

Product datasheet for **MC216979**

Zfp207 (NM_001130169) Mouse Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Zfp207 (NM_001130169) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Zfp207
Synonyms:	8430401D15Rik; BuGZ; Zep; Znf207
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)



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Fully Sequenced ORF: >MC216979 representing NM_001130169
 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCC**GCGATCGCC**

ATGGGTCGAAGAAGAAGAAGCAGCTCAAGCCGTGGTCTGGTATTGTAACAGAGATTTTGTATGATGAGA
 AGATCCTTATACAGCACAAAAAGCAAAGCATTAAATGCCATATTTGTCATAAGAAATTATACACAGG
 ACCTGGTTTAGCAATTCATTGCATGCAGGTCCATAAAGAAACCATAGATGCAGTACCAAATGCAATACCT
 GGGAGAACAGACATAGAGTTGGAATATATGGCATGGAAGGTATCCAGAGAAAGATATGGATGAAAGAC
 GGCGACTTCTTGAACAGAAAACACAAGAGAGTCAGAAAAAGAAACAACAAGATGATTCTGATGAATATGA
 TGATGATGAATCTGCAGCCTCAACTTCCTTTAGCCACAGCCTGTTCAACCTCAGCAAGTTATATCCCA
 CCAATGGCTCAGCCAGGACTGCCTCCAGTCCAGGGCCACCAGGAATGCCTCCAGGCATACCTCCATTGA
 TGCCAGGTGTTCCCTCCCCTGATGCCAGGCATGCCTCCAGTGCAGGAATGCCGCTGGATTGCATCA
 TCAGAGAAAATACACCCAGTCATTTTGGCGTGAAAACATAATGATGCCAATGGGTGGAATGATGCCACCT
 GGACCTGGAATACCACCTCTGATGCCAGGTATGCCACCAGGTATGCCCCACCTGTTCCAGCTCCTGGAA
 TTCTTCCAATGACTCAAGCACAGGCTGTTTCAGCACAGGTATTCTTAATAGACCACCTGCACCAACAGC
 AGCAGTACCTGCTCCACAGCCTCCAGTTACTAAGCCTCTTTTCCCAGTCTGGACAGATGGGGACACCT
 GTAACAAGCTCAAGTACAGCTTCATCCAATTCAGAAAGTCTGCTGCATCTTCTAAAGCTCTGTTTCCTA
 GCACAGCACAAGCTCAGGCAGCTGTCCAAGGACCTGTTGGTACAGATTTTAAGCCCTTAAATAGTACTCC
 TGCAGCAACAACACTACAGAACCCCAAGCCTACATTCCTGCTTATACACAGTCTACAGCGTCAACCACT
 AGTACAACAACAGTACTGCAGCAAAGCCAGCAGCTTCAATAACAAGTAAGCCTGCTACACTCACAAACA
 CCAGTGAACCAAGTAAGTTGATCCATCCAGATGAGGATATCACTGGAAGAAAGAAGGGCACAGTTACC
 TAAATATCAGAGAAATCTTCTCGACCAGGACAAACTCCAATTGGTAATCCACCAGTTGGACCAATTGGG
 GGTATGATGCCACCACAGCCAGGCTGCCACAGCAGGCAATGCGACCTCCAATGCCACCTCATGGTC
 AGTATGGTGGTCAATCAAGGCATGCCAGGTTATCTTCTGGCGCTATGCCACCGTATGGACAGGGACC
 ACCAATGGTGCCCTTACCAAGGTGGGCTCCTCGACCTCCAATGGGAATGAGACCTCCTGTAATGTGC
 CAAGGTGGCCGTTACTGA

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites: Sgfl-Mlul

ACCN: NM_001130169

Insert Size: 1488 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_001130169.1](#), [NP_001123641.1](#)

RefSeq Size: 2376 bp

RefSeq ORF: 1488 bp

Locus ID: 22680

UniProt ID: [Q9JMD0](#)

Cytogenetics: 11 B5

Gene Summary: Kinetochores- and microtubule-binding protein that plays a key role in spindle assembly. ZNF207/BuGZ is mainly composed of disordered low-complexity regions and undergoes phase transition or coacervation to form temperature-dependent liquid droplets. Coacervation promotes microtubule bundling and concentrates tubulin, promoting microtubule polymerization and assembly of spindle and spindle matrix by concentrating its building blocks (PubMed:26388440). Also acts as a regulator of mitotic chromosome alignment by mediating the stability and kinetochore loading of BUB3. Mechanisms by which BUB3 is protected are unclear: according to a first report, ZNF207/BuGZ may act by blocking ubiquitination and proteasomal degradation of BUB3. According to another report, the stabilization is independent of the proteasome (By similarity).[UniProtKB/Swiss-Prot Function] Transcript Variant: This variant (1) represents the longest transcript and encodes the longest isoform (1).