

# ***MRC PPU REAGENTS***

## **Standard Operating Procedure**

### **Preparation of ASK1 [2 - 1374]**

**Enzyme description:-** ASK1 [2 – 1374]

**Clone number:-** DU 4707

**Source:-** Recombinant

**Expression system:-** *E.coli*

**Tag:-** N-terminal GST

**Purification method:-** GSH Sepharose

**Calculated molecular mass:-**

Monoisotopic 181, 792.47 daltons

Average Mass 181, 907.61 daltons

[cysteines reduced, methionines have not been oxidised]

**Theoretical pI:-** 5.57

**Purity:-** 85 %

**Activation protocol:-** Constitutively active

**Enzyme storage buffer:-**

50 mM Tris-HCl pH 7.5, 270 mM Sucrose, 150 mM NaCl, 0.1 mM EGTA,  
0.1 % 2-mercaptoethanol, 0.02 % Brij-35, 1 mM benzamidine, 0.2 mM PMSF

**Storage temperature:-** -80 °C

**Assay:-** Standard filter binding assay

**Assay buffer:-**

50 mM Tris-HCl pH 7.5, 0.1 mM EGTA, 0.1 % 2-mercaptoethanol, 10 mM MgAc

**Substrate:-**

Myelin Basic Protein Final concentration: 0.3 mg/ml

# ***MRC PPU REAGENTS***

## **Clone Data Sheet**

### **ASK1 [2 – 1374]**

<b><u>Protein</u></b>	ASK1 [2 – 1374]
<b><u>Clone number</u></b>	DU 4707
<b><u>Species</u></b>	Human
<b><u>Accession number</u></b>	NM_005923
<b><u>Tags</u></b>	N-terminal GST
<b><u>Native sequence</u></b>	Amino acids S2 – T1374 (end) of human ASK1. Residue S238 of the fusion protein is equivalent to S2 of the native enzyme. The GST tag is located at residues 1 – 220.
<b><u>Protease cleavage</u></b>	PreScission ( <u>LEVLFQGPL</u> ) residues 221 - 229
<b><u>Cloning sites</u></b>	<i>Salt</i> 1 site of pGEX 6P-3

## *MRC PPU REAGENTS*

### Bacterially expressed protein

MSPILGYWKIKGLVQPTRLLLEYLEEKYEEHLYERDEGDKWRNKKFELGLEFPNLPYYIDG  
DVKLTQSMAIIRYIADKHNMLGGCPKERAIEISMLEGAVLDIRYGVSRIAYSKDFETLKVDF  
LSKLPPEMLKMFEDRLCHKTYLNGDHVTHPDFMLYDALDVVLYMDPMCLDAFPKLVCFKKRI  
EAIPQIDKYLKSSKYIAWPLQGWQATFGGGDHPPKSDLEVLFOGPLGSPNSRVD**STEADEG**  
**ITFSVPPFAPSGFCTIPEGGICRRGAAAVGEGEEHQLP PPPPGSFWNVESAAAPGIGCPA**  
**ATSSSSATRGRGSSVGGGSRRTTVAYVINEASQGQLVVAESEALQSLREACETVGATLETL**  
**HFGKLDGETTVLDRFYNADIAVVEMSDAFRQPSLFYHLGVRESFSMANNIILYCDTNSDS**  
**LQSLKEIICQKNTMCTGNYTFVPYMITPHNKVYCCDSSFMKGLTELMQPNFELLGPICLP**  
**LVDRFIQLLKVAQASSSQYFRESILNDIRKARNLYTGKELAAELARIRQRVDNIEVLTADI**  
**VINLLSYRDIQDYDSIVKLVETLEKLPTFDLASHHHVKFHYAFALNRRNLPGDRAKALDI**  
**MIPMVQSEGQVADMYCLVGRIYKDMFLDSNFTDTESRDHGASWFKKAFESEPTLQSGINY**  
**AVLLLAAGHQFESSFELRKVGKLSLLGKKGNLEKLQSYWEVGFFLGASVLANDHMRVIQ**  
**ASEKLFKLTTPAWYLKSI VETILYKHFVKLTTEQPVAKQELVDFWMDFLVEATKTDVTVV**  
**RFPVLILEPTKIYQPSYLSINNEVEEKTISIWHVLPDDKKGHEWNFSASSVRGVSISKFE**  
**ERCCFLYVLHNSDDFQIYFCTELHCKKFFEMVNTITEEKGRSTEEDCESDLLEYDYEYDE**  
**NGDRVVLGKGTYGIVYAGRDLSNQVRIAIKEIPERDSRYSQPLHEEIALHKHLKHKNIQY**  
**LGSFSENGFIKIFMEQVPGGSLSALLRSKWGPLKDNEQTIGFYTKQILEGLKYLHDNQIVH**  
**RDIKGDNVLINTYSGVLKISDFGTSKRLAGINPCTETFTGTLOYPEI IDKGPRGYGKAA**  
**DIWSLGTI IEMATGKPPFYELGEPQAAMFKVGMFKVHPEIPESMSAEAKAFILKCFEPDP**  
**DKRACANDLLVDEFLKVSSKKKKTKPKLSALSAGSNEYLRSISLPVPVLVEDTSSSEYGS**  
**VSPDTELKVDPPFSFKTRAKSCGERDVKGIRTLFLGIPDENFEDHSAPPSPEEKDSGFFMLR**  
**KDSERRATLHRILTEDQDKIVRNLMESLAHGAEPPKLKWEHITTLIASLREFVRSTDRKII**  
**ATTL SKLLELDFDSHGISOVQVVLFGFQDAVNKVLNRHNKPHWMFALDSIRKAVQTAI**  
**TILVPELRPHFSLASESDTADQEDLDVEDDHEEQPSNQTVRRPQAVIEDAVATSGVSTLSS**  
**TVSHDSQSAHRSLNVQLGRMKIETNRLLEELVRKEKELQALLHRAIEEKDQEKHLKLSQ**  
**PIEIPELPVFHLNSSGTNTEDSELTDWLRVNGAEDTISRFLAEDYTLLDVLYYVTRDDLK**  
**CLRLRGGMLCTLWKAIIDFRNKQT**

## *MRC PPU REAGENTS*

### Complete Nucleotide Sequence

ATGTCCCCTATACTAGGTTATTGGAAAATTAAGGGCCTTGTGCAACCCAC  
TCGACTTCTTTTGGAAATATCTTGAAGAAAAATATGAAGAGCATTTTGTATG  
AGCGCGATGAAGGTGATAAATGGCGAAACAAAAAGTTTGAATTGGGTTTG  
GAGTTTCCCAATCTTCCTTATTATATTGATGGTGATGTTAAATTAACACA  
GTCTATGGCCATCATACTGTTATATAGCTGACAAGCACAAACATGTTGGGTG  
GTTGTCCAAAAGAGCGTGCAGAGATTTCAATGCTTGAAGGAGCGGTTTTG  
GATATTAGATACGGTGTTCGAGAATTGCATATAGTAAAGACTTTGAAAC  
TCTCAAAGTTGATTTTCTTAGCAAGCTACCTGAAATGCTGAAAATGTTTCG  
AAGATCGTTTATGTCATAAAACATATTTAAATGGTGATCATGTAACCCAT  
CCTGACTTCATGTTGTATGACGCTCTTGATGTTGTTTTATAACATGGACCC  
AATGTGCCTGGATGCGTTCCCAAAATTAGTTTGTTTTTAAAAACGTATTG  
AAGCTATCCCACAAATTGATAAGTACTTGAAATCCAGCAAGTATATAGCA  
TGGCCTTTGCAGGGCTGGCAAGCCACGTTTGGTGGTGGCGACCATCCTCC  
AAAATCGGATCTGGAAGTTCTGTTCCAGGGGCCCTGGGATCCCCGAATT  
CCCGGGTGCACAGCACGGAGGCGGACGAGGGCATCACTTTCTCTGTGCCA  
CCCTTCGCCCCCTCGGGCTTCTGCACCATCCCCGAGGGCGGCATCTGCAG  
GAGGGGAGGAGCGGCGGGTGGCGAGGGCGAGGAGCACCAGCTGCCAC  
CGCCGCCCGGGCAGTTTCTGGAACGTGGAGAGCGCCGCTGCCCTGGC  
ATCGGTTGTCCGGCGGCCACCTCCTCGAGCAGTGCCACCCGAGGCCGGGG  
CAGCTCTGTTGGCGGGGGCAGCCGACGGACCACGGTGGCATATGTGATCA  
ACGAAGCGAGCCAAGGGCAACTGGTGGTGGCCGAGAGCGAGGCCCTGCAG  
AGCTTGCGGGAGGCGTGCGAGACAGTGGGCGCCACCCGGAAACCCTGCA  
TTTTGGGAAACTCGACTTTGGAGAAACCACCGTGCTGGACCGCTTTTACA  
ATGCAGATATTGCGGTGGTGGAGATGAGCGATGCCTTCCGGCAGCCGTCC  
TTGTTTTACCACCTTGGGGTGAGAGAAAGTTTCAGCATGGCCAACAACAT  
CATCCTCTACTGTGATACTAACTCGGACTCTCTGCAGTCACTGAAGGAAA  
TCATTTGCCAGAAGAATACTATGTGCACTGGGAACTACACCTTTGTTCTT  
TACATGATAACTCCACATAACAAAGTCTACTGCTGTGACAGCAGCTTCAT  
GAAGGGGTTGACAGAGCTCATGCAACCGAACTTCGAGCTGCTTCTTGGAC  
CCATCTGCTTACCTCTTGTGGATCGTTTTATTCAACTTTTGAAGGTGGCA  
CAAGCAAGTTCTAGCCAGTACTTCCGGGAATCTATACTCAATGACATCAG  
GAAAGCTCGTAATTTATACACTGGTAAAGAATTGGCAGCTGAGTTGGCAA  
GAATTCGGCAGCGAGTAGATAAATATCGAAGTCTTGACAGCAGATATTGTC  
ATAAATCTGTTACTTTCTACAGAGATATCCAGGACTATGATTCTATTGT  
GAAGCTGGTAGAGACTTTAGAAAACTGCCAACCTTTGATTTGGCCTCCC  
ATCACCATGTGAAGTTTCAATTATGCATTTGCACTGAATAGGAGAAATCTC  
CCTGGTGACAGAGCAAAGCTCTTGATATTATGATTCCCATGGTGCAAAG  
CGAAGGACAAGTTGCTTCAGATATGTATTGCC TAGTTGGTCGAATCTACA  
AAGATATGTTTTTGGACTCTAATTTACGGGACTGAAAGCAGAGACCAT  
GGAGCTTCTTGGTTCAAAAAGGCATTTGAATCTGAGCCAACACTACAGTC  
AGGAATTAATTATGCGGTCCCTCCTCCTGGCAGCTGGACACCAGTTTGAAT  
CTTCCTTTGAGCTCCGAAAGTTGGGGTGAAGCTAAGTAGTCTTCTTGGT  
AAAAAGGGAAACTTTGAAAAACTCCAGAGCTACTGGGAAGTTGGATTTTT  
TCTGGGGGCCAGCGTCC TAGCCAATGACCACATGAGAGTCATTCAAGCAT  
CTGAAAAGCTTTTTAAACTGAAGACACCAGCATGGTACCTCAAGTCTATT  
GTAGAGACAATTTAATATATAAGCATTTTGTGAAACTGACCACAGAACA

## *MRC PPU REAGENTS*

GCCTGTGGCCAAGCAAGAACTTGTGGACTTTTGGATGGATTTCCCTGGTCCG  
AGGCCACAAAGACAGATGTTACTGTGGTTAGGTTTCCAGTATTAATATTA  
GAACCAACCAAAATCTATCAACCTTCTTATTTGTCTATCAACAATGAAGT  
TGAGGAAAAGACAATCTCTATTTGGCACGTGCTTCCCTGATGACAAGAAAG  
GTATACATGAGTGGAAATTTTAGTGCCTCTTCTGTGAGGGGAGTGAGTATT  
TCTAAATTTGAAGAAAGATGCTGCTTTCTTTATGTGCTTCACAATTCTGA  
TGATTTCCAAATCTATTTCTGTACAGAACCTCATTGTAAAAAGTTTTTTG  
AGATGGTGAACACCATTACCGAAGAGAAGGGGAGAAGCACAGAGGAAGGA  
GACTGTGAAAGTGACTTGCTGGAGTATGACTATGAATATGATGAAAATGG  
TGACAGAGTCGTTTTAGGAAAAGGCACCTTATGGGATAGTCTACGCAGGTC  
GGACTTGAGCAACCAAGTCAGAATTGCTATTAAGGAAATCCAGAGAGA  
GACAGCAGATACTCTCAGCCCCGTCATGAAGAAATAGCATTGCATAAACA  
CCTGAAGCACAAAAATATTGTCCAGTATCTGGGCTCTTTCAGTGAGAATG  
GTTTCATTA AAAATCTTCATGGAGCAGGTC CCTGGAGGAAGTCTTTCTGCT  
CTCCTTCGTTCCAAATGGGGTCCATTA AAAGACAATGAGCAAACAATTGG  
CTTTTATACAAAGCAAATACTGGAAGGATTAAAATATCTCCATGACAATC  
AGATAGTTCACCGGGACATAAAGGGTGACAATGTGTTGATTAATACCTAC  
AGTGGTGTTCTCAAGATCTCTGACTTCGGAACATCAAAGAGGCTTGCTGG  
CATAAACCCCTGTACTGAAACTTTTACTGGTACCCTCCAGTATATGGCAC  
CAGAAATAATAGATAAAGGACCAAGAGGCTACGGAAAAGCAGCAGACATC  
TGGTCTCTGGGCTGTACAATCATTGAAATGGCCACAGGAAAACCCCATTT  
TTATGAACTGGGAGAACCACAAGCAGCTATGTTCAAGGTGGGAATGTTTA  
AAGTCCACCCTGAGATCCAGAGTCCATGTCTGCAGAGGCCAAGGCATTC  
ATACTGAAATGTTTTGAACCAGATCCTGACAAGAGAGCCTGTGCTAACGA  
CTTGCTTGTTGATGAGTTTTTTAAAAGTTTTCAAGCAAAAAGAAAAGACAC  
AACCTAAGCTTTCAGCTCTTTCAGCTGGATCAAATGAATATCTCAGGAGT  
ATATCCTTGCCGGTACCTGTGCTGGTGGAGGACACCAGCAGCAGCAGTGA  
GTACGGCTCAGTTTCACCCGACACGGAGTTGAAAGTGGACCCCTTCTCTT  
TCAAAACAAGAGCCAAGTCTCGGGAGAAAGAGATGTCAAGGGAATTCGG  
ACACTCTTTTTGGGCATTCAGATGAGAATTTTGAAGATCACAGTGCTCC  
TCCTTCCCCTGAAGAAAAGATTCTGGATTCTTCATGCTGAGGAAGGACA  
GTGAGAGGCGAGCTACCCTTCACAGGATCCTGACGGAAGACCAAGACAAA  
ATTGTGAGAAACCTAATGGAATCTTTAGCTCACGGGGCTGAAGAACCGAA  
ACTAAAATGGGAACACATCACACCCTCATTGCAAGCCTCAGAGAATTTG  
TGAGATCCACTGACC GAAAAATCATAGCCACCACACTGTCAAAGCTGAAA  
CTGGAGCTGGACTTCGACAGCCATGGCATTAGCCAAGTCCAGGTGGTACT  
CTTTGGTTTTTCAAGATGCTGTCAATAAAGTTCTTTCGGAATCATAACATCA  
AGCCGCACTGGATGTTTGCCTTAGACAGTATCATTCGGAAGGCGGTACAG  
ACAGCCATTACCATCCTGGTTCCAGAACTAAGGCCACATTTTCAGCCTTGC  
ATCTGAGAGTGATACTGCTGATCAAGAAGACTTGGATGTAGAAGATGACC  
ATGAGGAACAGCCTTCAAATCAAAGTGTCCGAAGACCTCAGGCTGTCATT  
GAAGATGCTGTGGCTACCTCAGGCGTGAGCACGCTCAGTTCTACTGTGTC  
TCATGATTTCCAGAGTGCTCACCGGTCCTGAATGTACAGCTTGGAAGGA  
TGAAAATAGAAACCAATAGATTACTGGAAGAATTGGTTCGGAAGAGAAA  
GAATTACAAGCACTCCTTCATCGAGCTATTGAAGAAAAGACCAAGAAAT  
TAAACACCTGAAGCTTAAGTCCCAACCCATAGAAATTCCTGAATTGCCTG  
TATTTTCATCTAAATTTCTTCTGGCACAAATACTGAAGATTCTGAACTTACC

***MRC PPU REAGENTS***

GACTGGCTGAGAGTGAATGGAGCTGATGAAGACACTATAAGCCGGTTTTT  
GGCTGAAGATTATACACTATTGGATGTTCTCTACTATGTTACACGTGATG  
ACTTAAAATGCTTGAGACTAAGGGGAGGGATGCTGTGCACACTGTGGAAG  
GCTATCATTGACTTTCGAAACAAACAGACTtgagtcgac