Certificate of Analysis: Lyophilized Microorganism Specification and Performance Upon Release

Specifications	
Microorganism Name: Campylobacter lari **Catalog Number:** 01132 **Lot Number:** 1132-07 **Reference Number:** ATCC® 35221™* **Purity:** Pure **Passage from Reference:** 2	**Expiration Date:** 2019/1/31 **Release Information:** **Quality Control Technologist:** Mary L Bowman **Release Date:** 2017/3/31

Performance

Macroscopic Features:	Medium:
Small to medium sized, circular in shape with entire edges. Colorless to grey to slightly tan/olive green, shiny	Chocolate
Microscopic Features:	**Method:**
Gram negative rods (long, thin)	Gram Stain (1)

ID System: MALDI-TOF

See attached ID System results document.

Other Features/ Challenges: Results

(1) Catalase (3% Hydrogen Peroxide): positive
(1) Oxidase (Kovacs): positive
Rapid Hippurate Hydrolysis: negative
Campy Isolation Agar (42 C, MA): Good growth
Campy Isolation Agar (35 C, MA): Good growth
Campy Isolation Agar (25 C, MA): No growth at 72 hours
Motility (Wet Mount): darting-positive

Amanda Kuperus
Quality Control Manager
AUTHORIZED SIGNATURE

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Bruker Daltonik MALDI Biotyper Classification Results

Meaning of Score Values

Range	Interpretation	Symbols	Color
2.00 - 3.00	High-confidence identification	(+++)	green
1.70 - 1.99	Low-confidence identification	(+)	yellow
0.00 - 1.69	No Organism Identification Possible	(-)	red

Meaning of Consistency Categories (A - C)

Category	Interpretation
(A)	**High consistency:** The best match is a high-confidence identification. The second-best match is (1) a high-confidence identification in which the species is identical to the best match, (2) a low-confidence identification in which the species or genus is identical to the best match, or (3) a non-identification.
(B)	**Low consistency:** The requirements for high consistency are not met. The best match is a high- or low-confidence identification. The second-best match is (1) a high- or low-confidence identification in which the genus is identical to the best match or (2) a non-identification.
(C)	**No consistency:** The requirements for high or low consistency are not met.

Sample Name: Campylobacter lari
Sample Description: 01132
Sample ID: 1132-07
Sample Creation Date/Time: 2017-03-23T14:49:30.657 MB
Applied MSP Library(ies): BDAL, Mycobacteria Library (bead method), Filamentous Fungi Library 1.0, Listeria

Sample Name	Sample ID	Organism (best match)	Score Value
A11 (+++) (A)	1132-07	Campylobacter lari	2.39

Comments:

N/A

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Certificate of Analysis: Lyophilized Microorganism Specification and Performance Upon Release

Specifications	
Microorganism Name: Escherichia coli **Catalog Number:** 0335 **Lot Number:** 335-219 **Reference Number:** ATCC® 25922™* **Purity:** Pure **Passage from Reference:** 3	**Expiration Date:** 2018/11/30 **Release Information:** **Quality Control Technologist:** Christine Condon **Release Date:** 2017/1/23

Performance

Macroscopic Features:	**Medium:**
2 colony types, both are gray & beta hemolytic: one is circular to irregular, convex, slightly erose edge & smooth; other is larger, irregular, low convex, erose edge & rough	SBAP
Microscopic Features:	**Method:**
Gram negative straight rod	Gram Stain (1)

ID System: MALDI-TOF

See attached ID System results document.

Other Features/ Challenges: Results

(1) Oxidase (Kovacs): negative

Beta-glucuronidase (E. coli Broth w/MUG): positive

(1) Ampicillin (10 mcg - Disk Susceptibility): 16 - 22 mm

(1) Gentamicin (10 mcg - Disk Susceptibility): 19 - 26 mm

(1) SXT (1.25/23.75 mcg - Disk Susceptibility): 23 - 29 mm

Amanda Kuperus

Quality Control Manager

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Bruker Daltonik MALDI Biotyper Classification Results

Meaning of Score Values

Range	Interpretation	Symbols	Color
2.00 - 3.00	High-confidence identification	(+++)	green
1.70 - 1.99	Low-confidence identification	(+)	yellow
0.00 - 1.69	No Organism Identification Possible	(-)	red

Meaning of Consistency Categories (A - C)

Category	Interpretation
(A)	**High consistency:** The best match is a high-confidence identification. The second-best match is (1) a high-confidence identification in which the species is identical to the best match, (2) a low-confidence identification in which the species or genus is identical to the best match, or (3) a non-identification.
(B)	**Low consistency:** The requirements for high consistency are not met. The best match is a high- or low-confidence identification. The second-best match is (1) a high- or low-confidence identification in which the genus is identical to the best match or (2) a non-identification.
(C)	**No consistency:** The requirements for high or low consistency are not met.

Sample Name: Escherichia coli
Sample Description: 0335
Sample ID: 335-219
Sample Creation Date/Time: 2017-01-12T15:07:53.975 CC
Applied MSP Library(ies): BDAL, Mycobacteria Library (bead method), Filamentous Fungi Library 1.0, Listeria

Sample Name	Sample ID	Organism (best match)	Score Value
E1(+++) (A)	335-219	Escherichia coli	2.508

Comments:

closely related to Shigella and not definitely distinguishable at the moment

Microbiologics®

Certificate of Analysis: Lyophilized Microorganism Specification and Performance Upon Release

Specifications

Microorganism Name: Salmonella enterica subsp. enterica serovar Typhimurium
Catalog Number: 0363
Lot Number: 363-265
Reference Number: ATCC® 14028™*
Purity: Pure
Passage from Reference: 3

Expiration Date: 2018/12/31
Release Information:
Quality Control Technologist: Christine Condon
Release Date: 2017/2/24

Performance

Macroscopic Features:
Medium, gray/white, circular, slightly irregular edges, convex colonies

Microscopic Features:
Gram negative straight rods

Medium:
SBAP

Method:
Gram Stain (1)

ID System: MALDI-TOF

See attached ID System results document.

Other Features/ Challenges: Results

(1) Oxidase (Kovacs): negative

Hektoen Enteric agar: good growth, blue-green colonies with black centers

(1) Salmonella O antiserum Factor O:4 (Included in group B): positive

(1) Salmonella O antiserum Factor O:5 (Included in group B): positive

(1) Salmonella O antiserum Factor O:12 (Included in group B): positive

Amanda Kuperus
Quality Control Manager
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Bruker Daltonik MALDI Biotyper Classification Results

Meaning of Score Values

Range	Interpretation	Symbols	Color
2.00 - 3.00	High-confidence identification	(+++)	green
1.70 - 1.99	Low-confidence identification	(+)	yellow
0.00 - 1.69	No Organism Identification Possible	(-)	red

Meaning of Consistency Categories (A - C)

Category	Interpretation
(A)	**High consistency:** The best match is a high-confidence identification. The second-best match is (1) a high-confidence identification in which the species is identical to the best match, (2) a low-confidence identification in which the species or genus is identical to the best match, or (3) a non-identification.
(B)	**Low consistency:** The requirements for high consistency are not met. The best match is a high- or low-confidence identification. The second-best match is (1) a high- or low-confidence identification in which the genus is identical to the best match or (2) a non-identification.
(C)	**No consistency:** The requirements for high or low consistency are not met.

Sample Name: Salmonella enterica subsp. enterica serovar Typhimurium
Sample Description: 0363
Sample ID: 363-265
Sample Creation Date/Time: 2017-02-10T10:48:31.181 CC
Applied MSP Library(ies): BDAL, Mycobacteria Library (bead method), Filamentous Fungi Library 1.0, Listeria

Sample Name	Sample ID	Organism (best match)	Score Value
F2 (+++) (A)	363-265	Salmonella sp	2.40

Comments:

Salmonella can only be identified on genus level.

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Certificate of Analysis: Lyophilized Microorganism Specification and Performance Upon Release

Specifications	
Microorganism Name: Klebsiella pneumoniae **Catalog Number:** 01005 **Lot Number:** 1005-22 **Reference Number:** ATCC® BAA-1705™* **Purity:** Pure **Passage from Reference:** 3	**Expiration Date:** 2019/6/30 **Release Information:** **Quality Control Technologist:** Mary L Bowman **Release Date:** 2017/7/17

Performance

Macroscopic Features:

Large, circular, convex, entire edge, light gray, glistening and mucoid. A second type that is flatter and less mucoid may be seen.

Medium:

SBAP

Microscopic Features:

Short, broad, straight, gram negative bacilli with rounded ends.

Method:

Gram Stain (1)

ID System: MALDI-TOF

See attached ID System results document.

Other Features/ Challenges: Results

(1) Oxidase (Kovacs): negative

Modified Hodge Test: positive

Amanda Kuperus

Quality Control Manager

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Bruker Daltonik MALDI Biotyper Classification Results

Meaning of Score Values

Range	Interpretation	Symbols	Color
2.00 - 3.00	High-confidence identification	(+++)	green
1.70 - 1.99	Low-confidence identification	(+)	yellow
0.00 - 1.69	No Organism Identification Possible	(-)	red

Meaning of Consistency Categories (A - C)

Category	Interpretation
(A)	**High consistency:** The best match is a high-confidence identification. The second-best match is (1) a high-confidence identification in which the species is identical to the best match, (2) a low-confidence identification in which the species or genus is identical to the best match, or (3) a non-identification.
(B)	**Low consistency:** The requirements for high consistency are not met. The best match is a high- or low-confidence identification. The second-best match is (1) a high- or low-confidence identification in which the genus is identical to the best match or (2) a non-identification.
(C)	**No consistency:** The requirements for high or low consistency are not met.

Sample Name: Klebsiella pneumoniae
Sample Description: 01005
Sample ID: 1005-22
Sample Creation Date/Time: 2017-07-13T12:15:26.004 MB
Applied MSP Library(ies): BDAL, Mycobacteria Library (bead method), Filamentous Fungi Library 1.0, Listeria

Sample Name	Sample ID	Organism (best match)	Score Value
C12 (+++) (A)	1005-22	Klebsiella pneumoniae	2.36

Comments:

N/A

Certificate of Analysis: Lyophilized Microorganism Specification and Performance Upon Release

Specifications **Microorganism Name:** Klebsiella pneumoniae **Catalog Number:** 01060 **Lot Number:** 1060-09 **Reference Number:** ATCC® BAA-2146™* **Purity:** Pure **Passage from Reference:** 2	**Expiration Date:** 2019/4/30 **Release Information:** **Quality Control Technologist:** Christine Condon **Release Date:** 2017/5/8

Performance

Macroscopic Features:	**Medium:**
Small to medium, grey, glossy, circular, convex, entire edge	SBAP
Microscopic Features:	**Method:**
Gram negative rods. Coccobacillary to medium forms may be present.	Gram Stain (1)

ID System: MALDI-TOF

See attached ID System results document.

Other Features/ Challenges: Results

(1) Oxidase (Kovacs): negative

Metallo-beta-lactamase test: postive

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Quality Control Manager

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Bruker Daltonik MALDI Biotyper Classification Results

Meaning of Score Values

Range	Interpretation	Symbols	Color
2.00 - 3.00	High-confidence identification	(+++)	green
1.70 - 1.99	Low-confidence identification	(+)	yellow
0.00 - 1.69	No Organism Identification Possible	(-)	red

Meaning of Consistency Categories (A - C)

Category	Interpretation
(A)	**High consistency:** The best match is a high-confidence identification. The second-best match is (1) a high-confidence identification in which the species is identical to the best match, (2) a low-confidence identification in which the species or genus is identical to the best match, or (3) a non-identification.
(B)	**Low consistency:** The requirements for high consistency are not met. The best match is a high- or low-confidence identification. The second-best match is (1) a high- or low-confidence identification in which the genus is identical to the best match or (2) a non-identification.
(C)	**No consistency:** The requirements for high or low consistency are not met.

Sample Name: Klebsiella pneumoniae
Sample Description: 01060
Sample ID: 1060-09
Sample Creation Date/Time: 2017-05-03T14:37:20.515 CC
Applied MSP Library(ies): BDAL, Mycobacteria Library (bead method), Filamentous Fungi Library 1.0, Listeria

Sample Name	Sample ID	Organism (best match)	Score Value
B1 (+++) (A)	1060-09	Klebsiella pneumoniae	2.59

Comments:

N/A

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Certificate of Analysis: Lyophilized Microorganism Specification and Performance Upon Release

Specifications	
Microorganism Name: Staphylococcus saprophyticus **Catalog Number:** 0134 **Lot Number:** 134-28 **Reference Number:** ATCC® BAA-750™* **Purity:** Pure **Passage from Reference:** 3	**Expiration Date:** 2018/10/31 **Release Information:** **Quality Control Technologist:** Mary L Bowman **Release Date:** 2016/12/23

Performance

Macroscopic Features: Medium, circular, convex, entire edge, white, glistening	**Medium:** SBAP
Microscopic Features: Gram positive cocci singly, in pairs and tetrads	**Method:** Gram Stain (1)

ID System: Vitek GP (1)	**Other Features/ Challenges: Results**
See attached ID System results document.	(1) Catalase (3% Hydrogen Peroxide): positive (1) Coagulase (rabbit plasma-tube): negative

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Quality Control Manager
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TESTING CERT #2655.01

Microbiologics

bioMerieux Customer: 1012555
System #: 521711

Laboratory Report

Printed Dec 23, 2016 07:33 CST
Printed by: mlb

Isolate Group: 134 28-1

Bench: MB

Card Type: GP Testing Instrument: 000017919CE3 (15220)

Bionumber: 020000014670231
Organism Quantity:

Comments:	

Identification Information	**Card:**	GP	**Lot Number:**	242399410	**Expires:**	Nov 20, 2017 12:00 CST	
	Completed:	Dec 22, 2016 16:32 CST	**Status:**	Final	**Analysis Time:**	5.00 hours	
Selected Organism	99% Probability		**Staphylococcus saprophyticus**				
	Bionumber:	020000014670231			**Confidence:**	Excellent identification	
SRF Organism							
Analysis Organisms and Tests to Separate:							
Analysis Messages:							
Contraindicating Typical Biopattern(s)							

Biochemical Details

2	AMY	-	4	PIPLC	-	5	dXYL	-	8	ADH1	-	9	BGAL	+	11	AGLU	-
13	APPA	-	14	CDEX	-	15	AspA	-	16	BGAR	-	17	AMAN	-	19	PHOS	-
20	LeuA	-	23	ProA	-	24	BGURr	-	25	AGAL	-	26	PyrA	-	27	BGUR	-
28	AlaA	-	29	TyrA	-	30	dSOR	-	31	URE	+	32	POLYB	-	37	dGAL	-
38	dRIB	-	39	ILATk	-	42	LAC	+	44	NAG	-	45	dMAL	+	46	BACI	+
47	NOVO	+	50	NC6.5	+	52	dMAN	+	53	dMNE	-	54	MBdG	-	56	PUL	-
57	dRAF	-	58	O129R	+	59	SAL	-	60	SAC	+	62	dTRE	+	63	ADH2s	-
64	OPTO	+															

Installed VITEK 2 Systems Version: 07.01
MIC Interpretation Guideline:
AES Parameter Set Name:

Therapeutic Interpretation Guideline:
AES Parameter Last Modified:

Microbiologics®

Certificate of Analysis: Lyophilized Microorganism Specification and Performance Upon Release

Specifications	
Microorganism Name: Penicillium chrysogenum **Catalog Number:** 0178 **Lot Number:** 178-20 **Reference Number:** ATCC® 10106™* **Purity:** Pure **Passage from Reference:** 3	**Expiration Date:** 2018/9/30 **Release Information:** **Quality Control Technologist:** Mary L Bowman **Release Date:** 2017/1/23

Performance

Macroscopic Features:	Medium:
Rapidly expanding floccose colonies, initially white, turning dark blue-green with age, exudes bright yellow pigment into medium.	Malt Extract Agar
Microscopic Features:	**Method:**
Hyaline septate mycelia that produce hyaline conidiophores. The conidiophores branch into brush-like penicillus. Spores are borne in long chains from terminal sterigmata.	Lactophenol Blue (1)

ID System: MALDI-TOF

See attached ID System results document.

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Quality Control Manager
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Bruker Daltonik MALDI Biotyper Classification Results

Meaning of Score Values

Range	Interpretation	Symbols	Color
2.00 - 3.00	High-confidence identification	(+++)	green
1.70 - 1.99	Low-confidence identification	(+)	yellow
0.00 - 1.69	No Organism Identification Possible	(-)	red

Meaning of Consistency Categories (A - C)

Category	Interpretation
(A)	**High consistency:** The best match is a high-confidence identification. The second-best match is (1) a high-confidence identification in which the species is identical to the best match, (2) a low-confidence identification in which the species or genus is identical to the best match, or (3) a non-identification.
(B)	**Low consistency:** The requirements for high consistency are not met. The best match is a high- or low-confidence identification. The second-best match is (1) a high- or low-confidence identification in which the genus is identical to the best match or (2) a non-identification.
(C)	**No consistency:** The requirements for high or low consistency are not met.

Sample Name: Penicillium chrysogenum
Sample Description: 0178
Sample ID: 178-20
Sample Creation Date/Time: 2017-01-20T16:41:08.578 CC
Applied MSP Library(ies): BDAL, Mycobacteria Library (bead method), Filamentous Fungi Library 1.0, Listeria

Sample Name	Sample ID	Organism (best match)	Score Value
D2 (+++) (A)	178-20	Penicillium chrysogenum	2.15

Comments:

N/A

Certificate of Analysis: Lyophilized Microorganism Specification and Performance Upon Release

Specifications	
Microorganism Name: Eikenella corrodens **Catalog Number:** 0189 **Lot Number:** 189-31 **Reference Number:** ATCC® BAA-1152™* **Purity:** Pure **Passage from Reference:** 3	**Expiration Date:** 2018/10/31 **Release Information:** **Quality Control Technologist:** Mary L Bowman **Release Date:** 2016/12/19

Performance

Macroscopic Features:	Medium:
Medium, circular, convex, entire edge, light tan, glistening, "bleach-like" odor; as culture ages, margins flatten out.	Chocolate
Microscopic Features:	**Method:**
Gram negative straight rod with rounded ends.	Gram Stain (1)

ID System: Vitek NH (1)

See attached ID System results document.

Other Features/ Challenges: Results

(1) Oxidase (Kovacs): positive
(1) Catalase (3% Hydrogen Peroxide): negative
MacConkey Agar (AER/35 C/48 Hr): no growth

Amanda Kuperus
Quality Control Manager
AUTHORIZED SIGNATURE

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TESTING CERT #2655.01

bioMerieux Customer: 1012555
System #: 521711

Laboratory Report

Printed Dec 16, 2016 09:35 CST
Printed by: mlb

Isolate Group: 189 31-1

Bench: MB

Card Type: NH Testing Instrument: 000017919CE3 (15220)

Bionumber: 0622002400
Organism Quantity:

Comments:	

Identification Information	**Card:**	NH	**Lot Number:**	245369522	**Expires:**	Jan 25, 2017 12:00 CST
	Completed:	Dec 15, 2016 17:31 CST	**Status:**	Final	**Analysis Time:**	6.00 hours
Selected Organism	97% Probability		**Eikenella corrodens**			
	Bionumber:	0622002400			**Confidence:**	Excellent identification
SRF Organism						
Analysis Organisms and Tests to Separate:						
Analysis Messages:						
Contraindicating Typical Biopattern(s)						
Eikenella corrodens	dMLT(16),					

Biochemical Details

1	ArgA	-	2	GGT	-	3	LysA	-	4	dGAL	-	5	LeuA	+	6	ELLM	+
7	PheA	-	8	ProA	+	10	PyrA	-	13	TyrA	-	15	APPA	+	18	dGLU	-
19	GLYG	-	20	dMNE	-	22	dMAL	-	28	SAC	-	33	NAG	-	36	URE	-
39	BGALi	-	40	ODC	+	41	AARA	-	45	PVATE	-	46	PHC	-	47	dMLT	+
51	MTE	-	52	IGLM	-	59	PHOS	-	61	dRIB2	-	62	OPS	-	64	dXYL	-

Installed VITEK 2 Systems Version: 07.01
MIC Interpretation Guideline:
AES Parameter Set Name:

Therapeutic Interpretation Guideline:
AES Parameter Last Modified:

Microbiologics®

Certificate of Analysis: Lyophilized Microorganism Specification and Performance Upon Release

Specifications	
Microorganism Name: Bacillus cereus **Catalog Number:** 0200 **Lot Number:** 200-30 **Reference Number:** ATCC® 14579™* **Purity:** Pure **Passage from Reference:** 3	**Expiration Date:** 2019/7/31 **Release Information:** **Quality Control Technologist:** Mary L Bowman **Release Date:** 2017/8/22

Performance

Macroscopic Features:	**Medium:**
Large, circular to irregular, flat, erose edge, gray, dull, beta hemolytic.	SBAP
Microscopic Features:	**Method:**
Straight, gram positive rod, with an ellipsoidal or spherical, terminal endospore	Gram Stain (1)

ID System: MALDI-TOF

See attached ID System results document.

Other Features/ Challenges: Results

(1) Catalase (3% Hydrogen Peroxide): positive

Parasporal crystals (Phase Contrast Microscopy): not present

Rhizoid colonies: not present

Amanda Kuperus

Quality Control Manager

AUTHORIZED SIGNATURE

Disclaimer: The last digit(s) of the lot number appearing on the product label and packing slip are merely a packaging event number. The lot number displayed on this certificate is the actual base lot number.

Note for Vitek®: Although the Vitek® panel uses many conventional tests, the unique environment of the card, combined with the short incubation period, may produce results that differ from published results obtained by other methods.

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ACCREDITED

TESTING CERT #2655.01

(1) These tests are accredited to ISO/IEC 17025:2005.

Bruker Daltonik MALDI Biotyper Classification Results

Meaning of Score Values

Range	Interpretation	Symbols	Color
2.00 - 3.00	High-confidence identification	(+++)	green
1.70 - 1.99	Low-confidence identification	(+)	yellow
0.00 - 1.69	No Organism Identification Possible	(-)	red

Meaning of Consistency Categories (A - C)

Category	Interpretation
(A)	**High consistency:** The best match is a high-confidence identification. The second-best match is (1) a high-confidence identification in which the species is identical to the best match, (2) a low-confidence identification in which the species or genus is identical to the best match, or (3) a non-identification.
(B)	**Low consistency:** The requirements for high consistency are not met. The best match is a high- or low-confidence identification. The second-best match is (1) a high- or low-confidence identification in which the genus is identical to the best match or (2) a non-identification.
(C)	**No consistency:** The requirements for high or low consistency are not met.

Sample Name: Bacillus cereus
Sample Description: 0200
Sample ID: 200-30
Sample Creation Date/Time: 2017-08-07T10:21:59.155 MB
Applied MSP Library(ies): BDAL, Mycobacteria Library (bead method), Filamentous Fungi Library 1.0, Listeria

Sample Name	Sample ID	Organism (best match)	Score Value
F4 (+++) (A)	200-30	Bacillus cereus	2.10

Comments:

Bacillus anthracis, cereus, mycoides, pseudomycoides, thuringiensis and weihenstephanensis are closely related and members of the Bacillus cereus group. In particular Bacillus cereus spectra are very similar to spectra from Bacillus anthracis. Bacillus anthracis is not included in the MALDI Biotyper database. For differentiation an adequate identification method has to be selected by an experienced professional. The quality of spectra (score) depends on the degree of sporulation: Use fresh material.

Microbiologics®

Certificate of Analysis: Lyophilized Microorganism Specification and Performance Upon Release

Specifications	
Microorganism Name: Shigella sonnei **Catalog Number:** 0303 **Lot Number:** 303-87 **Reference Number:** ATCC® 25931™* **Purity:** Pure **Passage from Reference:** 3	**Expiration Date:** 2019/1/31 **Release Information:** **Quality Control Technologist:** Christine Condon **Release Date:** 2017/3/7

Performance

Macroscopic Features:	Medium:
Two colony types (both are large and can spread); circular, convex, light gray, smooth and other type is irregular, raised, darker gray, and rough, (spreads more than other colony type).	SBAP
Microscopic Features:	**Method:**
Gram negative straight rod	Gram Stain (1)

ID System: Vitek GN (1)

See attached ID System results document.

Other Features/ Challenges: Results

(1) Oxidase (Kovacs): negative

Hektoen Enteric agar: good growth, colorless to blue-green colonies

Shigella Antiserum Poly Group D: positive

(1) Motility B Medium: negative

Amanda Kuperus

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A2LA ACCREDITED

TESTING CERT #2655.01

Microbiologics

bioMerieux Customer: 1012555
System #: 521711

Laboratory Report

Printed Mar 2, 2017 07:56 CST
Printed by: cmc

Isolate Group: 303 87-1

Bench: CC

Card Type: GN Testing Instrument: 000017919CE3 (15220)

Bionumber: 0405610140463610
Organism Quantity:

Comments:	
	GN QC OK

Identification Information	**Card:**	GN	**Lot Number:**	241399320	**Expires:**	Nov 19, 2017 12:00 CST
	Completed:	Feb 28, 2017 21:36 CST	**Status:**	Final	**Analysis Time:**	5.00 hours
Selected Organism	99% Probability		**Shigella sonnei**			
	Bionumber:	0405610140463610			**Confidence:**	Excellent identification

SRF Organism	

Analysis Organisms and Tests to Separate:

Analysis Messages:
Confirm by serological tests

Contraindicating Typical Biopattern(s)

Biochemical Details

2	APPA	-	3	ADO	-	4	PyrA	-	5	lARL	-	7	dCEL	-	9	BGAL	+
10	H2S	-	11	BNAG	-	12	AGLTp	-	13	dGLU	+	14	GGT	-	15	OFF	+
17	BGLU	-	18	dMAL	+	19	dMAN	+	20	dMNE	+	21	BXYL	-	22	BAlap	-
23	ProA	-	26	LIP	-	27	PLE	-	29	TyrA	+	31	URE	-	32	dSOR	-
33	SAC	-	34	dTAG	-	35	dTRE	+	36	CIT	-	37	MNT	-	39	5KG	-
40	lLATk	-	41	AGLU	-	42	SUCT	+	43	NAGA	-	44	AGAL	+	45	PHOS	+
46	GlyA	+	47	ODC	+	48	LDC	-	53	lHISa	-	56	CMT	+	57	BGUR	+
58	O129R	+	59	GGAA	-	61	lMLTa	-	62	ELLM	-	64	lLATa	-			

Installed VITEK 2 Systems Version: 07.01
MIC Interpretation Guideline:
AES Parameter Set Name:

Therapeutic Interpretation Guideline:
AES Parameter Last Modified: