

Product datasheet for **RC223966**

ERCC6L (NM_017669) Human Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	ERCC6L (NM_017669) Human Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	ERCC6L
Synonyms:	PICH; RAD26L
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
ORF Nucleotide Sequence:	>RC223966 representing NM_017669 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGATCGCC**

ATGGAGGCATCCCGAAGGTTTCCGGAAGCCGAGGCCTTGAGCCAGAGCAGGCTGCTCATTACCTAAGAT
ATGTGAAAGAGGCCAAAGAAGCAACTAAGAATGGAGACCTGGAAGAAGCATTTAACTTTTCAATTTGGC
AAAGGACATTTTCCCAATGAAAAAGTGTGAGCAGAATCCAAAAATACAGGAAGCCTTGGAGGAGTTG
GCAGAACAGGGAGATGATGAATTTACAGATGTGTGCAACTCTGGCTTGCTACTTTATCGAGAACTGCACA
ACCAACTCTTTGAGCACCAGAAGGAAGGCATAGCTTTCCTCTATAGCCTGTATAGGGATGGAAGAAAAGG
TGGTATATTGGCTGATGATATGGGATTAGGAAGACTGTTCAAATCATTGCTTTCCTTTCCGGTATGTTT
GATGCATCACTTGTGAATCATGTGCTGCTGATCATGCCAACCAATCTTATTAACACATGGGTAAGAAGAT
TCATCAAGTGGACTCCAGGAATGAGAGTCAAAACCTTTTCATGGTCCTAGCAAGGATGAACGGACAGAAA
CCTCAATCGGATTCAGCAAAGGAATGGTGTTATTATCACTACATACCAAAATGTTAATCAATAACTGGCAG
CAACTTTCAAGCTTTAGGGGCCAAGAGTTTGTGTGGGACTATGTCATCCTCGATGAAGCACATAAAATAA
AAACCTCATCTACTAAGTCAGCAATATGTGCTCGTGCTATTCTGCAAGTAATCGCCTCCTCCTCACAGG
AACCCCAATCCAGAATAATTTACAAGAACTATGGTCCCTATTTGATTTTGCTTGCAAGGGTCCCTGCTG
GGAACATTAATAAATTTAAGATGGAGTATGAAAATCCTATTACTAGAGCAAGAGAGAAGGATGCTACCC
CAGGAGAAAAAGCCTTGGGATTTAAAAATATCTGAAAACCTTAATGGCAATCATAAAACCTATTTTCTCAG
GAGGACTAAAGAAGACGTACAGAAGAAAAAGTCAAGCAACCCAGAGGCCAGACTTAATGAAAAGAAATCCA
GATGTTGATGCCATTTGTGAAATGCCTTCCCTTTCCAGGAAAAATGATTTAATTATTTGGATACGACTTG
TGCCTTTACAAGAAGAAATATACAGGAAATTTGTGCTTTAGATCATATCAAGGAGTTGCTAATGGAGAC
GCGCTCACCTTTGGCTGAGCTAGGTGTCTTAAAGAAGCTGTGTGATCATCCTAGGCTGCTGTCTGCACGG
GCTTGTGTTTGTAAATCTTGGGACATTCTGCTCAAGATGGAATGAGGGGAAGATTCCCCAGATG
TGGACCATATTGATCAAGTAAGTATGACACATTGATGGAAGAATCTGAAAAATGATATTCCTAATGGA
CCTACTTAAGAGGCTGCGAGATGAGGGACATCAAACTCTGGTGTCTTCTCAATCGAGGCAAAATTCTAAAC



[View online »](#)

ATCATTGAACGCCTCTTAAAGAATAGGCACTTTAAGACATTGCGAATCGATGGGACAGTTACTCATCTTT
 TGGAACGAGAAAAAGAATTAACTTATTCAGCAAAATAAGATTACTCTGTTTTCTGCTTACCACTCA
 AGTAGGTGGTGTCTGGTTAACATTAAGTGCAGCAACTAGAGTGGTCATTTTTGACCCTAGCTGGAATCCT
 GCAACTGATGCTCAAGCTGTGGATAGAGTTTACCGAATTGGACAAAAAGAGAATGTTGTGGTTTATAGGC
 TAATCACTTGTGGGACTGTAGAGGAAAAATATACAGAAGACAGGTTTTCAAGGACTCATTAATAAGACA
 AACTACTGGTGAAAAAAGAACCCTTTCCGATATTTTAGTAAACAAGAATTAAGAGAGCTCTTTACAATC
 GAGGATCTTCAGAACTCTGTAACCCAGCTGCAGCTTCAGTCTTTGCATGCTGCTCAGAGGAATCTGATA
 TAAAACTAGATGAACATATTGCCTACCTGCAGTCTTTGGGGATAGCTGGAATCTCAGACCATGATTGAT
 GTACACATGTGATCTGTCTGTAAAGAAGAGCTTGATGTGGTAGAAGAATCTCACTATATTCAACAAAGG
 GTTCAGAAAGCTCAATTCCTCGTTGAATTCGAGTCTCAAATAAAGAGTTCCTGATGGAACAACAAAGAA
 CTAGAAATGAGGGGGCTGGCTAAGAGAACCTGTATTTCTTCTCAACAAAGAAGAAATGCCCTAAATT
 GAATAAACACAGCCTCAGCCTTCACCTCTTCTAAGTACTCATCACTCAGGAAGAAGATATCAGTTCC
 AAAATGGCAAGTGTAGTCATTGATGATCTGCCAAAGAGGGTGAGAAACAAGATCTCTCCAGTATAAAGG
 TGAATGTTACCACCTGCAAGATGGTAAAGGTACAGGTAGTGTGACTCTATAGCTACTTTACCAAAGG
 GTTTGGAAGTGTAGAAGAACTTTGTAATACTCTTCATTGGGAATGGAAAAAGCTTTGCAACTAAAAAT
 GAAGCTGTACAAAAAGAGACATTACAAGAGGGGCTAAGCAAGAGGCACTGCAAGAGGATCCTCTGGAAA
 GTTTTAATTATGTACTTAGCAAATCAACCAAAGCTGATATTGGGCCAAATTTAGATCAACTAAAGGATGA
 TGAGATTTTACGTCAATTCCTTGGCCATTATTTCCATAACAAATGAAAGTCAAAATGCAGAAATCA
 AATGTATCCATTATTGAAATAGCTGATGACCTTCAGCATCCCATAGTGCAGTGCAGGATGCTCAAGCAA
 GTGAGGCCAAGTTGGAAGAGGAACCTTCAGCATCTTCACCACAGTATGCATGTGATTTCATCTTTTCTT
 GGAAGACTCAGCAGACAACAGACAAAATTTTCCAGTCAGTCTTTAGAGCATGTTGAGAAAGAAAATAGC
 TTGTGTGGCTCTGCACCTAATTCAGAGCAGGGTTTGTGCATAGCAAAACATGTCTCAGTTGGGAGTTTT
 CTGAGAAAGACGATGAACCAGAAGAAGTAGTAGTTAAAGCAAAATCAGAAGTAAAGCTAGAAGGATTGT
 TTCAGATGGCGAAGATGAAGATGATTCTTTTAAAGATACCTCAAGCATAAATCCATTCAACACATCTCTC
 TTTCAATTCTCATCTGTGAAACAATTTGATGCTTCAACTCCCAAAATGACATCAGTCCACCAGGAAGGT
 TCTTTTCATCTCAAAATCCAGTAGTGTAAATAAGTCTATGAACTCTAGAAGATCTCTGGCTTCTAGGAG
 GTCTCTTATTAATATGGTTTTAGACCACGTGGAGGACATGGAGGAAAGACTTGACGACAGCAGTGAAGCA
 AAGGGTCTGAAGATTATCCAGAAGAAGGGGTGGAGGAAAGCAGTGGCGAAGCCTCCAAGTATACAGAAG
 AGGATCCTTCCGAGAAACACTGTCTTCAGAAAACAAGTCCAGCTGGTTAATGACGTCTAAGCCTAGTGC
 TCTAGCTCAAGAGACCTCTCTTGGTGCCCTGAGCCTTTGTCTGGTGAACAGTTGGTTGGTTCTCCCCAG
 GATAAGGCGGCAGAGGCTACAAATGACTATGAGACTCTTGTAAGCGTGGAAGAAGCTAAAAGAGTGTG
 GAAAAATCCAGGAGGCCCTAAACTGCTTAGTTAAAGCGCTTGACATAAAAGTGCAGATCCTGAAGTTAT
 GCTCTTGACTTTAAGTTTGATAAGCACTTAATAACAAT

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Protein Sequence: >RC223966 representing NM_017669
 Red=Cloning site Green=Tags(s)

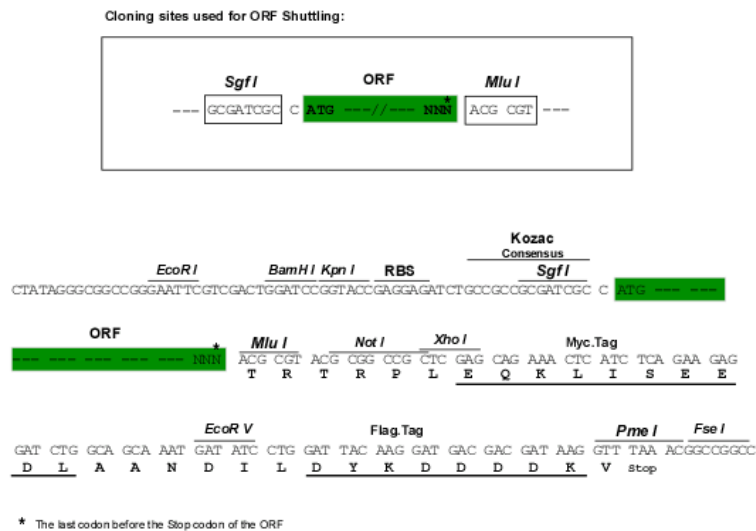
```
MEASRRFPEAEALSPEQAAHYLRVYKEAKEATKNGDLEEFKLFNLAKDIFPNEKVL SRIQKIQEAEEL
AEQGDDEFTDVCNSGLLLYRELHNQLFEHQKEGIAFLYSLYRDGRKGGILADDMGLGKTVQIIAFLSGMF
DASLVNHVLLIMPTNLINTWVKEFIKWTPGMRVKTFFHGPSKDERTRNLNRIQQRNGVITITYQMLINNQQ
QLSSFRGQEFVWDYVILDEAHKIKTSSTKSAICARAIPASNRLLLTGTPIQNNLQELWSLDFACQGSLL
GTLKTFKMEYENPITRAREKDATPGEKALGFKISENLMAIKPYFLRRTKEDVQKKKSSNPEARLNEKNP
DVDAICEMPSLSRKNDLIIWIRLVPLQEEIYRKFSVSLDHIKELLMETRSLAELGVKKLCDHPRLLSAR
ACCLLNLGTFSAQDNGEGEDSPDVHDIDQVTDLMEESGKMIFLMDLLKRLRDEGHQTLVFSQSRQILN
IIERLLKNRHFKTLRIDGTVTHLLEREKRINFQQNKDYSVFLTTQVGGVGLTLTAATRVVIFDPSWNP
ATDAQAVDRVYRIGQKENVVYRLITCGTVEEKIYRRQVFKDSLIRQTTGEKKNPFRYFSKQELRELFTI
EDLQNSVTQLQLQSLHAAQRKSDIKLDEHIAYLQSLGIAGISDHDLMYTCDLSVKEELDVEESHYIQQR
VQKAQFLVEFESQNKFEFLMEQQRTRNEGAWLREPVPSSSTKKKCPKLNKPQPSPLLSTHTQEEDISS
KMASVVIDDLPEKEGKQDLSSIKVNVTTLQDGKGTGSADSIATLPKGFSGVEELCTNSSLGMEKSFATKN
EAVQKETLQEGPKQEALQEDPLESFNYVLSKSTKADIGPNLDQLKDDEILRHCNPWPIISITNESQNAES
NYSIIIEIADDLSASHSALQDAQASEAKLEEEPSASSPQYACDFNLFLSDADNRQNFSSQSLEHVEKENS
LCGSAPNSRAGFVHSKTCLSWEFSEKDDPEEVVYKAKIRSKARRIVSDGEDEDDSKDTSSINPFNTSL
FQFSSVKQFDASTPKNDISPPGRFFSSQIPSSVNKSMNSRRSLASRRSLINMVLDHVEDMEERLDDSSSEA
KGPEDYPEEGVEESSGEASKYTEEDPSGETLSSENKSSWLMSTKPSALAQETSLGAPEPLSGEQLVGSPQ
DKAAEATNDYETLVKRGKELKECGKIQEALNCLVKALDIKSADPEVMLLTSLYKQLNNN
```

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Chromatograms: https://cdn.origene.com/chromatograms/mk8101_e06.zip

Restriction Sites: SgfI-MluI

Cloning Scheme:

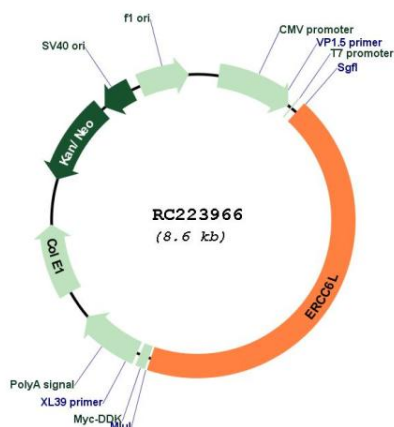


ACCN: NM_017669

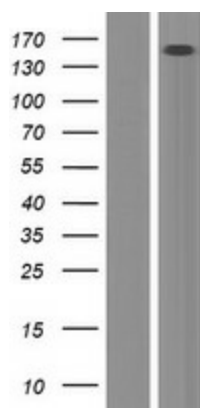
ORF Size: 3750 bp

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:	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:	NM_017669.4
RefSeq Size:	4224 bp
RefSeq ORF:	3753 bp
Locus ID:	54821
UniProt ID:	Q2NKK8
Cytogenetics:	Xq13.1
MW:	141.1 kDa
Gene Summary:	This gene encodes a member of the SWItch/Sucrose Non-Fermentable (SWI/SNF2) family of proteins, and contains a SNF2-like ATPase domain and a PICH family domain. One distinguishing feature of this SWI/SNF protein family member is that during interphase, the protein is excluded from the nucleus, and only associates with chromatin after the nuclear envelope has broken down. This protein is a DNA translocase that is thought to bind double-stranded DNA that is exposed to stretching forces, such as those exerted by the mitotic spindle. This protein associates with ribosomal DNA and ultra-fine DNA bridges (UFBs), fine structures that connect sister chromatids during anaphase at some sites such as fragile sites, telomeres and centromeres. This gene is required for the faithful segregation of sister chromatids during mitosis, and the ATPase activity of this protein required for the resolution of UFBs before cytokinesis. [provided by RefSeq, May 2017]

Product images:



Circular map for RC223966



Western blot validation of overexpression lysate (Cat# [LY413630]) using anti-DDK antibody (Cat# [TA50011-100]). Left: Cell lysates from untransfected HEK293T cells; Right: Cell lysates from HEK293T cells transfected with RC223966 using transfection reagent MegaTran 2.0 (Cat# [TT210002]).