

## *MRC PPU Reagents and Services*

### **Standard Operating Procedure**

#### **Preparation of Influenza B Virus M1 [1 – 248]**

**Enzyme description:-** IBV M1 [1 - 248]

**Clone number:-** DU 75470

**Source:-** Recombinant

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

**Tag:-** N-terminal GST

**Purification method:-** GSH Sepharose

**Calculated molecular mass:-**

Monoisotopic 54, 203.56 daltons

Average Mass 54, 239.01 daltons

[cysteines reduced, methionines have not been oxidised]

**Theoretical pI:-** 8.13

**Purity:-** 80 %

**Enzyme storage buffer:-**

50 mM Tris-HCl pH 7.5, 270 mM Sucrose, 150 mM NaCl, 0.1 mM EGTA, 0.5 mM TCEP

**Storage temperature:-** -70 °C

*MRC PPU Reagents and Services*

**Clone Data Sheet**

**Influenza B Virus M1 [1 – 248]**

<b><u>Protein</u></b>	IBV M1 [1 - 248]
<b><u>Clone number</u></b>	DU 75470
<b><u>Species</u></b>	Influenza B virus (IBV) strain B/Florida/04/2006
<b><u>Tags</u></b>	N-terminal GST
<b><u>Bacterially expressed protein</u></b>	<p>MSPILGYWKIKGLVQPTRLLLEYLEEKYEEHLYERDEGDKWRNKKFELG LEFPNLPYYIDGDVKLTQSMAIIRYIADKHNMLGGCPKERAEISMLEGA VLDIRYGVSR IAYS KDFETLKVDFLSKLP EMLKMFEDRLCHKTYLNGDH VTHPDFMLYDALDVVLYMDPMCLDAFPKLVCFKKRIEAI PQIDKYLKSS KYIAWPLQGWQATFGGGDHPPKSDLEVL FQGPLGS <b>MSLFGDTIAYLLSL</b> <b>TEDGEGKAELA EKLHCWFGGKEFDLDSALEWIKNKRC LTDIQKALIGAS</b> <b>ICFLKPKDQERKRRFITEPLSGMGT TATKKKGLILAERKMRRCVSFHEA</b> <b>FEIAEGHES SALLYCLMVMYLNPGNYSMQVKLGTLCALCEKQASHSHRA</b> <b>HSRAARSSVPGVRREMQMVSAMNTAKTMNGMGKGEDVQKLA EELQSNIG</b> <b>VLRSLGASQKN GEGIAKDVMEVLKQSSMGNSALVKKYL</b></p>
<b><u>Native sequence</u></b>	<p>Amino acids M1 – L248 (end residue) of IBV M1 protein. Residue M232 of the fusion protein is equivalent to M1 of the native enzyme. The GST tag is located at residues 1 – 220.</p>
<b><u>Protease cleavage</u></b>	PreScission ( <u>LEVLFQGP</u> ) residues 221 - 229