

## ***MRCPPU REAGENTS and SERVICES***

### **Standard Operating Procedure**

#### **Preparation of active MYCBP2 [4416 – 4678]**

**Enzyme description:-** MYCBP2 [4416 - 4678]

**Clone number:-** DU 28681

**Source:-** Recombinant

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

**Tag:-** N-terminal GST

**Purification method:-** GSH Agarose

**Calculated molecular mass:-**

Monoisotopic 56, 460.65 daltons

Average Mass 56, 498.26 daltons

[cysteines reduced, methionines have not been oxidised

**Theoretical pI:-** 6.20

**Purity:-** >80 %

**Activation protocol:-** Constitutively active

**Enzyme storage buffer:-**

50 mM HEPES pH 7.5, 10% glycerol, 150mM NaCl, 0.5 mM TCEP

**Storage temperature:-** -70 °C

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### **Clone Data Sheet**

#### **MYCBP2 [4416 - 4678]**

<b><u>Protein</u></b>	MYCBP2 [4416 - 4678]
<b><u>Clone number</u></b>	DU 28681
<b><u>Species</u></b>	Human
<b><u>Accession number</u></b>	O75592-1
<b><u>Tags</u></b>	N-terminal GST
<b><u>Bacterially expressed protein</u></b>	MSPILGYWKIKGLVQPTRLLLEYLEEKYEEHLYERDEGDKWRNKKFELG LEFPNLPIYYIDGDVKLTQSMAIIRYIADKHNLGGCPKERAESMLEGA VLDIIRGVSRAYSKDFETLKVDLFLSKLPEMLKMFEDRLCHKTYLNGDH VTHPDFMLYDALDVLYMDPMCLDAFPKLVCFKKRIEAIPQIDKYLKSS KYIAWPLQGWQATFGGGDHPPKSD <u>LEVLFQGP</u> LS <u>SATSLKQDADDMCM</u> <u>ICFTEALSAAPAIQLDCSHIFHLQCCRVL</u> ENRWLGRITFGFISCPIC KNKINHIVLKDLLDPIKELEYEDVVRKALMRLEYEGLHKSEAITTPGVRF YNDPAGYAMNRYAYYVCYKCRKAYFGGEARCDAEARGDDYDPRELICG ACSDVSRAQMCPKHGTDLEYKCRYCCSVAVFFCFGTTFCNACHDDFQ RMTSIPKEELPHCPAGPKGQLEGTECPLHVVHPPTGEEFALGCGVCRN AHTF
<b><u>Native sequence</u></b>	Amino acids S4416 – F4678 (end residue) of human MYCBP2. Residue S232 of the fusion protein is equivalent to S4416 of the native enzyme. The GST tag is located at residues 1 – 220.
<b><u>Protease cleavage</u></b>	PreScission site ( <u>LEVLFQGP</u> ) residues 221 – 228

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