

MRC PPU Reagents and Services

Standard Operating Procedure

Preparation of active PKN1 [460 - 942]

<u>Enzyme description:-</u>	PKN1 [460 – 942]
<u>Clone number:-</u>	DU 68070
<u>Source:-</u>	Recombinant
<u>Expression system:-</u>	Baculovirus expression vector system
<u>Tag:-</u>	N-terminal GST
<u>Purification method:-</u>	GSH Sepharose

Calculated molecular mass:-

Monoisotopic	81, 376.31 daltons
Average Mass	81, 428.17 daltons

[cysteines reduced, methionines have not been oxidised]

Theoretical pI:- 5.40

Purity:- >75 %

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, 0.2 mM PMSF, 1 mM Benzamidine.

Storage temperature:- -70 deg C

Assay buffer:-

50 mM Tris-HCl pH 7.5, 10 mM DTT, 10 mM magnesium acetate, 0.1 mM EGTA

Substrate:-

KKLNRTLVA Final concentration: 300 uM

MRC PPU Reagents and Services

Clone Data Sheet

PKN1 [460 – 942]

Protein PKN1 [460 – 942]

Clone number DU 68070

Species Human

Accession number Q16512-2

Tags N-terminal GST

**Baculovirus
expressed protein**

MSPILGYWKIKGLVQPTRLLEYLEEKYEEHLYERDEGDKWRNKKFEL
GLEFPNLPYYIDGDVKL TQSM IIRYIADKHNMLGGCPKERA EISM LE
GAVLDIRYGVSRIAYS KDFETLKVDFLSKLP EMLKMFEDRLCHKTYLN
GDHVTHPDFMLYDALDVVLYMDPMCLDAFPKLVCFKKRIEAI PQIDKY
LKSSKYIAWPLQGWQATFGGGDHPPKSDLEVL FQGPLGSPNSRVEMDN
ERHEVQLDMEPQGCLVAEVTFRNPVIERIPRLRRQKKIFSKQOGKAFO
RARQMNIDVATWVRLRLRRLIPNATGTGTFS PGASPGSEARTTGDISVE
KLNLGTSDSSPQKSSRDPPSSPSSLSSPIQESTAPELPSETQETPGP
ALCSPLRKSPLTLEDFKFLAVLGRGHFGKVLLSEFRPSGELFAIKALK
KGDIVARDEVESLMCEKRILAAVTSAGHPFLVNLFGCFQTP EHVCFVM
EYSAGGDLMLHIHSDVFSEPRAI FYSACVVLGQLFHEHKIVYRDLKL
DNLLLDTEGYVKIADFGLCKEGMGYGDR TSTFCGTPEFLAPEVLTDT S
YTRAVDWGLGVLLYEMLVGESPFPGDDEEEVFDSIVNDEVRYPRFLS
AEAIGIMRLLRRNPERRLGSSERDAEDVKKQPF FRTLGW EALLARRL
PPPFVPTLSGRTDVSNFDEEFTGEAP T LSPPRDARPLTAAEQAAFLDF
D

Native sequence Amino acids D460 – D942 (end residue C948) of human PKN1.
Residue D239 of the fusion protein is equivalent to M1 of the native
enzyme. The GST tag is located at residues 1 – 220.

Protease cleavage PreScission (LEVLFQGP) residues 221 - 228

Cloning sites *Xho*1 - *Not*1 into *Sal*1 - *Not*1 sites of pFastBac GST

MRC PPU Reagents and Services

Nucleotide
sequence of
insert

gacaatgagagggcatgAGGTGCAGCTGGACATGGAACCCCAGGGCTGC
CTGGTGGCTGAGGTCACCTTCCGCAACCCTGTCATTGAGAGGATTCCT
CGGCTCCGACGGCAGAAGAAAATTTTCTCCAAGCAGCAAGGGAAGGCG
TTCCAGCGTGCTAGGCAGATGAACATCGATGTCGCCACGTGGGTGCGG
CTGCTCCGGAGGCTCATCCCCAATGCCACGGGCACAGGCACCTTTAGC
CCTGGGGCTTCTCCAGGATCCGAGGCCCGGACCACGGGTGACATATCG
GTGGAGAAGCTGAACCTCGGCACTGACTCGGACAGCTCACCTCAGAAG
AGCTCGCGGGATCCTCCTTCCAGCCATCGAGCCTGAGCTCCCCATC
CAGGAATCCACTGCTCCCAGCTGCCTTCGGAGACCCAGGAGACCCCA
GGCCCCGCCCTGTGCAGCCCTCTGAGGAAGTCACCTCTGACCCTCGAA
GATTTCAAGTTCTTGGCGGTGCTGGGCCGGGGTCATTTTGGGAAGGTG
CTCCTCTCCGAATTCCGGCCCCAGTGGGGAGCTGTTCCGCATCAAGGCT
CTGAAGAAAGGGGACATTGTGGCCCCGAGACGAGGTGGAGAGCCTGATG
TGTGAGAAGCGGATATTGGCGGCAGTGACCAGTGCGGGACACCCCTTC
CTGGTGAACCTCTTCGGCTGTTTCCAGACACCGGAGCACGTGTGCTTC
GTGATGGAGTACTCGGCCGGTGGGGACCTGATGCTGCACATCCACAGC
GACGTGTTCTCTGAGCCCCGTGCCATCTTTTATTCCGCCTGCGTGGTG
CTGGGCCTACAGTTTCTTCACGAACACAAGATCGTCTACAGGGACCTG
AAGTTGGACAATTTGCTCCTGGACACCGAGGGCTACGTCAAGATCGCA
GACTTTGGCCTCTGCAAGGAGGGGATGGGCTATGGGGACCGGACCAGC
ACATTCTGTGGGACCCCCGAGTTCTTGGCCCCCTGAGGTGCTGACGGAC
ACGTCGTACACGCGAGCTGTGGACTGGTGGGGACTGGGTGTGCTGCTC
TACGAGATGCTGGTTGGCGAGTCCCATTCACAGGGGATGATGAGGAG
GAGGTCTTCGACAGCATCGTCAACGACGAGGTTTCGCTACCCCCGCTTC
CTGTCCGCCGAAGCCATCGGCATCATGAGAAGGCTGCTTCGGAGGAAC
CCAGAGCGGAGGCTGGGATCTAGCGAGAGAGATGCAGAAGATGTGAAG
AAACAGCCCTTCTTTCAGGACTCTGGGCTGGGAAGCCCTGTTGGCCCCG
CGCCTGCCACCGCCCTTTGTGCCACGCTGTCCGGCCGCACCGACGTC
AGCAACTTCGACGAGGAGTTACCGGGGAGCCCCCACACTGAGCCCCG
CCCCGCGACGCGCGGCCCTTACAGCCGCGGAGCAGGCAGCCTTCCTG
GACTTCGACTgagcggccgc