

## Product datasheet for **KN311043**

### Nlrp12 Mouse 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:	Nlrp12
Locus ID:	378425
Components:	<p><b>KN311043G1</b>, Nlrp12 gRNA vector 1 in pCas-Guide CRISPR vector (GE100002), Target Sequence: GGTAGGTAGACAGTCGATAG</p> <p><b>KN311043G2</b>, Nlrp12 gRNA vector 2 in pCas-Guide CRISPR vector (GE100002), Target Sequence: TACCTGGAAGAACTCGAGGC</p> <p><b>KN311043D</b>, donor DNA containing left and right homologous arms and GFP-puro functional cassette.</p> <p>Homologous arm and GFP-puro sequences:            pUC vector backbone in gray; Left arm sequence in blue; GFP-puro in green; Right arm in violet</p> <pre> AAGGCGAGTT ACATGATCCC CCATGTTGTG CAAAAAAGCG GTTAGCTCCT TCGGTCCTCC GATCGTTGTC AGAAGTAAGT TGGCCGCAGT GTTATCACTC ATGGTTATGG CAGCACTGCA TAATTCTCTT ACTGTCATGC CATCCGTAAG ATGCTTTTCT GTGACTGGTG AGTACTCAAC CAAGTCATTG TGAGAATAGT GTATGCCGGC ACCGAGTTGC TCTTGCCCGG CGTCAATACG GGATAATACC GCGCCACATA GCAGAACTTT AAAAGTGCTC ATCATTGGAA AACGTTCTTC GGGGCGAAAA CTCTCAAGGA TCTTACCGCT GTTGAGATCC AGTTCGATGT AACCCACTCG TGCACCCAAC TGATCTTCAG CATCTTTTAC TTTCACCAGC GTTTCTGGGT GAGCAAAAAC AGGAAGGCAA AATGCCGCAA AAAAGGGAAT AAGGGCGACA CGGAAATGTT GAATACTCAT ACTCTTCCTT TTTCAATATT ATTGAAGCAT TTATCAGGTT TATTGTCTCA TGAGCGGATA CATATTTGAA TGTATTTAGA AAAATAAACA AATAGGGGTT CCGCGCACAT TTCCCCGAAA AGTGCCACCT GACGTCTAAG AAACCATTAT TATCATGACA TTAACCTATA AAAATAGGCG TATCACGAGG CCCTTTCGGG TCGCGCGTTT CGGTGATGAC GGTAAAACC TCTGACACAT GCAGCTCCCG TTGACGGTCA CAGCTTGTCT GTAAGCGGAT GCCGGGAGCA GACAAGCCCG TCAGGGCGCG TCAGCGGGTG TTGGCGGGTG TCGGGGCTGG CTTAACTATG CGGCATCAGA GCAGATTGTA CTGAGAGTGC ACCATAAAAT TGTAACGTT AATATTTTGT TAAAATTCGC GTTAAATTTT TGTTAAATCA GCTCATTTTT TAACCAATAG GCCGAAATCG GCAAAATCCC TTATAATCA AAAGAAATAGC CCGAGATAGG GTTGAGTGTT GTTCCAGTTT GGAACAAGAG TCCACTATTA AAGAACGTGG ACTCCAACGT CAAAGGGCGA AAAACCGTCT ATCAGGGCGA TGGCCCACTA CGTGAACCAT CACCCAAATC AAGTTTTTTG GGGTCGAGGT GCCGTAAAGC ACTAAATCGG AACCCCTAAG GGAGCCCCCG ATTTAGAGCT TGACGGGGAA AGCCGGCGAA CGTGCGGAGA AAGGAAGGGA AGAAAGCGAA AGGAGCGGGC GCTAGGGCGC TGGCAAGTGT AGCGGTACAG CTGCGCGTAA CCACCACACC CGCCGCGCTT AATGCGCCGC TACAGGGCGC GACTATGGT TGCTTTGACG TATGCGGTGT GAAATACCGC ACAGATGCGT AAGGAGAAAA TACCGCATCA GCGCCATTTC GCCATTCAGG CTGCGCAACT GTTGGGAAGG GCGATCGGTG CGGGCTCTTT CGCTATTACG CCAGCTGGCG AAAGGGGGAT GTGCTGCAAG GCGATTAAGT TGGGTAACGC CAGGGTTTTC CCAGTCACGA CGTTGTAAAA CGACGGCCAG TGAATTGGAG GCTACAGTCA GTGGAGAGGA CTTTCACTGA CTGACTGACT GCGTCTCAAC           </pre>



View online »

TCTGGAATT	GAAGTCAGG	TCTCTGGAAG	AGTAGTCAGT	GCTCTGAAC	GCTAAGCCAT	CTCTCCAGCC
CGAACCTGGA	ACTGCTGGTC	CTCCCTCCTC	TACCTCAGGA	ATGCTGGCTT	TATAGGAGAG	TGCACCACTA
CATCTTCTCC	ATGCAGTCAG	TGGTGGAGCA	GAACCCAGGG	CTTCATTCTG	CCCACCGAGC	CACCTCCCCA
GCAGCCTTGC	ATATTCTTGG	AAGAGGCCAC	ATCTTCACAC	GTCCTTCTGA	GTTAGAAGGA	CACATAGAAA
GAGAGAAGAG	CGTGGTTGTT	TTGTCAAGGG	CGCTGGGAAA	ACAGCAGAAG	CTTGCTCAAT	GTGGGTCTATG
CTTACGGTTG	GATTACCTCA	TACCAACTTC	CACATCTTCA	AAGCAAACAC	TATTGTTGAC	AGGAGCAGGT
AGCAGAGAAG	CAATGGGTAG	GAGAGTATAA	TTTCTCGAAA	GATCTTGCCT	CTGTCCCAAT	CTTATTCTTA
TAACCTCTAC	CCCCTTTTGC	TTGTCAATGA	GGCCATGGCA	GCGTGTAGC	TACAGTGCTT	ATTGTCTTTT
TAGAGAGAGG	TTAGAGCATC	TGGCCCCGTT	ACTTTGTCCC	CCACTAGCAT	GGAGAGCGAC	GAGAGCGGCC
TGCCCCGCAT	GGAGATCGAG	TGCCGCATCA	CCGGCACCTT	GAACGGCGTG	GAGTTCGAGC	TGGTGGGCGG
CGGAGAGGGC	ACCCCCGAGC	AGGGCCGCAT	GACCAACAAG	ATGAAGAGCA	CCAAAGGCGC	CCTGACCTTC
AGCCCCCTAC	TGCTGAGCCA	CGTGATGGGC	TACGGCTTCT	ACCACTTCGG	CACCTACCCC	AGCGGTACG
AGAACCCCTT	CCTGCACGCC	ATCAACAACG	GCGGCTACAC	CAACACCCGC	ATCGAGAAGT	ACGAGGACGG
CGGCGTGCTG	CACGTGAGCT	TCAGCTACCG	CTACGAGGCC	GGCCGCGTGA	TCGGCGACTT	CAAGGTGATG
GGCACC GGCT	TCCCCGAGGA	CAGCGTGATC	TTCACCGACA	AGATCATCCG	CAGCAACGCC	ACCGTGGAGC
ACCTGCACCC	CATGGGCGAT	AACGATCTGG	ATGGCAGCTT	CACCCGACCC	TTCAGCCTGC	GCGACGCGCG
CTACTACAGC	TCCGTGGTGG	ACAGCCACAT	GCACTTCAAG	AGCGCCATCC	ACCCCAGCAT	CCTGCAGAAC
GGGGGCCCCA	TGTTGCGCTT	CCGCCGCGTG	GAGGAGGATC	ACAGCAACAC	CGAGCTGGGC	ATCGTGGAGT
ACCAGCACGC	CTTCAAGACC	CCGGATGCAG	ATGCCGCTGA	AGAAAGAGTT	TAAGAATTCC	GATCATATTC
AATAACCTT	AATATAACTT	CGTATAATGT	ATGCTATACG	AAGTTATTAG	GTCTGAAGAG	GAGTTTACGT
CCAGCCAAGC	TTAGGATCTC	GACCTCGAAA	TTCTACCGGG	TAGGGGAGGC	GCTTTTCCCA	AGGCAGTCTG
GAGCATGCGC	TTTAGCAGCC	CCGCTGGGCA	CTTGGCGCTA	CACAAGTGGC	CTCTGGCCTC	GCACACATTTC
CACATCCACC	GGTAGGCGCC	AACCGACTCC	GTTCTTTTGT	GGCCCCCTCG	CGCCACCTTC	TACTCTCTCC
CTAGTCAGGA	AGTTCCCCCC	CGCCCCGAGC	CTCGCGTCGT	GCAGGACGTG	ACAAATGGAA	GTAGCACGTC
TCACTAGTCT	CGTGCAAGTG	GACAGCACCG	CTGAGCAATG	GAAGCGGGTA	GGCCTTTGGG	GCAGCGGCCA
ATAGCAGCTT	TGCTCCTTCG	CTTTCTGGGC	TCAGAGGCTG	GGAAGGGGTG	GGTCCGGGGG	CGGGCTCAGG
GGCGGGCTCA	GGGGCGGGGC	GGGCGCCCGA	AGGTCTCCCG	GAGGCCCGGC	ATTCTGCACG	CTTCAAAAGC
GCACGTCTGC	CGCGCTGTTC	TCCTCTTCTT	CATCTCCGGG	CCTTTTCGAC	TGCATCCATC	TAGATCTCGA
GCAGCTGAAG	CTTACCATGA	CCGAGTACAA	GCCCACGGTG	CGCCTCGCCA	CCCGCGACGA	CGTCCCCAGG
GCCGTACGCA	CCCTCGCCGC	CGCGTTCCGC	GACTACCCCG	CCACGCGCCA	CACCGTCGAT	CCGGACCGCC
ACATCGAGCG	GGTCACCGAG	CTGCAAGAAC	TCTTCTCAC	GCRCGTGGG	CTCGACATCG	GCAAGGTGTG
GGTCGCGGAC	GACGGCGCCG	CGGTGGCGGT	CTGGACCACG	CCGGAGAGCG	TGAAGCGGG	GGCGGTGTTC
GCCGAGATCG	GCCC GCGCAT	GGCCGAGTTG	AGCGGTTCCC	GGCTGGCCGC	GCAGCAACAG	ATGGAAGGCC
TCCTGGCGCC	GCACCGGCC	AAGGAGCCCG	CGTGTTCTCT	GGCCACCGTC	GGCGTCTCGC	CCGACCACCA
GGGCAAGGGT	CTGGGCAGCG	CCGTCTGTCT	CCCCGGAGTG	GAGGCGGCCG	AGCGCGCCGG	GGTGCCCGCC
TTCTTGAGAG	CCTCCGCGCC	CCACAACCTC	CCCTTCTACG	AGCGGCTCGG	CTTACCGTC	ACCGCCGACG
TCGAGGTGCC	CGAAGGACCG	CGCACCTGGT	GCATGACCCG	CAAGCCCGGT	GCCTGACGCC	CGCCCCACGA
CCCGCAGCGC	CCGACCGAAA	GGAGCGCACG	ACCCCATGCA	TCGATGATAT	CAGATCCCCG	GGATGCAGAA
ATTGATGATC	TATTAACAAA	TAAAGATGTG	CACATAAATG	GAAGTTTTTC	CTGTCACTAT	TTGTTAAGAA
GGGTGAGAAC	AGAGTACCTA	CATTTTGAAT	GGAAGGATTG	GAGCTACGGG	GGTGGGGGTG	GGGTGGGATT
AGATAAATGC	CTGCTCTTTA	CTGAAGGCTC	TTTACTATTG	CTTTATGATA	ATGTTTCATA	GTTGGATATC
ATAATTTAAA	CAAGCAAAAC	CAAATTAAGG	GCCAGCTCAT	TCCTCCCACT	CATGATCTAT	AGATCTATAG
ATCTCTCTGT	GGATCATTGT	TTTTCTCTTG	ATTCCCACTT	TGTGGTTCTA	AGTACTGTGG	TTTCCAAATG
TGTCAGTTTC	ATAGCCTGAA	GAACGAGATC	AGCAGCCTCT	GTTCCACATA	CACCTTCATT	TCAGTATTGT
TTTGCCAAGT	TCTAATTCCA	TCAGAAGCTG	GTCGAGATCC	GGAACCCTTA	ATATAACTTC	GTATAATGTA
TGCTATACGA	AGTTATTAGG	TCCCTCGAAG	AGGTTCACTA	GGCGCGCCAT	TATTCCTGGG	GATTGCAGAG
GACCTGAGCC	AGGACAAAAT	TCCCTGGGGA	CGAATGGAGA	AGGCTGGTCC	TCTGAAATG	GCTCAGCTGA
TGGTGGCCCA	CATGGGGACA	AGGGAGGCTT	GGCTTCTGGC	TCTCAGCACC	TTTCAGAGGA	TTACAGGAA
GGACCTGTGG	GAGCGAGGAC	AGGGAGAAGA	CCTGGTGAGG	GGTAAGGAGG	GCAAGGGAGG	TACAGCCTGG
AACACATCCA	TTCTCTTATC	TTTACCATTG	TGAGCTGGAT	TCAAATTTCA	GCCAGTGGTC	TCTTTCCATG
TCTACTACAA	ATCCCCTCTT	TTTTTAAAAA	AGGACTTTGT	GGTGTATGTG	TGTGTGTGTG	TGTGTGTGTG
TGTGTGAGAG	AGAGAGAGAG	AGAGAGAGAG	AGAGAGAGAG	AATTTGCAAA	AGTCCATTGT	CTCCTTCTAC

AACTGGAATG GTTTTGGCCA TGTTCCTGAC CCTAATCCCA GTATTCTCAT ATGTGTACAT AATGTTGATC  
 ACACACACAC TCATGACTCC ACCTCCTGCT CCTCCCATGG ACCCCTTCCT CTCCCAAATA GTCCTCTTCC  
 TACTTTGATG TCTTTTTTATG TGAGAGACGA CTGACTGACT GACTGGAAAG AGGAAGGGCT GGAAGAGGAA  
 GGAGCTTGGC GTAATCATGG TCATAGCTGT TTCCTGTGTG AAATTGTTAT CCGCTCACAA TTCCACACAA  
 CATACGAGCC GGAAGCATAA AGTGTAAGC CTGGGGTGCC TAATGAGTGA GCTAACTCAC ATTAATTGCG  
 TTGCGCTCAC TGCCCGCTTT CCAGTCGGGA AACCTGTCGT GCCAGCTGCA TTAATGAATC GGCCAACGCG  
 CGGGGAGAGG CGGTTTGCGT ATTGGGCGCT CTTCCGCTTC CTCGCTCACT GACTCGCTGC GCTCGGTGCT  
 TCGGTGCGG CGAGCGGTAT CAGCTCACTC AAAGGCGGTA ATACGGTTAT CCACAGAATC AGGGGATAAC  
 GCAGGAAAGA ACATGTGAGC AAAAGGCCAG CAAAAGGCCA GGAACCGTAA AAAGGCCGCG TTGCTGGCGT  
 TTTTCCATAG GCTCCGCCCC CCTGACGAGC ATCACAACAAA TCGACGCTCA AGTCAGAGGT GGCGAAACCC  
 GACAGGACTA TAAAGATACC AGGCGTTTCC CCCTGGAAGC TCCCTCGTGC GCTCTCTGT TCCGACCCTG  
 CCGCTTACCG GATACCTGTC CGCCTTTCTC CTTTCGGGAA GCGTGGCGCT TTCTCATAGC TCACGTGTGA  
 GGTATCTCAG TTCGGTGTAG GTCGTTGCGT CCAAGCTGGG CTGTGTGCAC GAACCCCGG TTCAGCCCGA  
 CCGCTGCGCC TTATCCGGTA ACTATCGTCT TGAGTCCAAC CCGGTAAGAC ACGACTTATC GCCACTGGCA  
 GCAGCCACTG GTAACAGGAT TAGCAGAGCG AGGTATGTAG GCGGTGCTAC AGAGTTCTTG AAGTGGTGGC  
 CTAACACTAG CTACACTAGA AGAACAGTAT TTGGTATCTG CGCTCTGCTG AAGCCAGTTA CCTTCGAAAA  
 AAGAGTTGGT AGCTCTTGAT CCGGCAACAA AACCACCGCT GGTAGCGGTG GTTTTTTTGT TTGCAAGCAG  
 CAGATTACGC GCAGAAAAA AGGATCTCAA GAAGATCCTT TGATCTTTTC TACGGGGTCT GACGCTCAGT  
 GGAACGAAAA CTCACGTAA GGGATTTTGG TCATGAGATT ATCAAAAAGG ATCTTCACCT AGATCCTTTT  
 AAATTAATAA TGAAGTTTTA AATCAATCTA AAGTATATAT GAGTAACTT GGTCTGACAG TTACCAATGC  
 TTAATCAGTG AGGCACCTAT CTCAGCGATC TGTCTATTTT GTTCATCCAT AGTTGCCTGA CTCCCGCTCG  
 TGATAGATAAC TACGATACGG GAGGGCTTAC CATCTGGCCC CAGTGTGCA ATGATACCGC GAGAACCACG  
 CTCACCGGCT CCAGATTTAT CAGCAATAAA CCAGCCAGCC GGAAGGGCCG AGCGCAGAAG TGGTCTTGCA  
 ACTTTATCCG CCTCCATCCA GTCTATTAAT TGTGCGGGG AAGCTAGAGT AAGTAGTTCG CCAGTTAATA  
 GTTTGCGCAA CGTTGTTGCC ATTGCTACAG GCATCGTGGT GTCACGCTCG TCGTTTGTA TGGCTTCATT  
 CAGCTCCGGT TCCCAACGAT C

**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\_001033431

**UniProt ID:**

E9Q5R7

**Synonyms:**

Nalp12; PYPAF7

**Summary:**

Plays an essential role as an potent mitigator of inflammation (PubMed:26521018, PubMed:30559449). Primarily expressed in dendritic cells and macrophages, inhibits both canonical and non-canonical NF-kappa-B and ERK activation pathways (PubMed:30559449). Functions as a negative regulator of NOD2 by targeting it to degradation via the proteasome pathway (PubMed:30559449). In turn, promotes bacterial tolerance (PubMed:30559449). Inhibits also the DDX58-mediated immune signaling against RNA viruses by reducing the E3 ubiquitin ligase TRIM25-mediated 'Lys-63'-linked DDX58 activation but enhancing the E3 ubiquitin ligase RNF125-mediated 'Lys-48'-linked DDX58 degradation (By similarity). Acts also as a negative regulator of inflammatory response to mitigate obesity and obesity-associated diseases in adipose tissue (PubMed:30212649).[UniProtKB/Swiss-Prot Function]

## Product images:

