

Product datasheet for RN210904

Bex4 (NM 001037554) Rat Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: Bex4 (NM_001037554) Rat Untagged Clone

Tag: Tag Free Symbol: Bex4

Mammalian Cell Neomycin

Selection:

Vector:pCMV6-Entry (PS100001)E. coli Selection:Kanamycin (25 ug/mL)

Fully Sequenced ORF: >RN210904 representing NM_001037554

Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC

GCCGCGATCGCC

ACCTTGA

AGCGGACCGACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGATCTGGCAGCAAATGATATCC

TGGATTACAAGGATGACGACGATAAGGTTTAA

Restriction Sites: Sgfl-Rsrll

ACCN: NM_001037554

Insert Size: 357 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).



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Cytogenetics:

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 001037554.1, NP 001032643.1

Xq35

 RefSeq Size:
 723 bp

 RefSeq ORF:
 357 bp

 Locus ID:
 501624

 UniProt ID:
 Q3MKP9

Gene Summary: May play a role in microtubule deacetylation by negatively regulating the SIRT2 deacetylase

activity toward alpha-tubulin and thereby participate to the control of cell cycle progression

and genomic stability.[UniProtKB/Swiss-Prot Function]