

# *MRC PPU Reagents and Services*

## **Standard Operating Procedure**

### **Preparation of Nucleocapsid Protein [1 - 419] SARS CoV2**

**Enzyme description:-** N Protein [1 - 419]

**Clone number:-** DU 67728

**Source:-** Recombinant

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

**Tag:-** N-terminal MBP

**Purification method:-** Amylose Resin

**Calculated molecular mass:-**

Monoisotopic 89,809.39 daltons

Average Mass 89,864.61 daltons

[cysteines reduced, methionines have not been oxidised]

**Theoretical pI:-** 8.97

**Purity:-** 80 %

**Enzyme storage buffer:-**

50 mM Tris-HCl pH 7.5, 270 mM Sucrose, 150 mM NaCl, 0.1 mM EGTA,  
0.1 % 2-mercaptoethanol, 0.03 % Brij-35

**Storage temperature:-** -70 °C

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## Clone Data Sheet

### Nucleocapsid Protein [1 – 419] SARS CoV2

**Protein** N Protein [1 - 419]

**Clone number** DU 67728

**Accession number** QHD43423.2

**Tags** N-terminal MBP

**Bacterially  
expressed protein**

MKIEEGKLVIIWINGDKGYNGLAEVGGKKFEKDTGIKVTVEHPDKLE  
EKFPQVAATGDGPDIIFWAHDRFGGYAQSGLLAEITPDKAFQDKL  
YPFTWDAVRYNGKLIAYPIAVEALSLIYNKDLLPNPPKTWEEIPA  
LDKELKAKGKSALMFNLQEPYFTWPLIAADGGYAFKYENGYD IK  
DVGVDNAGAKAGLTFVLVDLIKKNHMNADTDYSIAEAAFNKGETAM  
TINGPWAWSNIDTSKVNYGVTVLPTFKGQPSKPFVGVLSAGINAA  
SPNKELAKEFLENYLLTDEGLEAVNKDKPLGAVALKSYYYEELVKD  
PRIAATMENAQKGEIMPNI PQMSAFWYAVRTAVINAASGRQTVDE  
ALKDAQTNSSNNNNNNNNNNLGD DDDDKVPEFLEVL FQG PLGSMS  
**DNGPQNQRNAPRITFGGPSDSTGSNQNGERSGARSKQRRPQGLPN**  
**NTASWFTALTQHGKEDLKFPRGQGVPIINTNSSPDDQIGYYRRATR**  
**RIRGGDGKMKDLSRWYFYLLGTGPEAGLPYGANKDGI IWVATEG**  
**ALNTPKDHIGTRNPANNAIIVLQLPQGTTL PKGFYAEGSRGGSQA**  
**SSRSSRSRNSRNSTPGSSRGTSPARMAGNGDAALALLLLDRL**  
**NQLESKMSGKGQQQGGQTVTKKSAEASKKPRQKRTATKAYNVTQ**  
**AFGRRGPEQTQGNFGDQELIROGTDYKHWQIAQFAPSASAFFGM**  
**SRIGMEVTPSGTWLTYTGAIKLDDKDPNFKDQVILLNKHIDAYKT**  
**FPPTPEPKDKKKKADETQALPQRQKKQQTVTLLPAADLDDFSKQL**  
**QQSMSSADSTQA**

**Native sequence** Amino acids M1 – A419 (end).  
Residue M404 of the fusion protein is equivalent to M1 of the native enzyme. The MBP tag is located at residues 1 – 367.

**Protease cleavage** Enterokinase (DDDDK) residues 384 – 388  
PreScission (LEVLFGQP) residues 393 – 400