

## **Product datasheet for TL320354**

## OriGene Technologies, Inc.

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## **FGFR1** Human shRNA Plasmid Kit (Locus ID 2260)

**Product data:** 

**Product Type:** shRNA Plasmids

**Product Name:** FGFR1 Human shRNA Plasmid Kit (Locus ID 2260)

Locus ID: 2260

Synonyms: H2, H3, H4, H5, CEK, FLG, FLT2, KAL2, BFGFR, C-FGR, CD331, N-SAM

Vector: pGFP-C-shLenti (TR30023)

E. coli Selection: Chloramphenicol (34 ug/ml)

**Mammalian Cell** 

Puromycin

Selection:

Format: Lentiviral plasmids

**Components:** FGFR1 - Human, 4 unique 29mer shRNA constructs in lentiviral GFP vector(Gene ID = 2260).

5µg purified plasmid DNA per construct

29-mer scrambled shRNA cassette in pGFP-C-shLenti Vector, TR30021, included for free.

RefSeq: NM 000604, NM 001174063, NM 001174064, NM 001174065, NM 001174066,

NM 001174067, NM 015850, NM 023105, NM 023106, NM 023107, NM 023108, NM 023109, NM 023110, NM 023111, NM 032191, NM 001354367, NM 001354368, NM 001354369, NM 001354370, NM 023110.1, NM 023110.2, NM 023105.1, NM 023105.2, NM 023107.1, NM 023107.2, NM 023108.1, NM 023108.2, NM 015850.2, NM 015850.3, NM 023106.1, NM 023106.2, NM 001174066.1, NM 001174064.1, NM 001174065.1, NM 001174063.1, NM 001174067.1, NM 023111.1, BC018128, BC018128.1, BC091494, BC015035, NM 023110.3,

NM 001174064.2, NM 023105.3, NM 023106.3, NM 001174066.2, NM 001174063.2,

NM 001174065.2, NM 015850.4

UniProt ID: P11362



## Summary:

The protein encoded by this gene is a member of the fibroblast growth factor receptor (FGFR) 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 binds both acidic and basic fibroblast growth factors and is involved in limb induction. Mutations in this gene have been associated with Pfeiffer syndrome, Jackson-Weiss syndrome, Antley-Bixler syndrome, osteoglophonic dysplasia, and autosomal dominant Kallmann syndrome 2. Chromosomal aberrations involving this gene are associated with stem cell myeloproliferative disorder and stem cell leukemia lymphoma syndrome. Alternatively spliced variants which encode different protein isoforms have been described; however, not all variants have been fully characterized. [provided by RefSeq, Jul 2008]

shRNA Design:

Performance Guaranteed:

These shRNA constructs were designed against multiple splice variants at this gene locus. To be certain that your variant of interest is targeted, please contact <a href="mailto:techsupport@origene.com">techsupport@origene.com</a>. If you need a special design or shRNA sequence, please utilize our <a href="mailto:custom shRNA service">custom shRNA service</a>.

OriGene guarantees that the sequences in the shRNA expression cassettes are verified to correspond to the target gene with 100% identity. One of the four constructs at minimum are guaranteed to produce 70% or more gene expression knock-down provided a minimum transfection efficiency of 80% is achieved. Western Blot data is recommended over qPCR to evaluate the silencing effect of the shRNA constructs 72 hrs post transfection. To properly assess knockdown, the gene expression level from the included scramble control vector must be used in comparison with the target-specific shRNA transfected samples.

For non-conforming shRNA, requests for replacement product must be made within ninety (90) days from the date of delivery of the shRNA kit. To arrange for a free replacement with newly designed constructs, please contact Technical Services at techsupport@origene.com. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data preferred).