

Product datasheet for **MC223611**

Gnas (NM_001077507) Mouse Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Gnas (NM_001077507) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Gnas
Synonyms:	5530400H20Rik; A930027G11Rik; C130027O20Rik; G; Ga; Galphas; Gn; Gnas1; Gnasxl; GPSA; Gs-; Gs-alpha; Gsa; GSP; N; Nes; Nesp; Nesp55; NespI; Oed; Oed-Sml; Oedsml; P; P1; P2; P3; PHP1A; PHP1B; POH; SCG; SCG6; XL
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Cell Selection:	Neomycin
Fully Sequenced ORF:	>MC223611 representing NM_001077507 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGATCGCC**

ATGGGCATGTTCAACTGCCTCCACGGCAATAATATGTCAGGACAACACGATATCCCCCTGAAGTCGGGG
AGCAGCCCGAGCAAGAACCTTTGGAAGCCCCAGGGCAGCTGCCCCCGGTGCTGGGGCTGGCCAGCCGA
AGAAATGGCGACCGAACCAGGACTCCGAACCGTCTAACAAATGAGCCCGTCCCCGACGAGACTGGCAGTGAG
ATCAGTGGACCCCGAAGACTCCTGACATCCAAAGCCCTGCCAGGCCTTCGAGGAAGTCCGAG
TGGGTGGAGACTACAGCCACCTCCGGAGGAAGCCATGCCATTCGAGACACAACAGCCAGCCTGGGAGA
TTTCTGGCCACCTGGAGCAGCCAGGACCATCTGGGACCCATCAGGCCTCAAGCCTCAACCCAGCG
ATTTTGGAGCCCGGGACCCCACTGGCGGAGCCAGGCCTGGGAGCCTATACCCCCACCAGAAGAAG
CTATGCCATTTGAGTTCAACGAGCCTGCCAGGAGACCATAGCCAGCCTCCCTTGCAAGTCCCAGACCT
TGCGCCAGGAGGTCCGGAAGCATTGGTCCCCAGAGCTTCTCCCGGAGCCCGGAACATCAGATTTGAA
AACGCTGGCTCCGAGAAGACTACAGCCCTCCCCCTGAAGAATCTGTGCCATTTAGGTGGTGGAGAAG
AATTCGGGGGGGATAGCCACCCCGAGGACTCCCGGAGTATCCCAAAATCGGCATTGGCGGGAGTT
CCCAGAGTCGCGGTCCCGAGTGCCTCTGCCTCGCTCCCGCCGAGAACGCGCCTCCCTCTGGGTCCGA
GGCGCCATTGACAGACCATTCGCGAGGCTGTGAGATCTCCTCCTAACTTCGATGCGACAGCCCCCGA
TGGAGATCACCAGACCCCTGCTTGAGATTGGCAGAGCCTCCATTGGGGTCGACGACGACCCGCTGTCAA
TATGGACAGCCCCCAATCGCAAGTGTGGCCGCCATCGAAGTCTCGGGAGCCCGAGTAAGAGCGGAG
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TGAAGGAAAAGTCCCTCTCCGGAGAGAGGGGACGGATCTTCCACCCAGCCTGAAGCAATGGATGCCAA
GCCAGCCCTGCTGCCAAGCCGTCTCTACCGATCTGATGCTGGAGCTCCTACGGATTCGCGATGCTC
ACAGATAGCCAGAGCGATGCCGGAAGACGGGACAGCCCGAGGAACGCCTTCAGATCTCCAGTCGGATC
CTGAAGAACTCGAAGAAGCCCGAGCTGTCCGCGCGATCTGACGGAGGGCAGCCCGAGTCGCCCCAGC



CACTCCTGCCGAGTCCGAGTCTGAAGGCAGCAGAGATCCAGCCGCCGAGCCAGCCTCCGAGGCAGTCCCT
 GCCACCACGGCCGAGTCTGCCTCCGGGGCAGCCCTGTACCCAGGTGGAGCCCGCAGCCGCGGCAGTCT
 CTGCCACCCTGGCGGAGCCTGCCGCCGGGCAGCCCTATCACCCCAAGGAGCCACTACCCGGGCAGT
 CCCCTCTGCTAGAGCCCATCCGGCCGCTGGAGCAGTCCCTGGCGCCCCAGCAATGTCAGCCTCTGCTAGG
 GCAGTCCCGCTAGGGCAGCCTATGCAGTCCACTGGTCTGGGGAGCCAGTCACTCTCAGTACTCCCG
 CCGCTCGGGCATCCCTTCCTGCCCGCCAGCAGTCCCGCCGGGCAGCCTCTGCTGCCCGGCAGTCCG
 TGCTGGCCGGTCAGCCTCTGCCGCCCCAGCAGGGCCCATTTAGACCCCCAGCCCGAGATCCAGGTT
 GCTGACCCGCCTACTCCGCGCCCTCCTCCGCGCCGACTGCCTGGCCTGACAAGTACGAGCGGGCCGAA
 GCTGCTGCAGGTACGAGGCATCGTCTGGCATCTGCGAGATCGAGTCTCCAGTGTAGTTCGGAAGAAGG
 GGCCACCGGTCTTCCAGTGGCTTCTGCGGCGAAACCGCCCTGGCCTGCCCGGAGCCACACGGTC
 GGGAGCAACCCAGTCCGCAACTTCTCACCCGAGCCTTCGGAAGTCTTCCGGTCTATCCGAGTGTACCC
 GATCACGATCCCTCAGCCCGGGAAGGCCAAGGATCCTATGGAGGAGAGGCGCAAACAGATGCGCAAAGA
 AGCCATTGAGATGCGAGAGCAGAAGCGCGCAGATAAGAAACGCAGCAAGCTCATCGACAAGCAACTGGAG
 GAGGAGAAGATGGACTACATGTGTACACACCGCCTGCTGCTTCTAGGTGCTGGAGAGTCTGGCAAAGCA
 CCATTGTGAAGCAGATGAGGATCCTGCATGTTAATGGGTTTAAACGGAGATAGTGAGAAGGCCACTAAAGT
 GCAGGACATCAAAAACAACCTGAAGGAGGCCATTGAAACCATTGTGGCCGCCATGAGCAACCTGGTGCC
 CCTGTGGAGCTGGCCAACCTGAGAACCAGTTCAGAGTGGACTACATTCTGAGCGTGATGAACGTGCCGA
 ACTTTGACTTCCACCTGAATTCTATGAGCATGCCAAGGCTCTGTGGGAGGATGAGGGAGTGCCTGCCTG
 CTACGAGCGCTCCAATGAGTACCAGCTGATTGACTGTGCCAGTACTTCTGGACAAGATTGATGTGATC
 AAGCAGGCCGACTACGTGCCAAGTGACCAGGACCTGCTTCGCTGCCGTCTGACCTCTGGAATCTTTG
 AGACCAAGTCCAGGTGGACAAAGTCAACTCCACATGTTGATGTGGGCGCCAGCGCATGAGCGCCG
 CAAGTGGATCCAGTCTCAATGATGTGACTGCCATCATCTTCGTGGTGGCCAGCAGCAGCTACAACATG
 GTCATTGGGAGGACAACAGACTAACCCGCTGCAGGAGGCTCTGAACCTTTCAAGAGCATCTGGAACA
 ACAGATGGCTGCGCACCATCTCTGTGATTCTTCTCCTCAACAAGCAAGCCTGCTTGTGAGAAAAGTCT
 CGCTGGCAAATCGAAGATTGAGGACTACTTTCCAGAGTTCGCTCGCTACACCACTCCTGAGGATGCGACT
 CCCGAGCCGGGAGAGGACCCACGCTGACCCGGCCAAAGTACTTCATTGGGATGAGTTTCTGAGAATCA
 GCACTGCTAGTGGAGATGGGCGCCACTACTGCTACCCTCACTTTACCTGCGCCGTGGACACTGAGAACAT
 CCGCCGTGCTTCAACGACTGCCGTGACATCATCCAGCGCATGCATCTCCGCAATACGAGCTGCTTAA

AGCGGACCGACGCGTACGCGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCC
 TGGATTACAAGGATGACGACGATAAGGTTTAA

- Restriction Sites:** Sgfl-RsrII
- ACCN:** NM_001077507
- Insert Size:** 3360 bp
- OTI Disclaimer:** Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).
- 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_001077507.2](#), [NP_001070975.1](#)

RefSeq Size: 3737 bp

RefSeq ORF: 3360 bp

Locus ID: 14683

UniProt ID: [Q6R0H7](#)

Cytogenetics: 2 97.89 cM

Gene Summary: This locus has a highly complex imprinted expression pattern. It gives rise to maternally, paternally, and biallelically expressed transcripts that are derived from four alternative promoters and 5' exons. Some transcripts contain a differentially methylated region (DMR) at their 5' exons, which is commonly found in imprinted genes and correlates with transcript expression. This gene has an antisense transcript. One of the transcripts produced from this locus, and the antisense transcript, are both paternally expressed noncoding RNAs, and may regulate imprinting in this region. In addition, one of the transcripts contains a second overlapping ORF, which encodes a structurally unrelated protein - Alex. Alternative splicing of downstream exons is also observed, which results in different forms of the stimulatory G-protein alpha subunit, a key element of the classical signal transduction pathway linking receptor-ligand interactions with the activation of adenylyl cyclase and a variety of cellular responses. Additional transcript variants have been found for this gene, but the full-length nature and/or biological validity of some variants have not been determined. [provided by RefSeq, Jun 2015]

Transcript Variant: This variant (5, also known as GNASXL) is paternally expressed. It has an alternate 5' exon, lacks an internal exon and uses an alternate splice site in the coding region, compared to variant 7. It encodes isoform f (also known as XLas), which has a longer and distinct N-terminus and lacks an internal segment, compared to isoform GNASL. Isoform f is the neuroendocrine-specific G-protein alpha subunit. **Sequence Note:** This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.