

## Standard Operating Procedure

### Preparation of active MSK1

|                                     |  |
|-------------------------------------|--|
| <b>Enzyme description:-</b>         | Active MSK1  |
| <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>    | 5 mg/L   |
| <b><u>Molecular mass:-</u></b>      | 95 kDa by SDS-PAGE   |
| <b><u>Purity:-</u></b>              | >90%   |
| <b><u>Contaminants:-</u></b>        | The preparation contains several minor degradation products. |

### **Activation protocol:-**

MSK1 (0.2 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 2 U/ml active GST-MAPK2/ERK2 at 30°C for 45 min. Following activation, the MSK1 is separated from the GST-MAPK2/ERK2 by incubation on GSH-agarose for 30 min (to remove most of the MAPK2/ERK2 followed by Ni-NTA chromatography (MSK1 binds to the resin by virtue of its His-tag). The re-purified active MSK1 is then eluted from the column and dialysed into enzyme storage buffer, snap frozen in liquid nitrogen prior to storage at -70°C.

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

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

**Storage temperature:-** -20°C

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## CLONE DATA SHEET – human MSK1

|   |   |
|---|---|
| <b><u>Protein</u></b>                       | Human MSK1  |
| <b><u>Accession no</u></b>                  | AF074393  |
| <b><u>Tags</u></b>                          | His6 and DYKDDDDK (FLAG) amino-terminal tags  |
| <b><u>Baculovirus-expressed protein</u></b> | MSYYHHHHHHHDYDIPTTENLYFQGAMGSATMDYK<br>DDDDKEEEGGSSGGAAGTSADGGDGGEQLLTVKH<br>ELRTANLTGHAEKVGIENFELLKVLGTGAYGKVFLV<br>RKISGHDTGKLYAMKVLKKATIVQKAKTTEHTRTER<br>QVLEHIRQSPFLVTLHYAFQTETKLHLILDYINGGEL<br>FTHLSQRERFTEHEVQIYVGEIVLALEHLHKLGIIRD<br>IKLENILLDSNGHVVLTDGFLSKEFVADETERAYSFC<br>GTIEYMAPDIVRGGDSGHDKAVDWWSLGVLMYEL<br>LTGASPFTVDGEKNSQAEISRRILKSEPPYPQEMSAL<br>AKDLIQRLLMKDPKRLGCGPRDADEIKEHLFFQKI<br>NWDDLAAKKVPAPFKPVIRDELVDVSNFAEEFTEMD<br>PTYSPAALPQSSEKLFQGYSFVAPSILFKRNAAVIDPL<br>QFHMGVERPGVTNVARSAMMKDSPFYQHYDLDLK<br>DKPLGEGSFSICRKCIVHKKSNQAFVAVKIISKRMEAN<br>TQKEITALKLCEGHPNIVKLHEVFHDQLHTFLVMEL<br>LNGGELFERIKKKKHFSETEASYIMRKLVS AVSHMH<br>DVGVVHRDLKPENLLFTDENDNLEIKIIDFGFARLKP<br>PDNQPLKTPCFTLHYAAPPELLNQNGYDESCDLWSL<br>GVILYTMLSGQVPFQSHDRSLTCTSAVEIMKKIKKGD<br>FSFEGEAWKNVSQEAKDLIQGLLTVDPNKRLKMSG<br>LRYNEWLQDGSQSSNPLMTPDILGSSGAAVHTCV<br>KATFHAFNKYKREGFCLQNVDKAPLAKRRKMKKT<br>STSTETRSSSESSHSSSHSHGKTTPTKTLQPSNPADS<br>NNPETLFQFSDSVA |
| <b><u>Native sequence</u></b>               | Residue 40 of the His <sub>6</sub> -tagged protein is equivalent to Glu 2 of MSK1. The fusion protein contains an N-terminal His <sub>6</sub> tag (residues 5-10) and Flag tag (DYKDDDDK, residues 32-39).  |
| <b><u>Protease cleavage site</u></b>        | ENLYFQ (rTEV protease) residues 18 – 23 of His <sub>6</sub> -tagged protein   |
| <b><u>Cloning sites</u></b>                 | A <i>Bam</i> HI site was introduced immediately 5' to the initiator ATG by PCR. The resulting <i>Bam</i> HI / <i>Kpn</i> I  |

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fragment was sub-cloned into the *Bam*HI/*Kpn*I sites of pFastBAC HTb.

### ORF in baculovirus

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AGATCTGCCACCATGGACTACAAGGACGACGATGACAAGGA
GGAGGAGGGTGGCAGCAGCGGCGGCGCCGCGGGGACCAGCG
CGGACGGCGGCGACGGAGGAGAGCAGCTCCTCACTGTCAAG
CACGAGCTGCGGACTGCTAATTTGACAGGACATGCTGAGAA
GGTGGGAATAGAAAATTTTGAGCTCCTGAAGGTCCTAGGAA
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GGCCATGATACTGGAAAAGCTGTATGCCATGAAAATTTTAA
AAAGGCAACAATCGTTCAAAGGCCAAAACCACAGAGCATA
CAAGGACAGAACGACAAGTCCTGGAACACATTAGGCAGTCG
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CAAACCTCATCTCATTTTAGATTATATAAATGGTGGTGAAC
TTTTTACTCATCTTTCTCAAAGAGAGCGTTTCCAGAGCAT
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TTGAGAATATTCTACTTGATTCTAATGGCCATGTGGTGCTG
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GCAGTTGACTGGTGGAGTTTGGGTGTTCTAATGTATGAATT
ACTAACTGGAGCATCTCCTTTCACTGTTGATGGAGAAAAAA
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CCTCCATATCCCCAAGAAATGAGTGCTTTAGCGAAAGACCT
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GCATGATGTTGGAGTGGTGCACAGGGATCTGAAACCTGAGA
ATTTATTGTTTACCAGATGAAAATGACAATTTGGAAATTA AA
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TCAGCCCCCTGAAGACTCCATGCTTACCCTTCATTATGCCG
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CCCCAGAGCTCTTGAATCAGAACGGCTACGATGAGTCCTGT  
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