

MRC PPU Reagents and Services

Standard Operating Procedure

Preparation of Influenza D Virus NP [1 – 552]

Enzyme description:- IDV NP [1 - 552]

Clone number:- DU 75213

Source:- Recombinant

Expression system:- *E.coli*

Tag:- N-terminal GST

Purification method:- GSH Agarose

Calculated molecular mass:-

Monoisotopic 88, 059.84 daltons

Average Mass 88, 116.43 daltons

[cysteines reduced, methionines have not been oxidised]

Theoretical pI:- 8.87

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

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Clone Data Sheet

Influenza D Virus NP [1 – 552]

<u>Protein</u>	IDV NP [1 - 552]
<u>Clone number</u>	DU 75213
<u>Species</u>	Influenza D virus (IDV) strain D/bovine/Oklahoma/660/2013
<u>Tags</u>	N-terminal GST
<u>Bacterially expressed protein</u>	<p>MSPILGYWKIKGLVQPTRLLLEYLEEKYEEHLYERDEGDKWRNKKFELG LEFPNLPYYIDGDVVKLTQSMAIIRYIADKHNMLGGCPKERAEISMLEGA VLDIRYGVSR IAYS KDFETLKVDFLSKLP EMLKMFEDRLCHKTYLNGDH VTHPDFMLYDALDVVLYMDPMCLDAFPKLVCFKKRIEAI PQIDKYLKSS KYIAWPLQGWQATFGGGDHPPKSDLEVL FQGPLGSMDS TKAQTPEEQRA KNAKTILENIQIYERMCDLFGVSEDDKLI IENSISIERMIRVVTDKKYQ DKKLKNAGSDLEKIANAGKVF CRLVESTAGKCSARLGMALKPNVEAVLT DVLGNELDRAAVLGKRMGFTAMFKSNLEEVLYQRGKNQLKKRNAAETFT LSQGASLEARFRPIMEKHLGVGT VVASIKNILASKKNGNYRNKMVRKPG GNRESWSPLEREISFLNKKLFPGPMRQLCKKFEYLNDQEKQLALNLMLD ASLILKPQVTHKMIMPWSMWLAVK KYAEMNKGPSLEDLAAYSGVRAFM AFNTACYMSKFTIGKGI VDAEIMENGNDKMQTLAMACFGLAYEDTGIV AAMISQPMKKRYQLRVGNFN PPEEGTIKGT SAGYFHKWAEFGNRLPFNS FGTGESKQISNSGVFAVQRPSTTNIQRLAELMARNTGETSDNFTQLVQK IREQVGT FADQKANLREFTGGYIYDITDVT KSNPKIPQLGGDSFFFEFT GSDVPRTGAKRRVGGADDVTPGTSQPKKRGRQGAGAESSMDIETVGED</p>
<u>Native sequence</u>	Amino acids M1 – D552 (end residue) of IDV NP protein. Residue M232 of the fusion protein is equivalent to M1 of the native enzyme. The GST tag is located at residues 1 – 220.
<u>Protease cleavage</u>	PreScission (<u>LEVLFQGP</u>) residues 221 - 229