

Product datasheet for **SC337472**

UHRF1 (NM_001290052) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	UHRF1 (NM_001290052) Human Untagged Clone
Tag:	Tag Free
Symbol:	UHRF1
Synonyms:	hNP95; hUHRF1; huNp95; ICBP90; Np95; RNF106; TDRD22
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)



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Fully Sequenced ORF: >NCBI ORF sequence for NM_001290052, the custom clone sequence may differ by one or more nucleotides

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ATGTGGATCCAGGTTTCGGACCATGGACGGGAGGCAGACCCACACGGTGGACTCGCTGTCCAGGCTGACCA
AGGTGGAGGAGCTGAGGCGGAAGATCCAGGAGCTGTTCCACGTGGAGCCAGGCTCAGAGGCTGTTCTA
CAGGGGCAAACAGATGGAGGACGGCCATACCCTCTTCGACTACGAGGTCCGCCTGAATGACACCATCCAG
CTCCTGGTCCGCCAGAGCCTCGTGCTCCCCACAGCACCAAGGAGCGGGACTCCGAGCTCTCCGACACCG
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TGACAGCAGGCCAGCCGATGAGGACATGTGGGATGAGACGGAATTGGGGCTGTACAAGGTCAATGAGTAC
GTCGATGCTCGGGACACGAACATGGGGCGTGGTTTGGAGCGCAGGTGGTCAGGGTGACGCGGAAGGCC
CCTCCCGGACGAGCCCTGCAGCTCCACGTCCAGGCCGGCGCTGGAGGAGGACGTCATTTACCACGTGAA
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GAGCGGCCGGGTGAAGGGAGCCCATGGTTGACAACCCATGAGACGGAAGAGCGGGCCGCTCTGCAAGC
ACTGCAAGGACGAGCTGAACAGACTCTGCCGGTCTGCGCCTGCCACCTGTGCGGGGCCGCGCAGGACCC
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AGGACCGCGAACAGTCTGTGATCAGAACTCACCAACACCAACAGGGCGCTGGCTCTCAACTGCTTTG
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ATGATGAGCCTGGCCCTTGGACGAAGGAGGGGAAGGACCGGATCAAGAAGCTGGGGCTGACCATGCAGTA
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GAGCAGCAGGAGGGGGCTTCGCGTCCCCAGGACGGGCAAGGGCAAGTGGAAAGCGGAAGTCCGGCAGGAG
GTGGCCCGAGCAGGGCCGGTCCCCGCGCCGACATCCAAGAAAACCAAGGTGGAGCCCTACAGTCTCAC
GGCCAGCAGAGCAGCCTCATCAGAGAGGACAAGGCAACGCCAAGCTGTGGAATGAGGTCTGGCGTCA
CTCAAGGACCGGCCGGCAGCGGACGCCGTTCCAGTTGTTTCTGAGTAAAGTGGAGGAGACGTTCCAGT
GTATCTGCTGTGAGGAGCTGGTGTTCGGGCCATCACGACCGTGTCCAGCACACGTTGCAAGGACTG
CCTGGACAGATCCTTTCGGGCACAGGTGTTCCAGCTGCCCTGCCTGCCGCTACGACCTGGGCCGACGAT
GCCATGCAGGTGAACAGCCTCTGCAGACCGTCTCAACCAGCTTTCCTCCGGCTACGGCAATGGCCGGT
GA
    
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Restriction Sites: SgfI-MluI

ACCN: NM_001290052

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:	<ol style="list-style-type: none">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_001290052.1 , NP_001276981.1
RefSeq Size:	3901 bp
RefSeq ORF:	2382 bp
Locus ID:	29128
UniProt ID:	Q96T88
Cytogenetics:	19p13.3
Protein Families:	Druggable Genome, Transcription Factors
Gene Summary:	<p>This gene encodes a member of a subfamily of RING-finger type E3 ubiquitin ligases. The protein binds to specific DNA sequences, and recruits a histone deacetylase to regulate gene expression. Its expression peaks at late G1 phase and continues during G2 and M phases of the cell cycle. It plays a major role in the G1/S transition by regulating topoisomerase IIalpha and retinoblastoma gene expression, and functions in the p53-dependent DNA damage checkpoint. It is regarded as a hub protein for the integration of epigenetic information. This gene is up-regulated in various cancers, and it is therefore considered to be a therapeutic target. Multiple transcript variants encoding different isoforms have been found for this gene. A related pseudogene exists on chromosome 12. [provided by RefSeq, Feb 2014]</p> <p>Transcript Variant: This variant (5) differs in the 5' UTR, compared to variant 1. Variants 1, 3, 4 and 5 all encode isoform 1. 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.</p>