

## **Product datasheet for SC317746**

## RNF70 (PJA1) (NM\_022368) Human Untagged Clone

## **Product data:**

**Product Type:** Expression Plasmids

Tag: Tag Free

Symbol: RNF70

Synonyms: PRAJA1; RNF70

Mammalian Cell Neomycin

Selection:

Vector: pCMV6-Entry (PS100001)

E. coli Selection: Kanamycin (25 ug/mL)

Fully Sequenced ORF: >SC317746 representing NM\_022368.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

GCTCGTTTAGTGAACCGTCAGAATTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGACTCGTCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

ATGGGTCAGGAATCTAGCAAGCCTGTATGGCCCAATCCAACAGGAGGGTATCAGTCCAATACAGGTAGG CAGAGAAAGACGAACTCCGAAGTCCCAATGCACAGATCAGCCCCCAGTCAAACCACCAAGAGGAGCCGA AATGAGGACTATTCCAGTACTTCCAGGTGGAGGGATACCGCCAATGACAATGAGGGCCACTCGGATGGC CTGGCAAGAAGAGGGGAGAGGCGAGAGTTCAAGTGGCTATCCCGAGCCAAAGTACCCTGAAGACAAACGG GAAGCGAGGAGTGACCAAGTGAAACCAGAAAAGGTGCCGAGACGACGACGACCATGGCCGACCCTGAC TTCTGGACGCACAGTGATGATTACTACAAATACTGCGACGAAGACTCTGACAGTGACAAAGAGTGGATT GCTGCTCTGCGTCGGAAATATCGAAGCCGAGAGCAAACCCTGTCCTCCAGTGGCGAAAGCTGGGAGACT CTGCCGGGGAAAGAAGAGCGGGAACCTCCACAGGCTAAGGTGAGTGCCAGCACTGGCACCAGCCCTGGC CCCGGTGCTAGTGCCAGTGCCGGGGCTGGCGCGGGGCCAGTGCTGGCAGCAATGGCAGCAATTACCTT GAAGAAGTTCGAGAACCATCTCTTCAGGAAGAGCAGGCATCCCTGGAAGAAGGAGAAATTCCTTGGCTC CAGTACCATGAGAATGACAGTAGCAGTGAGGGGGGATAATGATTCTGGTCACGAGTTGATGCAACCTGGG GTATTCATGCTGGATGGAAACAACAACCTTGAAGATGACTCCAGTGTGAGCGAAGACCTAGAAGTGGAT TGGAGCCTCTTTGATGGATTTGCAGATGGGTTAGGAGTGGCTGAAGCCATTTCCTATGTGGACCCTCAG TTCCTCACCTACATGGCACTTGAAGAACGCCTGGCCCAGGCAATGGAAACTGCCCTTGCGCACTTGGAG ATCCTGGTCACTGAAGATCATGGCGCAGTTGGTCAGGAGATGTGCTGCCCCATCTGCTGTAGCGAATAT 

**ACGCGTACGCGGCCGCTC**GAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT

TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC



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EU: info-de@origene.com CN: techsupport@origene.cn Restriction Sites: Sqfl-Mlul

**ACCN:** NM\_022368

Insert Size: 1368 bp

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).

OTI Annotation: This TrueClone is provided through our Custom Cloning Process that includes sub-cloning into

OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected

reference without frameshifts, and is delivered as lyophilized plasmid DNA.

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.

Note: Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um

filter is required.

**RefSeq:** <u>NM\_022368.4</u>

RefSeq Size: 2316 bp

RefSeq ORF: 1368 bp

**Locus ID:** 64219

UniProt ID: Q8NG27

Cytogenetics: Xq13.1

Domains: RING

Protein Families: Druggable Genome

**MW:** 50.4 kDa



## Gene Summary:

This gene encodes an enzyme that has E2-dependent E3 ubiquitin-protein ligase activity. This enzyme belongs to a class of ubiquitin ligases that include a RING finger motif, and it can interact with the E2 ubiquitin-conjugating enzyme UbcH5B. This gene is located in an area of chromosome X where several X-linked cognitive disability disorders have been associated, and it has also been found as part of a contiguous gene deletion associated with craniofrontonasal syndrome, though a direct link to any disorder has yet to be demonstrated. Alternative splicing results in multiple transcript variants. [provided by RefSeq, May 2010] Transcript Variant: This variant (3) lacks an in-frame segment of the 5' coding region, compared to variant 1, resulting in an isoform (c) that is shorter than isoform a.