

Product datasheet for SC107725

RNF41 (NM_194359) Human Untagged Clone

Product data:

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|---------------------------|---|
| Product Type: | Expression Plasmids |
| Product Name: | RNF41 (NM_194359) Human Untagged Clone |
| Tag: | Tag Free |
| Symbol: | RNF41 |
| Synonyms: | FLRF; NRDP1; SBB103 |
| Mammalian Cell Selection: | None |
| Vector: | <u>pCMV6-XL6</u> |
| E. coli Selection: | Ampicillin (100 ug/mL) |
| Fully Sequenced ORF: | >OriGene ORF within SC107725 sequence for NM_194359 edited (data generated by NextGen Sequencing) |

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ATGGGGTATGATGTAACCCGTTTCCAGGGGGATGTTGACGAAGATCTTATCTGCCCTATT
TGCAGTGGAGTCTTGGAGGAGCCAGTACAGGCACCTCATTGTGAACATGCTTTCTGCAAC
GCCTGCATCACCCAGTGGTTCTCTCAGCAACAGACATGTCCAGTGGACCGTAGTGTGTG
ACGGTCGCCCATCTGCGCCAGTACCTCGGATCATGCGGAACATGTTGTCAAAGCTGCAG
ATTGCCTGTGACAACGCTGTGTTCCGGCTGTAGTGCCGTTGTCCGGCTTGACAACCTCATG
TCTCACCTCAGCGACTGTGAGCACAACCCGAAGCGGCCTGTGACCTGTGAACAGGGCTGT
GGCCTGGAGATGCCCAAAGATGAGCTGCCCAACCATAACTGCATTAAGCACCTGCGCTCA
GTGGTACAGCAGCAGACACGCATCGCAGAGCTGGAGAAGACGTCAGCTGAACACAAA
CACCAGCTGGCGGAGCAGAAGCGAGACATCCAGCTGCTAAAGGCATACATGCGTGCAATC
CGCAGTGTCAACCCCAACCTTCAAGAACCTGGAGGAGACAATTGAATACAACGAGATCCTA
GAGTGGGTGAACCTCCCTTCAGCCAGCAAGAGTGACCCGCTGGGGAGGGATGATCTCGACT
CCTGATGCTGTGCTCCAGGCTGTAATCAAGCGCTCCCTGGTGGAGAGTGGCTGTCCTGCT
TCTATTGTCAACGAGCTGATTGAAAATGCCACGAGCGTAGTGGCCCCAGGGTCTGGCC
ACACTAGAGACTAGACAGATGAACCGACGCTACTATGAGAACACGTGGCCAAGCGCATC
CCTGGCAAGCAGGCTGTTGTCGTGATGGCCTGTGAGAACCAGCACATGGGGGATGACATG
GTGCAAGAGCCAGGCCTTGTGATGATATTTGCGCATGGCGTGAAGAGATATAA

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Clone variation with respect to NM_194359.2



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Gene Summary:

This gene encodes an E3 ubiquitin ligase. The encoded protein plays a role in type 1 cytokine receptor signaling by controlling the balance between JAK2-associated cytokine receptor degradation and ectodomain shedding. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jun 2011]

Transcript Variant: This variant (3) differs in the 5' UTR compared to variant 1. Variants 1, 3 and 4 encode the same protein. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.