

## Product datasheet for MC224887

### Zfp142 (NM\_029888) Mouse Untagged Clone

#### Product data:

|                           |                                                                                 |
|---------------------------|---------------------------------------------------------------------------------|
| Product Type:             | Expression Plasmids                                                             |
| Tag:                      | Tag Free                                                                        |
| Symbol:                   | Zfp142                                                                          |
| Synonyms:                 | 9330177B18Rik; BB154236; mKIAA0236                                              |
| Mammalian Cell Selection: | Neomycin                                                                        |
| Vector:                   | pCMV6-Entry (PS100001)                                                          |
| E. coli Selection:        | Kanamycin (25 ug/mL)                                                            |
| Fully Sequenced ORF:      | >MC224887 representing NM_029888<br>Red=Cloning site Blue=ORF Orange=Stop codon |

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCCGCGATCGCC

ATGACAGACCTGTGTTGGCATCCCAGCTGGCTAATGGCACTGGGGAGATGGATGGACTGTGCTCTGAAC  
TATTGTTGATTCCCCACCTCTCTCAAACCATGGAATCCTGGGCCCTGTCCAGAATACCTGTGCTTCTGG  
GGAGCTTGCACCTTTGCCTGCTGACCCAGGCTGCCTGCTGGTAGAGGCCACAGCTACTGAAGAGGGTCCA  
GGGAACATGGAGATCATTGTGGAAGCAGTAACCTGGAACCTGTGCCCCAGGAGCTCCTGAAGAGACCTCAG  
AGTCTGGCTGTGTGTTCTCTGCTGAAGATCGCAAGGGTCTGCAGAATCACCTGAGACAGACCCACAAGGC  
AGTTCTGTGCCCTGTTTATTCCGGGGCTGCTCCCTGCTTTTTGGGAGTCAGCAGGGAATGGAGCTGCAT  
CGGCAGGCCATTACCCTTTCCACTGCAGCCATTGCAGCTTCATGGGCTCCAACGTCAAATATTCCGGC  
AGCATCAGCGGAGCCATGGAGCCAGTGCACGGGAGAACTTTCTGCTGCTGTTCAGGGCCTTCCATCCCA  
GGAGCTACTGCCAGCTGCCAACTGCCTCCGGGACACAGAGAACCTTCTGAGGAAGCAAGTACACCTTTG  
CCTGGGCAGGAGTCAGCTGAAGAGGAGGATGCAGAGGAAGAGGAGAGCGTCACCCAAAAAGACTCCCAGA  
AAGTCATGGATAAAAGCCAAGGGGCTCAGCAGTTGGAAGGGCATGTGGGTTCAGGCACTGAGTCTCTCTT  
CAAGACACACATGTGCCAGAATGCAAGCGCTGCTTTAAGAAGCGGACCCATTTGGTAGAGCACCTACAT  
CTGCACTTTCCAGACCCAGCCTCCAGTGTCTTAAGTCCAGAAAGTTCTTACCAGCAAGAGCAAGCTCA  
AGACCCATCTGCTGCGGGAGCTGGGCGAGAAGGCCACCGATGTCCTCTGTGCCACTACAGTGCAGTAGA  
GAGGAATGCACTTAACCGCCACATGGCCAGCATGCACGAGGATATTTCCAATCTACTCAGACACCTAT  
GCCTGTCCCGTCTGTCTGTGAGGAATTCGCTCAGCCAAGCACTCAAGGAGCACCTCAAAGGCCACACAG  
CAGCAGCTGCAGCAGAACCATTACCCCTACACTGCTTTTCAAGGAGGCTGCACATATGTAGCACCTGACCG  
TAAGGCCTTCTTAAAGCACCTAAAGGAGATCCATGGAGTACGGGCTGTGGAGTGCCGCCATCACTCGTGT  
CCCATGCTTTTTGCCACAGCTGAAGCCATGGAAGCCCAACCAAGGCCATTATGCTTTTCCACTGTCCCC  
ACTGTGACTTTGCCTGCTCCAATAAGCACTTGTCCGAAACATAAGAAACAAGGTCAACCCAGGCAGTGA  
AGAGCTACGGTGACCTTCTGTCCCTTTGCCACCTTCAACCCAGTAGCCTACCAGGACCACGTTGGCAAG  
ATGCATGCTTATGAGAAGATCCACCACTGCTCTGAATGTAACCTTTGCCACTGCCACAAAAGGGTGCTTA



TCCGACACATGCTGCTGCATACTGGAGAGAAGCCTCACAATGTGAGCTCTGTGACTTCACATGCCGAGA  
TGTTAGCTACCTATCGAAGCACATGCTGACCCACTCCAACACCAAGGATTACATGTGTACTGAATGTGGC  
TATGTCACCAAGTGAAGCACTACCTGAGTGTGCACATGCCGAAACATGCAGGGGATCTCAGATACCAAT  
GCAACCAGTGCTCCTACCGCTGCCATCGAGCTGATCAGCTGAGCAGCCACAACTACGGCACCAAGGGCAA  
GTCCCTCATGTGTGAGGTGTGCTTTGCTGCAAGCGGAAGTATGAGCTACAGAAGCACATGGCTTCC  
CAGCACCACCCTGGCACACCTGCCCACTCTACCCCTGCCGCTATTGTAGCTATCAGAGCCGCCACAAGC  
AGGCCCTCCTGAGCCATGAGAACTGCAAGCACACCCATCTCCGTGAGTTTCACTGCGCCCTCTGTGACTA  
CCGTACCTTTAGCAATACCACACTCTTCTCCACAAGCGTAAGGTCCATGGCTACATGCCAGGGGACCAG  
GTTTGGCAGTTCTGCAATGCTAGCCAAGAGTTGGAGGGGGCTAGGCAGTGCCTGGCACCTCCTTCAGACT  
CAGGACCTTCAAGCCAGCTGTCTGCCAGCCTGAGAGAGAAGACCGTGAACATGAAATTGTGGCTAACTC  
CAATATGGACCAGGCCCTGCCAGAGACAAATGAGGAGGCCAGCCCCAAAAGACAAGATGGCATTGAGGCT  
CCCCAAGAGGATGATCAGGTTGACAGCCCCAGCCTGGGGGAGGTGGAAGAGGGTGGCTGCACACTACACT  
TGGAGGCCCTGAGAGTAGAGCTAGAACCTGAGACTGAGCCACTACCCCTGAGGAGCTCACCAGAACAGC  
CACTGTGGAGTTGAGGCCCTCTGGATCCCTCAGGACCATGGGAACAGAAAGACCAGGTGGACTAGAAGAG  
CCAGCACTGTCCAGCTTTGACAGTATTGAGACGCTGCCTTGGTTGCTGAAGAAGAGCCGTTGTGGAGA  
AGCTAGCCTCTGAGCCTCCAGAAATCCCCTAATTTCTGAAGAAGCACCTAATACATTCAAGGCAGCTCT  
GACTGCTGAGACTGTACCGTTGCCCTCCATTTCTGAGTCAGAGTCGCTACTTAAGGCCATGAGGAGACAG  
GACAAAGAACAAGCAGAAGCGTTGGTGTGGAGGGCCGGGTACAAATGGTCGTGATCCAGGGAGAAGGAA  
GAGCCTTCCGCTGCCACACTGTCCTTTTACTCTGCGGGGAAAAAGCCCTAACTGCACTCCAAATC  
GGGGTGCCAAAGGTCGCCGAGAGCCCCCTGCTATGCCCTGAGTGTGGAGCTAGCTTTAAGCAACAGCGGGC  
CTCAGCACCACATGATGAAGAAATGCCCTGTTCTTCTCAAAAAGAACAAGGCTTTGCCAAACCTGTCT  
CTCCCACTTACATCCTCAGCTCCAGATAACCAAGCCTCACAGGATGCTGAAAGTAGGAAGCCCCACC  
TTTACCATCAAAGGTAGAGCTTTGCTTCCAAAAGATGCTCCTTCTGACCTCCTGGGGGCCAGGAGTA  
GAGGAACCTCTTCCACACCCCTCTGACTTCCCAACCTCTCCACCAGAAAACCTTACCACAGGGACCT  
CTGAGAAGTTCCACTTTGAGCAAGGCAAGTTTCACTGCAGCTCGTGTACATTCTTTGTTCTCGGCTTTC  
TTCTATTACCTCTCATGTGACTGAAGGCTGTAGAGGGGGCGTGGCCAGAAAAGAAAGCGTGGGCGCCCC  
CAGACCCATGCAGTTGTGCTTCTCTTAACAATGGGACTCTACCTCCTGAATACTGGAAGTACAGAGT  
CCAGTCTTAGTGATGGGGACACAGCTGTGGTTTCAAGCAAAAAGGGTGCCCTCTTCTCTGCCCCACATG  
TCCCTTTAGCTGTGAGCAGGAAGGACCCTGAGGACTCACCAGACCCAGGGCTGCCCTTAAGTCTGGA  
GATCTGCACTGTGGTCTCTGCCATTCACTGCTCCTGCTGCTGCCCTGAGGCTCCATCAAAAACGGA  
GGCATCCCACTGCATCCCAGCCTCTGGCCCCGGCCCTCCTGCAGTGTGGGACTGTGGCTTTACCTG  
CAAACAAGCCGGTGCCTTCAAGCAGCATCGGCGGCTCAAGCATGAGGGTGTGAAGCCACATCAGTGCCCT  
TTCTGTGACTTTTCTACTACTAGACGGTACCGGTTAGAGGCCACCAAGTCTCGACACACAGGTGTTGGCC  
GCATCCCTTGCACTCTGCCCCTCAGACATTTGGTACCAACTCAAACTGCGCTTGCACTCACTACAGAGT  
ACATGACAAAACACCCACTCACTTCTGTCCACTGTGTGACTATAGTGGTTATCTTCGTATGACATCACT  
CGCCATGTCAACAGCTGCCACAGGCCCTTCTCTTTTCTGCACTCAGTGTGAAGCCCAAGTTTAGCT  
CAGAAACAGCACTCAAGCAGCATGCTCTGCGACGACACCCAGAACCCACACCTCCTTCTCTGGCTGTCC  
TGTAGAGGTACCGAAGGCCCTTGCAGTGTCCCACTGTGGCCTGCTCTGTCCAGTCTGCCAGCCTG  
CGAGGACACACCCGTAACAGCACCAAGGCTGGAGTGTGGGGCTGCCAGGAGTCTTCCCTAACCGTC  
CAGCTCTTGATGAGCATCGGAGGCAACATCATTTACGCCACCGCTGCCAGCTCTGCAGCTTTGCCGCCC  
GGAGCGAGTAGGCCTGGTGAAGCACTACCTGGAACAGCACGAGGAGTCTTCAACAGCCCCCTCAGATGGG  
GATGCTGGCCAGCCCTCTCTTGTGCCCTTTTGTGACTTTGATGTGCCATCAGTTGGTGTGATC  
ACCATGTAAGGGACATGGAGGTACCCGGCTTTACAAGTGTACTGACTGTGCTATAGCACCACAAAATCG  
GCAGAAGATCACCTGGCACAGCCGATTCACACTGGGGAAAAGCCTTACCACTGTCACTCTCGGCCTAT  
GCCTGTGCTGACCCTTCCGCTCAAGTACCACATGCGGATCCATAAGGAGGAACGGAAGTACCTATGCC  
CTGAGTGTGGGTACAAGTGCAAGTGGGTCAACCACTCAAATACCACATGACCAAGCATACAGGACTGAA  
GCCTTACCAATGTCTGAGTGTGAGTACTGCACCAACCGGGCTGACGCACTGCGGGTACACCGGAGACA  
CGGCACAGAGAAGCCCGGCTTTCATGTGTGAGCAGTGTGGCAAGGCATTCAAGACACGCTTCTGCTGC  
GCACCCACCTCCGAAAGCACAGTGAAGGCCAAGCCCTATGTTTGAATGTTTGCCACCGTGCTTCCGCTG  
GGCTGTGGACTGCGACATCATGCTTTACCCATACCGATCGCCACCCATTCTTCTGCCCTCTGCAGC  
TACAAGGCCAAGCAAAAGTTCCAGGTGGTCAAGCATGTGCGTAGGCACCCCTGACCAGGCCGACCCCA  
ACCAGGGTGTGGCAAGACCCCACTCCACAGTGCATCTACATGATGTGAAGCTGGAAGATCCCAG  
CCCTCTGCTCCTCCGCTCCCTCTACTGGACCTGAGGGCTGA

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAATGATATCCTGGATT  
ACAAGGATGACGACGATAAGGTTTAA

|                               |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         |
|-------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <b>Restriction Sites:</b>     | Sgfl-MluI                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               |
| <b>ACCN:</b>                  | NM_029888                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               |
| <b>Insert Size:</b>           | 5223 bp                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |
| <b>OTI Disclaimer:</b>        | 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).                                                                                                                                                                          |
| <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>Note:</b>                  | Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.                                                                                                                                                                                                                                                                                                                                                                                                |
| <b>RefSeq:</b>                | <a href="#">NM_029888.3</a> , <a href="#">NP_084164.1</a>                                                                                                                                                                                                                                                                                                                                                                                                                                                               |
| <b>RefSeq Size:</b>           | 6192 bp                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |
| <b>RefSeq ORF:</b>            | 5223 bp                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |
| <b>Locus ID:</b>              | 77264                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   |
| <b>UniProt ID:</b>            | <a href="#">G5E869</a>                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  |
| <b>Cytogenetics:</b>          | 138.54 cM                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               |
| <b>Gene Summary:</b>          | <p>May be involved in transcriptional regulation.[UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (2) uses an alternate in-frame splice junction compared to variant 1. The resulting isoform (2) has the same N- and C-termini but is shorter compared to isoform 1.</p>                                                                                                                                                                                                                        |