



Detection of Delta+ Variant and Inferred Identification of Delta Variant Using Advanta SARS-CoV-2 Mutation Assay Panel for Research and Community Surveillance

Increased Prevalence of Delta and Decline of Epsilon in Global Epidemiology

The spike protein mutation L452R, historically linked to the Epsilon variant of SARS-CoV-2, more recently is identified as a key mutation in the Delta and Delta+ variants. While the Epsilon variant has declined below 1% of cases in the United States, its origin and primary location of existence, the Delta variant has increased beyond 20% and is on the verge of becoming the dominant variant globally.¹ Based on these epidemiological statistics, it is clear that detection of the L452R mutation can be used to infer the presence of the Delta variant for surveillance purposes.

The Advanta™ SARS-CoV-2 Mutation Assay Panel (Table 1) can be used to detect the L452R mutation associated with the Delta variant and the K417N mutation associated with the Delta+ variant. Combined with prevailing epidemiology, this assay (when run with additional reagents not provided) can be a quick, cost-effective tool for community surveillance of the fast-growing Delta and Delta+ variants.

Assay Design for L452R

The detection of L452R is conducted in a two-step process (Figure 1): 1. One-step RT-preamplification with

forward and reverse primers that produces a 99 bp amplicon product, and 2. qPCR-based detection of L452 wild-type (CUG) or L452R mutant (CGG) that produces a 90 bp amplicon product. The qPCR assay uses either a mutant or wild-type-specific forward detection primer, a common detection probe with FAM-quencher, and a common reverse primer. The specificity of the assay relies on two target-specific primers and the probe.

Specific Detection of L452R

In order to demonstrate specific detection of L452R, Twist Synthetic SARS-CoV-2 RNA Controls, Wuhan-Hu-1 (wild type; Control 2; PN 102024), and India/CT-ILSGS00361/2021 Kappa variant (mutant to isolate for L452R; Control 18 (B.1.617.1); PN 104338) were tested using the Advanta SARS-CoV-2 Mutation Assay Panel with the 192.24 GE IFC (integrated fluidic circuit) with 20 replicates each. The test also included assays for N1, N2, and RNase P in quadruplicate for COVID-19. The heat map of the Ct value is shown in Figure 2. The N1, N2, and RNase P were positive for both wild-type and Kappa variant for all the assay replicates. The no template control (NTC), negative control (NC), and positive control were tested in duplicate, and controls met the expected results. The results from Biomark™ Pathogen Detection Software are compiled in Table 1 and aligned with expected detection outcome based on the current assays in the panel.

Mutation (where first reported)	B.1.1.7 (UK)	B.1.351 (S. Africa)	P.1 (Brazil)	B.1.427 B.1.429 (US-CA)	B.1.526 (US-NY)	B.1.617.2 (India)	B.1.617.2.1 (India)
WHO identification	Alpha	Beta	Gamma	Epsilon	Iota	Delta	Delta+
K417N		✓				Not detected	✓
K417T			✓			Not detected	Not detected
L452R				✓		✓	✓
E484K		✓	✓		✓	Not detected	Not detected
N501Y	✓	✓	✓			Not detected	Not detected
Δ69/70	✓					Not detected	Not detected

Table 1. Mutations detected by the Advanta SARS-CoV-2 Mutation Assay Panel

