

Development of a High Multiplex Respiratory Infectious Disease Detection Panel Using Next Generation qPCR Technology

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Abstract

Quantitative PCR using FRET-based real-time detection is restricted to very few targets within a single reaction. The ICEPlex® System offers an automated, high multiplexing solution. The ICEPlex System integrates PCR thermal cycling with dynamic, sequential amplicon separation by capillary electrophoresis (CE), and multi-color quantitative detection. Unlike probe-based qPCR, the ICEPlex System directly measures amplicon accumulation through incorporation of labeled primers.

Here we demonstrate that the ICEPlex system is capable of simultaneous detection and discrimination of important respiratory infectious agents belonging to Influenza and Respiratory Syncytial Virus (RSV). Primera Dx Multiplex Respiratory panel consists of primers that can detect and differentiate seasonal influenza strains, swine influenza, novel influenza strains, as well as RSV A and RSV B. The Multiplex Respiratory panel also contains primers that can detect drug resistant strains of Influenza. In addition, the assay consists of an internal control, extraction control, and a calibration control.

The ICEPlex System is an automated bench top instrument that has overcome many of the limitations of traditional real-time PCR based multiplexing. Using this high multiplex respiratory panel, we demonstrate the performance of the ICEPlex System in detection and discrimination of seasonal influenza strains, swine influenza, novel influenza strains, and drug resistant mutant strains.

Technology

The ICEPlex System is a fully automated real time PCR platform that combines an amplification module (thermocycler) and a detection module (a capillary electrophoresis cartridge, two solid state lasers with excitation maximum at 488 nm and 639 nm and a spectrophotometer with CCD camera). All ICEPlex System reagents are kept on board of the platform enabling an easy consumable maintenance (Figure 1).

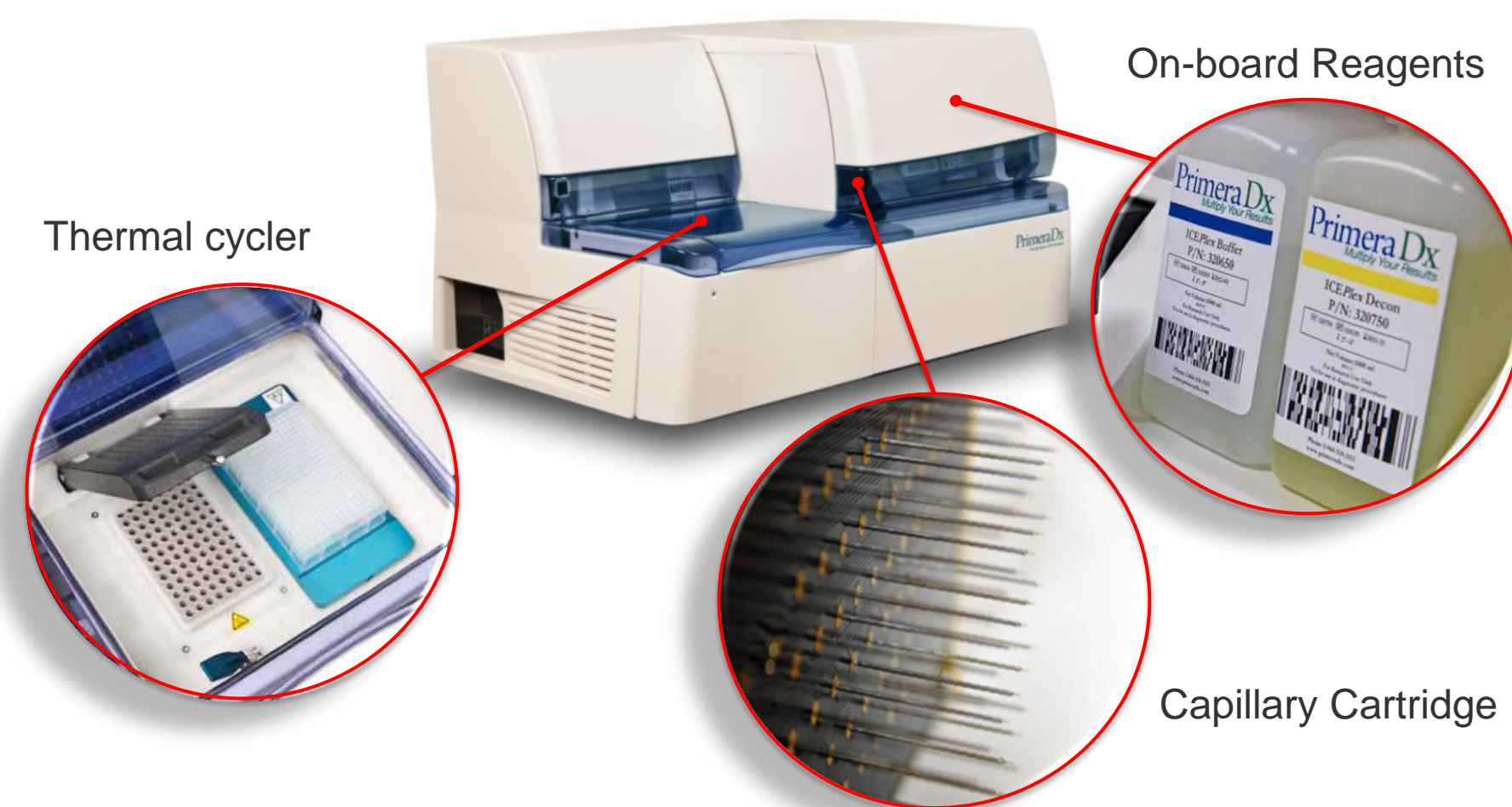


Figure 1. Components of the ICEPlex® System

The ICEPlex System generates fluorescently labeled PCR products (amplicons) which are separated based on their different sizes by capillary gel electrophoresis (CE). Accumulation of the fluorescent amplicons is monitored in real time by in house developed software that converts the fluorescent signal into amplification curves and calculates cycle thresholds (C_s) for all PCR targets. The combination of PCR and CE enables simultaneous detection and quantification of multiplex targets in single reaction with accuracy similar to real-time PCR methods (Figure 2).

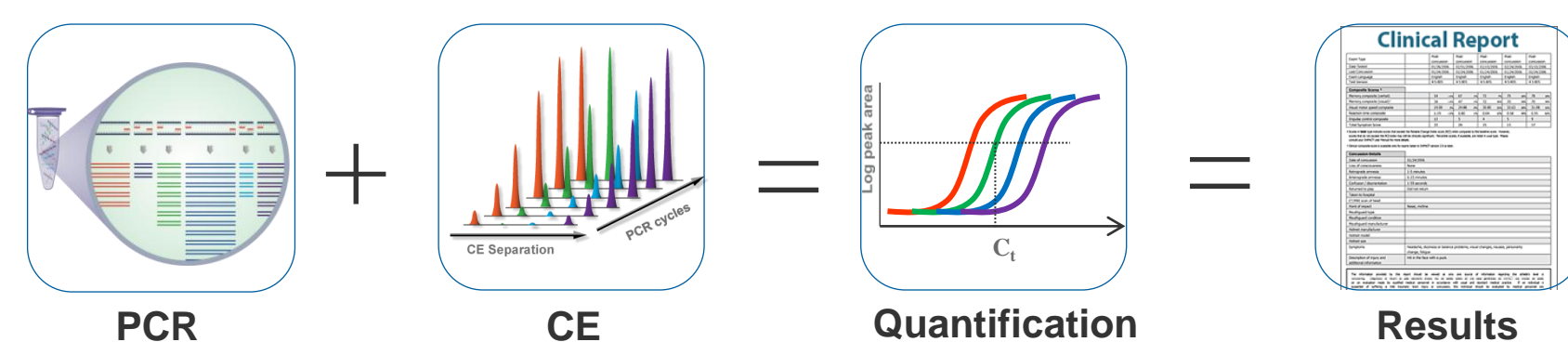


Figure 2. Multiplex real-time PCR detection on the ICEPlex® System

Introduction

Influenza virus causes highly contagious respiratory infection that can easily spread and cause considerable morbidity and mortality. It is major public health concern. Main causal agents of Influenza infection are Influenza virus A and B, and most of the Influenza pandemics are associated with Influenza A. Influenza A can infect human, animal, and birds, and can recombine to create novel Influenza strains that can be resistant to Oseltamivir (Tamiflu) and Zanamivir (Relenza). Another important viral agent that causes acute respiratory tract infection especially in young children and infants is respiratory syncytial virus (RSV), A and B. Here, we developed a multiplex respiratory panel that simultaneously and rapidly detects and discriminates Influenza A and B, Influenza A subtypes of public concerns and seasonal and pandemic importance, drug resistant (Oseltamivir) mutations, and RSV A and B.

Materials and methods

Primer design: Unique region of the genome for each target in the assay was identified by bioinformatics analysis followed by Basic Local Alignment Search Tool (BLAST) from the National Center for Biotechnology Information (Bethesda, MD). Geneious™ Pro software (Auckland, New Zealand) was used to carry-out ClustalW analysis (European Bioinformatics Institute, Cambridge, UK) of unique sequence, primer design on the consensus sequences (Steve Rozen and Helen J. Skaletsky (2000) Primer 3 on the WWW for general users and for biologist programmers), and primer-primer interaction among different primers (Cross-Hyb, Primera Dx, Inc., Mansfield, MA).

Genomic DNA templates: Synthetic oligonucleotides (IDT, Inc., Coralville, IA) that carry out the unique sequences for the different targets were used as templates to develop the panel. Influenza A, B, RSV A and B, A/H1, A/H3, N1-human, and N2 were tested using the RNA (data not shown).

PCR setup and amplification conditions: Multiplex PCR amplifications were performed in the 96 wells of a standard PCR plates. The PCR reactions contained: 1X Qiagen® Multiplex PCR Master Mix and 1 U of HotStarTaq® DNA Polymerase from Qiagen (Germantown, MD), 0.2 μM of gene specific forward and reverse primer pairs (IDT, Inc., Coralville, IA), of which one primer is labeled with FAM-dye (table 1) and 0.35X of the Universal Calibrators (PrimeraDx, Inc., Mansfield, MA). Twenty five microliter multiplex PCR reactions were subjected to thermocycling on the ICEPlex System.

PCR amplification conditions were as follow:

- 98° C for 10 minutes
- 16 cycles at 60° C for 20 sec., 72° C for 30 sec. and 96° C for 20 sec.
- 20 cycles at 64° C for 110 sec., 72° C for 150 sec., 96° C for 10 sec.

Organisms	Primers	Sequences
Influenza A	Forward	/56-FAM/TCATCCCCTCAGGCCCCCTC
	Reverse	GGGCACGGTGAGCGTGAACA
Influenza B	Forward	/56-FAM/AGTGGGGAACACACCTTCGGCG
	Reverse	GTCCTCTGGTGCCTTTCCGC
Influenza A/H1	Forward	/56-FAM/ATATACCCCAAGCCAAGTTCATGGCC
	Reverse	TGCCCCACAGCAGGACT
Influenza A/H3	Forward	/56-FAM/CACATACGGGGCCTGTCCCA
	Reverse	ATACCATCCACATTCCTCCCAACC
Influenza A/H5	Forward	/56-FAM/ACCATCGGGGAGTGCCCAAA
	Reverse	ATA TAA ATACCATTCCCTGCCATCCTCCC
Influenza A/H7	Forward	/56-FAM/TGTGGGGAAGTGCCCTGGT
	Reverse	ATCGATGAGGCCCTCCATCC
Influenza A/H9	Forward	/56-FAM/ACAAGGGGCTTGGCCCTATCA
	Reverse	GGGGCATAATCACCACCCGC
Influenza A/N1-human	Forward	/56-FAM/ATGGGGCCGCTGTACAAA
	Reverse	TCACTGTGCCAGTGTCTGGGT
Influenza A/N2	Forward	/56-FAM/AGGCTTCCGCTGGTGGGA
	Reverse	GGGTCGGATAAGGGTCTACCAC
SNP I117V	Forward	TTCCAAGGGGATGTGCCTGTTGA
	Reverse	/56-FAM/ TGTGAGGGCTTCTGCTTTGACAGTC
SNP I314V	Forward	CGTGCTATATAAATAGGCGAGATTAGAGAT
	Reverse	/56-FAM/GCCCTCTCTCGTTATTAGGATCCC
RSVA	Forward	/56-FAM/ACCTCACCAGAACCCAGC
	Reverse	GGCGTTGTTTGTGGTGGG
RSVB	Forward	/56-FAM/ACCACAGCATCCGAGCCCT
	Reverse	ATAACTCCATGGTATTTCGCCCCAG

Table 1. Gene specific primers for the different PCR targets in the multiplex respiratory panel

Results

Simultaneous detection of Influenza A and B, and Influenza A strain variants in single PCR reaction

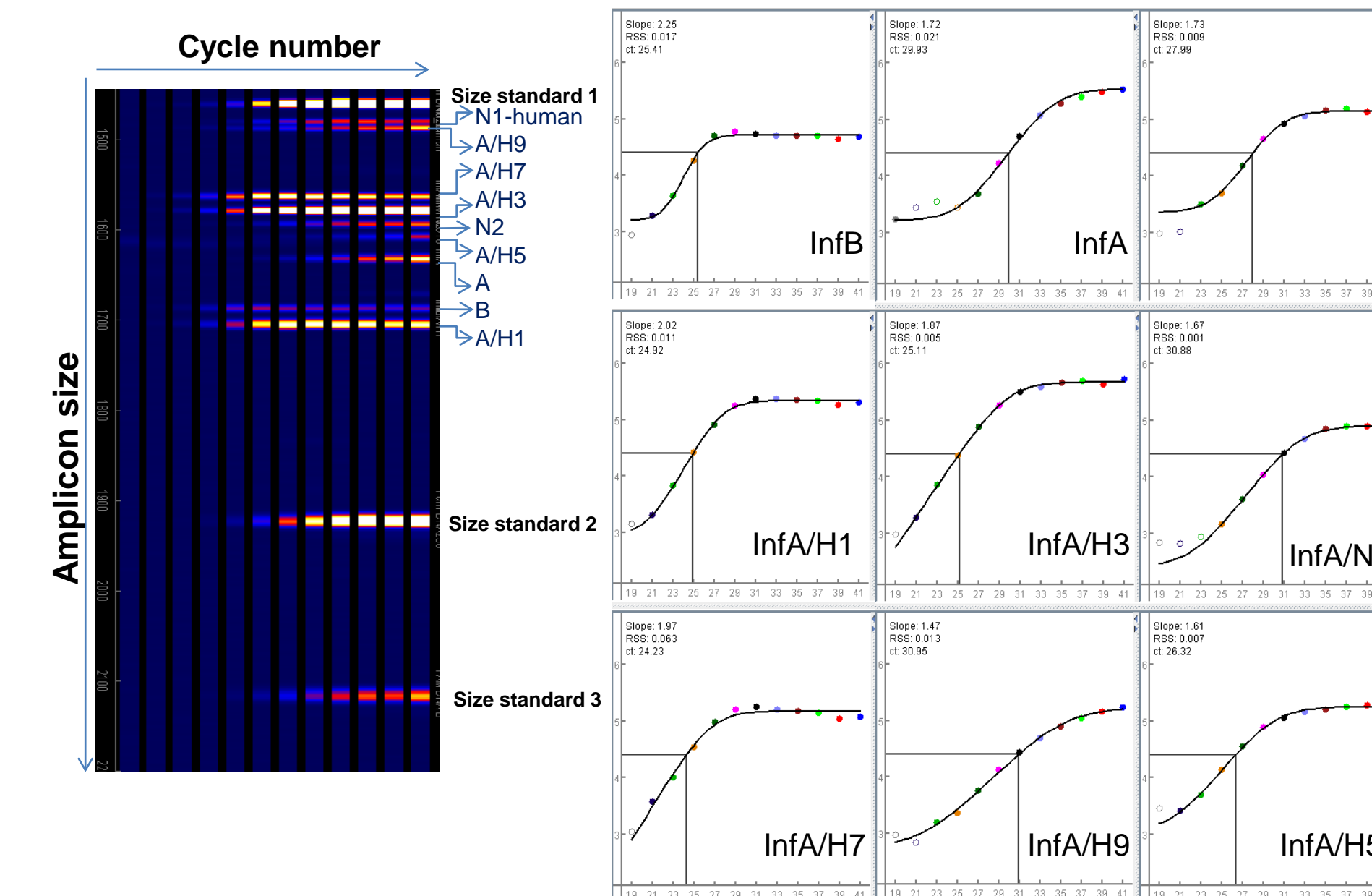


Figure 3. The multiplex respiratory panel can detect and discriminate Influenza A, B, and Influenza A strain variants.

The Multiplex Respiratory panel can simultaneously detect and differentiate RSV A and RSV B

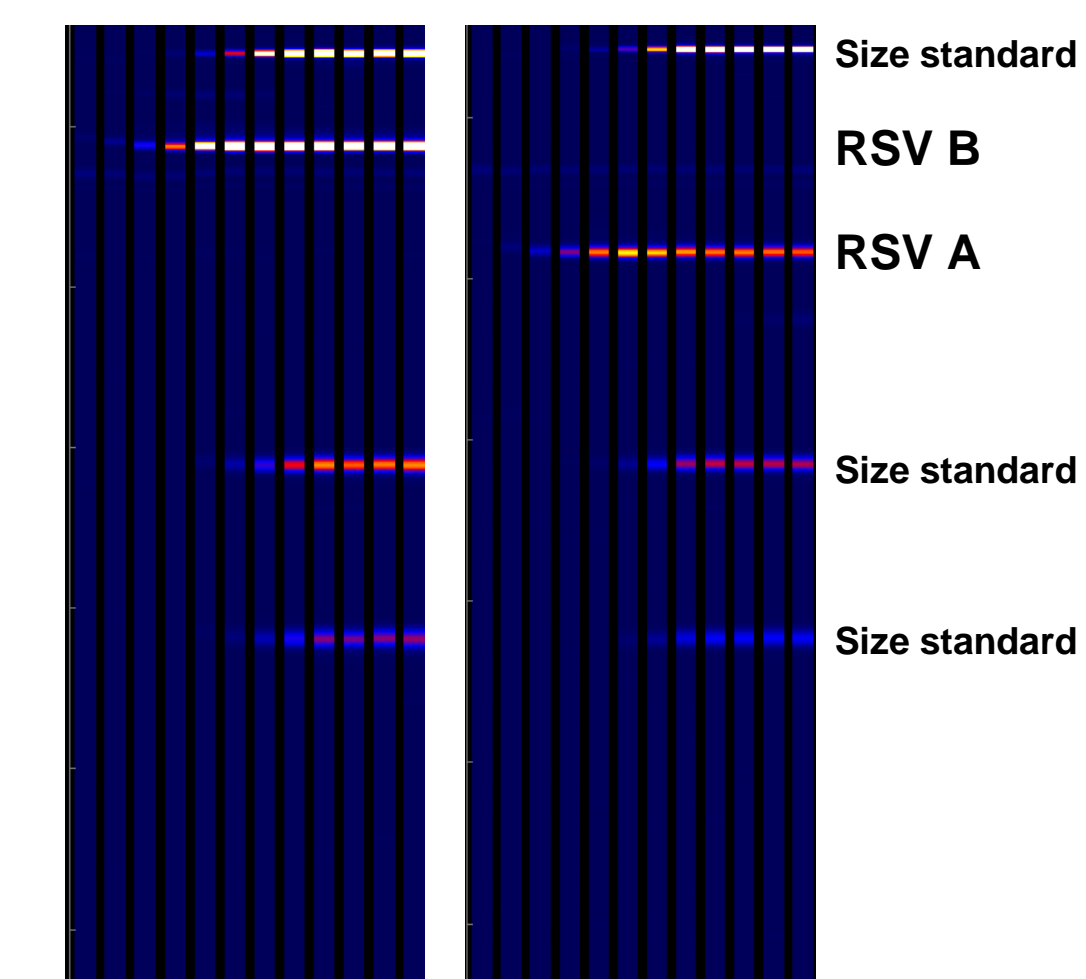


Figure 4. The Multiplex Respiratory panel can detect and discriminate respiratory syncytial virus A and B strains.

PDx Multiplex Respiratory panel can detect and discriminate Tamiflu drug resistant SNP

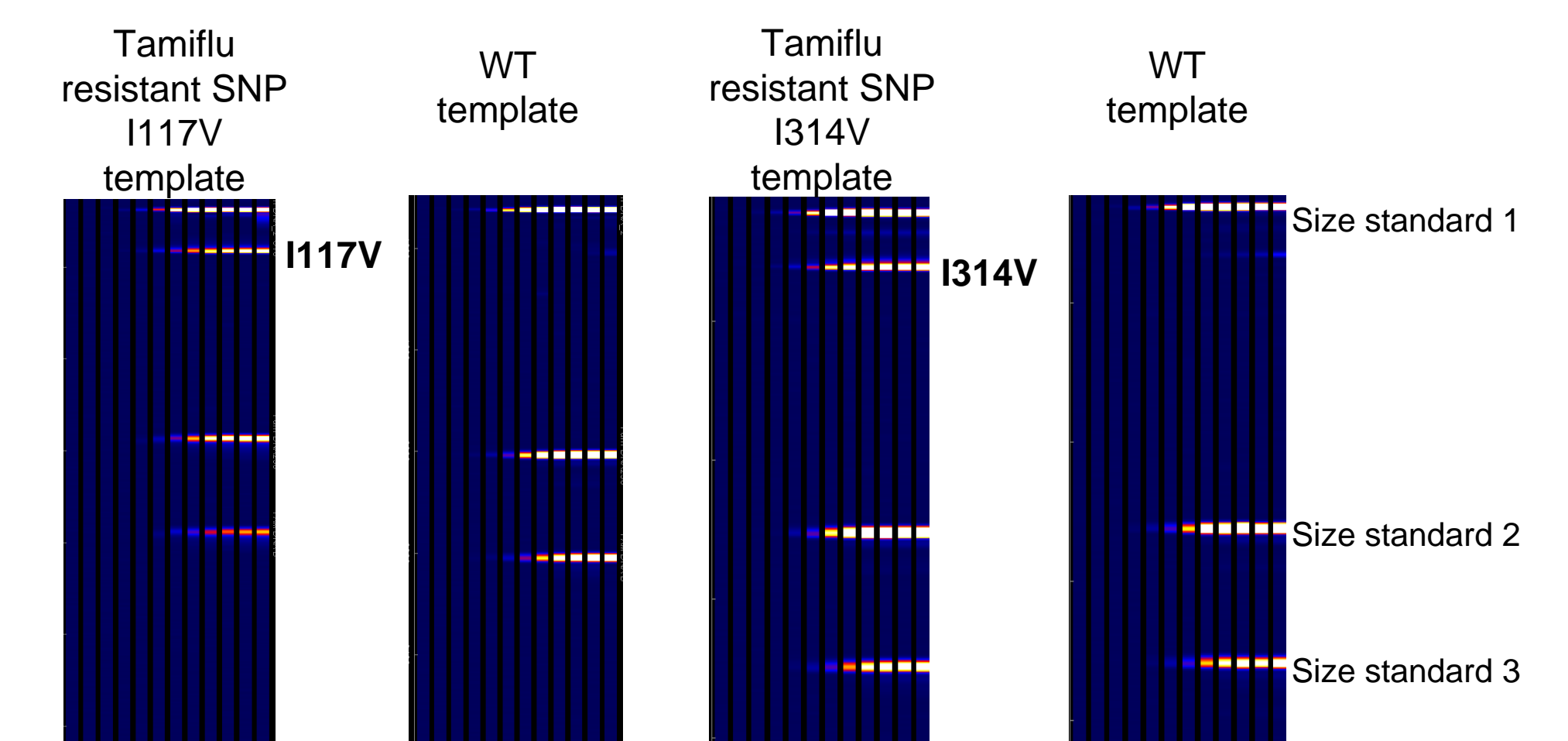


Figure 5. PDx Multiplex Respiratory panel can detect and discriminate Tamiflu drug resistant SNP, I117V and I314V, and thus helps in personalized medicine.

The PrimeraDx Multiplex Respiratory panel can detect Tamiflu resistant N2 subunit of avian and swine

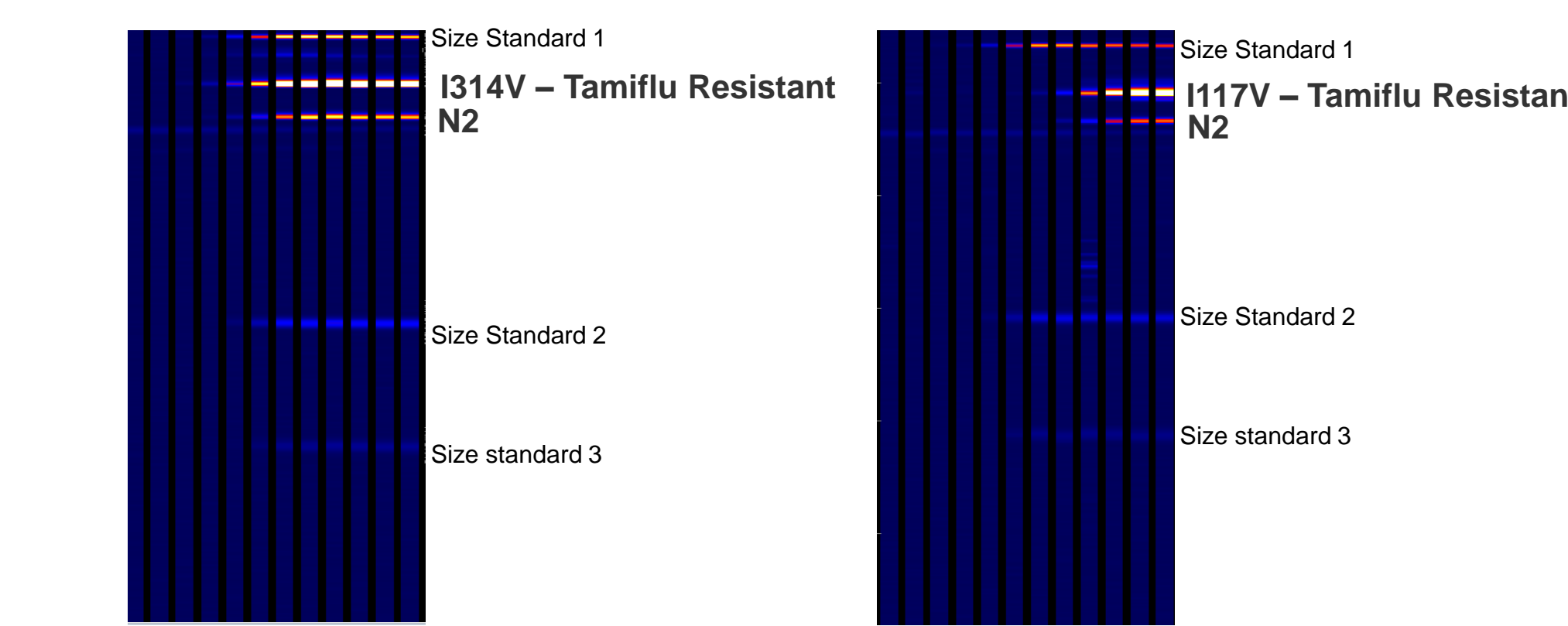


Figure 6. The Multiplex Respiratory panel simultaneously detects and discriminate SNP variant and the subtype

Conclusions

1. Pilot study demonstrates ICEPlex ability to detect multiple respiratory infectious agents, Influenza A and B, RSV A and B, Influenza A subtyping, and identify drug resistant SNP variants.
2. Primers have been tested against gram-negative bacteria, gram-positive bacteria, and fungal genomic DNA mixes for any cross-reactivity. The primers showed specificity with their templates.
3. The multiplex respiratory panel run time is less than three hours.

ICEPlex system and PDx Multiplex Respiratory panel have not been approved by the FDA for IVD. This information is for demonstration purpose only.