCAGC
 AAAAGGCCAG GAACCGTAAA AAGGCCGCGT TGCTGGCGTT TTTCCATAGG CTCCGCCCC CTGACGAGCA
 TCACAAAAAT CGACGCTCAA GTCAGAGGTG GCGAAACCCG ACAGGACTAT AAAGATACCA GGCGTTTCCC
 CCTGGAAGCT CCCTCGTGCG CTCTCCTGTT CCGACCCTGC CGCTTACCGG ATACCTGTCC GCCTTCTCC
 CTTCCGGAAG CGTGGCGCTT TCTCATAGCT CACGCTGTAG GTATCTCAGT TCGGTGTAGG TCGTTCGCTC
 CAAGCTGGGC TGTGTGCACG AACCCCGT TCAGCCCGAC CGCTGCGCCT TATCCGTAA CTATCGTCTT
 GAGTCCAACC CGGTAAGACA CGACTTATCG CCACTGGCAG CAGCCACTGG TAACAGGATT AGCAGAGCGA
 GGTATGTAGG CGGTGCTACA GAGTCTTGA AGTGGTGGCC TAACTACGGC TACACTAGAA GAACAGTATT
 TGGTATCTGC GCTCTGCTGA AGCCAGTTAC CTTCGAAAA AGAGTTGGTA GCTCTTGATC CGGCAAACAA
 ACCACCGCTG GTAGCGGTGG TTTTTTTGTT TGCAAGCAGC AGATTACGCG CAGAAAAAAA GGATCTCAAG
 AAGATCCTTT GATCTTTTCT ACGGGGTCTG ACCTCAGTG GAACGAAAAC TCACGTAAAG GGATTTTGGT
 CATGAGATTA TCAAAAAGGA TCTTACCTA GATCCTTTTA AATTAATAAT GAAGTTTTAA ATCAATCTAA
 AGTATATATG AGTAAACTTG GTCTGACAGT TACCAATGCT TAATCAGTGA GGCACCTATC TCAGCGATCT
 GTCTATTTTC TTCATCCATA GTTGCCTGAC TCCCCGCTG GTAGATAACT ACGATACGGG AGGGCTTACC
 ATCTGGCCCC AGTGCTGCAA TGATACCGCG AGAACACGC TCACCGGCTC CAGATTTATC AGCAATAAAC
 CAGCCAGCCG GAAGGGCCGA GCGCAGAAGT GGTCTGCAA CTTTATCCGC CTCCATCCAG TCTATTAATT
 GTTGCCGGGA AGCTAGAGTA AGTAGTTCGC CAGTTAATAG TTTGCGCAAC GTTGTGCCA TTGCTACAGG
 CATCGTGTG TCACGCTCGT CGTTTGGTAT GGCTTCATTC AGCTCCGTT CCAACGATC

GE10003, 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_001289591](#), [NM_001289592](#), [NM_028261](#)

UniProt ID:

[Q3TBT3](#)

Synonyms: 2610307O08Rik; ERIS; Mita; MPYS; STING

Summary: Facilitator of innate immune signaling that acts as a sensor of cytosolic DNA from bacteria and viruses and promotes the production of type I interferon (IFN-alpha and IFN-beta) (PubMed:18818105, PubMed:19433799, PubMed:19776740, PubMed:26229117, PubMed:26669264). Innate immune response is triggered in response to non-CpG double-stranded DNA from viruses and bacteria delivered to the cytoplasm (PubMed:18818105, PubMed:19433799, PubMed:19776740, PubMed:26229117, PubMed:26669264). Acts by binding cyclic dinucleotides: recognizes and binds cyclic di-GMP (c-di-GMP), a second messenger produced by bacteria, and cyclic GMP-AMP (cGAMP), a messenger produced by CGAS in response to DNA virus in the cytosol (PubMed:21947006, PubMed:23722158, PubMed:23258412, PubMed:23519410, PubMed:23910378). Upon binding of c-di-GMP or cGAMP, TMEM173/STING oligomerizes, translocates from the endoplasmic reticulum and is phosphorylated by TBK1 on the pLxIS motif, leading to recruitment and subsequent activation of the transcription factor IRF3 to induce expression of type I interferon and exert a potent anti-viral state (PubMed:25636800). In addition to promote the production of type I interferons, plays a direct role in autophagy (PubMed:30568238). Following cGAMP-binding, TMEM173/STING buds from the endoplasmic reticulum into COPII vesicles, which then form the endoplasmic reticulum-Golgi intermediate compartment (ERGIC) (By similarity). The ERGIC serves as the membrane source for WIPI2 recruitment and LC3 lipidation, leading to formation of autophagosomes that target cytosolic DNA or DNA viruses for degradation by the lysosome (By similarity). The autophagy- and interferon-inducing activities can be uncoupled and autophagy induction is independent of TBK1 phosphorylation (By similarity). Autophagy is also triggered upon infection by bacteria: following c-di-GMP-binding, which is produced by live Gram-positive bacteria, promotes reticulophagy (PubMed:29056340). Exhibits 2',3' phosphodiester linkage-specific ligand recognition: can bind both 2'-3' linked cGAMP (2'-3'-cGAMP) and 3'-3' linked cGAMP but is preferentially activated by 2'-3' linked cGAMP (PubMed:26300263). The preference for 2'-3'-cGAMP, compared to other linkage isomers is probably due to the ligand itself, which adopts an organized free-ligand conformation that resembles the TMEM173/STING-bound conformation and pays low energy costs in changing into the active conformation (By similarity). May be involved in translocon function, the translocon possibly being able to influence the induction of type I interferons (By similarity). May be involved in transduction of apoptotic signals via its association with the major histocompatibility complex class II (MHC-II) (PubMed:18559423).[UniProtKB/Swiss-Prot Function]

Product images:

