



The Combined Utility of the BIOFIRE® FILMARRAY® Research Configurations for Influent Wastewater Surveillance.

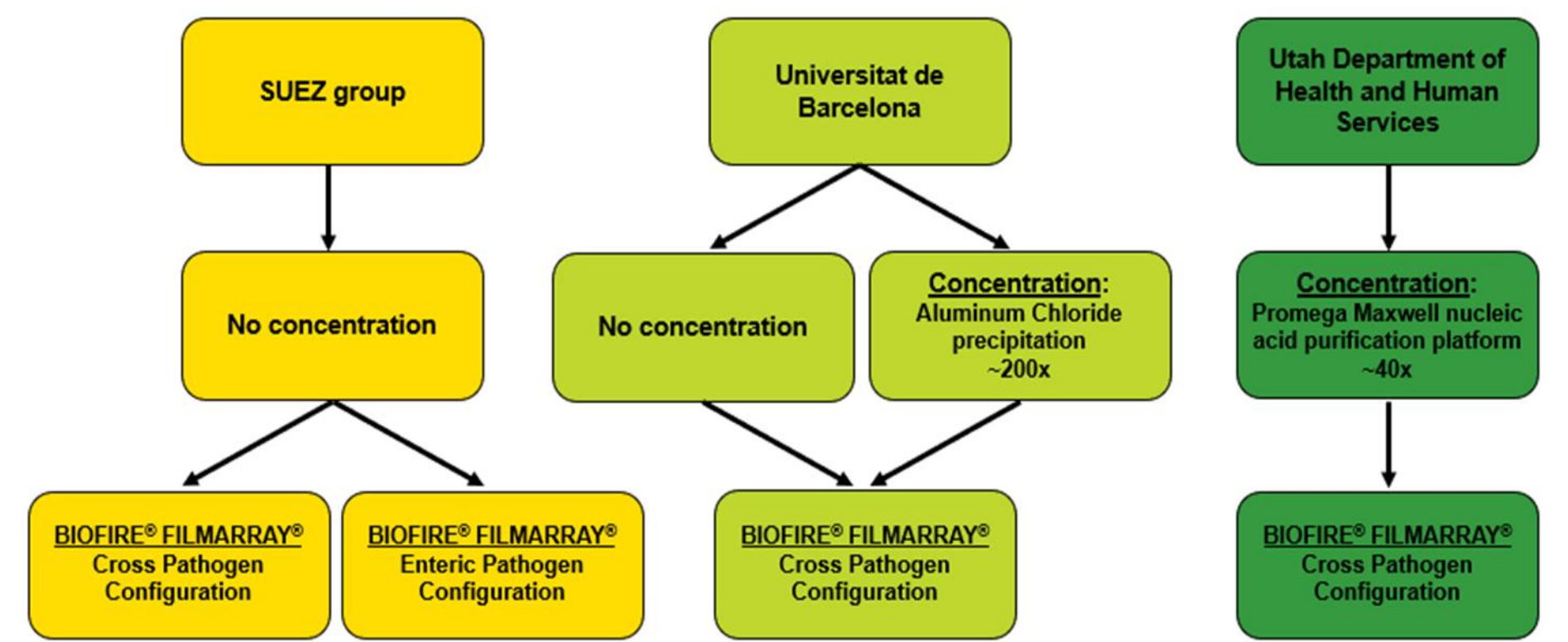
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INTRODUCTION

Wastewater-based surveillance is the intentional and consistent measurement of health markers in a wastewater or wastewater-derived sample. The BIOFIRE® FILMARRAY® qPCR system uses diagnostic panels designed to detect many human pathogens from a single patient sample of various sample types. The system includes integrated sample preparation, nucleic acid extraction, nested multiplex PCR, and melt curve analysis.

METHODS

To assess the performance of the BIOFIRE® system, two research configurations of existing BIOFIRE® respiratory, pneumonia, and enteric assays were used to monitor influent wastewater. Three study sites were set up throughout 2023. These sites included the Universitat de Barcelona (UB), SUEZ, and the Utah Department of Health and Human Services (UDHHS). These assays were chosen to assess their ability to detect the presence of analytes such as SARS-CoV-2, Influenza A/B, and Respiratory Syncytial Virus (RSV) in wastewater. Other assays including those for seasonal viruses, bacteria, fungi, and protozoa were also assessed in this study. Testing was performed with concentrated and unconcentrated influent wastewater samples following each site's standard preparation methods. For UDHHS, Crossing point (Cp) analysis was performed for Influenza and RSV using the BIOFIRE® FIREWORKS™ software (figure 4) to generate trend results which were then compared to clinical incidence rates in Utah (figure 5).



The BIOFIRE® FILMARRAY® System Wastewater Research Configurations Target Assays

Research Configuration 1 - Cross Pathogen / AMR Target List		Research Configuration 2 - Enteric Pathogen Target List	
Bacteria <i>Acinetobacter Calcoaceticus-baumannii complex</i> <i>Bordetella parapertussis</i> <i>Bordetella pertussis</i> <i>Chlamydia pneumoniae</i> <i>Enterobacter cloacae complex</i> <i>Escherichia coli</i> <i>Haemophilus influenzae</i> <i>Klebsiella aerogenes</i> <i>Klebsiella oxytoca</i> <i>Klebsiella pneumoniae group</i> <i>Legionella pneumophila</i> <i>Moraxella catarrhalis</i> <i>Mycoplasma pneumoniae</i> <i>Pseudomonas aeruginosa</i> <i>Protexis spp.</i> <i>Serratia marcescens</i> <i>Staphylococcus spp.</i> <i>Staphylococcus aureus</i> <i>Streptococcus agalactiae</i> <i>Streptococcus pneumoniae</i> <i>Streptococcus pyogenes</i> <i>Stenotrophomonas maltophilia</i>	Viral Adenovirus Seasonal Coronavirus (Non-SARS-CoV-2) Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) Human Metapneumovirus (hMPV) Human Rhinovirus/Enterovirus (HRV/VEV) Influenza A virus (FluA) Influenza B virus (FluB) Middle Eastern Respiratory Virus (MERS) Parainfluenza virus (PIV) Respiratory syncytial virus (RSV)	Bacteria <i>Campylobacter</i> <i>Clostridiaceae (Clostridium) difficile (toxin A/B)</i> <i>Clostridiaceae (Clostridium) difficile</i> <i>Plesiomonas shigelloides</i> <i>Salmonella</i> <i>Yersinia enterocolitica</i> <i>Vibrio spp.</i> <i>Vibrio cholerae</i>	Viral Adenovirus F40/41 Astrovirus Norovirus GI/GII Rotavirus A Sapovirus (I, II, IV, and V)
Antimicrobial Resistance Genes (AMR): Carbapenemases: ESBL, CTX-M, IMP KPC NDM OXA-48-Like VIM Methicillin resistance: mecA/C and MREJ (MRSA)		DIARRHEAGENIC ESCHERICHIA COLI/SHIGELLA: <i>Enteropathogenic E. coli (EPEC)</i> <i>Enterotoxigenic E. coli (ETEC) list</i> <i>Shiga-like toxin-producing E. coli (STEC) stx1/stx2</i> <i>E. coli O157</i> <i>Shigella/Enteroinvasive E. coli (EIEC)</i>	
		Parasites <i>Cryptosporidium</i> <i>Cyclospora cayentanensis</i> <i>Entamoeba histolytica</i> <i>Giardia lamblia</i>	

SUEZ RESULTS



SUEZ - Cross Pathogen Configuration				SUEZ - Cross Pathogen Configuration				SUEZ - Cross Pathogen Configuration				
Type	Assay Name	Positive Detections (n=25)	Percent Positive Detection	Type	Assay Name	Positive Detections (n=25)	Percent Positive Detection	Type	Assay Name	Positive Detections (n=25)	Percent Positive Detection	
Viral	Adenovirus	25	100%	Bacterial	<i>Acinetobacter</i>	25	100%	AMR	CTX-M	25	100%	
	Seasonal CoV	3	12%		<i>C. pneumoniae</i>	1	4%		IMP	25	100%	
	FluA	3	12%		<i>E. cloacae</i>	24	96%		KPC	21	84%	
	FluB	0	0%		<i>E. coli</i>	25	100%		MREJ	24	96%	
	hMPV	0	0%		<i>H. influenzae</i>	0	0%		NDM	25	100%	
	HRV/VEV	25	100%		<i>K. aerogenes</i>	21	84%		OXA-48-like	25	100%	
	MERS	0	0%		<i>K. oxytoca</i>	25	100%		VIM	25	100%	
	PIV	5	20%		<i>K. pneumoniae</i>	25	100%					
	RSV	4	16%		<i>L. pneumophila</i>	4	16%					
	SARSCoV2	25	100%		<i>M. catarrhalis</i>	0	0%					
			<i>M. pneumoniae</i>	0	0%							
			<i>P. aeruginosa</i>	23	92%							
			<i>P. rotens spp.</i>	24	96%							
			<i>S. agalactiae</i>	22	88%							
			<i>S. aureus</i>	22	88%							
			<i>S. maltophilia</i>	24	96%							
			<i>S. marcescens</i>	22	88%							
			<i>S. pneumoniae</i>	0	0%							
			<i>S. pyogenes</i>	0	0%							

Table 1. BIOFIRE® FILMARRAY® Cross Pathogen Configuration SUEZ positivity results for non-concentrated 24-hour composite influent wastewater samples collected August – November 2023 (n=25).

SUEZ - Enteric Pathogen Configuration				SUEZ - Enteric Pathogen Configuration				SUEZ - Enteric Pathogen Configuration			
Type	Assay Name	Positive Detections (n=28)	Percent Positive Detection	Type	Assay Name	Positive Detections (n=28)	Percent Positive Detection	Type	Assay Name	Positive Detections (n=28)	Percent Positive Detection
Viral	Adenovirus Type F	28	100%	Bacterial	<i>Aeromonas spp.</i>	27	96%	Parasites	<i>C. cayentanensis</i>	0	0%
	Astrovirus	27	96%		<i>Campylobacter spp.</i>	22	79%		<i>Cryptosporidium spp.</i>	4	14%
	Norovirus	27	96%		<i>C. difficile</i>	6	21%		<i>E. histolytica</i>	2	7%
	Rotavirus A	26	93%		<i>P. shigelloides</i>	9	32%		<i>G. lamblia</i>	28	100%
	Sapovirus	28	100%		<i>Salmonella</i>	8	29%		<i>Enterogreggative (EAGC)</i>	28	100%
					<i>Shigella spp.</i>	14	50%		<i>Enteropathogenic (EPEC)</i>	28	100%
					<i>V. cholerae</i>	9	32%		<i>Enterotoxigenic (ETEC)</i>	28	100%
			<i>Vibrio spp.</i>	13	46%	<i>Shiga-like toxin-producing (STEC)</i>	27	96%			
			<i>Y. enterocolitica</i>	21	75%	<i>STEC + O157</i>	20	71%			

Table 2. BIOFIRE® FILMARRAY® Enteric Pathogen Configuration SUEZ positivity results for non-concentrated 24-hour composite influent wastewater samples collected August – November 2023 (n=28).

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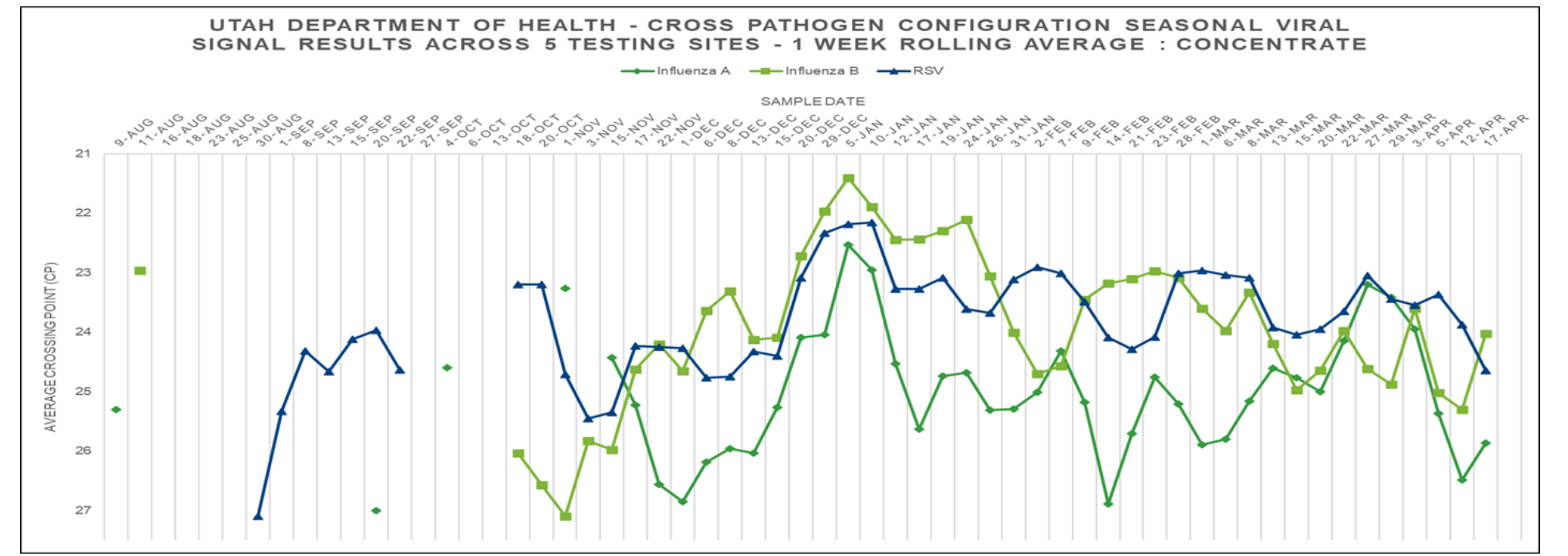


Figure 4. BIOFIRE® FILMARRAY® Cross Pathogen Configuration UDHHS seasonal viral results 1 week rolling average across 5 wastewater collection sites for concentrated 24-hour composite influent wastewater samples Salt Lake County collected August 2023 – April 2024 (n=288).

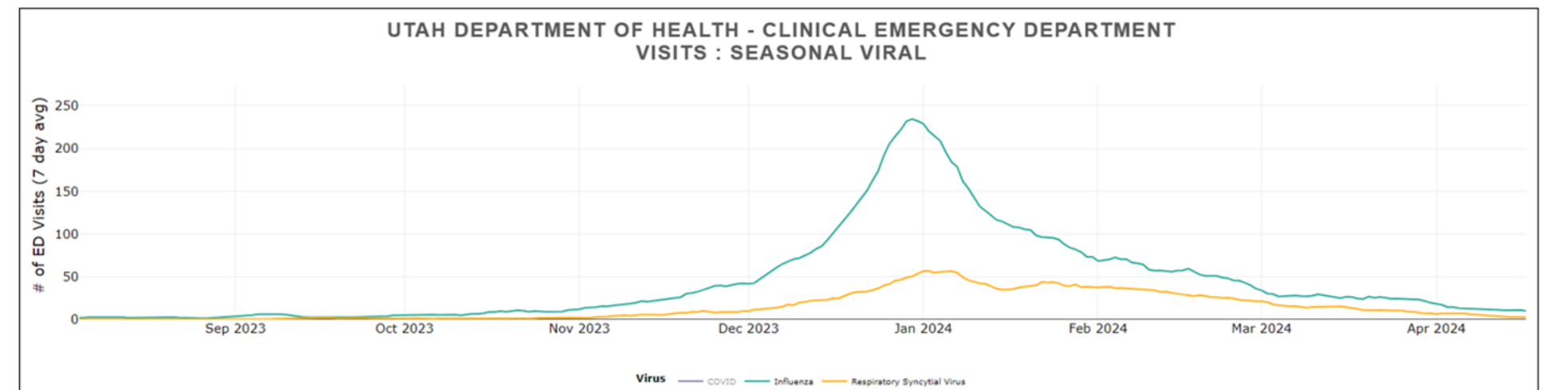


Figure 5. Statewide emergency department visits attributed to influenza and RSV reported by the Utah department of health and human services (<https://coronavirus.utah.gov/case-counts/>)

UNIVERSITAT DE BARCELONA

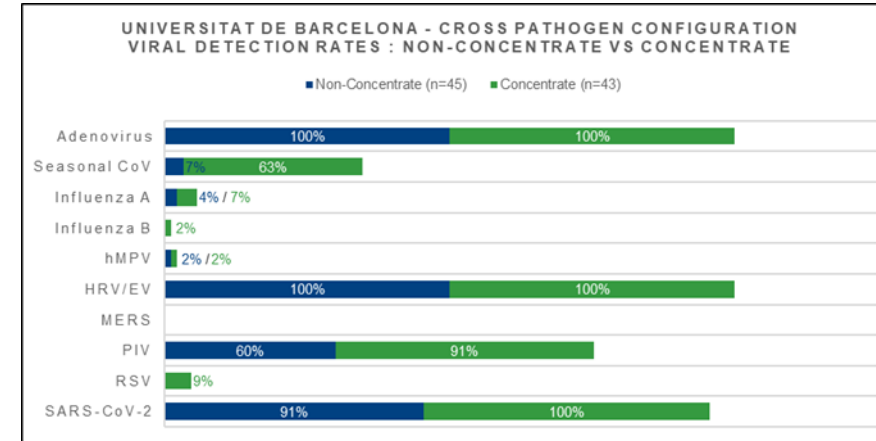


Figure 1. BIOFIRE® FILMARRAY® Cross Pathogen Configuration UB viral detection results for non-concentrated and concentrated 24-hour composite influent wastewater samples collected April – July 2023.

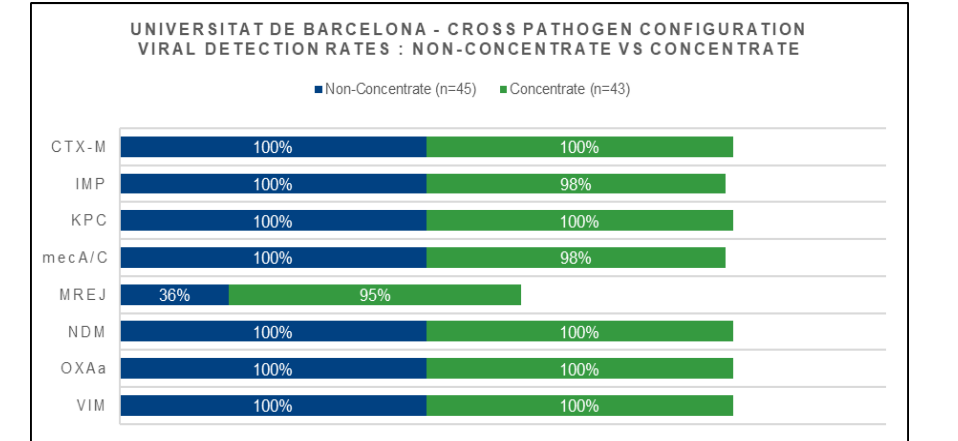


Figure 2. BIOFIRE® FILMARRAY® Cross Pathogen Configuration UB AMR detection results for non-concentrated and concentrated 24-hour composite influent wastewater samples collected April – July 2023.

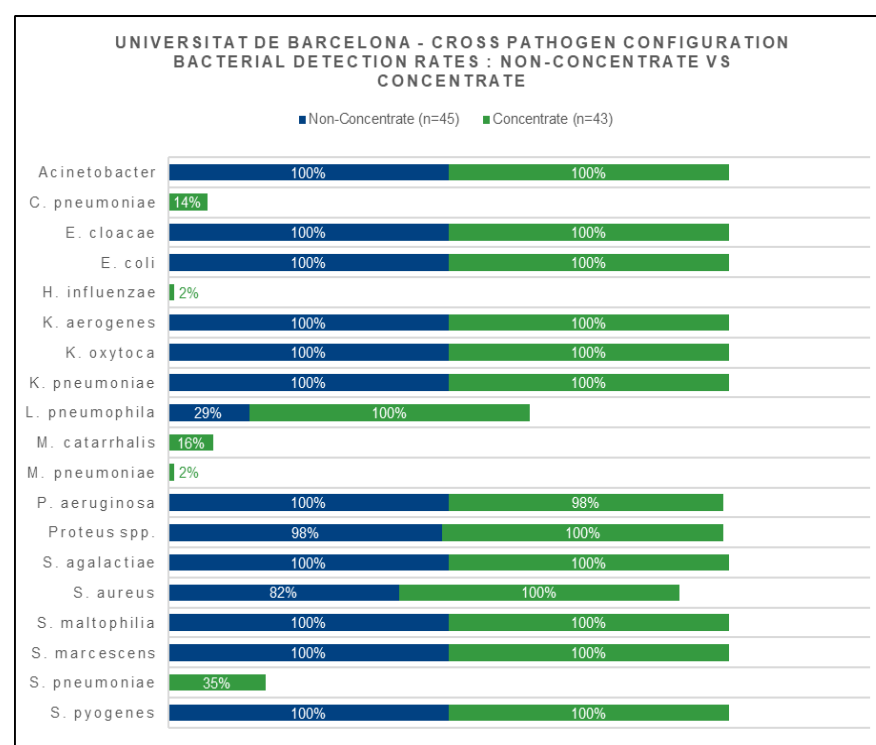


Figure 3. BIOFIRE® FILMARRAY® Cross Pathogen Configuration UB bacterial positivity results for non-concentrated and concentrated 24-hour composite influent wastewater samples collected April – July 2023.

AMR Gene Result	Applicable Bacteria
mecA/C and MREJ	<i>Staphylococcus aureus</i> <i>Acinetobacter calcoaceticus-baumannii</i> Complex
CTX-M	<i>Enterobacter cloacae</i> complex <i>Escherichia coli</i> <i>Klebsiella oxytoca</i> <i>Klebsiella pneumoniae</i> group <i>Protexis spp.</i>
IMP	<i>Pseudomonas aeruginosa</i> <i>Serratia marcescens</i>
KPC	<i>Enterobacter cloacae</i> complex <i>Escherichia coli</i> <i>Klebsiella oxytoca</i> <i>Klebsiella pneumoniae</i> group <i>Protexis spp.</i> <i>Serratia marcescens</i>
NDM	
OXA-48-like	
VIM	

Table 3. AMR gene targets and associated organisms.

Universitat de Barcelona Sample Sites	Population
Tarragona	145,832
Girona	153,162
Besòs	1,829,978
Prat de Llobregat	1,181,412
Terrassa	250,023

Table 4. Population per testing site UB.

GLOBAL RESULT SUMMARY

For influent wastewater samples across all testing sites, SARS-CoV-2 was detected in approximately 90% of unconcentrated samples and in 99% of concentrated samples tested. Influenza and RSV were mainly detected throughout respiratory season. For site Utah, increasing wastewater signal was observed for Flu and RSV from mid-October peaking on January 5th corresponding with clinical incident rates for the state. Environmental presence of these Viral analytes continued through April. Regular detections were observed for Adenovirus, HRV/VEV, seasonal coronaviruses (non-SARS-CoV-2), Parainfluenza virus, Norovirus, Rotavirus, Astrovirus, Sapovirus, *Bordetella parapertussis*, *Giardia lamblia*, Diarrheagenic *E. coli/Shigella*, and *Yersinia enterocolitica*. Furthermore, non-listed bacterial enteric pathogen assays showed high positivity rates in influent wastewater samples, as expected.

CONCLUSIONS

BIOFIRE® research configurations showed robust analyte detection with influent wastewater samples. These feasibility results suggest that the BIOFIRE® assays can be used for the intended purpose of wastewater-based surveillance.

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