Development of a Laboratory Verification Protocol for Concurrent Detection of Bacterial, Fungal, and Antimicrobial Resistance Genes in a Multiplex Syndromic Joint Infection Panel M. Cronin¹; T. Fadgen¹; B. Hoelcle²; R. Young¹; L. Ogden¹; J. Green¹; S. Thatcher¹ **IDWeek 2021** ¹BioFire Diagnostics, LLC, Salt Lake City, UT, USA; ²ZeptoMetrix[®], LLC, Buffalo, NY, USA #640

Background

Performance verification is a critical component of implementing a diagnostic test in a clinical lab and can be time consuming and costly. A verification protocol and organism panel were developed in collaboration with ZeptoMetrix[®], LLC to verify all analyte detections for the BioFire[®] Joint Infection (JI) Panel*. The proposed BioFire Joint Infection Panel detects 31 potential pathogens and 8 potential antimicrobial resistance (AMR) genes associated with joint infections from synovial fluid specimens.

*Investigational Use Only. Not for use in diagnostic procedures. Not available for sale. Under review by US FDA.



Methods

A protocol was developed using pilot NATtrolTM controls from ZeptoMetrix[®], synovial fluid, and the BioFire[®] FilmArray[®] 2.0 and the BioFire[®] FilmArray[®] Torch Systems. Control materials were tested in the presence of synovial fluid from pooled human donors. Synovial fluid was characterized for Joint Infection Panel targets by screening the specimen on the BioFire Joint Infection Panel prior to starting the verification procedure. The 32 targets required for all analyte detections were divided into 5 pools of 6 to 7 analytes and then tested over multiple days on several systems.

Laboratory Verification Workflow for the BioFire Joint Infection Panel





Verification Results for the BioFire Joint Infection Panel															
Organism and Resistance Genes			Summary								Summary				
			# Positives	# Possible Positives	# Negatives	# Modules	% Positivity	Organism and Resistance Genes			# Positives	# Possible Positives	# Negatives	# Modules	% Positivity
Pool 1	Candida		8	8	33	8	100		Angerococcus prevotii/ vaging	lis	8	8	33	8	100
	Candida albicans		8	8	33	8	100								100
	Enterococcus faecalis	vanB	8	8	33	8	100	4	Cutibacterium avidum/ granulosum		8	8	33	8	100
	Enterococcus faecium	vanA	8	8	33	8	100		Escherichia coli	IMP	8	8	33	8	100
	Finegoldia magna		8	8	33	8	100	Pc	Kingella kingae		8	8	33	8	100
	Morganella morganii		8	8	33	8	100		Proteus spp.		8	8	33	8	100
	Streptococcus spp.		8	8	24	8	100		/IP		8	8	33	8	100
	Streptococcus agalactiae		8	8	33	8	100		Clostridium perfringens		8	8	33	8	100
	Streptococcus pyogenes		8	8	33	8	100		KPC-2 (KPC) Z138 (CTX-N OXA-48 like) Z460 (CTX-N NDM)	KPC-2 (KPC)					
	vanA/B		8	8	33	8	100	Pool 5		Z138 (CTX-M &					
Pool 2	Enterobacter cloacae complex Haemophilus influenzae Peptoniphilus		8	8	33	8	100			OXA-48 like)	8	8	33	8	100
			8	8	33	8	100			Z460 (CTX-M &					
			8	8	33	8	100			NDM)					
	Peptostreptococcus anaerobius		8	8	33	8	100		Salmonella spp.		8	8	33	8	100
	Serratia marcescens		8	8	33	8	100		Stanbulgegegus lus dun ancis		0	0	22	0	100
	Staphylococcus aureus	mecA/C + MREJ	8	8	33	8	100		Staphylococcus lugaunensis		8	ð	33	8	100
	mecA/C + MREJ		8	8	33	8	100		CTX-M		8	8	33	8	100
Pool 3	Bacteroides fragilis		9	9	32	8	100		КРС		8	8	33	8	100
	Citrobacter		9	9	32	8	100		NDM		8	8	33	8	100
	Klebsiella aerogenes		9	9	32	8	100		OXA-48 like		8	8	33	8	100
	Neisseria gonorrhea		9	9	32	8	100								
	Parvimonas micra		8	9	33	8	89								
	Pseudomonas aeruginosa	VIM	9	9	32	8	100								
	Streptococcus spp.		9	9	24	8	100								
	Streptococcus pneumoniae		9	9	24	8	100								
	VIM		9	9	24	8	100								

A total of 41 BioFire Joint Infection tests were performed using the protocol developed and pilot control material (ZeptoMetrix[®] NATJIP-BIO).

- Expected positives: 263/264 (99.6%)
- Expected negatives: 1008/1008 (100%)
- Antibiotic resistance markers correctly identified when a correlated bacteria was present: 65/65 (100%)
- One missed detection (P. micra) was recovered upon re-testing and may be due sample handling of pilot verification materials, or manufacturing variability in the prototype test pouches.

Low-level organism may be present in matrix-document and proceed with testing

Step 2: Combine inactivated organisms and matrix to create a verification pool

Step 3: Ensure sample is mixed we prior to removing a sample for testing

Step 4: Follow Quick Guide instructions for pouch

Step 5: Run BioFire® Joint Infection Panel

Step 6: Document results







- days.
- Time to complete:
 - minutes.
 - system run time.
- 10 replicates per pool.



 al Fluid Screening Gram Positive Bacteria Gram Negative Bacteria Yeast SF#1 reported Strepted SF#2 reported Enterod cloacae complex SF#3 and SF#4 were repared targets Data presented here units Synovial fluid with JI pared 	ed human BioFire Joint reparing es were ytes ococcus spp. obacter negative for ased SF #4 anel
#2SF #3SF #4detections may be use	ed-but
novial Fluid Sample reported.	may be

Conclusions

Efficient performance verification may be achieved by combining 32 organisms/8 AMR into 5 pools and can be completed with 20 test runs in 4

• A single BioFire Joint Infection test run was completed in about 55

• Verification workflow of 20 tests completed in 18 hours 18 minutes of

• The pooling scheme provides multiple positive/negative detections for every BioFire Joint Infection target and sufficient material for running as many as

The protocol and controls serve as a useful tool for providing reliable detections of targets over multiple days, operators and systems and offers a flexible solution for supporting verification needs.

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