

## Product datasheet for **SC215190**

### FGFR2 (NM\_001144916) Human 3' UTR Clone

#### Product data:

Product Type:	3' UTR Clones
Product Name:	FGFR2 (NM_001144916) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	FGFR2
Synonyms:	BBDS; BEK; BFR-1; CD332; CEK3; CFD1; ECT1; JWS; K-SAM; KGFR; TK14; TK25
ACCN:	NM_001144916
Insert Size:	1555 bp



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**Insert Sequence:** >SC215190 3'UTR clone of NM\_001144916  
 The sequence shown below is from the reference sequence of NM\_001144916. The complete sequence of this clone may contain minor differences, such as SNPs.  
 Blue=Stop Codon Red=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
CCACACATAAACGGCAGTGTTAAAACATGAATGACTGTGTCTGCCTGTCCCAAACAGGACAGCACTGG
GAACCTAGCTACTGAGCAGGGAGACCATGCCTCCCAGAGCTTGTGTCTCCACTTGTATATATGGAT
CAGAGGAGTAAATAATTGAAAAAGTAATCAGCATATGTGTAAAGATTTATACAGTTGAAAACCTGTAAAT
CTTCCCAGGAGGAGAAGAAGGTTTCTGGAGCAGTGGACTGCCACAAGCCACCATGTAACCCCTCTCAC
CTGCCGTGCGTACTGGCTGTGGACCAGTAGGACTCAAGGTGGACGTGCGTTCTGCCTTCCTTGTAAAT
TTGTAATAATTGGAGAAGATTTATGTCAGCACACACTTACAGAGCACAAATGCAGTATATAGGTGCTGG
ATGTATGTAATATATTTCAAATTTATGTATAAATATATATTATATATTTACAAGGAGTTATTTTTGTAT
TGATTTTAAATGGATGTCCCAATGCACCTAGAAAATTGGTCTCTCTTTTTTAATAGCTATTTGCTAAA
TGCTGTTCTTACACATAATTTCTTAATTTTCACCGAGCAGAGGTGAAAAATACTTTTGCTTTCAGGGA
AAATGGTATAACGTTAATTTATTAATAAATTTGGTAATATACAAAACAATTAATCATTTATAGTTTTTTT
TGTAATTTAAGTGGCATTCTATGCAGGCAGCACAGCAGACTAGTTAATCTATTGCTTGGACTTAACTA
GTTATCAGATCCTTTGAAAAGAGAATATTTACAATATATGACTAATTTGGGGAAAATGAAGTTTTGATT
TATTTGTGTTTAAATGCTGCTGTCAGACGATTGTTCTTAGACCTCCTAAATGCCCATATTTAAAGAAC
TCATTCATAGGAAGGTGTTTCATTTTGGTGTGCAACCCTGTCATTACGTCAACGCAACGTCTAACTGGA
CTTCCCAAGATAAATGGTACCAGCGTCTCTTAAAAGATGCCTTAATCCATTCTTGAGGACAGACCTT
AGTTGAAATGATAGCAGAATGTGCTTCTCTGGCAGCTGGCCTTCTGCTTCTGAGTTGCACATTAATC
AGATTAGCCTGTATTCTCTTCAGTGAATTTGATAATGGCTTCCAGACTCTTTGGCGTTGGAGACGCCT
GTTAGGATCTTCAAGTCCCATCATAGAAAAATTGAAACACAGAGTTGTTCTGCTGATAGTTTTGGGGATA
CGTCCATCTTTTTAAGGGATTGCTTTCATCTAATTCTGGCAGGACCTCACAAAAGATCCAGCCTCATA
CCTACATCAGACAAAATATCGCCGTTGTTCTTCTGTACTAAAGTATTGTGTTTTGCTTTGGAAACACC
CACTCACTTTGCAATAGCCGTGCAAGATGAATGCAGATTACACTGATCTTATGTGTTACAAAATTGGAG
AAAGTATTTAATAAAAACCTGTTAATTTTTATACTGACAATAAAAAATGTTTCTACAGATATTAATGTTAA
CAAGACAAAATAAATGTCACGCAACTTATTTTTTAA
ACGCGTAAGCGGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTTTCGATTCCACCGCCGCTTCTATGAAAGG
  
```

**Restriction Sites:** SgfI-MluI

**OTI Disclaimer:** Our molecular clone sequence data has been matched to the sequence identifier above as a point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences , e.g., single nucleotide polymorphisms (SNPs).

**Components:** The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The package also includes 100 pmols of both the corresponding 5' and 3' vector primers in separate vials.

**RefSeq:** [NM\\_001144916.2](#)

**Summary:**

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]

**Locus ID:**

2263

**MW:**

60.1