

# University of Dundee

## Standard Operating Procedure

### Preparation of active JNK1 $\alpha$ 1 / SAPK1c

<b><u>Enzyme description:-</u></b>	Active JNK1 $\alpha$ 1 / SAPK1c
<b><u>Source:-</u></b>	Recombinant
<b><u>Expression system:-</u></b>	Baculovirus expression vector system (BEVS)/Insect cells
<b><u>Tag:-</u></b>	His(6)
<b><u>Purification method:-</u></b>	Ni <sup>2+</sup> -NTA agarose.
<b><u>Expression level:-</u></b>	2-3 mg/L
<b><u>Molecular mass:-</u></b>	43 kDa by SDS-PAGE
<b><u>Purity:-</u></b>	>90%
<b><u>Contaminants:-</u></b>	No major contaminating proteins as judged by SDS-PAGE.

### **Activation protocol:-**

JNK1 $\alpha$ 1/ SAPK1c (0.086 mg/ml - 2  $\mu$ M) is activated in 50mM Tris/HCl pH 7.5, 0.1mM EGTA, 0.1 %  $\beta$ -mercaptoethanol, 0.1 mM sodium vanadate, 10 mM magnesium acetate, 0.1 mM ATP with active GST-MKK4 (13  $\mu$ g/ml - 200 nM) and GST-MKK7 $\beta$  (15  $\mu$ g/ml - 200 nM) at 30°C for 40 min. Following activation, the enzyme is separated from the GST-MKK4 and GST-MKK7 $\beta$  by Ni-NTA agarose chromatography (JNK1 $\alpha$ 1/ SAPK1c binds to the column by virtue of its His-tag). The re-purified active JNK1 $\alpha$ 1/ SAPK1c is then eluted from the column and dialysed into enzyme storage buffer and fractions are pooled and snap frozen in liquid nitrogen prior to storage at -70°C.

### **Enzyme storage buffer:-**

50 mM Tris/HCl pH 7.5, 270 mM sucrose, 150 mM NaCl, 0.1 mM EGTA, 0.1 %  $\beta$ -mercaptoethanol, 0.02% Brij-35, 0.2 mM PMSF, 1 mM Benzamidine.

**Storage temperature:-** Aliquot, snap freeze and store at -70°C.

# University of Dundee

## CLONE DATA SHEET –Human JNK1 $\alpha$ 1/ SAPK1c

<b><u>Protein</u></b>	Human JNK1 $\alpha$ 1 / SAPK1c
<b><u>Accession number</u></b>	L26318
<b><u>Tags</u></b>	His(6)
<b><u>Baculovirus-expressed protein</u></b>	MHHHHHMSRSKRDNNFYSVEIGDSTFTVLK RYQNLKPIGSGAQGIVCAAYDAILERNVAIKK LSRPFQNTQTHAKRAYRELVLMKCVNHKNIIGL LNVFTPQKSLEEFQDVYIVMELMDANLCQVIQ MELDHERMSYLLYQMLCGIKHLHSAGIIHRDL KPSNIVVKS DCTLKILDFGLARTAGTSFMMTP YVTRYRAPEVILGMGYKENVDLWSVGCIM GEMVCHKILFPGRDYIDQWNKVIEQLGTPCPE FMKKLQPTVRTYVENRPKYAGYSFEKLFDPVL FPADSEHNKLKASQARDLLSKMLVIDASKRIS VDEALQHPYINVWYDPSEAEAPPKIPDKQLD EREHTIEEWKELIYKEVMDLEERTKNGVIRGQP SPLAQVQQ
<b><u>Native sequence</u></b>	Met8 of the fusion-protein is Met1 of JNK1 $\alpha$ 1. / SAPK1c There is a His(6)-Tag at residues 2-7.
<b><u>Protease cleavage site</u></b>	None
<b><u>Cloning sites</u></b>	Nde1 / XhoI sites of modified pFastBAC 1.
<b><u>JNK1<math>\alpha</math>1/SAPK1C NUCLEOTIDE SEQUENCE</u></b>	ATGCACCATCACCATCACCATATGAGCAGAAGCAAGCGTGACAAC AATTTTTATAGTGTAGAGATTGGAGATTCTACATTCACAGTCCTG AAACGATATCAGAATTTAAAACCTATAGGCTCAGGAGCTCAAGGA ATAGTATGCGCAGCTTATGATGCCATTCTTGAAAGAAATGTTGCA ATCAAGAAGCTAAGCCGACCATTTGAGAATCAGACTCATGCCAAG CGGGCCTACAGAGAGCTAGTTCTTATGAAATGTGTTAATCACAAA AATATAATTGGCCTTTTGAATGTTTTACACCACAGAAATCCCTA GAAGAATTTCAAGATGTTTACATAGTCATGGACCTCATGGATGCA AATCTTTGCCAAGTGATTGAGATGGAGCTAGATCATGAAAGAATG TCCTACCTTCTCTATCAGATGCTGTGTGGAATCAAGCACCTTCAT TCTGCTGGAATTATTCATCGGGACTTAAAGCCCAGTAATATAGTA GTAAAATCTGATTGCACTTTGAAGATTCTTGACTTCGGTCTGGCC

## University of Dundee

AGGACTGCAGGAACGAGTTTTATGATGACGCCTTATGTAGTGACT  
CGCTACTACAGAGCACCCGAGGTCATCCTTGGCATGGGCTACAAG  
GAAAACGTGGATTTATGGTCTGTGGGGTGCATTATGGGAGAAATG  
GTTTGCCACAAAATCCTCTTTCCAGGAAGGGACTATATTGATCAG  
TGGAATAAAGTTATTGAACAGCTTGGAACACCATGTCCTGAATTC  
ATGAAGAACTGCAACCAACAGTAAGGACTTACGTTGAAAACAGA  
CCTAAATATGCTGGATATAGCTTTGAGAAACTCTTCCCTGATGTC  
CTTTTCCCAGCTGACTCAGAACACAACAAACTTAAAGCCAGTCAG  
GCAAGGGATTTGTTATCCAAAATGCTGGTAATAGATGCATCTAAA  
AGGATCTCTGTAGATGAAGCTCTCCAACACCCGTACATCAATGTC  
TGGTATGATCCTTCTGAAGCAGAAGCTCCACCACCAAAGATCCCT  
GACAAGCAGTTAGATGAAAGGGAACACACAATAGAAGAGTGGAAA  
GAATTGATATATAAGGAAGTTATGGACTTGGAGGAGAGAACCAAG  
AATGGAGTTATACGGGGGCAGCCCTCTCCTTTAGCACAGGTGCAG  
CAGtag