

# Influenza B virus VICTORIA lineage TaqMan 2018 / FluB-VICT-TM2018 Influenza B virus VICTORIA lineage TaqMan 2018 / FluB-VICT-TM2018

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## Abstract

This protocol was designed and developed at this laboratory.

The protocol specifically aims to amplify strains of Influenza B VICTORIA virus lineage and not strains of the YAMAGATA virus lineage or other virus species. The assay targets the haemagglutinin (HA) region and is designed as a qualitative lineage-typing test for human cases of seasonal influenza virus type B infections.

FluB-VICT-TM2018 is ideally used alongside its companion protocol, 'Influenza B virus YAMAGATA lineage TaqMan 2018' (FluB-YAMA-TM2018), which aims to target influenza B virus YAMAGATA lineage strains exclusively. The two assays perform best as UNIPLEX protocols; a drop in sensitivity was observed when combined in a DUPLEX format.

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## Before start

- If using a different brand or model of real-time thermocycler, check the concentration of ROX is adequate.
- Method assumes the user is familiar with the thermocycler and software used to run the protocol and with PCR in general.

## Materials

SuperScript™ III Platinum™ One-Step qRT-PCR Kit 11732088 by Life Technologies

## Protocol

### Oligonucleotide sequences

#### Step 1.

| Name              | Sequence 5'-3'   |
|-------------------|--|
| FluB-HA-lineageF  | ACCAG <b>R</b> GGGAACTATGCCC                               |
| FluB-HA-lineageR1 | CCD <b>G</b> ATGT <b>R</b> AYAGG <b>Y</b> YTG <b>R</b> CYT |
| FluB-HA-lineageR2 | CCGGATGT <b>D</b> ACAGGT <b>Y</b> TGAC <b>Y</b> T          |
| FluB-HA-lineageR3 | CCAGATGTA <b>A</b> YAGGTCT <b>K</b> AYTT                   |

|                    |   |
|--------------------|---|
| FluB-HA-lineageR4  | CCD <b>D</b> GATGTAACAGGTCTG <b>R</b> CYT                         |
| FluB-HA-Vict-Probe | CY5 - CAGACCAAAATGCAC <b>R</b> GGGA <b>A</b> HATACC- <i>BHQ-3</i> |

## Reagents

### Step 2.



#### REAGENTS

SuperScript™ III Platinum™ One-Step qRT-PCR Kit 11732088 by Life Technologies

## Reaction set-up

### Step 3.

The assay has been used on both a Rotor-Gene 6000 and a Rotor-Gene Q real-time thermocycler

Prepare sufficient mix for the number of reactions.

Include a suitable 'dead volume' as necessary if using a robotic dispenser.

### MIX PREPARATION

| Reagent   | Volume (µl) x1 | Final reaction concentration |
|---|----------------|------------------------------|
| Nuclease-free water                             | 3.46           | N/A                          |
| FluB-HA-lineageF 200pmol/µl                     | 0.07           | 700nM                        |
| FluB-HA-lineageR1 200pmol/µl                    | 0.05           | 500nM                        |
| FluB-HA-lineageR2 200pmol/µl                    | 0.05           | 500nM                        |
| FluB-HA-lineageR3 200pmol/µl                    | 0.05           | 500nM                        |
| FluB-HA-lineageR4 200pmol/µl                    | 0.05           | 500nM                        |
| FluB-HA-Vict-Probe 100pmol/µl                   | 0.03           | 150nM                        |
| 2X Reaction Mix <sup>1</sup>                    | 10             | 1X                           |
| MgSO <sub>4</sub> 50mM                          | 0.8            | 5mM                          |
| SuperScript® III/Platinum® Taq Mix <sup>1</sup> | 0.4            | 1X                           |
| ROX Reference Dye (25µM)                        | 0.04           | 0.05µM                       |
| Template  | 5              | N/A                          |
| <b>TOTAL</b>                                    | <b>20</b>      |                              |

<sup>1</sup>Superscript™III Platinum™ One-step qRT-PCR kit

- Dispense 15µL to each reaction well.
- Add 5µL of template (extracted RNA, controls or NTC [nuclease-free water] ).
- Total reaction volume is 20µL

## Amplification

### Step 4.

#### CYCLING CONDITIONS

|      |                    |     |
|------|--------------------|-----|
| 50°C | 5min               | 1X  |
| 95°C | 2min               | 1X  |
| 95°C | 3sec               | 40X |
| 60°C | 30sec <sup>1</sup> |     |

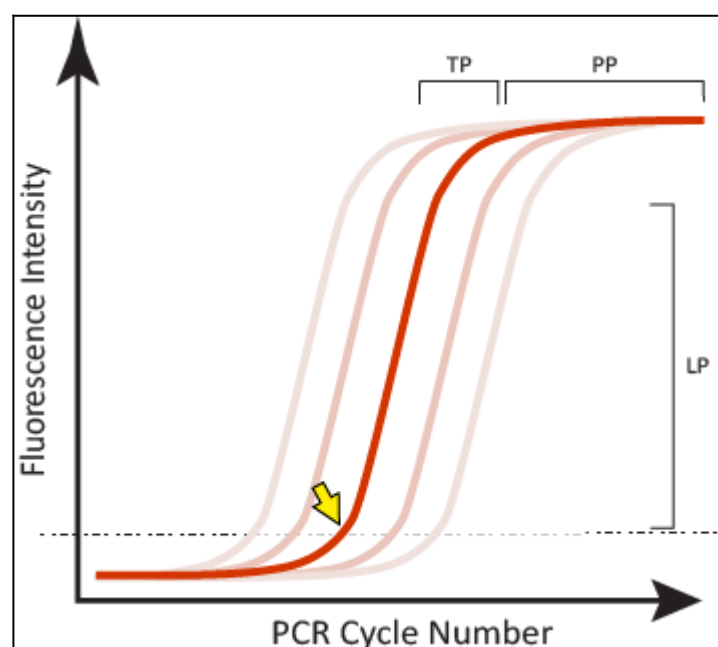
<sup>1</sup>Florescence acquisition step

## Result Analysis

### Step 5.

The definition used for a satisfactory positive result from a real-time fluorogenic PCR should include each of the following:

1. A **sigmoidal curve** – the trace travels horizontally, curves upward, continues in an exponential rise and followed by a curve towards a horizontal plateau phase
2. A **suitable level of fluorescence** intensity as measured in comparison to a positive control (y-axis)
3. A **defined threshold ( $C_T$ ) value** which the fluorescent curve has clearly exceeded (Fig.1 arrow), which sits early in the log-linear phase and is  $<40$  cycles
4. A flat or non-sigmoidal curve or a curve that crosses the threshold with a  $C_T >40$  cycles is considered a negative result.
5. NTCs should not produce a curve



**Figure 1.** Examples of satisfactory sigmoidal amplification curve shape when considering an assay's fluorescent signal output. The crossing point or threshold cycle ( $C_T$ ) is indicated (yellow arrow); it is the value at which fluorescence levels surpass a predefined (usually set during validation, or arbitrary) threshold level as shown in this normalized linear scale depiction. LP-log-linear phase of signal generated

during the exponential part of the PCR amplification; TP-a slowing of the amplification and accompanying fluorescence signal marks the transition phase; PP-the plateau phase is reached when there is little or no increase in fluorescent signal despite continued cycling.

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