

Product datasheet for **MR230655**

Btrc (NM_001286465) Mouse Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Btrc (NM_001286465) Mouse Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	Btrc
Synonyms:	b-TrCP; beta-TrCP; Beta-Trcp1; E3RS-IkappaB; E3RSIkappaB; Fbw1a; FWD1; HOS; Slimb
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Cell Selection:	Neomycin



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ORF Nucleotide Sequence:

>MR230655 representing NM_001286465
 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCC**CGCATCGCC**

ATGGACCCGGCAGAGCGGTGCTGCAGGAGAAAGCGCTTAAGTTTATGTGCTCTATGCCAGGTCTCTGT
 GGCTGGGCTGCTCCAGCCTGGCGGACAGCATGCCTTCGCTGCGATGCCTGTATAACCCAGGGACTGGCGC
 ACTCACAGCTTTCCAGAATTCCTCAGAGAGAGAAGACTGTAATAATGGCGAACCCCTAGGAAGATAATA
 CCAGAGAAGAATTCACCTAGACAGACTTACAACAGCTGTGCCAGGCTTTCATAAACCAAGAGACAGTAT
 GTCTAACAAAGCACTGCTATGAAGACTGAAAATTGTGTGGCCAAAGCCAACTTGCCAATGGCACTTCCAG
 CATGATTGTGCCAAGCAGCGGAACTCTCAGCAAGCTATGAGAAGGAAAAGGAGCTGTGTGTCAAGTAT
 TTTGAGCAGTGGTCAGAGTCTGATCAAGTGAATTTGTAGAACACCTTATATCCCAATGTGCTACTACC
 AGCATGGGCACATCAACTCCTACCTAAAACCTATGCTGCAGAGGGATTCATAACTGCCTGCCAGCAGC
 GGGTCTGGACCACATCGCTGAGAACATTCTGTACTTGGACGCCAAGTCACTGTGTGCTGCTGAGCTC
 GTGTGCAAGGAATGGTACCGCTGACGTCGGACGGCATGCTGTGAAAAAGCTCATCGAGAGGATGGTCA
 GGACGGACTCTCTGTGGCGAGGCCTGGCAGAGCGCAGAGGCTGGGGACAGTACTTATCAAAAAACAACC
 TCCTGATGAGAACGCTCCTCCAACTCCTTTTATAGAGCGCTTTATCCTAAAATCATACAAGACATTGAG
 ACAATAGAGTCCAATTGGAGATGTGGGCGACATAGTTTACAGAGAATCCACTGCCGGAGTGAAACAAGTA
 AAGGGGTTTACTGTTTACAGTACGACGACCAGAAGATAGTCAGCGGCCTTCGAGACAACACCATCAAGAT
 CTGGGATAAAAGCACACTGGAATGCAAGCGGATTCACGGGCCACACGGGCTCCGTCTGTGTCTGCAG
 TACGATGAGAGGGTGTATCATCACAGGCTCCTCAGACTCCACCGTCAGAGTGTGGGATGTAATGCAGGTG
 AGATGCTAAACACATTGATTCACACTGTGAAGCCGTTCTGCACCTGCGCTTCAATAATGGCATGATGGT
 GACCTGTTCCAAAGACCGTTCATCGCTGTGTGGGATATGGCTTCCCAACTGACATCACCTCAGGAGG
 GTGCTGGTGGGACACCGAGCTGCGGTCAATGTTGTAGACTTTGATGACAAGTACATCGTTTCTGCCTCTG
 GAGATAGAACCATAAAGGTGTGGAACACAAGTACCTGTGAATTCGTAAGGACCCTAAATGGGCACAAGCG
 TGGCATCGCCTGTTTGCAGTACAGAGACAGGCTGGTGGTGGAGCGGCTCCTCTGACAACACCATCAGGCTG
 TGGGACATAGAGTGTGGAGCATGCCTGCGAGTGTGGAGGGCCATGAGGAGTTGGTACGCTGCATTTCGAT
 TTGATAACAAAAGGATAGTGAGCGGAGCCTATGATGGGAAAATTAAGTGTGGGATCTTATGGTGTCTT
 GGACCCGCGTGTCCAGCAGGACTCTCTGTCTGCGGACACTTGTGGAGCATTCTGGAAGAGTTTCCGC
 CTCAGTTTGTGATGAATTCAGATTGTGAGTACATGATGACACAATTCTCATCTGGGACTTCTCTGA
 ATGATCCAGCTGCTCACGCTGAACCGCCCCGCTCCCCTTCTCGGACATACACTACATCTCCAGA

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Protein Sequence:

>MR230655 representing NM_001286465
 Red=Cloning site Green=Tags(s)

MDPAEAVLQEKALKFMCSPRSLWLGCSLADSMPSLRCLYNPGTGALTAFQNSSEREDCNNGEPPRKII
 PEKNSLRQTYNSCARLCLINQETVCLTSTAMKTENCVAKAKLANGTSSMIVPKQRKLSASYEKEKELCVKY
 FEQWSESDQVEFVEHLISQMCHYQHGHIINSYLPMLQRDFITALPARGLDHIAENILSYLDAKSLCAAEL
 VCKEYWRVTS DGMLWKKLIERMVRTDSLWRGLAERRGWGQYLFKNKPPDENAPPNSFYRALYPKIIQDIE
 TIESNWRCGRHSLQRIHCRSETS KGVYCLQYDDQKIVSGLRDNTIKIWDKSTLECKRILTGHTGSVLCLO
 YDERVITGSSDSTVRVWDVNAGEMLNTLIHHCEAVLHLRFNNGMMVTCSKDRSIAVWDMASPTDITLRR
 VLVGHRAAVNVDFDDKIVYSASGDRITKVVNTSTCEFVRTLNHGKRGIAQLQYRDRLVVS GSSDNTIRL
 WDIECGACLRVLEGHEELVRCIRFDNKRIVSGAYDGKIKVWDLMAALDPRAPAGTLCLRTLVEHSGRVFR
 LQFDEFQIVSSSHDDTILIWDFLNDPAAHAEPSPSRITYTISR

TRTRPLEQKLI SEEDLAANDILDYKDDDDKV

Restriction Sites:

SgfI-MluI

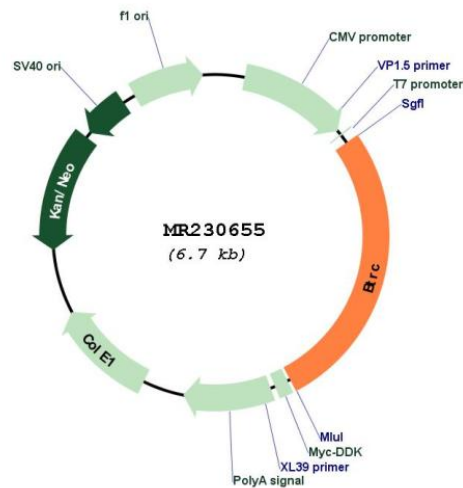
Cloning Scheme:

Cloning sites used for ORF Shuttling:



* The last codon before the Stop codon of the ORF

Plasmid Map:



ACCN: NM_001286465

ORF Size: 1815 bp

OTI Disclaimer:	The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. More info
OTI Annotation:	This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.
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_001286465.1 , NP_001273394.1
RefSeq Size:	6340 bp
RefSeq ORF:	1818 bp
Locus ID:	12234
UniProt ID:	Q3ULA2
Cytogenetics:	19 C3
MW:	69.4 kDa

Gene Summary:

Substrate recognition component of a SCF (SKP1-CUL1-F-box protein) E3 ubiquitin-protein ligase complex which mediates the ubiquitination and subsequent proteasomal degradation of target proteins. Recognizes and binds to phosphorylated target proteins (PubMed:10097128, PubMed:16371461, PubMed:18782782, PubMed:9859996, PubMed:9990853, PubMed:21911472). SCF(BTRC) mediates the ubiquitination of phosphorylated NFKB, ATF4, CDC25A, DLG1, FBXO5, PER1, SMAD3, SMAD4, SNAI1 and probably NFKB2. SCF(BTRC) mediates the ubiquitination of CTNNB1 and participates in Wnt signaling (By similarity). SCF(BTRC) mediates the ubiquitination of NFKBIA, NFKBIB and NFKBIE; the degradation frees the associated NFKB1 to translocate into the nucleus and to activate transcription (PubMed:9859996, PubMed:10097128). Ubiquitination of NFKBIA occurs at 'Lys-21' and 'Lys-22' (PubMed:9859996, PubMed:10097128). SCF(BTRC) mediates the ubiquitination of CEP68; this is required for centriole separation during mitosis (By similarity). SCF(BTRC) mediates the ubiquitination and subsequent degradation of nuclear NFE2L1 (PubMed:21911472). Has an essential role in the control of the clock-dependent transcription via degradation of phosphorylated PER1 and PER2 (PubMed:18782782). May be involved in ubiquitination and subsequent proteasomal degradation through a DBB1-CUL4 E3 ubiquitin-protein ligase (By similarity). Required for activation of NFKB-mediated transcription by IL1B, MAP3K14, MAP3K1, IKBKB and TNF (By similarity). Required for proteolytic processing of GLI3 (PubMed:16371461). Mediates ubiquitination of REST, thereby leading to its proteasomal degradation (By similarity).[UniProtKB/Swiss-Prot Function]