

## Product datasheet for **MG225449**

### Unc13c (NM\_001081153) Mouse Tagged ORF Clone

#### Product data:

Product Type:	Expression Plasmids
Tag:	TurboGFP
Symbol:	Unc13c
Synonyms:	1500037O19Rik; AU019458; D9Ert414e; Munc13-3; Unc13h3
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-AC-GFP (PS100010)
E. coli Selection:	Ampicillin (100 ug/mL)

**ORF Nucleotide Sequence:** >MG225449 representing NM\_001081153  
Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTACTATAGGGCGCCGGGAATTCGTGACTGGATCCGGTACCGAGGAGATCTGCC  
GCCGCGATCGCC

ATGGTGGCCAGTCTCTTTAAGAGCTTGATTTAGCTTATATACATAAGCTTTGCAAAGGAATGTTTACAA  
AGAAATTTGGGAAATACAACAAAAAGAAAGAGAATCGTCAGCAGAAAAAGATCAAGATTTTCCAAGTGC  
TGGCCACACCAACCCCTAAATTGTCAAATGCCCTGAAAAGCACTGTAAAAAGATTGCAAAGTGTCTCA  
TCTACCCGGAATCTCCATAGAAGACGAAGAGGGTCAAAAGACTTCTCCCTGTACCCACATTCAGTT  
ACCGAGTTGCTATCGCCAATGGCCTGCAAACAGCCGTGACCAACAGCGATGAGGATCTGCTCCAGGAGCT  
GTCTTCCATCGAGAGCTCGTACTCTGAATCCTTCAATGAGCTAAGGAGCAGCACGGAAAAACAGTCCAA  
TCAACACACACCATGCCTGTCCGACGTAACAGGAAGAGCTCGAGCAGTTTGGCACCTTCTGAGGGCAGCT  
CAGATGGGGAGCGCACCTCCACACCTTAAACTGGGTGCTCTACGAAACTGAGGAAATGGAAGAAGAG  
CCAAGAGTGCCTCTCCTCAGACTCGGAGTTGAGCACCGTCAAGAAAACCTGGGGCATAGAAGCAAATCA  
TTAGATAGAAGTGGCCGAACCCAAAGACAAATGTCTTAGAGCCAGGGTTCAGTCTCCTTGCTGCATTA  
GCCAAACCCACGATGTCATGGAATGATCTTTAAAGAGCTTCAGGGTATAAGTCAGATCGAAACAGAAT  
TTCCGAACTCCGAGGGCACGTGAATGCTCTCAAGTACTCTATCGATGAAATCTCCAGCAGTGTGGAGGTA  
GTCCAGAGCGAGATCGAACAGCTTCAACAGGGTTTGTCCAGGCTCGAAGGGAAACTCGAGACATCCACG  
ATTACATTAACACTTGGGTACATGAGGAGTAAGGTAAGCCTGAGATTTTAAATGTGCCTGAAGAAAG  
ACACGAGTACGTGGAAAGTGTGGTGTACCAAAATCTTATAGATAAAATGGGTTTCTCAGATGTACCGAAT  
GCTATTAATTTGAATTTGCTCAGAGGATAGGGCAGCAAAGGGACTGCCAAAATGCAAACCTCGACCCA  
TCCTAGTGTACTTTGAAACCCCTCAGCAAAGGGATTCTGTCTTAAAGAAATCGTATAAGCTGAAAGGTAC  
AGGCATCGGGATCTCCACAGATATTCTAATTATGATATCAGAGAACGGAAAGAAAAAGGGGATTGCGCA  
TCTTCCAGACATACGAAAGCATGGATATGAAGTTATCCACGCCAGAGCCAAAGCCAAGAAGAAGCCTT  
GGCTGTACCCCAATGACAGTACAGAGAGCTGGAGTACAGACCTCAGCAGAAGCAGCTATGCTGACTCTCC  
GGCAAAGGGAAGTTCTTCTAAATCCAGCTCAAAGTACACAGTGTGCTCGGTCCAAGAAACAAAGCAGCTAAC  
AGCAGAATCACAGAAGTCAGACTACAATAAAGCGCCCTCCCAACCGCCAGCATCAAGTACCCAGAAA



AGCAAACACCTCACTATGTGGAGGCCACACCACCACTCTGGCATTACAGAGTGACTTCTTCACTCTTAA  
ACTTAGTCGCTCCGAATCGGATTTCTCCAAATTGTGTCAATCTTATTCAGAGGATTTTTCTGAAAGCCAG  
TTTTTCTGTGCAACTAATGGGAGCTCCCTCCTTCTCGCTGACCGGGAACCTTGGCAGAGGAAACAGG  
AGGGCATGCCTGCCTTGTACCACCGTCTTCAGGACCAGGGTTTGGATGAGACTATTCCAGCAGTGCCAGG  
GCAGGCTGAAATTGAGAATACAGAAACAGTAGACAGCGGTATGAGCAACAGCATGGTGTGCATCTGGA  
GACCGAAGCAATTACAGCGTTCTCAGCTCTCATTGCACGAGGACCTGTCTCCATGGAAGGAGTGGAATC  
AAGCAGGGCAAGGAAGTATGATGTTGGCTTAGATTCACTACCCAGGAACCTTTTGACTACGATACCAA  
CAGTTTGTCTGACCAACAGCTGGATCTCTTAGCAAAGACCTAGATGACCTTGGCAAGTGCCACAGCGAC  
CTGCAAGATGACTCTGAAAGCTATGATTTAACCAGGATGACAACCTCTCTCCATGCCCTGGATTGGATA  
ATGAACCACAAGGCCAGTGGTTGGCCAGTACGACTCTTATCAGGAAGCCAATTCTAATGACCTGTATCC  
CAACCAAGCCATCCATCCATGATGATAGAAGTCAAAGCGAACTGCAAAGTGATGATTCAGAGGGGGCC  
CAGCCCAAGTCTGGCAGTGGCTAAGCATTGACCTTTCTGATAAAGACTTTTAAAGTCCCAAAATTTG  
GATCTACTCTTCAGAGAGCAAAGTACGCCCTGGAAGTGGTGTGGAACAAGAGCACACAGAGTCTCAGTGG  
GTGTGAGGACAGCGGCTCTTCTTAATGGGGAGATTTCCGACATTATCACAATCCACCAGCAAGGAGTCA  
AGTACCACCTTGACTCTGATATCTACACGGAACCTATTACTATAAAGCCGAGGAAGAGGAGGACTACT  
GTGAACCAGTGGCTGACAGTGAACAGATTATGTTGAAGTGTGGAGCAGGTCCTAGCTAAGCTAGAAAA  
CAGGACTAGTATCACTGAAGTAAACGAACACATTAAGACTATGACCACCTTCTATGAAACTCCTTAC  
GAGACCCCGCAAGATGAGGGCTATGATGGCAGGCTGATGACATCATTAGTGAAGGAGAAGTGGAAACCT  
TAAATGAACCAGCAGTTGAAATGGAATTAGCAGAAGTGAAGCAAAACCTCCCTGCAGAGTCACTGGA  
GGTCAATGAAGCCAAAGAGAATCAGGCCTTCTTCAAAGAAGCAGCTTAAAGGGCTATAAAGAGCAGATG  
GCAGAGCTGGAAGAGAAGATCCTGGCTGGAGATAGCAGCTCTATGGATGAAAAGGCTCGAATAGTAAGTG  
GCAATGATTTGGATGCTTCCAAATCTCCGCACTCCAGGTGTTCCGGTGGGGCTGGGCGTGGACTGTATGG  
TATTGATAGCATGCCTGACCTCCGAGGAAAAAGACATTGCCTATTGTGCGGGATGTGGCCATGACCTG  
GCTGCCCGGAAATCTGGTCTCTCCCTTGCTATGTTATCAGGACATCACTGAATAATGAGGAGCTGAAAA  
TGCATGTCTTCAGGAAGACCTTGCAGGCATTGATCTACCCCATTTCTTCCACCCTCCCAAACTTCGA  
GGTCTGGACGGCTACCACCCACCTACTGCTATGAATGTGAAGGGCTCCTGTGGGGCATCGCAAGGCAA  
GGCATGAAGTGTCTGGAGTGTGGGTGAAATGCCACGAGAAGTGTGAGGACCTGCTCAACGCCGACTGTC  
TGCAGCGAGCAGCAGAGAAGAGCTCTAAGCACGGTGCAGGAGCAAGACCCAGACTATCATCACAGCCAT  
GAAAGAAAGAAATGAAGATCAGGGAGAGAAACAGACCAGAGGTGTTTGAAGTCATTGAGGAAATGTTCCAG  
ATCTCAAAGAGGATTTTGTGCAGTACACAAGGCTGCCAAGCAGAGCGTGTGGATGGGACGTGCAAGT  
GGTCTGCAAAAATACCATTACAGTGGTTCCGCACAAGGCTTGCAGGCAAAAGACAAAACAGGGTCCAG  
CGACCCATATGTGACAGTTCAAGTTGGCAAAAATAAAGAAGAACGAAAATCTTTTGGAAACTTGAAT  
CCAGTGTGGGACGAGAAGTCTTCTTTGAGTGTCAACTCGACAGATCGAATCAAGGTGAGAGTATGGG  
ATGAAGATGATGACATTAATCCAGAGTTAAGCAGCACTTCAAGAAGGAGTCCGATGACTTTCTGGGACA  
GACGATTGTGGAAGTCAAGACTCTAAGTGGTGGATGGATGCTGGTACAACCTAGAGAAAACGGACAGAT  
AAGTCACTGTGTCTGGGGCCATCCGACTAAAAATCAATGTGGAGATAAAGGGAGAGAAGAGAAGTCCGAC  
CATACCATATAACAATCACTTGTTTACACGAGAACCTGTTCCATTACCTGACGGAAGTGAAGTCTAATGG  
CAGTGTAAAAATACCAGAGGTCAAGGGCGATGAAGCCTGGAAGGTTTTCTTTGATGATGCGTCTCAGGAA  
ATAGTGGATGAATTTGCCATGCGTTATGGGGTTGAGTCCATTTATCAAGCTATGACGCACTTTTCATGTC  
TGTCTTCTAAGTACATGTGCCCGGGTGTCCCTGCAAGTATGAGCGCCCTGCTGGCGAACATCAATGCCTT  
CTATGCTCATAACAATGCTCCTCACTAACGTGCAGGTTTCTGCCTCTGATCGCTTTGCTGCTACAACTTT  
GGGAGGGAAAAATTCATAAACTACTGGACCAGTTGCATAACTCCTTAAGGATTGACCTGTCAAAGTACA  
GGGAAAATTTCCAGCCAGCAACAGTGAAGACTACAGGACCTAAGTCCACTGTGGACCTGTTGACCAG  
CATCACCTTCTCCGGATGAAGGTCTGGAGCTGCAGAGCCCCCAAGGCCAGTGCAGTTGTGAAGGAC  
TGTGTGAGGCTTGTCTGGATTCCACATACAAGTATATCTTCGACAACCTGCCATGAACTCTATTCTCAGC  
TGATAGACCCGAGCAAGAAGCAGGATGTCCCTCGAGAAGACCAGGGACCAACCACCAAGAACCTGGACTT  
CTGGCCACAGTTGATTACACTGATGGTCACTATTATTGATGAGGATAAACTGCCTACACCCTGTTCTC  
AATCAGTTTCCACAAGAACTAAACATGGGGAAGATAAGTGCAGAAATCATGTGGTCTCTTTTTGCTCTCG  
ATATGAAATATGCTCTAGAGGAACATGAGAAGCAACGCTTATGCAAGAGCACCGATTATATGAATTTGCA  
TTTTCAAAGTGAATGGTTTTATAATGAGTATGTACGAGAACCTCCTGCCTTCAAGGATGCTGTTCCAGAA  
TATTCCTGTGGTTTGGACCTTTTGTATGCAATGGCTAGATGAAAATGAAGATGTGCCATGGAGTTCC  
TTCATGGAGCACTGGGAAGAGACAAAAAAGATGGATTCCAGCAGACATCGGACCACGCCCTTTCTCCTG  
TTCTGTGGTTGATGCTTTGCTCAGCTTAACCAGAGCTTTGAGATCATTAAAGAACTGGAATGCCCAAT  
CCTGAAGCCCTATCACACTAATGAGACGATTTGCAAAGACCATCAATAAGGTTCTGGTCCAGTACGCTG  
CGATCGTATCAAAAGACTTCACTTCACTGCGATAAAGGAGACAGTGCCTTGCATATTGATGAACAACAT  
CCAACAACCTGCGGTCCAGCTGAAAAAATGTTTGAATCCATGGGTGGGAAAGAGCTAGACCCTGAAGCT

AGCACTATTCTGAAAGAACTTCAAATCAAACCTCAACGGGGTTCTGGATGCACTCAGCATCACTTATGGAG  
AAAGTTTCCAACCTACAATTGAAGAGTGTATAAAGCAGATGGTGCCGAGCTGAATCAGATGCGAGCCAA  
CGGGAACAGCACAGCGAATAAGAACAGTGCAGCGATGGATGCAGAGATTGTGCTGAGGCCGCTCATGGAC  
TTCTTGGATAAGACATTAAGTCTTTCAGCAAAAATTTGTGAGAAAACGGTCCTGAAGCGAGTTCTGAAGG  
AGCTCTGAAAATTAGTACTTAACAAGATAGAGAAGCAGATTGTTCTCCCTCCACTGACAGACCAAACGGG  
GCCTCAGATGATCTTCATTGCTGCTAAAGATCTTGGACAGTTATCCAACTGAAGGAGCATATGATTGGA  
GAAGATGCCAAGGGTCTGACACCAAGACAATGTGCAATAGTGGAGGTGGTCTTGCCACTATAAAGCAAT  
ATTTTCATGCAGGAGGAAATGGCCTGAAGAAGATTTTCTGGAGAAAAGCCAGATCTTCATTCCCTGAG  
ATACGCTCTCAGCCTCTACACAAAACCTACTGATGCTTTGATCAAGAAGTTCATAGAGACACAGGGCTCA  
CAGAGTCGTTCCCTCAAAGATGCTGTAGGCCAGATCTCTGTCCATGTGGATGTCACCACCACCCAGGGA  
CTGGAGACCATAAAGTCACTGTGAAAGTAATTGCGATCAATGACCTAAACTGGCAGACCACCGCCATGTT  
CCGCCCTTTGTGGAAGTGTGATGCTGGGACCCAGCCTAGGGGACAAGAAGAGAAAACAAGGGACAAAG  
ACAAAAAGCAACACATGGTCACCGAAGTACAATGAAACCTTCCAGTTCATTCTGGGTAAACGAGAACCGCC  
CAGGAGCCTATGAACTTCACTCTCGTTAAGGACTACTGCTTTGCCCGAGAAGATCGAATAATTGGAAT  
GACAGTCATTGAGTACAGAACATAGCAGAGAAAAGGGAGCTACGGGCATGGTACCCTCTTCTGAAAAAT  
CTCTCTATGGATGAAACTGGTTTAACCATCCTCAGAATACTCTCTCAGAGGACCAGCGACGATGTCGCTA  
AAGAGTTTGTGAGACTTAAGTCAGAAACGAGGTCTATTGAAGAGAGTGCC

ACGCGTACGCGGCCGCTCGAG - GFP Tag - GTTTAA

**Protein Sequence:**

>MG225449 representing NM\_001081153  
 Red=Cloning site Green=Tags(s)

MVASLFKSLILAYIHKLCCKGMFTKKLGNTTKKKENRQKKDQDFPTAGHTKPPKLSNALKSTVKKIAKCS  
 STRNFSIEDEEGHKDFSLSPTFYSYRVAIANGLQTAVTNSDEDLLQELSSIESSYSEFNFELRSSTENQVQ  
 STHTMPVRRNRKSSSLAPSEGSSDGERTLHTLKL GALRKLKRWKKSQECVSSDSELSTVKKTWGIRSKS  
 LDRTARNPKTNVLEPGFSSSGCISQTHDVMEMIFKELQGISQIETELSELRGHVNALKYSIDEISSSVEV  
 VQSEIEQLRTGFVQARRETRDIHDYIKHLGHMGSKVSLRFLNVPEERHEVYVESVYQILIDKMGFSDVNP  
 ATKIEFAQRIGQQRDCPNAKPRPILVYFETPQQRDSVLKKSYPKGTGIGISTDILTVDIRERKEKGVLP  
 SSQTYESMDMKLSTPEPKAKNAWLPNDSDRELESDLRSRYADSPAKGSSSKSSKSHSARSKNKAAN  
 SRTSQKSDYNKRPSQPPASSTPEKQTPHYVEATPPLWHSQSDFFTLKLSRSEDFSKLCQSYSEDFSESQ  
 FFCRTNGSSLLSSSDRELWQRKQEGMPALYHRLQDQGLDETIPAVPGQAEIENTETVDSGMSNSMYCASG  
 DRSNYSQSLSLHEDLSPWKEWNQAGQGTDDVGLDSSTQEPFDYDTNSLSDQQDLSSKDLDDL GKCHSD  
 LQDDSESYDLTQDDNSSPCPLDNEPQGWVQYDSYQANSNDL YPNQSHPSMMYRSQSELQSDDSEGA  
 QPKSWHSRLSIDLSDKTFKFKFGSTLQRAKSALEVVWNKSTQSLSGCEDSGSSLMGRFRTL SQSTANES  
 STTLDSDIYTEPYYYKAEEDDYCEPVADETDYVEVMEQVLA KLENRTSITEVNEHIKDYDHPYSYETPY  
 ETPQDEGYDQADDIISEGELETLENEPAVEMELAEDENQNLPAESLEVMKPKRIRPSFKEAALRAYKKQM  
 AELEEKILAGDSSSMDEKARIVSGNDLDASKFSALQVFGGAGRGL YGIDSMPDLRRKKTLP IVRDVAMTL  
 AARKSGLSLAMVIRTSLNNEELKMHVFRKTLQALIYPISSTTPHNFEVWTATPTTYCEGELLWG IARQ  
 GMKCLECGVKCHEKCQDLLNADCLQRAAEKSSKHGAEDKTQTIITAMKERMKIRERNRPEVFEIQEMFQ  
 ISKEDFVQYTKAAKQSVLDGTSKWSAKITITVVSQAQLQAKDKTGSSDPYVTVQVGNKRRTKTFIGNLN  
 PVWDEKFFFECHNSTDRIVRVWDEDDDIKSRVKQHFKKESDDFLGQTI VEVRTLSGEMDVWYNLEKRTD  
 KSAVSGAIRL KINVEIKGEEKVAPYHIQYTCLEHNFHYL TEVKSNGSVK IPEVKGDEAWKVFDDASQE  
 IVDEFAMRYGVESIYQAMTHFSCLSSKYMCPGVPVMSALLANINAFYAHTTVSTNVQVSASDRFAATNF  
 GREKFIKLLDQLHNSLRIDL SKYRENF PASNSERLQDLKSTVDLLTSITFFRMKVLELQSPPKASAVVKD  
 CVRACL DSTYKYIFDNCHELYSQLIDPSKKQDVPREDQGP TTKNLDWFQQLITLMVTI IDEDKTAYTPVL  
 NQFPQELNMGKISAEIMWSL FALDMKYALEEHEKQRLCKSTDYMNLFHKVKWFYNEYVRELPAFKDAVPE  
 YSLWFEPFVMQWLDENEDVSMFLHGALGRDKKDG FQQTSDHALFSCSVVDVFAQLNQSF EIIKKLECPN  
 PEALSHLMRRFAKTINKVLVQYAAIVSNDFSSYCDKETVPCILMNNIQQLRVQLEKMFESMGGKELDPEA  
 STILKELQIKLNGVLDAL SITYGESFQLTIEECIKQMGAE LNQMRANGNSTANKNSAAMDAEIVLRPLMD  
 FLDKTL SLSAKICEKT VLRVLEKELWKLVLNKIEKQIVL PPLTDQTGPQMIFIAAKDLGQLSKLKEHMIR  
 EDAKGLTPRQCAIVEVVLATIKQYFHAGGNLKKNFLEKSPDLHSLRYALSLYTQTTDALIKKF IETQGS  
 QSRSSKDAVGQISVHVDVTTTPGTGDHKVTVKVIAINDLNWQTAMFRPFVEVCMLG PSLGDKKRRKQGTK  
 TKSNTWSPKYNETFQF ILGNENRPGAYELHL SVKDYCFAREDR IIGMTVIQLQNI AEKGSYGAWYPLLKN  
 LSMDETGLTILRILSQRTSDDVAKEFVRLKSETRSIEESA

TRTRPLE - GFP Tag - V

**Restriction Sites:**

Sgfl-MluI



RefSeq Size: 8081 bp

RefSeq ORF: 6633 bp

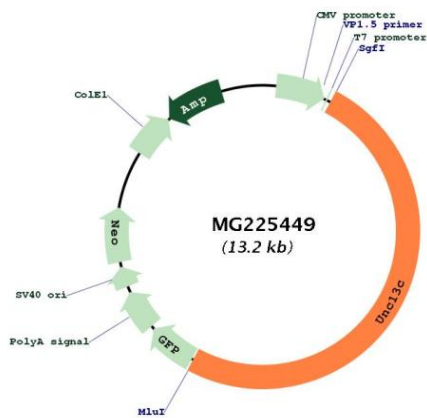
Locus ID: 208898

UniProt ID: [Q8K0T7](#)

Cytogenetics: 9 40.8 cM

**Gene Summary:** May play a role in vesicle maturation during exocytosis as a target of the diacylglycerol second messenger pathway. May be involved in the regulation of synaptic transmission at parallel fiber - Purkinje cell synapses.[UniProtKB/Swiss-Prot Function]

**Product images:**



Circular map for MG225449