

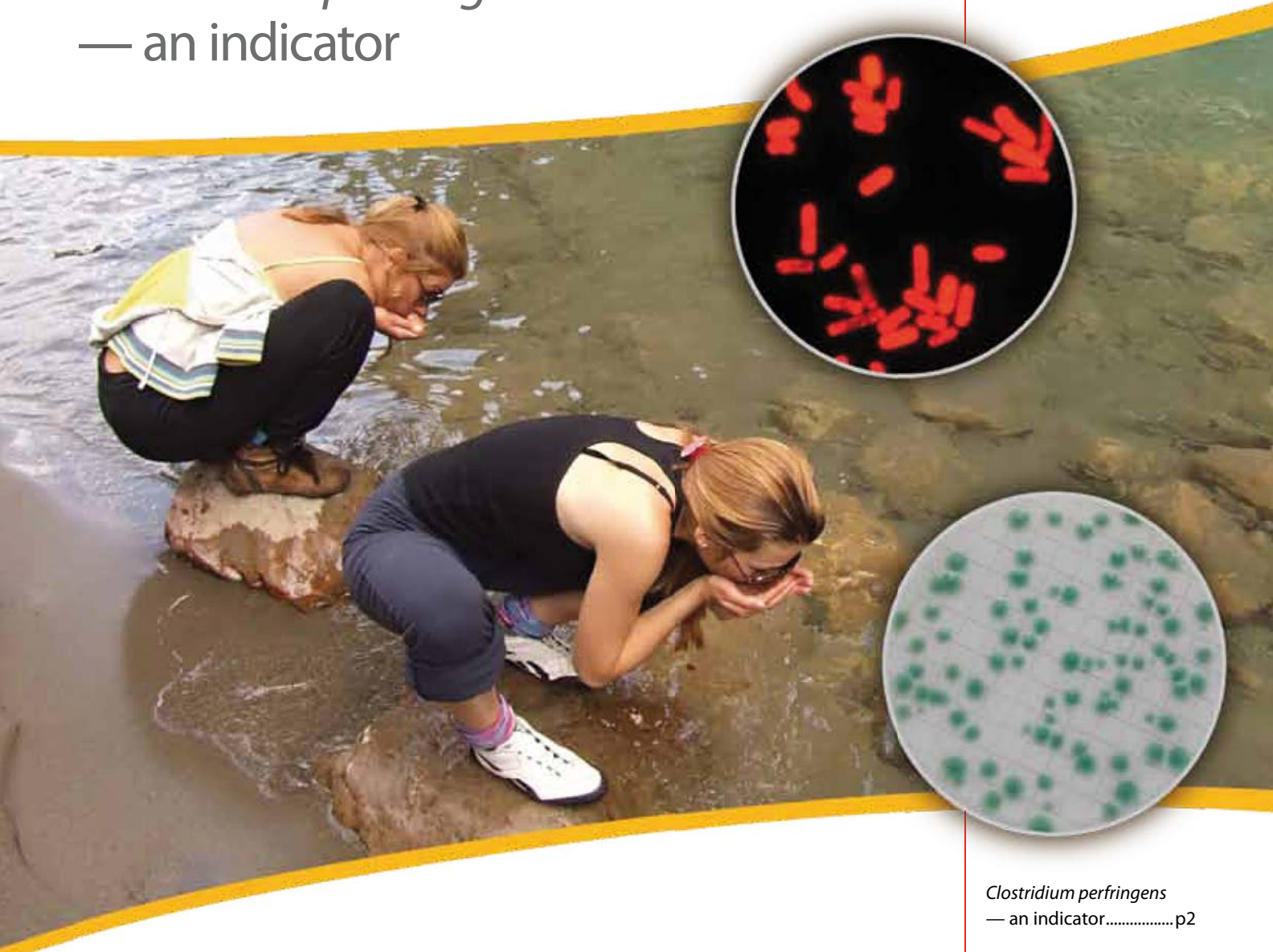


Microbiology Focus

Volume 3.1, 2011



Clostridium perfringens — an indicator



Surface water a source of *Clostridium perfringens*
(source *C. perfringens* FISH image: BioVisible BV, The Netherlands)

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— an indicator.....p2

Burkholderia cepacia
complex: multilingual
bacteriap6



Clostridium perfringens — an Indicator

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Clostridium perfringens is an effective indicator of faecal contamination. Unlike *E. coli*, *C. perfringens* can withstand processing treatments and unfavorable growth conditions.

Clostridium perfringens is found in undercooked or improperly sterilized canned foods (germination of endospores) and in water (surface water). The natural contamination source is human and animal faeces transmitted into food products primarily by water. *C. perfringens* produces an extensive range of invasins and exotoxins. The enterotoxins cause the undesirable, predominantly meat-associated, food poisoning, as well as wound and surgical infections that lead to gas gangrene.

C. perfringens plays a subsidiary role in water examination [6]. *Clostridia* are spore builders and are resistant to heating, chlorination and other stress factors. In contrast to vegetative cells like coliforms (*E. coli*, enterococci), which are less resistant, *C. perfringens* has the advantage of a robust capability to survive [8]. Therefore, while faecal contamination is detected primarily by coliforms as an indicator, which could disappear after a processing step, *C. perfringens* remains present. The organism is not a hazard in water; rather, it is problematic when the water comes in contact with food.

In consideration of the aforementioned facts, it is obvious that detection and identification of *C. perfringens* is an important step toward the control and eradication of this potent pathogen. *C. perfringens* is an anaerobic, Gram-positive, spore-forming rod-shaped (see Figure 7, pg 4) bacterium [3, 5]. Some characteristic enzymes of *C. perfringens* are hemolysins (β -hemolysis), lecithinase, extracellular proteases, lipases (phospholipase-C), phosphatase, collagenase, hyaluronidase, saccharolytic enzymes and enzymes to reduce sulphite to sulphide. These enzymes are also used as detection and differentiation targets. It is also notable that *C. perfringens* is a non-motile bacterium, and it is the most important of the sulphite reducing clostridia [5].

Also, *C. perfringens* normally grows at 44 °C, whereas some other clostridia are inhibited at this temperature. This property is used in ISO methods to give the medium more selectivity [4].

Early detection of *Clostridium* in food is important to control outbreaks. To facilitate detection, Fluka has introduced a new chromogenic media, CP *ChromoSelect* Agar, for enumeration and differentiation of *Clostridium sp.*, in particular *Clostridium perfringens*, in aqueous samples (Figure 1).

For the standard detection of *C. perfringens*, mCP and TSC agar have been recommended [6, 7]. However, there are problems associated with each of these media. CP *ChromoSelect* Agar is more reliable and easier to handle than m-CP and TSC agars. The color does not diffuse in the agar and confirmation is not required since the green coloration is specific for *C. perfringens* [9]. CP *ChromoSelect* Agar avoids the disadvantages of m-CP agar, including the presence of ammonia, which prevents subculturing the *C. perfringens* colonies, the overly-selective nature of m-CP agar, and the evanescence of the red color of colonies after the addition of ammonia, which makes further confirmation impossible. In addition, the recovery of *C. perfringens* was rejected by ISO in favor of methods based on TSC agar [1, 8]. CP *ChromoSelect* Agar also eliminates the excessive and variable blackening of the peripheral colonies encountered with TSC agar, which makes colony counting at lower dilutions difficult and leads to false positives (Figure 2). It is also more reliable at high bacteria counts, where the TSC agar can produce false negatives because of an interference with the other enzymatic mechanisms from acid production and oxygen contact (Figure 3). TSC detects all

Figure 1: Drinking water sample cultured on CP *ChromoSelect* Agar. *C. perfringens* appears as distinct green colonies.

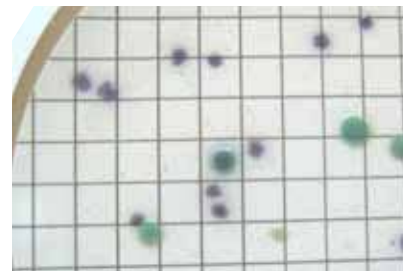


Figure 2: Comparison of m-CP, TSC and CP *ChromoSelect* Agar (from left to right)



Figure 3: *C. perfringens* ATCC 10873 on CP *ChromoSelect* Agar (above) and TSC agar (below). Note the false negatives on the TSC agar.

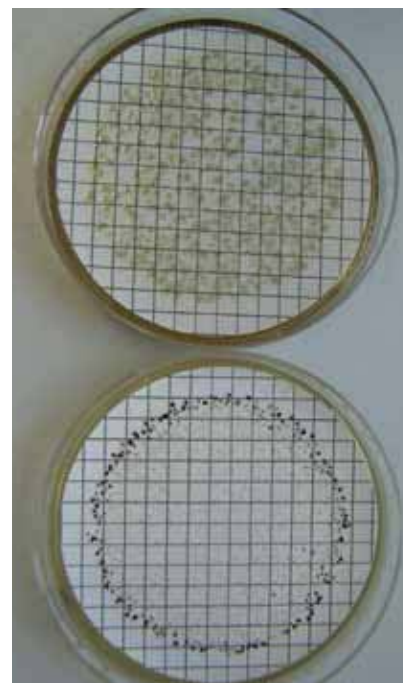




Figure 4: TSC agar + fluorogenic substrate = TSCF agar

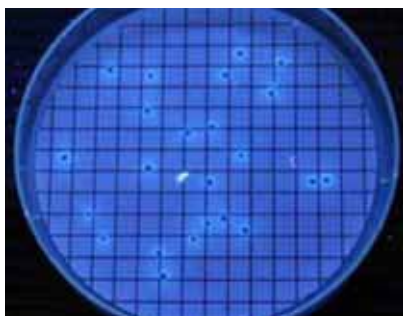
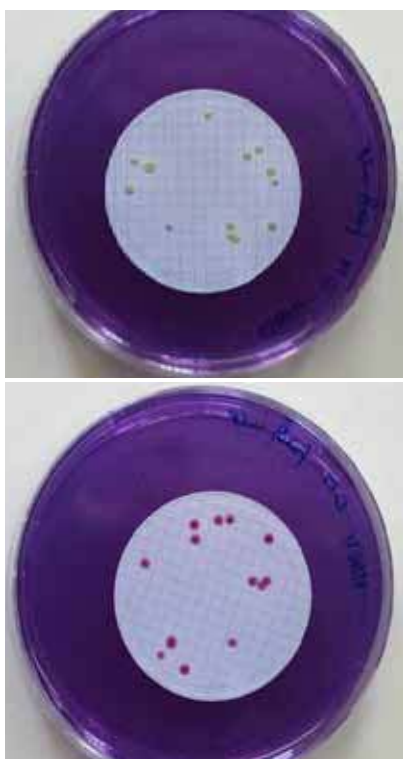


Figure 5: *C. perfringens* on mCP Agar before (above) and after (below) ammonia hydroxide vapors



sulphite-reducing clostridia, however, and not only *C. perfringens*. TSC agar can be improved with the addition of MUP, a fluorogenic substrate which helps to confirm the presence of phosphatase. This modification is designated as TSCF (Figure 4).

In the present study, three media types (mCP [Figure 5], TSCF and CP ChromoSelect Agar) were evaluated for recovery of *C. perfringens* in various surface water samples. Using a membrane filtration technique on 139 water samples, 131 samples (94.2%) were found to be presumably positive for *C. perfringens* in at least one of the culture media. Green colored colonies on CP ChromoSelect agar (Figure 1) were counted as presumably *C. perfringens* isolates.

Out of 483 green colonies on CP ChromoSelect Agar, 96.9 % (465 strains, indole negative) were identified as *C. perfringens*, 15 strains (3.1%) were indole positive and were identified as *C. sordelli*, *C. bifermentans* or *C. tetani*. Only 3 strains (0.6 %) gave false positive results and were identified as *C. fallax*, *C. botulinum*, and *C. tertium* (Table 1). Variance analysis of the data obtained shows no statistically significant differences in the counts obtained between media employed in this work (Figure 6 and Table 2).

Table 1: All green colonies isolated from CP ChromoSelect Agar and identified with API system (n=483)

strains	Indole reaction	n
<i>C. perfringens</i>	-	465 (96.3%)
<i>C. tertium</i>	-	1 (0.2%)
<i>C. botulinum</i>	-	1 (0.2%)
<i>C. fallax</i>	-	1 (0.2%)
<i>C. bifermentans</i>	+	2 (0.4%)
<i>C. sordelli</i>	+	12 (2.5%)
<i>C. tetani</i>	+	1 (0.2%)

Figure 6: Comparison between TSCF agar and the mCP and CPC (CP ChromoSelect Agar) media for enumerating strains of *C. perfringens* in water samples.

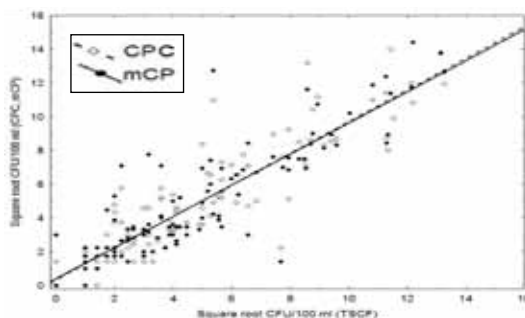


Table 2: Analysis of variance of *C. perfringens* spore counts obtained on all media

Effect	Sum of squares	df	Mean square	F ratio	p value
Media	0.392	2	0.196	0.190	0.827
Residual	235.578	228	1.033		

In general, the identification of typical and atypical colonies isolated from all media demonstrated that CP *ChromoSelect* Agar was the most useful medium for *C. perfringens* recovery in water samples [2].

In addition to its advantages over m-CP and TSC agars, CP *ChromoSelect* Agar is an ideal growth media. It contains only vegetable peptones and, together with yeast extract, it is an excellent source of nitrogen, carbon, amino acids and vitamin B complex. Sucrose acts as the fermentable carbohydrate and reducing agents lower the redox potential of the media. Diverse salts provide the required ions for enzymatic reactions. Buffering agents stabilize the pH within the ideal growth range. Inhibitors D-cycloserine and polymyxin B give the medium its selectivity, while further selectivity is achieved by incubation under anaerobic conditions at 44 °C. Various promoters and substrates protect injured cells to improve recovery rate and enhance growth. The chromogenic enzyme substrates in the CP *ChromoSelect* Agar provide the differentiation, particularly for *C. perfringens* (Table 3). A negative indol reaction (Kovac's Reagent) confirms the presence of *C. perfringens*.

Table 3: *Clostridium* sp. cultural characteristics in CP *ChromoSelect* Agar

Organisms (ATCC)	Growth	Colony Appearance
<i>Clostridium perfringens</i> (13124)	+++	Green
<i>Clostridium bifementans</i> (638)	+++*	Dark blue with violet halo
<i>Clostridium sporogenes</i> (8534)	-	-
<i>Clostridium sordelli</i> (9714)	++	Dark green with halo (change to red with Kovac's Reagent)
<i>Enterococcus faecalis</i> (29212)	++	Violet
<i>Escherichia coli</i> (25922)	-	-
<i>Pseudomonas aeruginosa</i> (27853)	-	Colorless
<i>Staphylococcus aureus</i> (25923)	-	-
<i>Bacillus subtilis</i> (6051)	-	-
<i>Salmonella typhimurium</i> (DSM 554)	++	Violet

* Growth at 40 °C, but no growth at 44 °C

Table 4: Typical Media for *Clostridium perfringens*

Nonselective Media	Cat. No.	Description
AEA Sporulation Broth (Base), modified Supplements: 4 g/l raffinose (Fluka 83400) + 930 mg/l sodium carbonate (Fluka 71351) + 42.6 mg/l cobalt chloride (Sigma-Aldrich 60818) + 20 mg/l sodium ascorbate (Sigma 11140)	17170	For early sporulation of <i>Clostridium perfringens</i> from foods.
Brewer thioglycollate medium	B2551	Used for testing the sterility of biological products.
Clostridial Nutrient Medium*	27546	For the cultivation and enumeration of clostridia and other anaerobes as well as facultative microorganisms in food, clinical and other material acc. to Hirsch & Grinsted.
Cooked Meat Broth	60865	For the primary cultivation of aerobic, microaerophilic and anaerobic bacteria from clinical specimens.
Reinforced Clostridial Agar	91365	Used for the cultivation and enumeration of Clostridia.
Nonselective Differential Media	Cat. No.	Description
Clostridial Differential Broth*	27544	For the MPN enumeration of all clostridia in food and other material acc. to Gibbs and Freame.
Gelatin Iron Agar	G0289	Used for detecting gelatin liquefaction and hydrogen sulphide production.
Lactose Gelatin Medium (Base)* Supplement: 120 g/l gelatine* 180 Bloom (Fluka 48722)	61348	For the detection of lactose and gelatine metabolizing microorganisms (<i>Cl. perfringens</i>).
Meat Liver Agar	46379	For the cultivation of anaerobic microorganisms.
Motility Nitrate Medium* Supplement: 5 ml/l glycerol (Sigma 49767)	14305	Selective medium for motile nitrate-utilizing microorganisms (<i>Ps. aeruginosa</i> , <i>Cl. perfringens</i>).
Nutrient Gelatin	70151	Nutrient Gelatin is recommended for the determination of gelatin-liquefying microorganisms as well as for the enumeration of proteolytic organisms in water by the plate count test.
Sulfite Iron Agar* Supplement: 1.4 g/l ferrous sulfate (Fluka 44980)	86128	For the detection and enumeration of <i>Clostridium</i> species in meat and meat products.

Did you know...

Clostridium perfringens has one of the highest reproduction rates in the bacteria kingdom. Under optimal conditions, the cell doubles every 10 minutes!

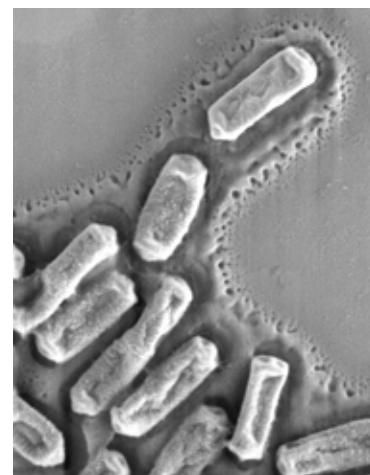


Figure 7: Scanning electron micrograph of *C. perfringens* grown on a silicon wafer (source: S. Melville, Department of Biological Sciences, Virginia Tech University)



Table 4: Typical Media for Clostridium perfringens (Contd.)

Selective Differential Media	Cat. No.	Description
CP ChromoSelect Agar Supplements: 2 vials/l M-CP selective Supplement I (Fluka 51962) or 2 vial/l of Perfringens T.S.C. Supplement (P9352)	12398	A chromogenic Agar for enumeration detection and differentiation of <i>Clostridium sp.</i> , in particular <i>Clostridium perfringens</i> , in aqueous samples.
m-CP Agar Base Supplements: 2 vial/l m-CP Selective Supplement I (Fluka 51962) + 2 vial/l m-CP Selective Supplement II (Fluka 82265)	75605	Recommended by the Directive of the Council of the European Union 98/83/EC for isolation and enumeration of <i>Clostridium perfringens</i> from water sample using membrane filtration technique.
Perfringens Agar Base Supplements: 50 ml/l Egg Yolk Emulsion (Fluka 17148) + 2 vial/l of Perfringens S.F.P. Supplement (53436) or 2 vial/l of Perfringens T.S.C. Supplement (P9352)	39727	Used for the presumptive identification and enumeration of <i>Clostridium perfringens</i> from food.
SPS Agar*	85627	Medium for detection, isolation and enumeration of <i>Clostridium perfringens</i> and <i>Clostridium botulinum</i> in food acc. to Angelotti et al. (1962).
SPS Agar, modified	17231	For the selective isolation and enumeration of <i>Clostridium perfringens</i> from foods.
TSC Agar* Supplements: TSC Agar: 400 mg/l D-cycloserine (Sigma 30020) or 2 vial/l Perfringens T.S.C. Supplement (Fluka P9352) and 50 ml/l Egg Yolk Emulsion (Fluka 17148) TSC Agar egg yolk free: 400 mg/l D-cycloserine (Fluka 30020) or 2 vial/l Perfringens T.S.C. Supplement (Sigma/Fluka P9352) TSC Fluorescence Agar: 5.5 g/l rehydrated TSC Agar supplement (Fluka 80548) SFP Agar (Shahidi-Ferguson Perfringens Agar): 2 vial/l of Perfringens S.F.P. Supplement (Fluka P9477) and 50ml/l Egg Yolk Emulsion (Fluka 17148)	93745	For the isolation and enumeration of vegetative forms as well as spores from <i>Clostridium perfringens</i> in food, clinical specimens and other material acc.to Harmon et al. (1971).
TSN Agar*	93735	Highly selective medium for the detection and enumeration of <i>Clostridium perfringens</i> in food and other material.

* not sold in USA

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The *Burkholderia cepacia* Complex: A Group of Multilingual Bacteria

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In the world of microorganisms, some members of the genus *Burkholderia* are multilingual, utilizing at least three different communication systems.

Over the past 20 years, it has become evident that bacteria exist not only as individual cells; in addition, they often coordinate their activities and act in a concerted manner similar to that of multicellular organisms. Interactions between cells are brought about by cell-to-cell communication systems that depend on the production and detection of small diffusible signal molecules. This phenomenon, which has been termed 'quorum sensing' (QS) [1], provides bacteria with a regulatory mechanism to express certain phenotypic traits in a population density-dependent manner. To date, various signal molecules belonging to different chemical classes have been described.

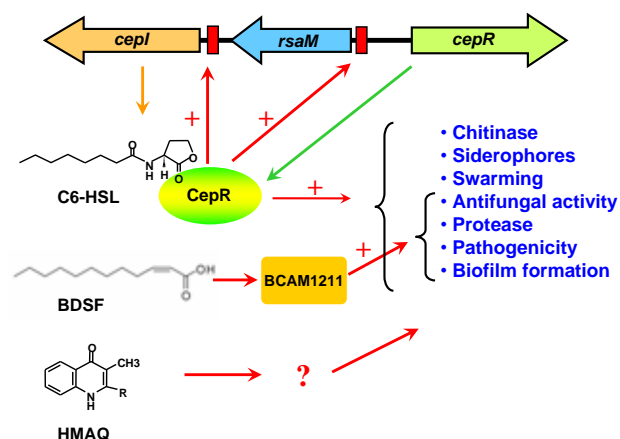
Some bacteria utilize more than just one communication system and therefore can be considered multilingual. One well-investigated example are the bacteria belonging to the *Burkholderia cepacia* complex (Bcc). The Bcc currently comprises a group of 17 formally-named bacterial species that, although closely related, are phenotypically diverse. Strains of the Bcc are ubiquitously distributed in nature and have been isolated from soil, water, the rhizosphere of plants, industrial settings, hospital environments, and from infected humans. Some Bcc strains have emerged as problematic opportunistic pathogens in patients with cystic fibrosis as well as in immunocompromised individuals [2].

All Bcc species investigated so far produce N-octanoylhomoserine lactone (C8-HSL), a member of the N-acyl homoserine lactone (AHL) signal molecule family, by the aid of the AHL synthase Ceph. As the cell density increases, C8-HSL accumulates in the growth medium until a critical threshold concentration is attained. At this point, C8-HSL binds to its cognate LuxR-type receptor protein CepR, which, in turn, leads to the induction or repression of target genes (Figure 1). Previous work has identified several QS-regulated functions in strains of the genus *Burkholderia*, including the production of extracellular proteases, chitinases, siderophores and the fungicide pyrrolnitrin, swarming motility, biofilm formation and pathogenicity [3]. Moreover, some Bcc strains harbor additional AHL-based QS systems that utilize different AHL species to either fine-tune the CepR QS system or regulate separate sets of genes.

In addition to AHLs, Bcc strains were demonstrated to produce diffusible signal factors (DSFs). While most strains synthesize cis-2-dodecenoic acid, which was named the *Burkholderia* diffusible signal factor (BDSF), some Bcc species also synthesize other DSF family molecules. Subsequent studies have shown that BDSF plays a role in the regulation of bacterial virulence and biofilm formation [4]. BDSF is synthesized by the gene product of Bcam0581 and appears to be sensed by the histidine sensor kinase BCAM0227. Interestingly, the AHL- and BDSF-dependent QS systems appear to act in conjunction in the regulation of *B. cenocepacia* virulence, as a set of the AHL-controlled virulence genes is also positively regulated by BDSF [4, 5]. Furthermore, mutation of Bcam0581 results in substantially retarded energy production and exhibits impaired growth in minimal medium, highlighting the dual roles of the BDSF QS system in physiology and infection.

Finally, Bcc strains were also shown to produce a third class of signal molecules, namely 4-hydroxy-2-alkylquinolines (HAQs). Recent work has shown that *Burkholderia ambifaria* (a member of the Bcc) HAQs contain predominantly an unsaturated aliphatic side chain and are typically methylated (at the 3 position), hence their designation as 4-hydroxy-3-methyl-2-alkylquinolines (HMAQs) [6]. Interestingly, inactivation of HAQs' production of *B. ambifaria* has led to an increased AHL production and, consequently, affected phenotypes that are under the control of QS in *B. ambifaria*, namely antifungal activity, siderophore production, and proteolytic activity.

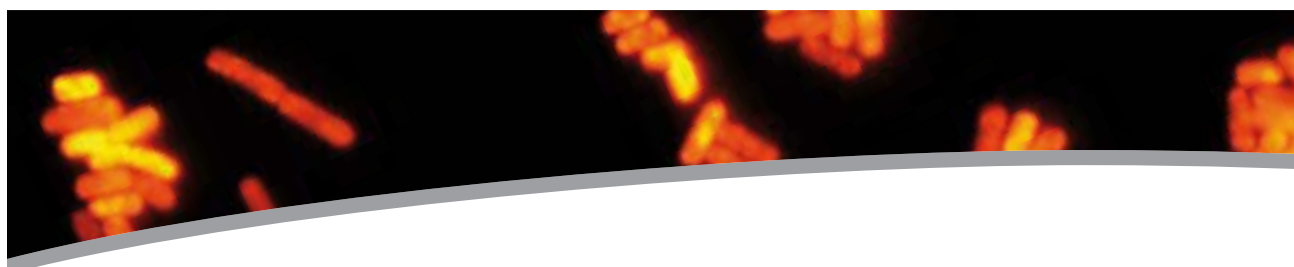
Figure 1: Bcc strains can utilize up to three different cell-to-cell communication systems that rely on different classes of signal molecules: N-acyl-homoserine lactones (particularly C8-HSL), cis-2-dodecenoic acid, the *Burkholderia* diffusible signal factor (BDSF), and 4-hydroxy-3-methyl-2-alkylquinolines (HMAQs).



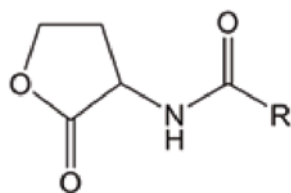
At present, it is unclear why Bcc strains synthesize multiple signal molecules. However, it should be kept in mind that these bacteria inhabit various habitats and thus may encounter a large diversity of bacteria. It is therefore tempting to speculate that the utilization of different communication systems will enable the bacteria to interact with a wide range of indigenous bacteria through chemical cross-talk. These interspecies interactions may be important for the fitness of the bacteria in their natural environment.

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Quorum Sensing Substances

Cat. No.	Signalling Compounds	Substance
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10942	C4-HCT	N-Butyryl-DL-homocysteine thiolactone
09945	C4-HSL	N-Butyryl-DL-homoserine lactone
09926	C6-HSL	N-Hexanoyl-DL-homoserine lactone
10939	C7-HSL	N-Heptanoyl-DL-homoserine lactone
10940	C8-HSL	N-Octanoyl-DL-homoserine lactone
17248	C10-HSL	N-Decanoyl-DL-homoserine lactone
17247	C12-HSL	N-Dodecanoyl-DL-homoserine lactone
10937	C14-HSL	N-Tetradecanoyl-DL-homoserine lactone
42052	DSF	cis-11-Methyl-2-dodecanoic acid
53727	3-Hydroxy-C12-HSL	N-(3-Hydroxydodecanoyl)-DL-homoserine lactone
51481	3-Hydroxy-C14-HSL	N-(3-Hydroxytetradecanoyl)-DL-homoserine lactone
K3255	3-Oxo-C6-HSL	N-(3-Oxohexanoyl)-DL-homoserine lactone
K3007	3-Oxo-C6-HSL	N-(3-Oxohexanoyl)-DL-homoserine lactone
O1639	3-Oxo-C8-HSL	N-(3-Oxooctanoyl)-DL-homoserine lactone
O1764	3-Oxo-C8-HSL	N-(3-Oxooctanoyl)-L-homoserine lactone
O9014	3-Oxo-C10-HSL	N-(3-Oxodecanoyl)-L-homoserine lactone
O9139	3-Oxo-C12-HSL	N-(3-Oxododecanoyl)-L-homoserine lactone
O9264	3-Oxo-C14-HSL	N-(3-Oxotetradecanoyl)-L-homoserine lactone
07077	pC-HSL	N-(p-Coumaroyl)-L-homoserine lactone
94398	PQS (Pseudomonas quinolone signal)	2-Heptyl-3-hydroxy-4(1H)-quinolone
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