

Product datasheet for **RC234792**

FGFR2 (NM_023029) Human Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	FGFR2 (NM_023029) Human Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	FGFR2
Synonyms:	BBDS; BEK; BFR-1; CD332; CEK3; CFD1; ECT1; JWS; K-SAM; KGFR; TK14; TK25
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)



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ORF Nucleotide Sequence:

>RC234792 representing NM_023029
 Red=Cloning site Blue=ORF Green=Tags(s)

CTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCCGGCGC
 GCC

ATGGTCAGCTGGGGTCGTTTCATCTGCCTGGTCGTGGTACCATGGCAACCTTGTCCCTGGCCCGCCCT
 CCTTCAGTTTAGTTGAGGATACACATTAGAGCCAGAAGATGCCATCTCATCCGGAGATGATGAGGATGA
 CACCGATGGTGGGAAGATTTTGTGAGTGAAGACAGTAACAACAAGAGAGCACCATACTGGACCAACACA
 GAAAAGATGAAAAGCGGCTCCATGCTGTGCCTGCGGCCAACACTGTCAAGTTTCGCTGCCAGCCGGGG
 GGAACCAATGCCAACCATGCGGTGGTGAAAAACGGGAAGGAGTTTAAGCAGGAGCATCGCATTGGAGG
 CTACAAGTACGAAACCAGCACTGGAGCCTCATTATGAAAGTGTGGTCCCATCTGACAAGGAAATTAT
 ACCTGTGTAGTGGAGAATGAATACGGTCCATCAATCACACGTACCACCTGGATGTTGTGGAGCGATCGC
 CTCACCGCCCATCCTCAAGCCGGACTGCCGCAAAATGCCTCCACAGTGGTCGGAGGAGACGTAGAGTT
 TGTCTGCAAGGTTTACAGTGTGCCAGCCCCACATCCAGTGGATCAAGCACGTGAAAAAGAACGGCAGT
 AAATACGGGCCCGACGGGCTGCCCTACCTCAAGGTTCTCAAGGCCCGGTGTTAACACCACGGACAAGG
 AGATTGAGGTTCTCTATATTCGGAATGTAACTTTTGAGGACGCTGGGGAATATACGTGCTTGGCGGGTAA
 TTCTATTGGGATATCCTTCACTCTGCATGTTGACAGTTCTGCCAGCGCCTGGAAGAGAAAAGGAGATT
 ACAGCTTCCCAGACTACCTGGAGATAGCCATTTACTGCATAGGGGTCTTCTTAATCGCCTGTATGGTGG
 TAACAGTCATCCTGTGCCGAATGAAGAACACGACCAAGAAGCCAGACTTCAGCAGCCAGCCGGCTGTGCA
 CAAGCTGACCAACCGTATCCCCTGCGGAGACAGGTAACAGTTTCGGCTGAGTCCAGCTCCTCCATGAAC
 TCCAACACCCCGCTGGTGAAGATAACAACACGCCTCTCTTCAACGGCAGACACCCCATGCTGGCAGGGG
 TCTCCGAGTATGAACTTCCAGAGGACCCAAAATGGGAGTTTCAAGAGATAAGCTGACACTGGGCAAGCC
 CCTGGGAGAAGGTTGCTTTGGGCAAGTGGTCATGGCGGAAGCAGTGGGAATTGACAAAGACAAGCCCAAG
 GAGGCGGTACCCTGGCCGTGAAGATGTTGAAAGATGATGCCACAGAGAAAGACCTTTCTGATCTGGTGT
 CAGAGATGGAGATGATGAAGATGATTGGGAAACACAAGAATATCATAAATCTTCTTGGAGCCTGCACACA
 GGATGGGCCTCTCTATGTCATAGTTGAGTATGCCTCTAAAGGCAACCTCCGAGAATACCTCCGAGCCCGG
 AGGCCACCCGGGATGGAGTACTCTATGACATTAACCGTGTTCCTGAGGAGCAGATGACCTTCAAGGACT
 TGGTGTGATGCACCTACCAGCTGGCCAGAGGCATGGAGTACTTGGCTTCCAAAAATGTATTCATCGAGA
 TTTAGCAGCCAGAAATGTTTTGGTAACAGAAAACAATGTGATGAAAATAGCAGACTTTGGACTCGCCAGA
 GATATCAACAATATAGACTATTACAAAAGACCACCAATGGGCGGCTTCCAGTCAAGTGGATGGCTCCAG
 AAGCCCTGTTTGTAGAGTATACACTCATCAGAGTGTGCTGGTCTTCCGGGTGTTAATGTGGGAGAT
 CTTCACTTTAGGGGGCTCGCCCTACCCAGGGATTCCCCTGGAGGAACTTTTTAAGCTGCTGAAGGAAGGA
 CACAGAATGGATAAGCCAGCCAACTGCACCAACGAACTGTACATGATGATGAGGGACTGTTGGCATGCAG
 TGCCCTCCAGAGACCAACGTTCAAGCAGTTGGTGAAGACTTGGATCGAATTCTCACTCTCACAAACCA
 TGAGGAATACTTGGACCTCAGCAACCTCTCGAACAGTATCACCTAGTTACCCTGACACAAGAAGTTCT
 TGTTCTTCAGGAGATGATTCTGTTTTTCTCCAGACCCCATGCCTTACGAACCATGCCTTCTCAGTATC
 CACACATAAACGGCAGTGTTAAAAACA

ACGCGTACGCGCGCCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Protein Sequence: >RC234792 representing NM_023029
Red=Cloning site Green=Tags(s)

MVSWGRFICLVVVTMATLSLARPSFSLVEDTTLEPEDAISSGDDDDTDGAEDFVSENSNNKRAPYWTNT
 EKMEKRLHAVPAANTVKFRCPAGGNPMPMTMRWLKNGKEFKQEHRIGGYKVRNQHWSLIMESVVP SDKGNY
 TCVVENEYGSINHTYHLDVVERSHPRPILQAGLPANASTVVGDDVEFVCKVYSDAQPHIQWIKHVEKNGS
 KYGPDGLPYLKVLKAAGVNTT DKEIEVL YIRNVTFEDAGEY TCLAGNSIGISFHSAWLTVLPAPGREKEI
 TASP DYLEIAIY CIGVFLIACMVVTVILCRMKNTTKKPDFSSQPAVHKLTKRIPLRRQVTVSAESSSMN
 SNTPLVRITTRLSTADTPMLAGVSEYELPEDPKWEPDRDKLTLGKPLGEGCFGQVYVMAEAVGIDKDKPK
 EAVTVAVKMLKDDATEKDLSDLVSEMEMMKMIGKHNIINLLGACTQDGPLYVIVEYASKGNLREYLRAR
 RPPGMEYSYDINRVPEEQMTFKDLVSCTYQLARGMEYLASQKCIHRDLAARNVLTENNVMKIADFLGAR
 DINNIDYKKTNGRLPVKWMPEALFDRVYTHQSDVWSFGVLMWEIFTLGGSPYPGIPVEELFKLLKEG
 HRMDKPANCTNELYMMMRDCWHAVPSQRPTFKQLVEDLDRILTLTTNEEYLDLSQPLEQYSPSPDTRSS
 CSSGDDSVFSPDPMPYEPCLPQYPHINGSVKT

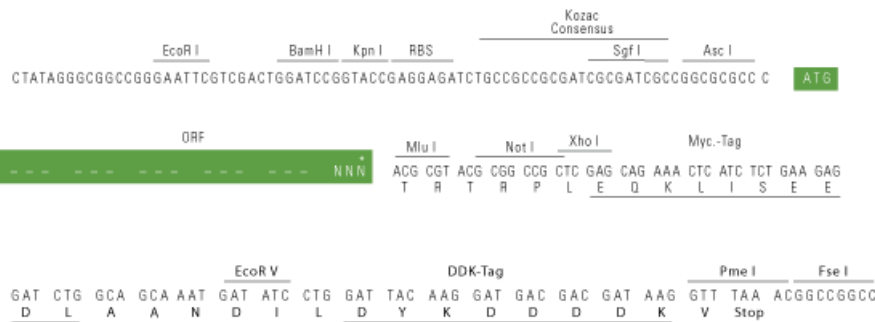
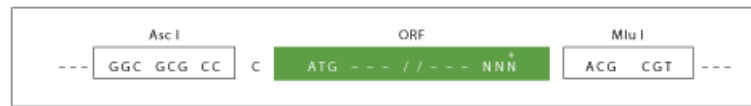
TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Restriction Sites:

AscI-MluI

Cloning Scheme:

Cloning sites used for ORF Shuttling:



* The last codon before the Stop codon of the ORF

ACCN: NM_023029

ORF Size: 2196 bp

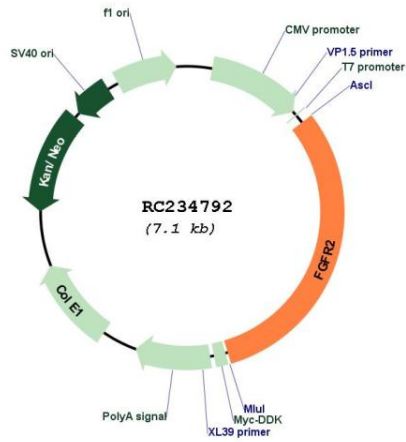
OTI Disclaimer: The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially 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_023029.2 , NP_075418.1
RefSeq Size:	3890 bp
RefSeq ORF:	2199 bp
Locus ID:	2263
UniProt ID:	P21802
Cytogenetics:	10q26.13
Domains:	pkinase, TyrKc, S_TKc, ig, IGc2, IG
Protein Families:	Druggable Genome, Protein Kinase, Secreted Protein, Transmembrane
Protein Pathways:	Endocytosis, MAPK signaling pathway, Pathways in cancer, Prostate cancer, Regulation of actin cytoskeleton
MW:	82.7 kDa
Gene Summary:	<p>The protein encoded by this gene is a member of the fibroblast growth factor receptor family, where amino acid sequence is highly conserved between members and throughout evolution. FGFR family members differ from one another in their ligand affinities and tissue distribution. A full-length representative protein consists of an extracellular region, composed of three immunoglobulin-like domains, a single hydrophobic membrane-spanning segment and a cytoplasmic tyrosine kinase domain. The extracellular portion of the protein interacts with fibroblast growth factors, setting in motion a cascade of downstream signals, ultimately influencing mitogenesis and differentiation. This particular family member is a high-affinity receptor for acidic, basic and/or keratinocyte growth factor, depending on the isoform. Mutations in this gene are associated with Crouzon syndrome, Pfeiffer syndrome, Craniosynostosis, Apert syndrome, Jackson-Weiss syndrome, Beare-Stevenson cutis gyrata syndrome, Saethre-Chotzen syndrome, and syndromic craniosynostosis. Multiple alternatively spliced transcript variants encoding different isoforms have been noted for this gene. [provided by RefSeq, Jan 2009]</p>

Product images:



Circular map for RC234792