

## Assay Profile

## LNA-based qPCR assays for the genotyping of SARS-CoV-2 variants of concern (VOCs)

A robust design pipeline with several parameters for optimal primer-probe combinations was wet-lab tested for the mutations. The specificity and sensitivity of the assays were validated using the QIAprep&amp<sup>®</sup> Viral RNA UM Kit. The user should order primers and probes from a partnering oligo provider. QIAGEN can provide recommendations (in this case, our trusted partner **biomers.net**), but it is up to the end-user to validate the assays in their workflow. Assay validation should be carried out to ensure that the laboratory workflow performs adequately for the circulating viruses.



	Bio-Rad CFX	ABI QuantStudio	QIAGEN Rotor-Gene Q
IVT copies/ reaction		Ct	
105	23.4	23.6	23.4
104	26.7	27.4	27.0
10 <sup>3</sup>	30.0	30.6	30.1
10 <sup>2</sup>	33.2	34.4	33.1
10 <sup>1</sup>	37.4	37.7	36.4

Specific and sensitive detection of the mutation T478K. In green: amplification curves from dilution series  $(10^5-10^1 \text{ cp/rxn}, \text{ in duplicates})$  of an in-vitro transcript (IVT) bearing the SARS-CoV-2 mutation T478K. In red: amplification curve of an IVT of the corresponding WT sequence from SARS-CoV-2 at  $10^7 \text{ cp/rxn}$ . In black (overlapping with the red line) is the NTC.



	Bio-Rad CFX	ABI QuantStudio	QIAGEN Rotor-Gene Q
IVT copies/ reaction		Ct	
105	25.1	25.6	25.1
104	28.3	29.3	28.5
10 <sup>3</sup>	31.6	33.4	32.0
10 <sup>2</sup>	35.2	36.7	35.6
101	38.5	41.3	39.1

Specific and sensitive detection of the mutation E484K. In green: amplification curves from dilution series ( $10^5 - 10^1 \text{ cp/rxn}$ , in duplicates) of an in-vitro transcript (IVT) bearing the SARS-CoV-2 mutation E484K. In red: amplification curve of an IVT of the corresponding WT sequence from SARS-CoV-2 at  $10^7 \text{ cp/rxn}$ . In black (overlapping with the red line) is the NTC.

## Sample to Insight



	Bio-Rad CFX	ABI QuantStudio	QIAGEN Rotor-Gene Q
IVT copies/ reaction		Ct	
105	24.3	24.4	24.2
104	27.6	27.8	27.6
10 <sup>3</sup>	31.0	31.1	30.9
10 <sup>2</sup>	34.3	34.8	34.1
10 <sup>1</sup>	38.0	39.3	38.1

Specific and sensitive detection of the mutation E484Q. In green: amplification curves from dilution series ( $10^5-10^1$  cp/rxn, in duplicates) of an in-vitro transcript (IVT) bearing the SARS-CoV-2 mutation E484Q. In red: amplification curve of an IVT of the corresponding WT sequence from SARS-CoV-2 at  $10^7$  cp/rxn. In black (overlapping with the red line) is the NTC.



	Bio-Rad CFX	ABI QuantStudio	QIAGEN Rotor-Gene Q
IVT copies/ reaction		Ct	
105	24.5	24.4	24.12
104	27.8	27.8	27.25
10 <sup>3</sup>	31.2	31.1	31.01
10 <sup>2</sup>	34.7	34.4	33.99
10 <sup>1</sup>	37.6	37.6	38.21

Specific and sensitive detection of the mutation N501Y. In green: amplification curves from dilution series ( $10^5-10^1$  cp/rxn, in duplicates) of an in-vitro transcript (IVT) bearing the SARS-CoV-2 mutation N501Y. In red: amplification curve of an IVT of the corresponding WT sequence from SARS-CoV-2 at  $10^7$  cp/rxn. In black (overlapping with the red line) is the NTC.



	Bio-Rad CFX	ABI QuantStudio	QIAGEN Rotor-Gene Q
IVT copies/ reaction		Ct	
105	22.9	22.6	22.72
104	26.2	25.8	25.28
10 <sup>3</sup>	29.5	29.1	28.98
10 <sup>2</sup>	33.0	32.4	32.65
10 <sup>1</sup>	36.6	35.7	35.83

Specific and sensitive detection of the mutation L452R. In green: amplification curves from dilution series ( $10^5-10^1$  cp/rxn, in duplicates) of an in-vitro transcript (IVT) bearing the SARS-CoV-2 mutation L452R. In red: amplification curve of an IVT of the corresponding WT sequence from SARS-CoV-2 at  $10^7$  cp/rxn. In black (overlapping with the red line) is the NTC.



	ABI Bio-Rad CFX QuantStudio		QIAGEN Rotor-Gene Q
IVT copies/ reaction		Ct	
105	22.4	22.30	23.0
104	25.7	25.63	25.9
10 <sup>3</sup>	29.0	29.02	29.3
10 <sup>2</sup>	32.4	32.31	33.2
10 <sup>1</sup>	35.4	35.50	36.1

**Specific and sensitive detection of the mutation P681R.** In green: amplification curves from dilution series  $(10^5-10^1 \text{ cp/rxn}, \text{ in duplicates})$  of an in-vitro transcript (IVT) bearing the SARS-CoV-2 mutation P681R. In red: amplification curve of an IVT of the corresponding WT sequence from SARS-CoV-2 at  $10^7 \text{ cp/rxn}$ . In black (overlapping with the red line) is the NTC.



	Bio-Rad CFX	ABI QuantStudio	QIAGEN Rotor-Gene Q
IVT copies/ reaction		Ct	
105	21.47	21.67	21.57
104	24.80	24.65	24.82
10 <sup>3</sup>	27.83	28.26	27.80
10 <sup>2</sup>	31.62	31.52	31.49
10 <sup>1</sup>	34.24	35.64	35.04

Specific and sensitive detection of the deletion mutation HV 69/70. In green: amplification curves from dilution series  $(10^5-10^1 \text{ cp/rxn}, \text{ in duplicates})$  of an in-vitro transcript (IVT) bearing the SARS-CoV-2 deletion mutation HV 69/70. In red: amplification curve of an IVT of the corresponding WT sequence from SARS-CoV-2 at  $10^7 \text{ cp/rxn}$ . In black (overlapping with the red line) is the NTC.



	Bio-Rad CFX	ABI QuantStudio	QIAGEN Rotor-Gene Q
IVT copies/ reaction		Ct	
105	23.18	23.17	24.17
104	26.37	26.54	27.13
10 <sup>3</sup>	29.64	29.87	29.53
10 <sup>2</sup>	32.77	33.08	32.85
10 <sup>1</sup>	35.70	36.34	37.46

Specific and sensitive detection of the mutation P681H. In green: amplification curves from dilution series ( $10^5-10^1$  cp/rxn, in duplicates) of an in-vitro transcript (IVT) bearing the SARS-CoV-2 mutation P681H. In red: amplification curve of an IVT of the corresponding WT sequence from SARS-CoV-2 at  $10^7$  cp/rxn. In black (overlapping with the red line) is the NTC.

	Control genes			
– Mutation assays	N1	N2	Egene	RdRp
N501Y	✓	✓	1	✓
E484K		✓	1	1
E484Q		✓	1	1
P681R	✓	✓		
L452R	1	✓	1	1
T478K		✓		
K417N	✓	✓	1	1
K417T		✓	1	
T20N	✓	✓	1	
Del HV 69/70	✓	✓	1	1
P681H	✓	✓	1	

Genotyping assay compatibility with control genes in a duplex reaction

To order all your oligos quickly and easily, visit biomers.net at https://www.biomers.net/en/products/Catalog\_Products/Primers-for-QIAGEN-SARS-CoV-2-Assays.

To order the QIAprep&amp Viral RNA UM Kit, visit QIAGEN at www.qiagen.com/qiaprepamp-viral-rna-um-kit.

Download a Quick-Start Protocol for the SARS-CoV-2 LNA qPCR Assays at www.qiagen.com/covid-genotyping-QSP.

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The assays described here are intended for molecular biology applications. These assays are not intended for the diagnosis, prevention, or treatment of a disease.

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