

Genomic RNA from Enterovirus D68, US/KY/14-18953

Catalog No. NR-49137

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Contributor:

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Manufacturer:

BEI Resources

Product Description:

Genomic RNA was isolated from a preparation of cell lysate and supernatant from human rhabdomyosarcoma cells infected with enterovirus D68 (EV-D68), US/KY/14-18953.

EV-D68, US/KY/14-18953 was isolated from a nasopharyngeal swab taken from a human in Kentucky, USA, in August, 2014.¹ The complete genome of EV-D68, US/KY/14-18953 has been sequenced (GenBank: [KM851231](https://www.ncbi.nlm.nih.gov/nuclom/KM851231)).

NR-49137 has been qualified for PCR applications by amplification of an approximately 1100 nucleotide sequence. Recommended dilutions for successful RT-PCR amplification are indicated on the Certificate of Analysis for each lot.

Material Provided:

Each vial contains 100 µL of viral genomic RNA in TE buffer (10 mM Tris-HCl, 1 mM EDTA, pH 7.0). The viral genomic RNA is in a background of cellular nucleic acid and carrier RNA. The vial should be centrifuged prior to opening.

Packaging/Storage:

NR-49137 was packaged aseptically in screw-capped plastic cryovials. The product is provided frozen on dry ice and should be stored at -60°C or colder immediately upon arrival. Freeze-thaw cycles should be minimized.

Citation:

Acknowledgment for publications should read “The following reagent was obtained through BEI Resources, NIAID, NIH: Genomic RNA from Enterovirus D68, US/KY/14-18953, NR-49137.”

Biosafety Level: 1

Appropriate safety procedures should always be used with this material. Laboratory safety is discussed in the following publication: U.S. Department of Health and Human Services, Public Health Service, Centers for Disease Control and

Prevention, and National Institutes of Health. Biosafety in Microbiological and Biomedical Laboratories. 5th ed. Washington, DC: U.S. Government Printing Office, 2009; see www.cdc.gov/biosafety/publications/bmbl5/index.htm.

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References:

1. Brown, B. A., et al. “Seven Strains of Enterovirus D68 Detected in the United States during the 2014 Severe Respiratory Disease Outbreak.” Genome Announc. 2 (2014): e01201-14. PubMed: 25414503.

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