

FGFR4 Blocking Peptide (N-term) Synthetic peptide Catalog # BP20781a

# Specification

# FGFR4 Blocking Peptide (N-term) - Product Information

Primary Accession Other Accession P22455 Q03142

# FGFR4 Blocking Peptide (N-term) - Additional Information

Gene ID 2264

**Other Names** Fibroblast growth factor receptor 4, FGFR-4, CD334, FGFR4, JTK2, TKF

Target/Specificity The synthetic peptide sequence is selected from aa 143-158 of HUMAN FGFR4

#### Format

Peptides are lyophilized in a solid powder format. Peptides can be reconstituted in solution using the appropriate buffer as needed.

**Storage** Maintain refrigerated at 2-8°C for up to 6 months. For long term storage store at -20°C.

**Precautions** This product is for research use only. Not for use in diagnostic or therapeutic procedures.

## **FGFR4 Blocking Peptide (N-term) - Protein Information**

Name FGFR4

Synonyms JTK2, TKF

### Function

Tyrosine-protein kinase that acts as a cell-surface receptor for fibroblast growth factors and plays a role in the regulation of cell proliferation, differentiation and migration, and in regulation of lipid metabolism, bile acid biosynthesis, glucose uptake, vitamin D metabolism and phosphate homeostasis. Required for normal down- regulation of the expression of CYP7A1, the rate-limiting enzyme in bile acid synthesis, in response to FGF19. Phosphorylates PLCG1 and FRS2. Ligand binding leads to the activation of several signaling cascades. Activation of PLCG1 leads to the production of the cellular signaling molecules diacylglycerol and inositol 1,4,5-trisphosphate. Phosphorylation of FRS2 triggers recruitment of GRB2, GAB1, PIK3R1 and SOS1, and mediates activation of RAS, MAPK1/ERK2, MAPK3/ERK1 and the MAP kinase signaling pathway, as well as of the AKT1 signaling pathway. Promotes SRC-dependent phosphorylation of the matrix protease MMP14 and its lysosomal degradation. FGFR4 signaling is down-regulated by receptor internalization and degradation; MMP14 promotes internalization and degradation of FGFR4.



Mutations that lead to constitutive kinase activation or impair normal FGFR4 inactivation lead to aberrant signaling.

### **Cellular Location**

Cell membrane; Single-pass type I membrane protein. Endosome. Endoplasmic reticulum. Note=Internalized from the cell membrane to recycling endosomes, and from there back to the cell membrane

### **Tissue Location**

Expressed in gastrointestinal epithelial cells, pancreas, and gastric and pancreatic cancer cell lines

# **FGFR4 Blocking Peptide (N-term) - Protocols**

Provided below are standard protocols that you may find useful for product applications.

### <u>Blocking Peptides</u>

## FGFR4 Blocking Peptide (N-term) - Images

## FGFR4 Blocking Peptide (N-term) - Background

Tyrosine-protein kinase that acts as cell-surface receptor for fibroblast growth factors and plays a role in the regulation of cell proliferation, differentiation and migration, and in regulation of lipid metabolism, bile acid biosynthesis, glucose uptake, vitamin D metabolism and phosphate homeostasis. Required for normal down-regulation of the expression of CYP7A1, the rate-limiting enzyme in bile acid synthesis, in response to FGF19. Phosphorylates PLCG1 and FRS2. Ligand binding leads to the activation of several signaling cascades. Activation of PLCG1 leads to the production of the cellular signaling molecules diacylglycerol and inositol 1,4,5-trisphosphate. Phosphorylation of FRS2 triggers recruitment of GRB2, GAB1, PIK3R1 and SOS1, and mediates activation of RAS, MAPK1/ERK2, MAPK3/ERK1 and the MAP kinase signaling pathway, as well as of the AKT1 signaling pathway. Promotes SRC-dependent phosphorylation of the matrix protease MMP14 and its lysosomal degradation. FGFR4 signaling is down-regulated by receptor internalization and degradation; MMP14 promotes internalization and degradation of FGFR4. Mutations that lead to constitutive kinase activation or impair normal FGFR4 inactivation lead to aberrant signaling.

## FGFR4 Blocking Peptide (N-term) - References

Partanen J.M., et al.EMBO J. 10:1347-1354(1991). Ron D., et al.J. Biol. Chem. 268:5388-5394(1993). Takaishi S., et al.Biochem. Biophys. Res. Commun. 267:658-662(2000). Kostrzewa M., et al.Mamm. Genome 9:131-135(1998). Ezzat S., et al.J. Clin. Invest. 109:69-78(2002).