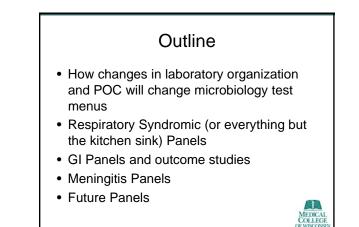
Syndromic Multiplex Panels – What's New, How Will We Use Them and What is the Impact on the Microbiology Laboratory

We Practice What We Teach

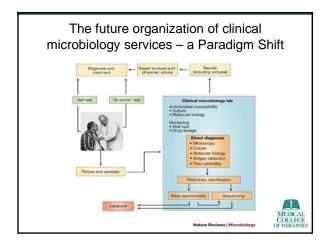
Nathan A Ledeboer Associate Professor of Pathology Medical College of Wisconsin

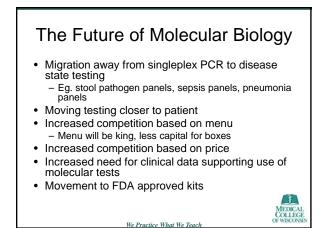
Medical Director, Microbiology and Molecular Pathology Wisconsin Diagnostic Laboratories and Froedtert Hospital

> MEDICAL COLLEGE OF WISCONSIN



We Practice What We Teach



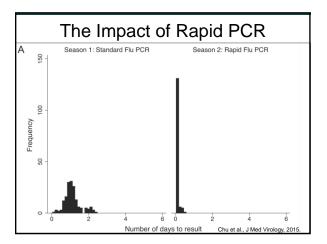


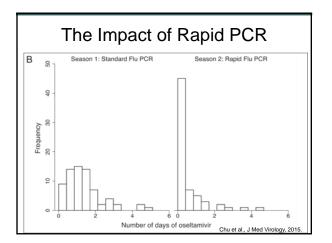
## Performance of POC Molecular Influenza Tests

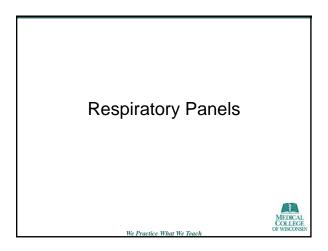
- Binnicker et al., JCM 2015
  - Compared Liat (Roche, Indianapolis, IN) to laboratory based Simplexa (Focus, Chantilly, VA)
  - Enrolled 197 respiratory swabs
  - Found Liat was 99.2% and 100% sensitive for Flu A and B, respectively
  - Found Liat was 100% specific compared to Simplexa

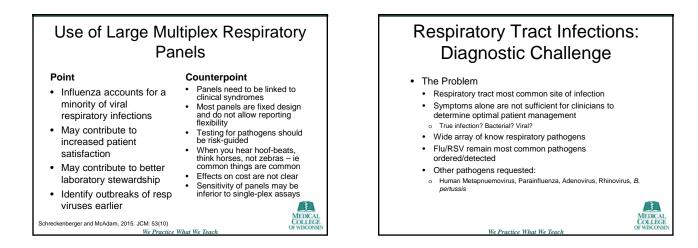
We Practice What We Teach

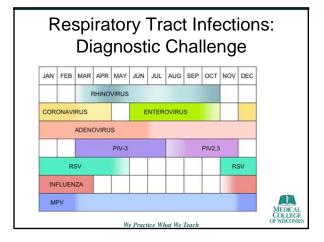








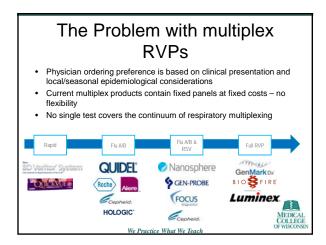


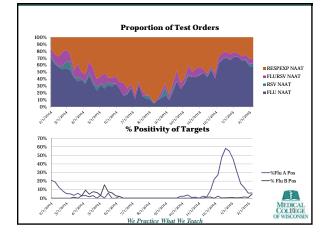


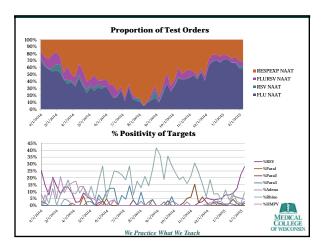


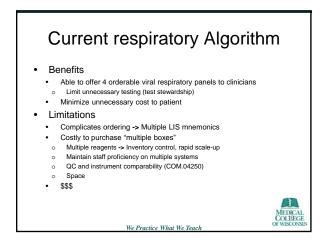
issaya	Manufacturor	Methodology	Preextraction required	Viruses reported <sup>®</sup>
FilmArray RP=	BioFire Diagnostics	Endpoint melt curve analysis	No	AdV; CoV HKU1, NL63; influenza virus A (H1/2009, H1, H3); influenza virus B; MPV; PIV1, -2, -3, -4; RSV; RhV/EV
eSensor RVP	GenMark Dx	Voltammetry	Yes	AdV (C, B/E); influenza virus A (H1/2009, H1, H3); influenza virus B; MPV; PIV1, -2, -3; RSV (A/B); RhV
xTAG RVPv1	Luminex Molecular Diagnostics	Fluorescence-labeled bead array	Yes	AdV; influenza virus A (H1, H3); influenza virus B; MPV; PIV1, -2, -3; RSV (A/B); RhV/EV
xTAG RVP fast	Luminex Molecular Diagnostics	Fluorescence-labeled bead array	Yes	AdV; influenza virus A (H1, H3); influenza virus B; MPV; RSV; RhV/EV

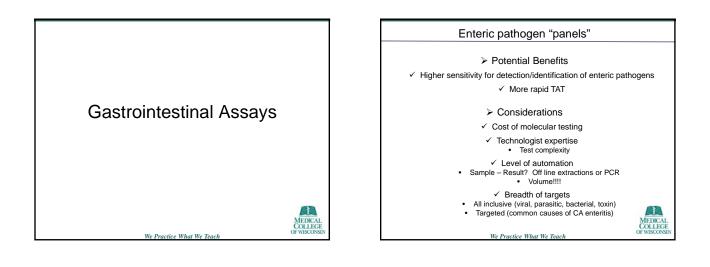
IVIUIUDIE	ex Assav	s for De	tection c	of Respira	atory Virus	ses
	,	% Sensitivity (95% C	l) of:		,	
Virus	No. of true-positive specimens (n = 300			XTAG		
	specimens tested)	FimArray RP	eSensor RVP	RVPv1	RVP fast	
AdV	35	57.1 (40.8, 72.0)	100 (88.2, 100)	74.3 (57.8, 86.0)	82.9 (66.9, 92.3)	
Influenza virus						
A	30	86.24 (68.8, 95.1)	100 (86.5, 100)	100 (86.5, 100)	86.7 (69.7, 95.3)	
A H1/09	16	73.3= (47.6, 89.5)	100 (77.3, 100)	100 (77.3, 100)	81.3 (58.2, 94.2)	
A H3	14	100 (74.9, 100)	100 (74.9, 100)	92.9 (66.5, 99.9)	78.6 (51.7, 93.2)	
в	22	77.3 (56.2, 90.3)	100 (82.5, 100)	95.5 (78.5, 99.9)	45.5 (26.9, 65.4)	
MPV	26	96.2 (79.6, 99.9)	100 (84.8, 100)	100 (84.8, 100)	100 (84.8, 100)	
PW						
1	14	100 (74.9, 100)	100 (74.9, 100)	100 (74.9, 100)	NA	
2	13	92.3 (64.6, 99.9)	100 (73.4, 100)	100 (73.4, 100)	NA	
3	13	100 (73.4, 100)	100 (73.4, 100)	100 (73.4, 100)	NA	
RSV						
A	22	88.4 (65.8, 96.1)	100 (82.5, 100)	86.4 (65.8, 96.1)	86.4 (65.8, 96.1)	
в	14	100 (74.9, 100)	100 (74.9, 100)	92.9 (66.5, 99.9)	85.7 (58.8, 97.2)	
RhV/EV	43	83.7 (69.7, 92.2)	90.7 (77.8, 96.9)	93.0 (80.7, 98.3)	93.0 (80.7, 98.3)	M

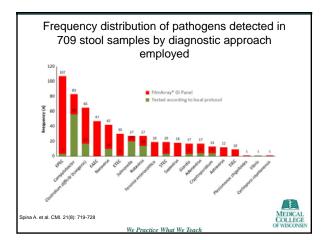


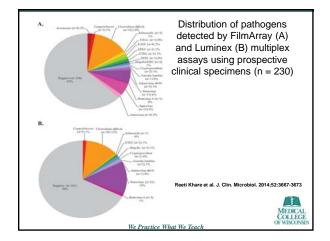




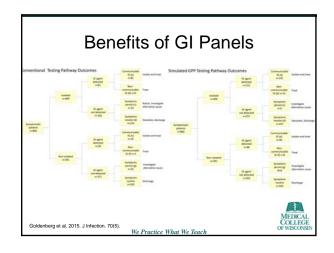








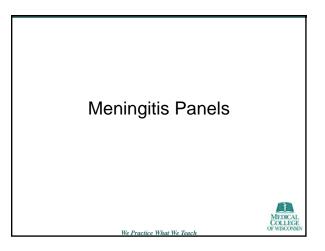
Target	FilmArray Lumine				ines			FilmArray		Luminex		
	TN	TP	FP	EN	TN	TP	5P	FN	Sensitivity (% 195% CID	Specificity (% (95% CID)	Sensitivity (% 195% CID	Specificity (% 195% CII)
Aeromonas spp.	248		0	16					23.8 (10.2, 47.1)	100 (98.2, 100)		
Campylobacter spp.	240	28	1	1	241	23	0	6	96.6 (81.4, 99.9)	99.6 (97.4, 100)	79.3 (61.3, 90.5)	100 (98.1, 100)
Clostridium difficile	229	33	5	3	229	33	4	3	91.7 (77.4, 97.9)	97.9 (95.0, 99.2)	91.7 (77.4, 97.9)	98.3 (95.5, 99.5)
Plesiomonas shigelloides	265	2	0	0					100 (29, 100)	100 (98.3, 100)		
Salmonalla spo.	246		0	0	246	20	0	4	100 (83.7, 100)	100 (98.1, 100)	83.3 (63.5, 93.9)	100 (98.1, 100)
Yersinia enterocolítica	242	27	1	0	243	13	0	14	100 (85.2, 100)	99.6 (97.5, 100)	48.1 (30.7, 66)	100 (98.1, 100)
Vibrio spp.	270	0	0	0					ND	ND		
Vibrio cholerae	270	0	0	0	270	0	0	0	ND	ND	ND	ND
EAEC	256	0	0	0					ND	ND		
EPEC	243	0	0	0					ND	ND		
ETEC	267	0	0	0	270	0	0	0	ND	ND	ND	ND
STEC	240	28	2	0	241	27	1	1	100 (85.7, 100)	99.2 (96.8, 100)	95.4 (80.8, 99.9)	99.6 (97.5, 100)
E col/0157	258	11	0	0	257	10	1	1	100 (70, 100)	100 (98.2, 100)	90.9 (60.1, 99.9)	99.6 (97.6, 100)
Shigella/EIEC	258	10	1	1	259	9	0	2	90.9 (60.1, 99.9)	99.6 (97.6, 100)	81.8 (51.1, 96)	ND
Cryptosporidium spp.	264	5	0	0	265	5	0	0	100 (51.1, 100)	100 (98.3, 100)	100 (51.1, 100)	100 (98.3, 100)
Cyclospora cavetanensis	269	0	0	0					ND	ND		
Entamoeba histolytica	269	0	0	1	268	1	0	0	0 (0, 83.3)	100 (98.3, 100)	100 (16.8, 100)	100 (98.3, 100)
Giardia Iamblia	263	7	0	0	263	7	0	0	100 (60, 100)	100 (98.3, 100)	100 (60, 100)	100 (98.3, 100)
Adenovirus 40/41	258	9	2	1	259	8	0	2	90 (57.4, 99.9)	99.2 (97.1, 100)	80 (47.9, 95.4)	100 (98.2, 100)
Norovirus GI/GII	225	41	0	3	194	41	32	3	93.2 (81.3, 98.6)	100 (98.0, 100)	93.2 (81.3, 98.6)	85.8 (80.7, 89.8)
Rotavirus	255	14	1	0	254	13	1	1	100 (74.9, 100)	99.6 (97.6, 100)	92.9 (66.5, 99.9)	99.6 (97.6, 100)
Sapovirus	237	28	0	3					90.3 (74.3, 97.4)	100 (98.1, 100)		
Astrovirus	257	11	0	0					100 (70, 100)	100 (98.2, 100)		



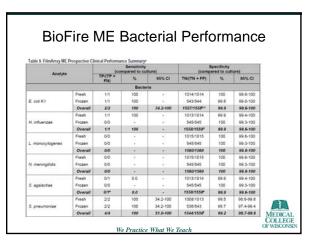
		0000	Sav	ingo		
Isolation days	3.0 days	2.5 days	2.0 days	1.5 days	1.0 days	
Total isolation days GPP testing pathway		1581	1447			
Total isolation cost under GPP testing pathway	£151,602	£139,754	£127,944	£116,057	£104,209	
Total laboratory testing costs for GPP	£56,243	£56,243	£56,243	£56,243	£56,243	
Total costs for GPP testing pathway	£207,845	£195,997	£184,187	£172,300	£160,452	
Total costs for conventional testing pathway	£228,661	£228,661	£228,661	£228,661	£228,661	
Net savings using GPP testing pathway	£20,816	£32,664	£44,474	£56,361	£68,209	

diarrheal stool sp	pecimens that		FilmArray GI Pane e for <i>C. difficile</i> an esting	
Pathogen on FilmArray GI Panel		testing positive for indi- hat were originally neg-		
	C. difficile, N = 142	Rotavirus≞, N = 16	All negative patients, N = 158	
Norovirus	9 (2)	1	10	
Rotavirus	8 (2)		8	
EPEC	8 (3)		8	
EIEC/Shigella	2 (2)	1	3	
EAEC	2		2	
ETEC	2 (1)		2	
Astrovirus		2 (1)	2	
Salmonella		1	1	
Cryptosporidium	1		1	
Aeromonas	1		1	
C. difficile	2	1 (1)	3	
Adenovirus		1	1	
Total pathogens	35/142 (24.6%)	7/16 (43.8%)	42/158 (26.6%)	<b>FB</b>
Total patients	29/142 (20.4%)	6/16 (37.5%)	35/158 (22.2%)	MEDICAL COLLEGE
Rand et al. 2015. DMID. 82(2).	We Practic	e What We Teach		OF WISCONSIN

Advantagés	Disadvantages
aboratory perspective Potential for lower cost with larger volumes Potential for enhanced efficiency and higher throughput Rapid reporting on engative samples and reductions in turnaround times Bundled test that can be applied to all faces samples	Contently significantly more expensive with low volumes: Calmer confirmation still required for autibated susceptibility studing ± strain typing Domand for festing may grow due to perception of rapidity of result. Need to retain their individual tests for creatin situations, e.g., hospitalised patients
ndividual patient management Rapid diagnosis Enhanced sensitivity for STEC, and C. difficile in particular Less dependent on clinical details or clinician request Results less affected by concurrent antibiotic therapy	Does not alter management for vast majority of patients Over diagnosis and low-loved positives of uncertain significance, which may result in unnecessary treatment uncertainty of the stream of the stream of the stream of the stream protocol of the stream of the stream of the stream of the Notatila to increase antibiotic presenting No antibiotic succeptibility result
nfection prevention and control and public health Rapid detection of outbreaks More reliable detection of pathogene potentially missed by culture, e.g., subiol fermenting STEC Increased detection of EP Shorter period of isolation for patients with negative results More sophisticated cohering of patients according to pathogen	No culture isolate available for strain typing Limited to organisms included in panel, which may not account for local epidemiolog Not suitable for terms to so dc. dcamance Loss of opportunity to passively screen for MDROs on stool



	lethods for FilmArray ME Pane			
FilmArray Analyte	Comparator Method	Comparator Test Location		
H. influenzae		Source Laboratory		
L. monocytogenes	CSF bacterial culture			
N. meningitidis				
S. agalactiae				
S. pneumoniae				
CMV				
EV				
HSV-1				
HSV-2	Two PCR assays with			
HHV-6	bi-directional sequencing <sup>a</sup>	BioFire Laboratory		
HPeV				
VZV	-			



Analyte		(compared	e Percent Ag to PCR with sequencing	bi-directional	Negative Percent Agreement (compared to PCR with bi-directiona sequencing)			
		TP/(TP + FN)	%	95% CI	TN/(TN + FP)	%	95% CI	
			Virus	os				
	Fresh	2/2	100	34.2-100	1010/1013	99.7	99.1-99.5	
CMV	Frozen	1/1	100	20.7-100	544/544	100	99.3-100	
	Overall	3/3	100	43.9-100	1554/15574	99.8	99.4-99.5	
EV	Fresh	43/44	97.7	88.2-99.6	965/971	99.4	98.7-99.7	
	Frozen	1/2	50.0		542/543	99.8	99.0-100	
	Overall	44/46*	95.7	85.5-98.8	1507/1514	99.5	99.0-99.8	
HSV-1	Fresh	1/1	100	-	1013/1014	99.9	99.4-100	
	Frozen	1/1	100		543/544	99.8	99.0-100	
	Overall	2/2	100	34.2-100	1556/1558	99.9	99.5-100	
HSV-2	Fresh	6/6	100	61.0-100	1008/1009	99.9	99.4-100	
	Frozen	4/4	100	51.0-100	540/541	99.8	99.0-100	
	Overall	10/10	100	72.2-100	1548/1550	99.9	99.5-100	
HHV-6	Fresh	13/15	86.7	62.1-96.3	997/1000	99.7	99.1-99.9	
	Frozen	5/6	83.3	43.6-97.0	535/536	99.8	99.0-100	
	Overall	18/21 <sup>k</sup>	85.7	65.4-95.0	1532/1536*	99.7	99.3-99.9	
	Fresh	9/9	100	70.1-100	1003/1006	99.7	99.1-99.9	
HPeV	Frozen	0/0			545/545	100	99.3-100	
	Overall	9/9	100	70.1-100	1548/1551	99.8	99.4-99.9	
	Fresh	3/3	100	43.9-100	1010/1012	99.8	99.3-99.9	
VZV	Frozen	1/1	100		543/544	99.8	99.0-100	
	Overall	4/4	100	51.0-100	1553/1556m	99.8	99.4-99.9	

