

## Product datasheet for **RN217514**

### Nfasc (NM\_001160315) Rat Untagged Clone

#### Product data:

**Product Type:** Expression Plasmids  
**Product Name:** Nfasc (NM\_001160315) Rat Untagged Clone  
**Tag:** Tag Free  
**Symbol:** Nfasc  
**Synonyms:** NF  
**Vector:** pCMV6-Entry (PS100001)  
**E. coli Selection:** Kanamycin (25 ug/mL)  
**Cell Selection:** Neomycin  
**Fully Sequenced ORF:** >RN217514 representing NM\_001160315  
Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCC**CGGATCGCC**

ATGGCCAGGCAGCAGGCGCCACCCTGGGTCCACGTAGCCCTCATCCTCTTCTCCTCAGCCTCGGAGGGG  
CCATTGAGATCCGATGGATCTGACCCAACCCCAACGATCACCAAGCAGTCGGTGAAGGACCACATCGT  
GGACCCCGAGATAACATCCTGATTGAATGTGAAGCTAAAGGGAACCCCGCCCCAGTTTCCACTGGACT  
CGCAACAGCAGGTTCTTCAACATTGCCAAGGACCCACGGGTGTCCATGAGGAGGAGGTCTGGGACCTTGG  
TGATCGACTTCCGAGTGGTGGCGGCCTGAGGAGTACGAAGGGGAGTACCAGTGCTTTGCCCGGAACAA  
ATTCGGACAGCTCTTAGCAACCGCATCCGCTGCAGGTGTCAAATCTCCCTGTGGCCCAAGGAAAC  
CTAGACCCCGTCTGGTCAAGAGGGTGGCCCTTAACCTGCAGTGCAACCCCCACCTGGCCTCCCAT  
CCCCGTCTCTGGATGAGCAGTCCATGGAGCCATCACCCAGGACAAGCGTGTCTCCAGGGTCA  
CAACGGGACCTGACTTCTCCAACGTCATGCTGCAGGACATGCAGACCGACTACAGTGAATGCACGC  
TTCCACTTCAACACACCATTCAGCAGAAGAATCCCTTACCCTCAAAGTCCCTACCACCCGAGGAGTTG  
CGGAAAGAACACCTAGCTTCAATGTATCCCCAGGCACGTCAAGCAGTCAGATGGTACTGCGCGGCATGGA  
CCTGCTGCTGGAGTGCATTGCCTCTGGGTCCCAACACCAGATATTGCATGGTACAAGAAAGGTGGGGAC  
CTCCCATCTGACAAGGCCAAGTTCGAGAATTTTAAACAAGGCTCTGCGCATCACCAATGTCTCTGAAGAGG  
ACTCTGGGAGTATTTCTGCCTGGCCTCCAACAAGATGGGCAGCATCCGGCACACGATCTCGGTGAGAGT  
AAAGGCTGCCCCATACTGGCTGGATGAGCCCAAGAACTTGATCTGGCTCCTGGTGAAGATGGGAGGCTG  
GTGTGTCGAGCCAATGGGAACCCGAAGCCGACCGTCCAGTGGTGGTGAATGGAGACCTTTGCAATCGG  
CACCAACCAACCCGAGGTGGCCGGAGACACTATCATCTCCGGGACACTCAGATCAGCAGCAG  
GGCAGTGTACCAGTGAACACATCCAACGAACATGGCTACCTGCTGGCCAATGCCTTCGTCAGCGTATTA  
GATGTACCCCTCGGATGCTGTCTCCCCGAACAGCTCATCAGGGTGTCTTTACAACCGGACCGGAC  
TGGACTGTCCGTTCTTTGGGTCTCCATTCCAACACTCCGATGGTTTAAAGATGGGAAGGAAGCAACCT  
GGATGGTGGTAACTACCAGTCTACGAAAATGGCAGCCTGGAAAATCAAGATGATTCGCAAAGAGGACCAA  
GGCATCTACACCTGTGTGGCCACCAACATCCTGGGAAAAGCTGAAAATCAAGTCCGCTGGAGGTCAAAG



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ACCCCACCAGGATCTACAGGATGCCTGAAGACCAGGTGGCCAAGAGGGGCACCACAGTGCAGCTGGAGTG  
 CCGTGTGAAGCATGACCCCTCCTTGAAACTCACAGTCTCCTGGCTGAAGGATGACGAGCCACTCTACATT  
 GGAAACAGGATGAAAAAGGAAGATGACTCCCTGACCATCTTCGGAGTGGCAGAGCGGGACCAAGGCAGTT  
 ACACGTGCATGGCCAGCACCCGAGCTGGACCAGGACCTGGCAAAGGCTACCTCACTGTTCTAGGGCGGCC  
 AGACCGACCCAGGGACCTGGAGCTCACTGACCTGGCGAAAGGAGTGTGAGGCTGACCTGGATCCCGGGG  
 GATGATAACAACAGTCCCATCACAGACTACGTTGTCCAGTTTGAAGAGGACCAGTCCAGCCAGGAGTCT  
 GGCATGACCACTCCAAGTTCAGGAGTGTCAACTCAGCCGCTCCATCTGTCCCGTATGTCAACTA  
 TCAGTTCGAGTCATCGCTGTCAACGAGGTTGGGAGCAGCCACCCAGCCTTCCATCCGAGCGGTACCGA  
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 AGATCACATGGACGCCTATGAATGCTACCTCCGCCTTTGGCCCAACCTGCGCTACATTGTCAAGTGGCG  
 ACGGAGAGAAACCCGAGAGACTTGAACAATGTCACCGTGTGGGCTCTCGCTATGTGGTGGGCAGACC  
 CCTGTCTACGTACCCTATGAGATCCGAGTCCAGGCTGAAAATGACTTTGGGAAAGGCCCGGAGCCTGAAA  
 CCGTCATTGGGTACTCGGGGAAGATTATCCAGGGCTGCACCCACTGAAGTTAAATCCGAGTCTGAA  
 CAGCACAGCCATCAGCCTTCAGTGAACCGCTACTCTGACACGGTCCAGGGCCAGCTCAGAGAGTAT  
 CGAGCTTACTACTGGAGGAAAGCAGTTTGTGAAGAACCTGTGGTGTCTCAGAAGAGACAGCAGGCCA  
 GCTTCCCTGGCGACCCGCCCGGGGCTGGTGGTCCGCTGTCCCTACAGTAACCTACAAGCTGGAGAT  
 GGTGTGGTCAATGGGAGAGGTGATGGGCTCGAAGTGAACCAAGGAATTCACCACCCCGAAGGAGTA  
 CCCAGTGGCCCGAGCGTTTCAGAGTCCGACAGCCCAACCTGGAGACCATCAACCTGGAATGGGATCACC  
 CAGAGCACCCCAACGGGATCCTGATTGGATACACGCTCAGATACGTGCCCTTTAATGGAACCAACTGGG  
 AAAGCAGATGGTGGAAAACCTTCTCTCCCAATCAGACCAAGTTCCTCGGTGCAGAGAGCAGACCCCGTCC  
 CGTTACCGCTTCTCCCTCAGTCCAGGACGCAGGTGGGCTCTGGAGAAGCAGCCACGGAGGAGTCCCAA  
 CACCTCCAAATGAAGCTACTCCAACCTCAGCTTACACCAACAACCAGACTGACATCGCCACCCAGGGCTG  
 GTTCATCGGGCTCATGTGTGCCATTGCCCTTCTGGTGTGATCCTGCTTATCGTCTGCTTATCAAGAGG  
 AGTCGTGGCGCAAGTACCCAGTGCAGAGAAAAGAAGGATGTCCCTTGGGCCCTGAAGACCCCAAAGAAG  
 AAGATGGTTCATTTGACTACAGTACAGGAGACAACAAGCCCTGCAGGGCAGCCAGACATCTCTGGACGG  
 CACCATCAAGCAGCAGGAGAGTACGACAGCCTAGTGGACTACGGCAGGGTGGCAGGGCCAGTTCAAC  
 GAAGATGGCTCCTTTATTGGCCAGTACTGTGAGAAAGGACAAGGAGGAGACCGAGGGCAATGAGAGCT  
 CAGAGGCCACATCTCCAGTCAATGCCATCTATCCCTGGC**TGA**

**ACGCGT**ACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT  
 ACAAGGATGACGACGATAAGGTTTAA

- Restriction Sites:** SgfI-MluI
- ACCN:** NM\_001160315
- Insert Size:** 3474 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).
- OTI Annotation:** Clone contains native stop codon, and expresses the complete ORF without any c-terminal tag.
- 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\\_001160315.1](#), [NP\\_001153787.1](#)

**RefSeq Size:** 9785 bp

**RefSeq ORF:** 3474 bp

**Locus ID:** 116690

**Cytogenetics:** 13q13

**Gene Summary:**

This gene encodes an L1 family immunoglobulin cell adhesion molecule with multiple IGcam and fibronectin domains. The protein functions in neurite outgrowth, neurite fasciculation, and organization of the axon initial segment (AIS) and nodes of Ranvier on axons during early development. Both the AIS and nodes of Ranvier contain high densities of voltage-gated Na<sup>+</sup> (Nav) channels which are clustered by interactions with cytoskeletal and scaffolding proteins including this protein, gliomedin, ankyrin 3 (ankyrin-G), and betaIV spectrin. This protein links the AIS extracellular matrix to the intracellular cytoskeleton. This gene undergoes extensive alternative splicing, and the full-length nature of some variants has not been determined.

[provided by RefSeq, May 2009]

**Transcript Variant:** This variant (4) represents use of an alternate promoter and 5' UTR, and has multiple differences in the coding region but maintains the reading frame, compared to variant 1. The resulting isoform (4) includes the third fibronectin type 3 (FNIII) repeat, lacks the mucin-like domain, and lacks two segments in the N-terminal and central regions, compared to isoform 1. **Sequence Note:** The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on alignments.