

## Product datasheet for MR226679

### Ranbp2 (NM\_011240) Mouse Tagged ORF Clone

#### Product data:

|                          |   |
|--------------------------|---|
| Product Type:            | Expression Plasmids   |
| Product Name:            | Ranbp2 (NM_011240) Mouse Tagged ORF Clone                                   |
| Tag:                     | Myc-DDK   |
| Symbol:                  | Ranbp2  |
| Synonyms:                | A430087B05Rik; AI256741; NUP358   |
| Vector:                  | pCMV6-Entry (PS100001)  |
| E. coli Selection:       | Kanamycin (25 ug/mL)  |
| Cell Selection:          | Neomycin  |
| ORF Nucleotide Sequence: | >MR226679 representing NM_011240<br>Red=Cloning site Blue=ORF Green=Tags(s) |

TTTTGTAATACGACTCACTATAGGGCGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCCGCGATCGCC

ATGAGGCGTAGCAAGGCCGACGTGGAGCGGTACATCGCCTCGGTGCAGGGCTCCGCCCGTCGCCCCGAG  
AGAAGTCAATGAAAGGATTCTATTTTGC AAAGCTGACTATGAAGCTAAAGAATATGATCTTGCTAAAAA  
ATACATTTCTACGTATATTAATGTGCAAGAGAGACCCTAAAGCCACAGATTTTGGGCTTCTTTAT  
GAAGTCGAAGAAAATATAGACAAAGCTGTAGAATGTTATAAGCGTTCAGTGGAATTA AACCCAACGCAAA  
AAGATCTTGTTGAAGATTGCAGAATTGCTTTGTAAAAATGATGTGACTGATGGAAGAGCAAAACTG  
GGTTGAAAGAGCAGCCAACTTTTCCAGGAAGTCCTGCCATTTATAAGTTAAAGGAACAACTTTTAGAT  
TGTAAGGTGAAGATGGATGGAATAAACTTTTGACTTGATTCAGTCAGAACTTTATGCAAGACCTGATG  
ATATTCATGTAATATCCGACTAGTAGAGCTTTACCGTTCAAATAAAAGATTGAAGGATGCTGTGGCCCA  
CTGCCATGAGGCAGACAGGAACACAGCTTTACGTTCAAGTTTGAATGGAATTTATGTGTTGTACAGACT  
CTTAAGGAATATCTGGAGTCTCTGCAGTGTGGATTCTGATAAAAAGTACTTGGAGAGCAACCAATAAG  
ACTTACTTCTAGCCTATGCTAACCTTATGCTACTTACACTTTCCACCAGAGATGTGCAGGAAGGTAGAGA  
ACTATTGAAAGCTTTGATAGTGCTCTTACGCTGTGAAGTCTTCTGTGGGTGGAATGATGAGCTATCA  
GCTACTTTCTAGAAAACAAAAGGACATTTCTACATGCATGTTGGTTCTCTGCTTTTGAAGATGGGTGAGC  
AGAGTGACATCCAGTGGCGAGCTCTTTCTGAGCTGGCTGCATTGTGCTATCTTGTAGCATTTTCAGTTCC  
CAGACCAAGGTTAAATTAATAAAAGGAGAAGCTGGACAAAATCTCCTGGAAACGATGGCCACGATCGC  
CTTAGCCAGTCAGGCACATGTTGTTAACTTAAGTCGTGGCAAGCAAGACTTTCTAAAGGAAGTTGTGG  
AATCTTTTGTAAACAAAAGCGGCCAGTCTGCTCTCTGTGATGCTCTGTTTCCAGTCAGTCATCCAAGGA  
GAGATCCTTTCTGGGAATGATGATATTGAAACCTTGATGGACAAGTCCAGACCCTGATGACTTGGCT  
AGATATGATACTGGTGCCGTTTCGAGCACATAATGGTAGTCTTTCAGCACCTTACCTGGCTTGGCTTACAAT  
GGAATTCAGTGTCTACTTTACCAGCAATTCGAAAATGGTTAAAACAGCTCTTCCACCATTTCCTCAGGA  
GACGTC AAGGCTTGAACAAAATGCACCTGAATCAATCTGTATTTAGATCTTGAAGTATTCTTACTTGG  
GTTATATATACCAGTCACTTACAATTAAGGAGAAGTGAATTTCTCACCACACTTCTATCAACCATTGT



[View online »](#)

GTTTGCCACTTCCTGTGTGCAGACAGCTTTGTACAGAGAGACAGAAGACGTGGTGGGATGCAGTTTGTAC  
TCTGATTCACCGAAAGGCGCTACCTGGAACCTCAGCAAAATTGCGACTTCTAGTTCAACGTGAAATAAAC  
AGTCTGAGAGGCCAGGAAAAGCACGGCCTTCAACCTGCTCTGCTTGTGCATTGGGCCAGAGTCTTCAAA  
AAACAGGCAGCAGTCTTAATTCTTTTTATGATCAACGGGAATACATAGGCAGAAGTGTTCATTATTGGAG  
GAAAGTTTTACCATTGTTGAAGATGATCAGAAAAGAAGAATAGTATTCCTGAACCTATTGATCCTCTGTTT  
AAACATTTCCATAGTGTGGATATTCAAGCATCTGAAATGGTGAATATGAAGAAGATGCACACATAACTT  
TTGCTATATTGGATGCTTTAATGGAAATATAGAAGATGCTATGACTGCTTTTTGAATCTATTA AAAAATGT  
AGTTTTCTATTGGAACTTGCACCTGATTTTCCACAGGAAAGCAGAAGACATTGAAAATGATGCCTTGCT  
CCTGAAGAACAAGAAGAATGCAAAAATTATCTGAGAAAGACTAGGGACTACCTGATAAGGATTTTAGATG  
ATAGTGATTCAAATACTTCTGTGGTTCAGAAAATGGCTGTTCCCTTGAGTCTGTAAAAGAGATGCTGAA  
CTCAGTTATGCAGGAACCTGAAGACTACAGTGAAGGAGTACTCTTTATAAAAAATGGTTGCTGGCGAAGT  
GCTGATTCAGAGTTAAAGCATTCAACACCATCTCCTACCAAAATTTCTCTATCACCAAGTAAAAGTTACA  
AGTATTCTCAAAGACACCACCTCGATGGGCAGAAGATCAAAAATCTTTACTGAAAATGATCTGCCAGCA  
AGTAGAGGCCATTAAGAAAGAGATGCAGGAATTGAAGCTAAATAGCAATAACTCTGCATCCCCTCATCGC  
TGCCCGCAGAGCCTTACGGACAAGACCAGCTCTGACGGATATCAGGGCTCACAACTTTTCACGGGG  
CTCCACTAACAGTTGCAACTACTGGCCCATCAGTATATTATAGTCAGTCACCTGCATATAACTCCCAGTA  
TCTTCTCAGACCAGCTGTAATGTAACCTCCCAAGGGTCCAGTTTATGGCATGAATAGGCTTCCACCC  
CAACAACATATTTATGCATATTCACAACAGATGCACACACCACAGTGCAAAAGTTTCATCTGCTTGTATGT  
TCTCTCAGGAGATGTATGGTCTCCCTTGCCTTTGAGTCTCTGCAACAGGAATTTCTATCACCTAGGGG  
TGATGATTATTTAATTACAATGTTCAACAGACAAGCACAATCCACCTTTGCCAGAACCAGGGTATTTT  
ACAAAGCCTCCACTTGTAGTCCAGCTTCAAGGTGAGCAGAATCAAAGTTTATAGAATTTGGAAAATCCA  
ATTTCTGTTCCAGCCATGCAGGGTGAAGTAATAAGGCCACCCTTGACAACACCAGCACATAACAACAGCC  
AATCCCTTTAAATTTAACTCAAATTTCAAATCAAATGATGGCGACTTCAATTTTATCACCACAGGTA  
GTGGCCAGCCTCCTTCAACAGCTTATAGCAATAGTGAAGTCTTCTAGGCTTCTGACTTCAGATAAGC  
CTTTACAAGGAGATGGCTATAGTGGACTAAAGCCAATATCTGGCCAAGCTAGTGGGTCTCGGAATACATT  
CAGTTTTGGAAGCAAAAATACACTTACAGAAAACATGGGCCAAAATCAGCAGAAGAATTTTGGTTTTCAT  
CGCAGTGATGACATGTTTGTCTTTCACGGTCCAGGGAAGTCTGTATTTACAACAGCTGCTCCGAGCTGG  
CAAATAAAAGTGCATGAGACAGATGGAGGAAGTGTCTATGGGGATGAGGAGGATGATGGCCCTCACTTTGA  
GCCTGTGGTACCTCTTCTGACAAGATAGAAGTAAAAACTGGCGAGGAAGATGAAGAAGAATTTCTTTTGC  
AATCGTGCAAAGTTGTTCCGTTTTGATGGTGAATCCAAAGAATGGAAGGAACGTGGGATAGGCAATGTAA  
AAATACTAAGGCATAAAACATCTGGTAAAATTCGCCTTCTGATGAGACGAGAGCAAGTATTAAGATCTG  
TGCAAATCATTACATAAGCCAGATATGAAACTGACGCCAAATGCTGGCTCAGATAGATCTTTTGTATGG  
CATGCCCTTGATTATGCAGATGAATTGCCAAAACAGAACAGCTTGAATTAGATTCAAAAACCTCTGAGG  
AAGCAGCACTATTTAAGTGTAAATTTGAAGAGGCCAGAACATTTTAAAAGCCTTAGGAACAAAACATC  
TACAGCACAAAATCATACTCTTGAATTTGAAAGGAGTCTGCAACTCAGGATAACAAGGATATTTGCAAA  
GCTGATGGTGGAACTTGAATTTGAAATCCAGATTGTAAGAAGGAAGGGCCTTACTGGAATTTGTAACA  
GCTGCTCCTTTAAGAATGCTGCAACTGTAAGAAATGTGTATCATGCCAGAATACAAACCAACAGTAA  
CAAAGAGCTCCTAGGCTCCTCATTAGTTGAAAATGGCTTTGCTCCTAAAACCTGGCCTGAAAAATGCTCAA  
GATCGATTTGCAACAATGACTGCAAATAAAGAAGGCCATTGGGACTGCAGTGTGTTTGTAGTAAGAATG  
AACCCACCGTCTCTAGGTGCATTGCATGTCAGAACACAAAAGTCTGCGTCTTCGTTTTGTTCAAACCTCTTT  
CAAATTTGGTCAGGGAGATCTTCTAAGCTGTTGACAGTGATTTTCAAGTCTGTTTTTTCTAAAAAGAA  
GGTCAGTGGGAATGCAGTGTATGCCTAGTCCGAAATGAAAGAAGCGCTAAAAAATGTGTTGCTTGTGAGA  
ACCCAGGAAAACAGTTCAAAGAATGGCATTGTAGTTTGTGCTCGGTGAAAAATGAAGCACATGCTATAAA  
ATGTGTTGCTTGTAAATCCTGTTACACCAAGTTTATCTACTGCACCTCCCTCTTTAAGTTTGGTACC  
TCAGAGATGAGTAAGCCTTTCAGGATTGGATTTGAGGGTATGTTCCGAAGAAAGAAGGACAGTGGGACT  
GTAGCTTGTGTTTTGTCGAAAATGAAGCCAGTGTACTACTGTATTGCTTGTGAGTATCCAAATAAGCA  
GAATCAGCCTACATCTTGTGATCAGCTCCTGCCTCTTCAGAGACAAGCAGGTCTCCAAAGAGTGGATTT  
GAAGGCTTATCCCAAAAAGGAAGGAGAGTGGGAGTGTGCTGTTTGTCTGTACAAAATGAGAGCTCTT  
CCCTAAAATGTGTGGCTTGTGAAGCCTCAAGCCAACCTATAAGCCTCATGAAGCTCCTTCAGCTTTCAC  
AGTGGGCTCAAAGTCAAGTCAAATGAATCTGCAGGAAGTCAAGTGGGAACAGAATTCAAAAGTAACTTT  
CCAGAAAAGAAATTTAAAGTTGGCATTTCAGAGCAGAAATTTAAATTTGGGCATGTGGATCAAGAAAAAA  
CACCTTCTTTGCCTTTCAGGGTGGCTCTAATACAGAATTAAGTCAATCAAGGATGGATTTAGTTTTTG

CATTCTGTATCTGCTGATGGGTTAAATTTGGCATTTCAGGAAAAGGAAATCAAGAGAAAAAGAGTGAA  
 AACATCTTAAAAATGACCCTAGTTTCCAAGCTCATGATACTAGTGGTCAGAAGAATGGTAGTGGTGGG  
 TCTTTGGTCAGACAAGCAGCACCTTTACCTTTGCAGATCTTGCAAAGTCAACATCAAGAGAAGGGTTTCA  
 ATTTGGCAAGAAAAGCCCTAATTTTAAAGGGATTTTCAGGTGCAGGAGAAAAATTATTCTCGTCACAAAGC  
 GGCAAAGTGGCTGAGAAAAGCAAAATACGTCTTCTGATCTCGAGAAAAGATGATGATGCATATAAGACTGAGG  
 ACAGTGATGACATCCATTTTGGCCAGTGTGAGATGCCAGAAAAGGTGGAAGTAAACAGGAGAAGA  
 AGATGAAAAAGTTTTGTATTCACAAAAGGTGAAATTTAGATTTGATGCTGAGATAAGTCAAGTGGAAA  
 GAAAGGGGTTTGGGGAATTTAAAAATTCTCAAAAATGAAGTCAATGGTAAACTGAGAATGCTGATGCGAA  
 GAGAACAAAGTGCTAAAAGTTTGTGCCAATCATTGGATAACCACTACAATGAACCTGAAGCCTCTCTCGGG  
 GTCAGATAGAGCATGGATGTGGCTAGCTAGTGATTCTCCGATGGTGACGCGAAACTGGAACAAGTGGCT  
 GCAAAGTTTAAAACGCCAGAGCTAGCTGAAGAATCAAGCAGAAAATTTGAGGAGTGTCAACGTCTTCTCT  
 TAGACATTCACCTTCAGACACCCATAAAGTGTAGATACTGGCAGGGCTGCTAAATTAACAGAGAGC  
 TGAGGAAATGAAGAGCGGCTGAAAGATTTCAAAAACCTTTTGGCAAATGATCAAGTAAAGGTCACCGAT  
 GAAGAGAATGCAAGTTCAGGTGCAGATGCTCCCAGTGCATCAGACACCACAGCCAAGCAAAATCCTGATA  
 ACACCGGGCCTGCCTTAGAATGGGATAACTATGACTTACGGGAGGACGCCTTGGATGACAGTGTGAGCAG  
 CAGCTCAGTTCATGCTTCTCCATTGGCAAAGCAGCCTGTGCGGAAAAACCTCTTCCGCTTTGGTGAGTCC  
 ACTACAGGATTTAACTTCAGTTTTAAATCTGCTCTGAGTCCATCTAAGTCTCCAGCCAAATTAACCCAGA  
 GTGGAGCTTCAGTTGGCACTGATGAGGAGTCTGATGTTACTCAAGAAGAAGAGAGAGATGGACAGTACTT  
 TGAACCTGTTGTTCTTTACCTGATCTAGTTGAAGTATCCAGTGGTGAGGAAAAATGAACAAGTGTTTTT  
 AGTCACAGAGCAAAACTTACAGATACGATAAAGATGTTGGCCAGTGGAAAGAAAGAGGCCATTGGAGATA  
 TAAAGATCTTACAGAATTATGATAATAACAAGTTCGCATCGTATGAGAAGGGACCAGGTATTAAGT  
 TTGTGCCAATCACAGAATAACTCCAGATGACTTTGCAAATGAAAGGGACTGAAAGAGTGTGGGTG  
 TGGACTGCATGTATTTGCAGATGGAGAAAGAAAAATAGAACATTTGGCTGTTCTGTTTTAACTACAGG  
 ATGTTGCAGACTCATTTAAGAAAAATTTTGTAGTAAGCAAAAACAGCCCAAGAAAAAGATCATTTGATAAC  
 ACCTCATGTTTCTCATCTGAGCACCCCTAGAGAGTCCCATGTGGCAAAAATGCTATTGCTGTTTTAGAA  
 GAAACCACAAGAGAAAGGACAGATTTAACTCAGGGAGATGAAGTAATAGATACAACCTCAGAAGCTGGAG  
 AAACATCTAGCACTTCTGAAACAACACCAAAAAGCAGTAGTTTTCTCCTCAAGTTTGTATTTGGCTCAGA  
 ATCTGTTAAAAGCATTTTTAGTAGTAAAAATCCAAGCCATTTGCATTTGGCAACAGTTCAGCCACTGGT  
 TCTTTGTTGGATTTAGTTTTAATGCACCTTTGAAAAATAGCAATAGTGAATGACTTCCAGAGTCCAGA  
 GTGGATCTGAAGGAAAAGTAAAACCCGACAAATGTGAGCTGCCACAGAAGTCTGACATCAAGCAATCTTC  
 TGATGGCAAAGTCAAAAATCTCAGTGCCTTTTCAAAGAGAAGTCTTCAACTAGTTATACATTTAAAACA  
 CCAGAAAAGGCACAAGAGAAGAGCAAACTGAAGATCTCCCTCAGATAATGATATTCTCATTGTGTATG  
 AACTCACTCCCACCCCTGAGCAGAAAGCCTGGCAGAGAAAATCTGCTTCTTCAACTTCTTTTGTTA  
 TAAGAATAGGCCTGGCTATGTTAGTGAAGAAGAGGAGGATGATGAAGATTATGAAATGGCTGTCAAGAAG  
 CTTAATGGAAAATCTATCTGGATGACTCAGAAAAACCTTGAAGAAGAAATCTAGCAGATAATGACAAGG  
 AATGTGTTATTGTTTGGGAAAAGGCCAACAGTTGAAGAAGAGCAAAAAGCAGATACTTTAAAGCTTCC  
 ACCTACATTTTCTGTGGAGTCTGCAGTACTGATGAGGACAATGGAATGGGAGGATTTTTCAGTCA  
 GAGCTTCGAAAAGTCTGTGAAGCTCAGAAATCCAGAAATGAGAAAGTACTGATAGAGTTGGTATTGAGC  
 ATATAGGTGAGACTGAAGTAACAACCCCTGTTGGCTGTAAGTCTGAAGAAGCTGATCTGATACCAAACA  
 CAGTAGCTCATCCCTGTTTCTGGGACTATGGACAAAACCTGTCGATTTGTCCACTAGAAAAGAAACTGAC  
 ATGGAATTCCAAAGCAAAGGGGAAAAACAAGCCTGTTTTGTTGGATTTGGAAGTGTACAGGCCTGTCAT  
 TTGCAAGCTTGGCTTCAAGTAATCTGGAGATTTTGTCTTTGGCTCTAAAGATAAAAAATTTCCAGTGGGC  
 AAATACTGGAGCAGTGTGTTTGGAAACACAGACAACAAGTAAAGGTGGCGAGGATGAAGACGGCAGTGT  
 GAAGATGTAGTTCATAATGAGGACATCCACTTTGAACCTATAGTCTCCTTACCAGAGGTAGAAGTGAAT  
 CTGGAGAGGAAGATGAAGAAGTTTTGTTCAAAGAGAGAGCCAACTTTATAGGTGGGACAGAGATGTCAG  
 TCAGTGGAAAGGAGAGAGGATTGGAGATATAAAGATTCTCTGCCATACAATGAAGAAGTATTATCGGATC  
 CTAAATGAGAAGAGACCAGTTTTTAAAGTGTGTGCAAACCATGTTATTACCAAAGCCATGGAATTTAAAC  
 CTTTAAATGTTTCAAACAATGCTTTAGTCTGGACTGCCTCCGATTATGCTGATGGAGAAGCAAAAGTAGA  
 ACAGCTTGCAGTGAATTAAGACAAAAGGAAATGACTGAGAGTTTTAAGAAAAAATTTGAAGAATGTCAA  
 CAAAATAATAAAAATCCAGAATGGACATACATCCCTTGCAGCAGAAATTAACAAGGACACCAATCCTG  
 TGGTGTTTTTGATGTTTGTGCGGATGGTGAACCTCTAGGACGGATAATCATGGAATTTTTTCAAATAT  
 CGTTCTCAAAGTGTGAGAAGTTCAGAGCATTATGCACTGGAGAGAAGGGCTTTGGTTTTAAGAAGTCC

ATTTTTCACCGAGTAGTTCAGATTTTATTTGCCAAGGGGAGATATTACCAAATACAATGGAACAGGTG  
GACAGTCCATTTATGGAGATAAATTTGACGATGAAAACTTTGATTTGAAACACACTGGTCCTGGCTTACT  
ATCCATGGCCAATTACGGCCAGAATACCAACAGTTCTCAATTTTTATAACACTGAAGAAAGCAGAACAT  
TTGGACTTTAAGCATGTAGTATTTGGGTTTGTTAAGGATGGCATGGATACTGTGAGAAAGATTGAATCAT  
TTGTTCTCCTAAAGGTCTGTTAGTAGAAGAATTTGTATCACAGAATGTGGACAGCTA

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT  
ACAAGGATGACGACGATAAGGTTTAA

**Protein Sequence:** >MR226679 representing NM\_011240  
 Red=Cloning site Green=Tags(s)

MRRSKADVERYIASVQGSAPSPREKSMKGFYFAKLYYEAKYDLAKKYISTYINVQERDPKAHRFLGLLY  
 EVEENIDKAVECYKRSVELNPTQKDLVLKIAELLCKNDVTDGRAKYWVERAAKL FPGSPA IYKLEQQLD  
 CKGEDGWNKLFDLIQSELYARPDDIHVNIRLVELYRSNKRLKDAVAHCHEADRNTALRSSLEWNL CVVQT  
 LKEYLES LQCLDSDKSTWRATNKDLLLAYANMLLTLSTRDVQEGRELLESFDSALQSVKSSVVGNDLS  
 ATFLETKGHFYMHVGSLLKMGQQSDIQWRALSELAALCYL VAFQVPRPKVKLIKGEAGQNLLETMAHDR  
 LSQSGHMLLNLSRGKQDFLKEVVESEFANKSGQSALCDALFSSQSSKERSFLGNDDIGNLDGQVPDPDDLA  
 RYDTGAVRAHNGSLQHLTWLGLQWNSLSTLPAIRKWLKQLFHHL PQETSRL ETNAPESICILDLEVFLLG  
 VIYTHLQLKEKCNHSHSYQPLCLPLPVCRLCTERQKTWWDVACTL IHRKALPGTSAKLRLLVQREIN  
 SLRGQEKHGLQPALLVHWAQSLQKTGSSLSNSFYDQREYIGRSVHYWRKVLPLLMIRKKNKSIPEPIDPLF  
 KHFFHSVDIQASEIGEYEEDAHI TFAILD AVNGNIEDAMTAFESIKNVVS YWNLALIFHRKAEDIENDALS  
 PEEQE ECKNYLRKTRDYLRILDDSDSNTSVVQKLPVPLESVKEMLNSVMQELEDYSEGGLYKNGCWRS  
 ADSELKHS TPSP TKYSLSPSKSYKYSKTPPRWAEDQNSLLKMICQQVEAIKKEMQELKLNNSNASPHR  
 WPAEYPGQDPADPGYQGSQTFHGAPLTVATTGPSVYYSQSPAYNSQYLLRPAANVTPTKGPVYGMNRLPP  
 QQHIYAYSQQMHTPPVQSSSACMFSQEMYGPPLRFESPATGILSPRGDDYFNYNVQQTSTNPPLPEPGYF  
 TKPPLVAHASRSAESKVI EFGKSNFVQPMQGEVIRPPLTPAHTTQPTPFKFNFSKNSNDGDFTFSSPQV  
 VAQPPSTAYSNSSELLGLLTS DKPLQGDGYSGLKPI SGQASGSRNTFSFGSKNTL TENMGPNQKNGGFH  
 RSDDMFAFHGPGKSVFTTAA SELANKSHETDGGSAHGDEEDDGPHFEPV VPLPDKIEVKTGEDEEEFFC  
 NRAKLFRFDGESKEWKERGI GNVKILRHKTSGKIRLLMRREQVLKICANH YISPDMLTPNAGSDRSFVW  
 HALDYADELPKPEQLAIRFKTP EEAALFKCKFEEAQNILKALGTNTSTAPNH TLRIVKESATQDNKDICK  
 ADGGNLNFEFQIVKKEGPHYWNCNSCSFKNAATAKKCVSQNTNPTS NKELLPPLVENGFAPKTGLENAQ  
 DRFATMTANKEGHWDSCVCLVRNEPTVSRCIACQNTKSASSFVQTSFKFGQGLPKSVSDSDFRSVFSKKE  
 GQWECSVCLVRNERSAKKCVACENPGKQFKEWHCSLCSVKNEAHAIKCVACNNPVTPSLSTAPPSFKGT  
 SEMSKPFRI GFEGMFAKKEGQWDCSLCFVRNEASATHCIAQYPNKQNPQPTSCVSAPASSETSRSKSGF  
 EGLFPKKEGEWECAVCSVQNESSSLKCVACEASKPTHKPHEAPSAFTVGSKSQSNE SAGSQVGFTEKSNF  
 PEKNFKVGI SEQKFKFGHVDQEKTPSFAFQGSNTEFKSIKDGFSFCIPVSADGFKFGIQEKGQEKKSE  
 KHLNDP SFQAHDTS GQKNGSGVVFQGTSTFTFADLAKSTSREGFQFGKKDPNFKFGSGAGEKLFSSQS  
 GKVAEKANTS SSDL EKDDDAYKTEDSDDIHFEFVVMPEKVELVTGEEDEKVLYSQRVKLFRFDAEISQWK  
 ERGLGNLKI LKNEVNGKLRMLMRREQVLKVCANH WITTTMNLKPLSGSDRAWMWLASDFSDGDAKLEQLA  
 AKFKTPELAE EFQKQFEECQRLLLDIPLQTPHKLVDTGRAAKLIQRAEEMKSGLKDFKFTLNDQVKVTD  
 EENASSGADAPSASDTTAKQNPNTGPALEWDNYDLREDALDSDSVSSSVHASPLASSPVRKNLFRFGES  
 TTGFNFSFKSALSPSKSPAKLNQSGASVGTDEESDVTQEEERDQYFEPV VPLPDLVEVSSGEENEQVVF  
 SHRAKLRYRDKDVGVQWKERGI GDIKILQNYDNKQVRIVMRRDQVLKLCANHRITPDMTLQTMKGTERRVW  
 WTACDFADGERKIEHLAVRFLQDVADSFKKIFDEAKTAQEKDSLITPHVSHLSTPRESPCGKIAIAVLE  
 ETTRETRD LTQGDEVIDTTSEAGETSSTSETTPKAVVSPPKFVFGSESVKSI FSSEKSKPFAFGNSSATG  
 SLFGFSFNAPLKNNSSEMTSRVQSGSEGKVKPKCELPQNSDIKQSSDGKVKNL SAFSKENSSTSYTFKT  
 PEKAQEKSKPEDLPSDNDILIVYELTPTPEQKALAEKLLLPSTFFCYKNRPGYVSEEEDEDEDYEMAVKK  
 LNGKLYLDDSEKPLEENLADNDKECVI VWEKKPTVEERAKADTLKLPPTFFCGVCSDTDEDNGNGEDFQS  
 ELRKVCEAQKSQNEKVTDRVGIEHIGETEVTNPVGCKSEEPDSDTKHSSSPVSGTMDKPVDLSTRKETD  
 MEFPSKGENKPVLF GFGSGTGLSFADLASSNSGDFAFGSKDKNFQWANTGAAVFGTQTTSKGGEDDGDSD  
 EDVVHNE DIHFEP IVSLPEVEVKS GEDEEVLFKERAKLYRWRDVSQWKERGI GDIKILWHTMKYYRI  
 LMRRDQVFKVCANHVITKAMELKPLNVSNNALVWTASDYADGEAKVEQLAVRFKTKEMTESFKKKFEECQ  
 QNI IKLQNGHTSLAAELSKDTPVVFDFVCADGEPLGRIMELFSNIVPQTAENFRALCTGEKGF GKNS  
 IFHRVVPDFICQGGDITKYNGTGGQSIYGDKFDDENFDLKHGTGGLLSMANYGQNTNSSQFFITLKA EH  
 LDFKHVVFVFKDGM DTVRKIESFGSPKGSVSRRICITECGQL

TRTRPLEQKLI SEEDLAANDILDYKDDDDKV

**Restriction Sites:** SgfI-MluI



|                               |   |
|-------------------------------|---|
| <b>Components:</b>            | 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).  |
| <b>Reconstitution Method:</b> | <ol style="list-style-type: none"><li>1. Centrifuge at 5,000xg for 5min.</li><li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li><li>3. Close the tube and incubate for 10 minutes at room temperature.</li><li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li><li>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.</li></ol>   |
| <b>RefSeq:</b>                | <u><a href="#">NM_011240.3</a></u> , <u><a href="#">NP_035370.2</a></u>   |
| <b>RefSeq Size:</b>           | 9477 bp   |
| <b>RefSeq ORF:</b>            | 9162 bp   |
| <b>Locus ID:</b>              | 19386   |
| <b>UniProt ID:</b>            | <u><a href="#">Q9ERU9</a></u>   |
| <b>Cytogenetics:</b>          | 10 29.34 cM   |
| <b>MW:</b>                    | 341.1 kDa   |
| <b>Gene Summary:</b>          | E3 SUMO-protein ligase which facilitates SUMO1 and SUMO2 conjugation by UBE2I. Involved in transport factor (Ran-GTP, karyopherin)-mediated protein import via the F-G repeat-containing domain which acts as a docking site for substrates. Binds single-stranded RNA (in vitro). May bind DNA. Component of the nuclear export pathway. Specific docking site for the nuclear export factor exportin-1 (By similarity). Sumoylates PML at 'Lys-490' which is essential for the proper assembly of PML-NB. Recruits BICD2 to the nuclear envelope and cytoplasmic stacks of nuclear pore complex known as annulate lamellae during G2 phase of cell cycle. Probable inactive PPIase with no peptidyl-prolyl cis-trans isomerase activity.[UniProtKB/Swiss-Prot Function] |