

# Verification of Analytical Bacterial Spectrum of Novodiag Bacterial GE+ and QIAstat-Dx GI Panels in a Hospital Setting

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# Novodiag



# QIAstat



Target	Method(s) used for routine testing	Target included on multiplex panel	
		QIAstat	Novodiag
<i>Campylobacter</i> <sup>b</sup>	Culture	✓	✓
<i>Salmonella</i> <sup>c</sup>	Culture	✓	✓
<i>Yersinia enterocolitica</i>	Culture	✓	✓
<i>Yersinia pseudotuberculosis</i>	Culture		✓
<i>Shigella</i> spp. <sup>d</sup>	Culture, in-house PCR	✓	✓
STEC	Culture, in-house PCR	✓	✓
EIEC	Culture, in-house PCR	✓	✓
EPEC	Culture, in-house PCR	✓	✓
ETEC	Culture, in-house PCR	✓	✓
EAEC	Culture, in-house PCR	✓	✓
<i>Clostridium difficile</i> <sup>e</sup> (tcdB)	Real-time PCR (GeneXpert)	✓	✓
<i>Plesiomonas shigelloides</i>	Culture	✓	
<i>Vibrio cholerae</i>	Culture	✓	✓
<i>Vibrio parahaemolyticus</i>	Culture	✓	✓
<i>Vibrio vulnificus</i>	Culture	✓	

Organisms (no. samples)	Species (no. samples)	Material (no. samples)	QIAstat Detection rate	Novodiag Detection rate
Campylobacter <sup>a</sup> (n=21)	C. jejuni (18)	Clinical samples (15)	14/15	15/15
		EQA, QCMD (GastroB18S-01 and 02)	2/2	2/2
		ATCC 33560	1/1	1/1
	C. coli (2)	EQA, QCMD (GastroB18S-03)	1/1	1/1
		ATCC 33559	1/1	1/1
	C. upsaliensis (1)	CCUG 23626	1/1	NI
Salmonella <sup>b</sup> (n=23)		Clinical samples (16)	12/16 <sup>c</sup>	10/16 <sup>c</sup>
		EQA, QCMD (GastroB18S-02)	1/1	0/1
		Clinical strains (4)	4/4	4/4
		ATCC 13076 and ATCC 14028	2/2	2/2
	Yersinia (n=16)	Y. enterocolitica (14)	7/7	6/7 <sup>d</sup>
		EQA, QCMD (GastroB18S-08)	1/1	0/1 <sup>d</sup>
		Clinical strains (5)	5/5	2/5 <sup>d</sup>
		ATCC 23715	1/1	0/1 <sup>d</sup>
		Y. pseudotuberculosis (2)	NI	2/2
Shigella spp. (n=13)	S. sonnei (7)	Clinical samples (6)	6/6	6/6
		ATCC 25931	1/1	1/1
	S. flexneri (4)	Clinical samples (2)	2/2	2/2
		EQA, QCMD (GastroB18S-07)	1/1	1/1
		ATCC 12022	1/1	1/1
	S. boydii (1)	NCTC 9359	1/1	1/1
	S. dysenteriae (1)	Clinical sample	1/1	1/1

# Vandbakterier

Organisms (no. samples)	Species	Material (no. samples)	QIAstat Detection rate	Novodiag Detection rate
Plesiomonas (n=3)	<i>P. shigelloides</i>	Clinical samples (1)  EQA, QCMD (GastroB18S-06)  ATCC 33560	1/1  1/1  1/1	NI
Vibrio (n=4)	<i>V. parahaemolyticus</i>  <i>V. vulnificus</i>  <i>V. cholerae</i>	Clinical strains (2)  Clinical strains (2)  NA	2/2  2/2	1/2  NI

# *Clostridioides (Clostridium) difficile*

Organism (no. samples)	Subtype (no. samples)	Material (no. samples)	QIAstat Detection rate	Novodiag Detection rate
<b><i>Clostridioides difficile</i></b> <b>(n=24)</b>	Toksin B (tcdB) (9)	Clinical samples (9)	9/9	8/9
	Toxin B (tcdB) plus binary toxin (cdtA/cdtB) (7)	Clinical samples (7)	7/7	3/7
	Toxin B (tcdB) plus binary toxin (cdtA/cdtB) and tcdCΔ117 (8)	Clinical samples (6) Clinical strains (1) EQA, QCMD (GastroB18S-04)	5/6 1/1 1/1	3/6 0/1 0/1

Organisms	Serotype	Targets	QIStat	Novodiag
EAEC;ETEC;EAEC	NA	eae, eltA, AggR	EPEC, LTST, AGGR	EPEC (eae), ETEC, EAEC
EAEC	NA	eae ( <b>lt,st?</b> )	(EPEC, LTST)	E. coli (eae)
ETEC	NA	estA	ETEC (LTST)	E. coli (ETEC)
EPEC	NA	eae	EPEC	E. coli (EPEC, eae)
STEC	O27:H30	stx2b	STEC (stx1/stx2)	EHEC stx2
EPEC	O111	eae	EPEC	E. coli (EPEC, eae)
EAEC	NA	AggR ( <b>eae?</b> )	EPEC, AGGR	E. coli (EPEC, eae) <b>missing aggR</b>
STEC	O26:H11	eae,stx1a	STEC (stx1/stx2)	EHEC (eae, stx1)
EIEC	NA	ipaH	EIEC/Shigella	Shigella spp. / EIEC
STEC	O26:H11	stx2a (eae pos SSI)	STEC (stx1/stx2)	EHEC,eae stx2
ETEC, EAEC	NA	eltA, estA aggR	ETEC, EAEC (LTST, AGGR)	E. coli (ETEC, EAEC)
ETEC	NA	eltA, estA	ETEC (LTST)	E. coli (ETEC)
STEC	O26:H11	eae,stx1a	STEC (stx1/stx2)	E. coli (eae, stx1)
EIEC	O96:H19	ipaH	EIEC/Shigella	Shigella spp. / EIEC
STEC	O157:H7	eae,stx1a;stx2c	STEC (stx1/stx2, O157:H7)	E. coli EHEC (eae,stx1,stx2)
STEC	O26:H11	stx1a	STEC (stx1, stx2, <b>0157: H7?</b> )	E. coli (eae, stx1)
STEC	O153, O178:H7!	stx1c ( <b>no eae primary</b> )	STEC (stx1/stx2)	E. coli EHEC (eae,stx1,stx2)
STEC	O145:H-	stx2a	STEC (stx1/stx2)	E. coli (EHEC, eae,stx2)
STEC	O157:H7	eae,stx1a, stx2a	STEC (stx1/stx2, STEC (O157:H7))	EHEC (eae, stx1,stx2)
STEC	O154:H31	stx1d	STEC (stx1/stx2)	EHEC (stx1)
STEC with estAp	OX187:O28;	stx2g,estAp	ETEC (lt/st), STEC (stx1/stx2)	EHEC (stx2), ETEC
STEC	O8:H9	stx2a stx2d	STEC (stx1/stx2)	EHEC (stx2)
STEC	O63:H6	stx2f	STEC (stx1/STX2)	EHEC (eae,stx2)
STEC	O145:H34	stx2f	STEC (stx1/stx2)	EHEC (eae,stx2)
STEC	O156:H4	stx2d	STEC (stx1/stx2)	EHEC (stx2)
STEC with eltA	O166:H15	stx2d eltA	STEC (stx1/stx2) ETEC (lt/st)	EHEC (stx2), ETEC
STEC	O9:H9	stx2e	STEC (stx1/stx2)	<b>Negative</b>
ETEC	O6:H16	eltA, estAh	ETEC (lt/st)	ETEC
EPEC	O157:H NA	eae	EPEC	<b>Negative</b>

<b>Conclusions</b>		
	QIAstat	Novodiag
<b>Hands-on-time</b>	Short	Short
<b>Turn-around-time</b>	+++	+++
<b>Amplification curves/ct value</b>	yes	no
<b>Reports</b>	STEC	STEC subtype
	Camp. spp	C. species level
<b>Technical failures</b>	++	+
<b>Multiplex, targets</b>	B+V+P	B (+P)
<b>Sensitivity, overall</b>	+++	++
<b>Salmonella</b>	++	+
<b>CD</b>	+++	+
<b>Yersinia</b>	+++ inkl. 1A	+++
<b>High throughput setting</b>	1 module +	4 modules +
<b>Versatility</b>	Resp. panel (CNS panel)	CARBA panel
<b>Price</b>	+++	+