

ABSTRACT

Thermophilic *Campylobacter* species produce the highest numbers of cases of bacteria associated human and animal enteritis annually. At the University of Cape Town an innovative isolation procedure has been used routinely for a number of years which has demonstrated the isolation of a large number of *Campylobacter* species which cannot be isolated using conventional high temperature (42°C) isolation methods. A range of new *Campylobacter* species have been identified as potential emerging pathogens as *C. coli* and *C. jejuni* are the most common species isolated. The alternative isolation procedure utilizes a membrane filtration step, modified hydrogen enriched atmosphere, no antibiotics and incubation at 37°C to isolate these emerging species. To enable large scale surveys to be performed rapidly and to generate a rapid confirmation of an emerging *Campylobacter* species colony has been isolated a new latex reagent has been developed, based on rabbit antisera raised against one of the established species. This new latex reagent, when used in combination with the established latex agglutination kit has been shown to detect all of the *Campylobacter* species, both emerging and established species. This new dual latex combination can therefore be used to facilitate more research into the clinical significance and environmental reservoirs of these emerging *Campylobacter* species. Initial studies using an extensive panel of 55 *Campylobacter* species isolates, comprising 21 different species and sub-species, have confirmed the wide specificity of this dual latex test. This dual latex test is currently being used to detect colonies being isolated from clinical, veterinary, food and environmental samples, including species from a range of our research, veterinary, food and environmental sources.

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PROCEDURE

Quality Control:
 The following check with the Positive Control should be performed each time the kit is used to confirm that the reagents are functioning correctly.
 1. A single 50µL drop of Positive Control should be dispensed on to three adjacent areas on the test slide.
 2. The latex reagent should be added to each of these areas as described in 'Test Procedure' below.
 3. Detentionation of a reagent should be expected if:
 • There is no reaction between the Test Latex Reagents and the Positive Control OR the reaction shows a significant loss of strength with time.
 • The Control Latex Reagent reacts with the Positive Control
 • A latex reagent becomes discoloured or forms lumps which do not disperse with gentle shaking.

Test Procedure:
 1. Bring all reagents to room temperature. Gently shake the latex reagents to ensure a homogeneous suspension.
 2. Dispense 50µL of isotonic saline on to each of three ovals of the agglutination slide.
 3. Add 1 drop (50µL) of Control Latex reagent to each of the three ovals of the agar surface. Mix the bacteria into each of the three drops of isotonic saline on the slide to form even suspensions.
 4. Add 1 drop (50µL) of Control Latex reagent to one of the bacterial suspensions on the slide. Similarly dispense 1 drop (50µL) of the two Test Latex reagents to the other two bacterial suspensions with the latex reagents using a mixing stick, starting with the Control Latex reagent. Spread the mixtures to the edges of the oval areas.
 5. Gently rock the slide, to keep the fluid suspensions in constant movement, for 2 minutes. Observe for agglutination.
 6. Discard the used mixing sticks and slides into a suitable disinfectant.
 7. Repeat the test results (see INTERPRETATION below)

INTERPRETATION
 An agglutination reaction is indicated by visible aggregation of the latex particles. The strength of the reaction may vary, and can be assessed according to the following guidelines.
 + reaction - fine, but readily discernible granularity against a milky background.
 ++ reaction - coarse clumping of particles around the periphery of the test oval, against a clear background.
 +++ reaction - coarse clumping of particles around the periphery of the test oval, against a clear background.

Campycheck. *Campylobacter* results should be interpreted as follows:

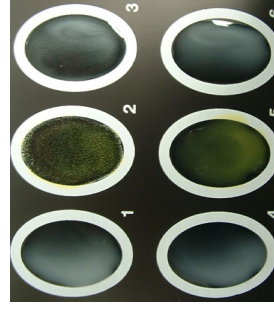
Thermophilic Latex	Species Latex	Control Latex	Interpretation
+	-	-	Thermophilic <i>Campylobacter</i>
+	+	-	Thermophilic <i>Campylobacter</i>
-	+	-	Non-Thermophilic <i>Campylobacter</i>
+ or -	+ or -	-	Auto-agglutination*

*An isolate that causes the Control Latex reagent to agglutinate cannot be tested by Campycheck. *Campylobacter*.

SUMMARY OF RESULTS ON CAMPYLOBACTER ISOLATES (ARS, USDA, Albany, CA)

Bacterial Species	Thermophilic Latex		Campylobacter spp. Latex		Control Latex		Detection Rate
	Positive	Negative	Positive	Negative	Positive	Negative	
<i>Campylobacter jejuni</i>	19	0	19	0	0	19	100%
<i>Campylobacter coli</i>	27	0	27	0	0	27	100%
<i>Campylobacter hyointestinalis</i>	22	0	22	0	0	22	100%
<i>Campylobacter upsaliensis</i>	8	0	8	0	0	8	100%
<i>Campylobacter lari</i>	0	12	12	0	0	12	100%
<i>Salmonella Thompson</i>	0	2	2	0	0	2	100%
<i>Escherichia coli O113</i>	0	1	0	1	0	1	0%
<i>Escherichia coli O128</i>	0	1	0	1	0	1	0%

LATEX AGGLUTINATION



PRODUCT PRESENTATION

THE CAMPYCHECK STRAIN SET

CAMPY CHECK	TAXON	STRAIN NUMBER	NOTES	SOURCE (clinical/geographic)
CC 1	<i>C. coli</i>	RM 2228	genome sequenced	Chicken USA
CC 2	<i>C. coli</i>	LM 33	hyolite genotype	Pig/DK
CC 3	<i>C. coli</i>	306.97		Human, diarrhoea/SA
CC 4	<i>C. concisus</i>	COUG 13144	type strain	Human, oral/USA
CC 5	<i>C. concisus</i>	COUG 19695	genotype 2	Human, diarrhoea/Belgium
CC 6	<i>C. concisus</i>	Liatovicia 336/86	genotype 1	Human diarrhoea/vomiting/SA
CC 7	<i>C. curvus</i>	SSI 18286	genotype 1	Human, oral/USA
CC 8	<i>C. curvus</i>	Liatovicia 526/92	genotype 2	Human, oral/SA
CC 9	<i>C. curvus</i>	Liatovicia 13A	genotype 3	Human, oral/SA
CC 10	<i>C. fetus subsp. fetus</i>	COUG32114		Human abortion, Sweden
CC 11	<i>C. fetus</i>	Ardh 1076		Human, loose stools/SA
CC 12	<i>C. fetus</i>	COUG 11287		Human, loose stools/SA
CC 13	<i>C. fetus subsp. venerealis</i>	Arm 4442		Human/France
CC 14	<i>C. fetus subsp. venerealis</i>	Arm 4442		Bovine/Australia
CC 15	<i>C. fetus subsp. venerealis</i>	Arm 516		Human, loose stools/SA
CC 16	<i>C. gracilis</i>	COUG 27220		Revised bovine fetus/N. Ireland, UK
CC 17	<i>C. gracilis</i>	COUG 13143		Human, periodontitis (USA)
CC 18	<i>C. helveticus</i>	COUG 30366		Human, periodontitis (USA)
CC 19	<i>C. helveticus</i>	COUG 34016		Cat diarrhoea/Switzerland
CC 20	<i>C. hominis</i>	NCTC CH001	type strain	Human faeces/UK
CC 21	<i>C. hyointestinalis subsp. hyointestinalis</i>	NCTC 9603	type strain	Human faeces/UK
CC 22	<i>C. hyointestinalis subsp. hyointestinalis</i>	LMG 6280	genotype 1	Human faeces, Belgium
CC 23	<i>C. hyointestinalis subsp. hyointestinalis</i>	LMG 6280	genotype 2	Human faeces, Belgium
CC 24	<i>C. hyointestinalis subsp. hyointestinalis</i>	176.96	genotype 3	Human neonate, diarrhoea/SA
CC 25	<i>C. hyointestinalis subsp. hyointestinalis</i>	234.95	genotype 4	Human infant, loose stools/SA
CC 26	<i>C. hyointestinalis subsp. bawneyi</i>	CHY 5	type strain	Pig stomach/UK
CC 27	<i>C. hyointestinalis subsp. bawneyi</i>	COUG 27651	genotype 0:17	Human/DK
CC 28	<i>C. jejuni subsp. jejuni</i>	SS 3254	serotype 0:17	Human faeces/Boggy/Germany
CC 29	<i>C. jejuni subsp. jejuni</i>	299.97		Human blood, Kwazulu/Natal/SA
CC 30	<i>C. jejuni subsp. jejuni</i>	NCTC 11668	genome sequenced	Human diarrhoea/UK
CC 31	<i>C. jejuni subsp. jejuni</i>	RM 1221	genome sequenced	Chicken/USA
CC 32	<i>C. jejuni subsp. jejuni</i>	SUS 4633	genome sequenced	Cattle/DK
CC 33	<i>C. jejuni subsp. jejuni</i>	47.37		Human faeces/USA
CC 34	<i>C. jejuni subsp. jejuni</i>	47.37		Infant, faeces/USA
CC 35	<i>C. jejuni subsp. jejuni</i>	NCTC 13004		Human faeces/UK
CC 36	<i>C. lariense</i>	DARDNI 6718D		Pig faeces/N. Ireland, UK
CC 37	<i>C. lari</i>	RM 2100 (NARTCO),		Human faeces/USA
CC 38	<i>C. lari</i>	COUG 22395	genome sequenced	Human diarrhoea/France
CC 39	<i>C. lari</i>	(LPTC)		Human diarrhoea/France
CC 40	<i>C. mucosalis</i>	COUG 1859	serotype A	Pig neonatal faeces/Scotland, UK
CC 41	<i>C. mucosalis</i>	COUG 23201	serotype B	Pig neonatal faeces/Scotland, UK
CC 42	<i>C. rectus</i>	COUG 20446	type strain	Human oral/USA
CC 43	<i>C. rectus</i>	COUG 11645	type strain	Human periodontitis/Sweden
CC 44	<i>C. shouboi</i>	COUG 30254	type strain	Human, oral/Japan
CC 45	<i>C. shouboi</i>	Liatovicia 86/92	bv. agglutinum	Infant diarrhoea/SA
CC 46	<i>C. shouboi</i>	LMG 11794	bv. pararegularis	Human faeces/Canada
CC 47	<i>C. shouboi</i>	90.37.2703	bv. rectalis	Human faeces/UK
CC 48	<i>C. upsaliensis</i>	COUG 19659	genome sequenced	Human faeces/USA
CC 49	<i>C. upsaliensis</i>	COUG 19607		Human faeces/Sweden
CC 50	<i>C. upsaliensis</i>	COUG 18470		Human urine/Sweden
CC 51	<i>C. ureolyticus</i>	Regahospital 98/80		Human faeces/DK
CC 52	<i>C. ureolyticus</i>	SS 71032		Human faeces/DK
CC 53	<i>C. ureolyticus</i>	192.07		Human faeces/DK
CC 54	<i>A. butzeri</i>	SS 71032		Human faeces/UK
CC 55	<i>A. butzeri</i>	192.07		Human faeces/UK
CC 56	<i>A. cyaneogriseus</i>	COUG 171801	type strain: subgroup 1	Human faeces/UK
CC 57	<i>A. cyaneogriseus</i>	SSI 70952		Human faeces/UK
CC 58	<i>A. cyaneogriseus</i>	COUG 171801		Human faeces/DK
CC 59	<i>A. sibirici</i>	BU 30CC 881		Poultry/UK

RESULTS ON CAMPYCHECK STRAIN SET (TEAGASC, EIRE)

Campycheck Code	Species	Original Microgen Latex	New Microgen Latex	Comments
CC1	<i>C. coli</i>	+	+	
CC2	<i>C. coli</i>	+	+	
CC3	<i>C. coli</i>	+	+	
CC4	<i>C. concisus</i>	+	+	
CC5	<i>C. concisus</i>	+	+	
CC6	<i>C. concisus</i>	+	+	
CC7	<i>C. curvus</i>	+	+	
CC8	<i>C. curvus</i>	+	+	
CC9	<i>C. curvus</i>	+	+	
CC10	<i>C. fetus</i>	+	+	
CC11	<i>C. fetus</i>	+	+	
CC12	<i>C. fetus</i>	+	+	
CC13	<i>C. fetus</i>	+	+	
CC14	<i>C. fetus</i>	+	+	
CC15	<i>C. fetus</i>	+	+	
CC16	<i>C. fetus</i>	+	+	
CC17	<i>C. fetus</i>	+	+	
CC18	<i>C. fetus</i>	+	+	
CC19	<i>C. fetus</i>	+	+	
CC20	<i>C. helveticus</i>	+	+	
CC21	<i>C. hominis</i>	+	+	Mixed by original Latex
CC22	<i>C. hyointestinalis</i>	+	+	Mixed by original Latex
CC23	<i>C. hyointestinalis</i>	+	+	Mixed by original Latex
CC24	<i>C. hyointestinalis</i>	+	+	
CC25	<i>C. hyointestinalis</i>	+	+	
CC26	<i>C. hyointestinalis</i>	+	+	
CC27	<i>C. hyointestinalis</i>	+	+	
CC28	<i>C. hyointestinalis</i>	+	+	
CC29	<i>C. jejuni</i>	+	+	
CC30	<i>C. jejuni</i>	+	+	
CC31	<i>C. jejuni</i>	+	+	
CC32	<i>C. jejuni</i>	+	+	
CC33	<i>C. jejuni</i>	+	+	
CC34	<i>C. jejuni</i>	+	+	
CC35	<i>C. jejuni</i>	+	+	
CC36	<i>C. lariense</i>	+	+	
CC37	<i>C. lari</i>	+	+	
CC38	<i>C. lari</i>	+	+	
CC39	<i>C. lari</i>	+	+	
CC40	<i>C. mucosalis</i>	+	+	
CC41	<i>C. mucosalis</i>	+	+	
CC42	<i>C. rectus</i>	+	+	Mixed by original Latex
CC43	<i>C. rectus</i>	+	+	Mixed by original Latex
CC44	<i>C. shouboi</i>	+	+	Mixed by original Latex
CC45	<i>C. shouboi</i>	+	+	Mixed by original Latex
CC46	<i>C. shouboi</i>	+	+	Mixed by original Latex
CC47	<i>C. shouboi</i>	+	+	Mixed by original Latex
CC48	<i>C. shouboi</i>	+	+	Mixed by original Latex
CC49	<i>C. upsaliensis</i>	+	+	
CC50	<i>C. upsaliensis</i>	+	+	
CC51	<i>C. upsaliensis</i>	+	+	
CC52	<i>B. ureolyticus</i>	+	+	Mixed by both Latexes
CC53	<i>B. ureolyticus</i>	+	+	
CC54	<i>A. butzeri</i>	+	+	
CC55	<i>A. butzeri</i>	+	+	
CC56	<i>A. cyaneogriseus</i>	+	+	
CC57	<i>A. cyaneogriseus</i>	+	+	Mixed by both Latexes
CC58	<i>A. cyaneogriseus</i>	+	+	
CC59	<i>A. sibirici</i>	+	+	

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