

## Lake Victoria marburgvirus (MARV) (strain Musoke-80) VP30 Protein (His & Myc)

### General Information

Synonyms: VP30;Transcriptional activator VP30;Minor nucleoprotein VP30

Protein Construction: 1-281 aa

Species: MARV

Expression Host: E. coli

Accession: P35258

Molecular Weight: 39.1 kDa (predicted)

AA Sequence: MQQPRGRSRTNRNHQVPTIYHETQLPSKPHYTNYPHPRARMSSTRSSAESSPTNHIPRARPPSTFNLSKPPPP  
PKDMCRNMKIGLPCADPTCNRDHDLDNLNRELLLLMARKMLPNTDKTFRSPQDCGSPSLSKGLSKDKQEQ  
TKDVLTLLENLGHILSYLHRSEIGKLDETSLRAALSLTCAGIRKTNRSINTMTLHMNHENLPQDQNGVIKQTYT  
GIHLDKGGQFEAALWQGWDKRSISLFVQAALYVMNNIPCESSISVQASYDHFILPQSQGKGQ

### QC Testing

Biological Activity: Activity has not been tested. It is theoretically active, but we cannot guarantee it. If you require protein activity, we recommend choosing the eukaryotic expression version first.

Purity: > 85% as determined by SDS-PAGE.

Endotoxin: < 1.0 EU/μg of the protein as determined by the LAL method.

Formulation: If the delivery form is liquid, the default storage buffer is Tris/PBS-based buffer, 5%-50% glycerol. If the delivery form is lyophilized powder, the buffer before lyophilization is Tris/PBS-based buffer, 6% Trehalose, pH 8.0.

### Preparation and Storage

Reconstitution:

Reconstitute the lyophilized protein in sterile deionized water. The product concentration should not be less than 100 μg/mL. Before opening, centrifuge the tube to collect powder at the bottom. After adding the reconstitution buffer, avoid vortexing or pipetting for mixing.

Stability & Storage:

Lyophilized powders can be stably stored for over 12 months, while liquid products can be stored for 6-12 months at -80°C. For reconstituted protein solutions, the solution can be stored at -20°C to -80°C for at least 3 months. Please avoid multiple freeze-thaw cycles and store products in aliquots.

Shipping:

In general, Lyophilized powders are shipping with blue ice. Solutions are shipping with dry ice.

### Protein Background

Acts as a transcription anti-termination factor immediately after transcription initiation, but does not affect transcription elongation. This function has been found to be dependent on the formation of an RNA secondary

structure at the transcription start site of the first gene.

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Tel:781-999-4286    E\_mail:info@targetmol.com    Address:34 Washington Street,Wellesley Hills,MA 02481