

Certificate of Analysis

FGFR1 (V561M), active

(Recombinant enzyme expressed in Sf21 insect cells)

Item # 14-734, 14-734-K, 14-734M

Parent Lot # D7BN045U

The data presented in this document apply to the parent lot shown above and to all pack sizes derived from subsequent vialling runs of this parent lot. An alphabetical suffix after the parent lot number is used to denote each vialling run.

Product Description: N-terminal GST-tagged, recombinant, human FGFR1 amino acids 456–765 containing the V561 mutation. Expressed by baculovirus in Sf21 insect cells. Purified using glutathione agarose. Co-ordinates encompass the protein kinase domain (478–767) according to SP P11362. The coordinates for recombinant human FGFR1 were described in Homann *et al*, J Biotech (2001); **86**:51-58. Blencke *et al* (2004) suggested mutations to a conserved threonine residue at the ATP binding site would result in inhibitor resistance. The amino acid valine 561 was mutated to a methionine in FGFR1 which corresponded to previously reported mutations found in Abl (T315) and EGFR (T766) which had been shown to confer resistance to selective inhibitors. Assay data for FGFR1 V561M had shown that this mutation had conferred resistance to PP58 (pyrido [2,3-d] pyrimidine tyrosine kinase inhibitor) compared with that of the wild type. (Blencke, S. *et al*, Chemistry & Biology (2004);**11**:691-701) Purity 100% by SDS-PAGE and Coomassie blue staining. MW = 62.3kDa.

Formulation: 1.2mg/ml of enzyme in 50mM Tris/HCl pH7.5, 300mM NaCl, 0.1mM EGTA, 0.03% Brij-35, 270mM sucrose, 1mM benzamidine, 0.2mM PMSF, 0.1% 2-mercaptoethanol. Frozen solution.

Storage and Stability: On receipt of material store at -70°C. Unopened reagent is stable for a minimum of 1 year from date of shipment when stored at recommended storage temperature. Avoid repeat freeze/thaw cycles. For maximum recovery of product, centrifuge original vial prior to removing the cap.

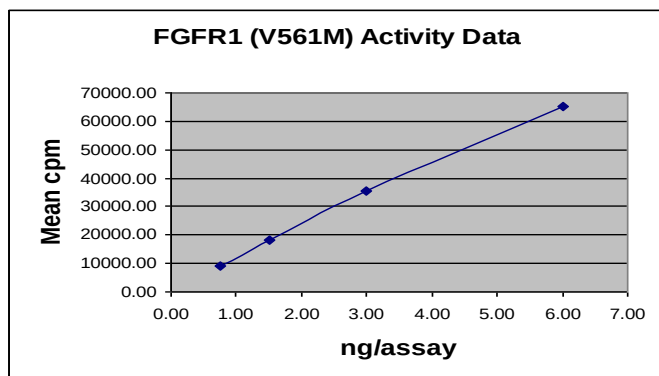
Handling Recommendations: Rapidly thaw the vial under cold water and immediately place on ice. Aliquot unused material into pre-chilled micro-centrifuge tubes and immediately snap-freeze the vials in liquid nitrogen prior to re-storage at -70°C.

Specific Activity (Parent lot# D7BN045U): 4741U/mg, where one unit of FGFR1 (V561M), active activity is defined as 1nmol phosphate incorporated into 500µM (GGEEEEYFELVKKKK) per minute at 30°C with a final ATP concentration of 100µM.

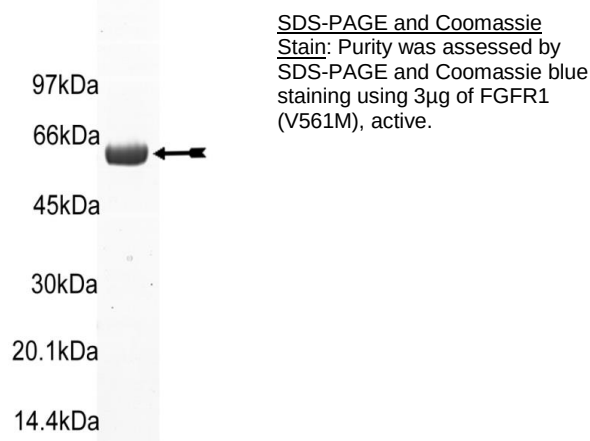
**FOR IN VITRO RESEARCH USE ONLY
NOT FOR USE IN HUMANS OR ANIMALS**

Quality Control Testing

Kinase Assay: 0.75–6.00ng of this lot of enzyme phosphorylated 500µM (GGEEEEYFELVKKKK) in the assay described on page two. Assay background was subtracted from the actual counts to yield the results shown below.



MS Tryptic Fingerprint: Confirmed product identity as FGFR1 (V561M) with the translated native sequence listed on page three.



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Kinase Assay Protocol

Stock Solutions:

1. **5 x Reaction Buffer:** 40mM MOPS/NaOH pH7.0, 1mM EDTA.
2. **(GGEEEEYFELVKKKK):** Use at a final assay concentration of 500µM. Prepare a 5mM stock and add 2.5µl of stock per assay point.
3. **FGFR1 (V561M), active:** Dilute with 20mM MOPS/NaOH pH7.0, 1mM EDTA, 0.01% Brij-35, 5% glycerol, 0.1% 2-mercaptoethanol, 1mg/ml BSA. Use 0.75–6.00ng per assay point.
4. **[γ -³³P]ATP:** 2.5 x magnesium acetate/[γ -³³P]ATP cocktail: 25mM MgAc and 0.25mM ATP to which is added [γ -³³P]ATP (specific activity approximately 500 - 800cpm/pmol as required.)

Assay Procedure (96 well plate format):

1. Add 5µl of 5 x reaction buffer per assay to wells.
2. Add 2.5µl of **(GGEEEEYFELVKKKK)**.
3. Add **2.5µl (0.75–6.00ng) FGFR1 (V561M), active**.
4. Add 10µl of diluted [γ -³³P]ATP mixture.
5. Add 5µl of dH₂O.
6. Incubate for 10 minutes at 30°C.
7. Stop the reaction by adding 5µl of 3% phosphoric acid.
8. Transfer a 10µl aliquot onto the appropriate area of a **P30 Filtermat**.
9. Wash the filtermat three times for 5 minutes with 75mM phosphoric acid.
10. Wash the filtermat once for 2 minutes with methanol.
11. Transfer the filtermat to a sealable plastic bag and add 4ml of scintillation cocktail.
12. Read in a scintillation counter. Compare cpm of enzyme samples with cpm of control samples that contain all assay components plus 1µl of 30% phosphoric acid.

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FGFR1 (V561M) Sequence Information

Protein	Human FGFR1 (V561M)
Tags	N-terminal GST
Native sequence	M231 of recombinant sequence is equivalent to M456 of native human FGFR1
Accession number	GenBank NM_000604

Recombinant FGFR1 (V561M) amino acid sequence:

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1  MSPILGYWKI  KGLVQPTRLL  LEYLEEKYEE  HLYERDEGDK  WRNKKFELGL  EFPNLPYYID
61  GDVKLTQSMa  IIRYIADKHN  MLGGCPKERA  EISMLEGAVL  DIRYGVSRIA  YSKDFETLKV
121 DFLSKLPEML  KMFEDRLCHK  TYLNGDHVTH  PDFMLYDALD  VVLYMDPMCL  DAFPKLVCFK
181 KRIEAIPOID  KYLKSSKYIA  WPLQGWQATF  GGGDHPPKSD  LEVLFQGPFE  MLAGVSEYEL
241 PEDPRWELPR  DRLVLGKPLG  EGCFGQVVLA  EAIGLDKDKP  NRVTKVAVKM  LKSDATEKDL
301 SDLISEMEMM  KMIGKHKNI  NLLGACTQDG  PLYVIMEYAS  KGNLREYLQA  RRPPGLEICY
361 NPSHNPEEQL  SSKDLVSCAY  QVARGMEYLA  SKKCIHRDLA  ARNVLVTEEN  VMKIADFGLA
421 RDIHHIDYYK  KTTNGRLPVK  WMAPEALFDR  IYTHQSDVWS  FGVLLWEIFT  LGGSPYPGVP
481 VEELFKLLKE  GHRMDKPSNC  TNELYMMMRD  CWHAVPSQRP  TFKQLVEDLD  RIVALTSNQE

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Recombinant FGFR1 (V561M) nucleotide sequence:

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1  atgtccccta  tactaggtta  ttggaaaatt  aagggccttg  tgcaaccac  tgcattctt
61  ttggaatatc  ttgaagaaaa  atatgaagag  catttgatg  agcgcgatga  aggtgataaa
121  tggcgaaaca  aaaagtttga  attgggtttg  gagtttcca  atcttcctta  ttatatgtat
181  ggtgatgta  aattaacaca  gtctatggcc  atcatacgtt  atatactga  caagcacaa
241  atgttgggtg  gttgtccaaa  agagcgtgca  gagatttcaa  tgcttgaagg  agcggttttg
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421  acatatatta  atggtgatca  tgtaacccat  cctgacttca  tggttgatga  cgctcttgat
481  gttgttttat  acatggaccc  aatgtgcctg  gatgcgttcc  caaaattagt  ttgttttaaa
541  aaacgtattg  aagctatccc  acaaattgat  aagtacttga  aatccagcaa  gtatatagca
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661  ctggaagttc  tggtccaggg  gcccgaaatt  atgctagcag  ggggtctctg  gtatgagctt
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901  tcagacctga  tctcagaaat  ggagatgatg  aagatgatcg  ggaagcataa  gaatatcatc
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1141 caggtggccc  gaggcattga  gtatctggcc  tccaagaagt  gcataaccgg  agacctggca
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1321 tggatggcac  ccgaggcatt  atttgaccgg  atctacaccc  accagagtga  tgtgtggtct
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1561 accttcaagc  agctggtgga  agacctggac  cgcacgtggt  ccttgacctc  caaccaggag
1621 taa

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