

Detection and Identification of Microorganisms using PathoKey® SQ FLEX Pathogen ID Assay



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INTRODUCTION

The PathoKey® SQ Pathogen ID Assay is an automated Next Generation Sequencing (NGS) based In-Vitro Diagnostic test intended for the detection and identification of microorganisms simultaneously.

The PathoKey® Assay allows for the selection of up to four different pathogen targets per sample via the use of specific primer pools for target amplification. Pathogen targets include bacteria, fungi, parasites, viruses and antimicrobial resistant genes. The assay has a Limit of Detection (LoD) of 10^3 – 10^4 CFU/mL or copies/mL and has been shown to work with clinical samples like whole blood, plasma, serum, nasopharyngeal swabs, saliva, urine, stool, FFPE, CSF, BAL and skin swabs. Additionally, the assay has a built-in positive control that acts as an external control to monitor workflow operation, a negative control to detect workflow contamination and an extraction control to confirm validity of the extraction process.

The PathoKey® Assay workflow utilizes an automated robotic liquid handling system (Sentosa® SX 101) for total nucleic acid extraction, PCR setup and NGS library preparation for up to 24 samples. Template preparation is done via emulsion PCR using Ion PGM Hi-Q View OT2 Kit on the Sentosa® ST401 system and sequencing is performed on the Ion Torrent NGS platform. Data interpretation is subsequently performed by the Sentosa® SQ Reporter software and a report containing the list of pathogens detected in each sample is generated. This minimizes hands-on time and human error while enabling high throughput.

WORKFLOW

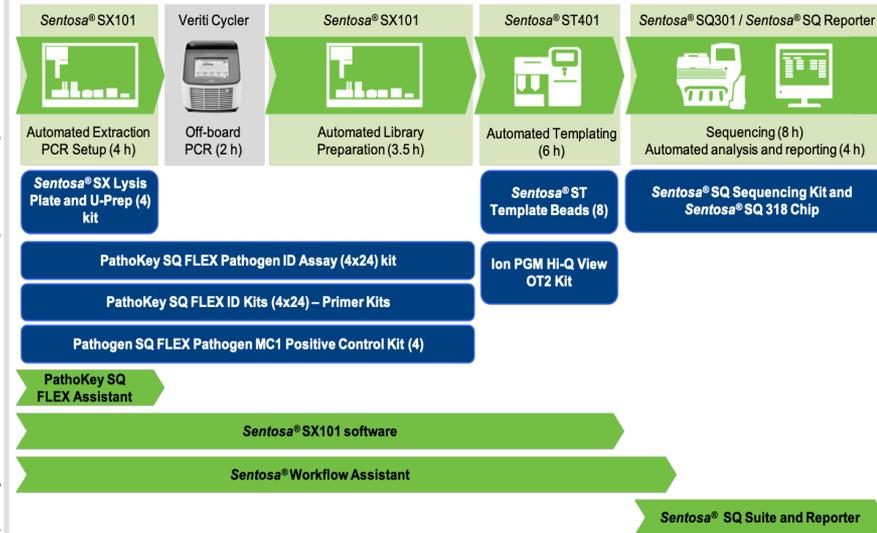


Figure 1. PathoKey® SQ Pathogen ID Assay Workflow

RESULTS



Analytical Sensitivity

The Limit of Detection (LoD) of the assay workflow for the bacteria, fungi, parasites, viruses and AMR bacteria strains tested was determined as 10^3 CFU/mL or copies/mL in biological matrices like nasopharyngeal and skin swabs, saliva, urine, whole blood, plasma, FFPE, BAL and CSF (Tables 1, 2 and 3).

Table 1. LoD for bacteria in nasopharyngeal swabs and plasma

Bacteria	Matrix	LoD (CFU/mL)
<i>Streptococcus pneumoniae</i>	Nasopharyngeal	10^3
<i>Stenotrophomonas maltophilia</i>	Nasopharyngeal	10^3
<i>Acinetobacter baumannii</i>	Nasopharyngeal	10^3
<i>Citrobacter freundii</i>	Nasopharyngeal	10^3
<i>Neisseria gonorrhoea</i>	Nasopharyngeal	10^3
<i>Vibrio parahaemolyticus</i>	Plasma	10^3
<i>Clostridium difficile</i>	Plasma	10^3
<i>Prevotella melaninogenica</i>	Nasopharyngeal	10^3
<i>Bacillus cereus</i>	Plasma	10^3
<i>Salmonella enterica</i>	Plasma	10^3
<i>Helicobacter pylori</i>	Nasopharyngeal	10^3

Table 2. LoD for fungi and parasites in various biological matrices

Fungi	Matrix	LoD (CFU/mL)
<i>Aspergillus sydowii</i>	Nasopharyngeal	10^3
<i>Pleurostoma richardsiae</i>	Skin	10^3
<i>Setosphaeria rostrata</i>	Nasopharyngeal	10^3
<i>Rhizopus microsporus</i>	Nasopharyngeal	10^3
<i>Rhodotprula mucilaginosa</i>	Skin	10^3
<i>Candida krusei</i>	Skin	10^3
<i>Cryptococcus neoformans</i>	CSF	10^3
<i>Candida albicans</i>	Nasopharyngeal	10^3
Parasites	Matrix	LoD (CFU/mL)
<i>Plasmodium falciparum</i>	Whole blood/ Plasma/FFPE	10^3
<i>Trypanosoma cruzi</i>	Whole blood/ Plasma/FFPE	10^3
<i>Acanthamoeba polyphaga</i>	Whole blood/ Plasma/FFPE	10^3

RESULTS, cont'd



Table 3. LoD for AMR bacteria and viruses in urine and saliva

AMR bacteria (genes)	Matrix	LoD (CFU/mL)
<i>Klebsiella pneumoniae</i> , K6 (SHV, SUL1)	Urine	10 ³
MRSA (ErmA, mecA)	Urine	10 ³
<i>Enterococcus faecium</i> (vanA)	Urine	10 ³
Viruses	Matrix	LoD (copies/mL)
Adenovirus	Saliva	10 ³
BK virus	Saliva	10 ³
Cytomegalovirus	Urine	10 ³
Herpes simplex virus 1	Saliva	10 ³
Herpes simplex virus 2	Urine	10 ³
Human herpesvirus 6	Saliva	10 ³
Human herpesvirus 7	Saliva	10 ³
Varicella-Zoster virus	Saliva	10 ³
Enterovirus	Urine	10 ³
JC virus	Saliva	10 ³

Clinical Evaluation

The assay was clinically evaluated with 41 patient urine samples, 43 urine DNA samples and 22 urine samples spiked with CDC AR Bank Isolates from POPLAR healthcare. The PathoKey® SQ Pathogen ID Assay was able to detect 91.6% of bacteria and 90.5% of AMR genes when compared to MicroPath ID qPCR technology platform (Tables 4 and 5). An example of the lab test report for one of POPLAR's sample is shown in Figure 2.

Table 4. Bacteria detection in POPLAR healthcare clinical samples

Bacteria detected with MicroID qPCR	Detected with PathoKey® Assay?
<i>Enterobacter cloacae</i>	Yes
<i>Enterococcus avium</i>	Yes
<i>Enterococcus faecalis</i>	Yes
<i>Enterococcus faecium</i>	Yes
<i>Escherichia coli</i>	Yes
<i>Klebsiella pneumoniae</i>	Yes
<i>Proteus mirabilis</i>	Yes
<i>Pseudomonas aeruginosa</i>	Yes
<i>Salmonella enterica</i> subsp. <i>enterica</i> serovar Concord	Yes
<i>Shigella flexneri</i>	Yes
<i>Streptococcus agalactiae</i>	Yes
<i>Ureaplasma urealyticum</i>	No

Table 5. AMR detection in POPLAR healthcare clinical samples

AMR genes detected with MicroID qPCR	Detected with PathoKey® Assay?
AAC	Yes
AADA	Yes
APH	Yes
CTX	Yes
dhfr	No
ErmA	Yes
ErmB	Yes
ErmC	Yes
gyrA	Yes
KPC	Yes
mecA	Yes
NDM	Yes
OXA	Yes
QnrA	No
QnrS	Yes
SHV	Yes
SUL1	Yes
TEM	Yes
TetM	Yes
vanA	Yes
VIM	Yes

Laboratory Test Report



Assay name : PathoKey™ SQ FLEX Pathogen ID Assay

Run name : 20210609_Microkey_R173 / 7YQXY

Barcode : SenlosaBC-22

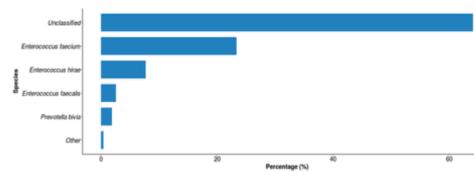
Sample Type : Unknown

Specimen ID : 20210609_Run173_S22

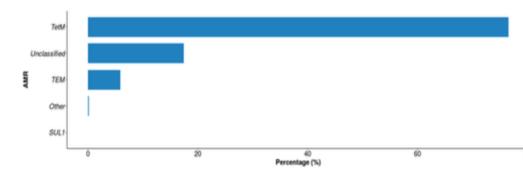
Date reported : 16-Jul-2021

Tested For : AMR, Bacteria

Species



AMR



No.	Species	Group	Read Count	Adjusted %
1	Enterococcus faecium	Bacteria	7428	23.34 %
2	Enterococcus faecalis	Bacteria	2467	7.67 %
3	Enterococcus faecalis	Bacteria	827	2.58 %
4	Prevotella bivia	Bacteria	658	1.89 %
5	Other	Various	228	0.41 %
6	Unclassified	Unknown	25406	64.11 %

No.	AMR	Group	Read Count	Adjusted %
1	TetM	AMR	25779	76.59 %
2	TEM	AMR	1978	5.88 %
3	Other	Various	45	0.12 %
4	SUL1	AMR	168	0.00 %
5	Unclassified	Unknown	5980	17.41 %

Figure 2. PathoKey® SQ Pathogen ID Assay Laboratory Test Report

CONCLUSION

We have developed a rapid and automated pathogen genotyping assay to enable the identification of various group of pathogens (bacteria, fungi, parasites, viruses and antimicrobial resistant genes) in a single sample which will contribute significantly to the study of host-microbiome research and greatly reduce the need to purchase multiple assay kits from different suppliers.

CONTACT

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