



Interpretation of Positive Blood Cultures When PCR Blood Culture Identification (BCID) Results are "Not Detected"

Nebraska Medicine currently uses a multi-plex PCR-based blood culture identification (BCID) system that is able to identify 19 potential pathogens growing in blood culture. BCID generally detects over 90% of the most common causative agents in bloodstream infections; however, when microbes not included on the panel are present in a blood culture, it returns a result of "Not Detected." This document aims to provide guidance in these scenarios supported by data collected at Nebraska Medicine from January 2018 to August 2019.

Table 1: Recommendations for treatment of patients with blood cultures growing organisms not detected on BCID

Gram Stain/Preliminary Culture Result	Likely Organism (% total BCID negative)*	Recommended Treatment
Gram-positive:		
Aerobe (most can also grow in anaerobic bottles)	Micrococcus sp. (18.1%) Coagulase-negative Staphylococcus (9.3%) Diphtheroids (7%)	None
Anaerobe bottle only	Peptostreptococcus sp. (4.4%) Lactobacillus sp. (2.6%) Clostridium sp. (2.6%)	None If therapy is desired: Metronidazole 500 mg PO q8h OR Penicillin G 4 million units IV q4h
Gram-negative:		
Aerobe (most can also grow in anaerobic bottles)	Acinetobacter sp. (1.8%) Stenotrophomonas maltophilia (1.6%) Pseudomonas fluorescens-putida group (1%)	Levofloxacin 750 mg IV/PO q24h
Anaerobe bottle only	Bacteroides fragilis group (9.3%) Fusobacterium sp. (4.7%)	Metronidazole 500 mg IV/PO q8h

^{*}A full list of isolated organisms can be found below in Table 2

Orange text = Cocci, Blue text = Bacilli (rods)

Gram-Positives

When BCID results as "Not Detected" but there is microbial growth, the organism is most frequently gram-positive (71%). Of the gram-positive results, the most common species isolated were *Micrococcus* sp., coagulase-negative *Staphylococcus*, and Diphtheroids (combined 48.5%). These organisms can grow in either aerobic or anaerobic bottles. They are usually considered contaminants and do not require treatment, but clinical judgment should be used. Instances in which these cultures may warrant treatment are when there are 2 out of 2 blood cultures positive or when the patient has a documented history of infection with the organism. Typically when multiple cultures are positive with these organisms, the cause is device-related and the primary treatment is removal of the device, when possible.

Anaerobic gram-positive organisms are more rarely isolated, with the most common being *Peptostreptococcus* sp., *Lactobacillus* sp., and *Clostridium* sp. These organisms will grow in an anaerobic bottle only. They are typically considered contaminants and require no treatment, but clinical judgment should be used. If therapy is desired, the recommended antibiotics are metronidazole 500 mg po q8h OR penicillin G 4 million units IV q4h. Either of these options should have activity against common gram-positive anaerobic species.

Gram-Negatives

When BCID results as "Not Detected" and the gram stain characteristics are negative, the organisms are more commonly anaerobes. Of these, *Bacteroides fragilis* group and *Fusobacterium* sp. are the most common. Although these organisms are not commonly seen in the blood, they are typically indicative of an underlying infection and are associated with high mortality rates, increasing the importance of appropriate therapy. Metronidazole is the preferred agent for these organisms because of its potent activity and lack of resistance. Penicillins with beta-lactamase inhibitors and carbapenems are also usually active.

Aerobic gram-negative rods that are "Not Detected" are rarer and typically include *Acinetobacter* sp., *Stenotrophomonas maltophilia*, and *Pseudomonas fluorescens-putida* group. These organisms are often associated with true infection and should be treated with an appropriate agent. Levofloxacin has reliable activity against all 3 of these as well as other aerobic gram-negatives, making it an appropriate empiric choice until identification. Of note, an updated BCID panel that detects Acinetobacter and *Stenotrophomonas maltophilia* is in development and will be coming to Nebraska Medicine when available.

Table 2: Organism Groups Identified by Culture and Not Detected by BCID

Organism Group Stratified by Gram Stain					
	n	% of Gram Stain	% of Total BCID Negative		
Gram Positive	274	100.0%	70.8%		
Micrococcus sp.	70	25.5%	18.1%		
Coagulase-negative Staphylococcus	36	13.1%	9.3%		
Diphtheroids	27	9.9%	7.0%		
Peptostreptococcus sp.	17	6.2%	4.4%		
Bacillus species, not anthracis	12	4.4%	3.1%		
Rothia sp.	10	3.6%	2.6%		
Clostridium sp.	10	3.6%	2.6%		
Lactobacillus sp.	10	3.6%	2.6%		
Viridans group Streptococcus	8	2.9%	2.1%		
Corynebacterium striatum group	7	2.6%	1.8%		
Abiotrophia/Granulicatella sp.	7	2.6%	1.8%		
Unable to identify	7	2.6%	1.8%		
Gemella sp.	7	2.6%	1.8%		
Enterococcus sp.	6	2.2%	1.6%		
Actinomyces sp.	5	1.8%	1.3%		
Aerococcus sp.	5	1.8%	1.3%		
Propionibacterium sp.	5	1.8%	1.3%		
Leuconostoc sp.	4	1.5%	1.0%		
Peptoniphilus sp.	3	1.1%	0.8%		
Parvimonas micra	3	1.1%	0.8%		
Eggerthella lenta	3	1.1%	0.8%		
Finegoldia magna	3	1.1%	0.8%		
Atopobium parvulum	1	0.4%	0.3%		

Staphylococcus aureus	Dermabacter hominis	1	0.4%	0.3%
Blautia producta 1 0.4% 0.3% Cutibacterium acnes 1 0.4% 0.3% Group C Streptococcus 1 0.4% 0.3% Eubacterium limosum 1 0.4% 0.3% Facklamia hominis 1 0.4% 0.3% Facklamia hominis 1 0.4% 0.3% Gram Negative 113 100% 29.2% Bacteroides fragilis group 36 31.9% 9.3% Fusobacterium sp. 18 15.9% 4.7% Acinetobacter sp. 7 6.2% 1.8% Prevotella sp. 6 5.3% 1.6% Stenotrophomonas maltophilia 6 5.3% 1.6% Unable to identify 5 4.4% 1.3% Pseudomonas fluorescens-putida group 4 3.5% 1.0% Pasteurella multocida 4 3.5% 1.0% Sphingomonas paucimobilis 3 2.7% 0.8% Morganella morganii 3 2.7% 0.8%	Staphylococcus aureus	1	0.4%	0.3%
Cutibacterium acnes 1 0.4% 0.3% Group C Streptococcus 1 0.4% 0.3% Eubacterium limosum 1 0.4% 0.3% Facklamia hominis 1 0.4% 0.3% Gram Negative 113 100% 29.2% Bacteroides fragilis group 36 31.9% 9.3% Fusobacterium sp. 18 15.9% 4.7% Acinetobacter sp. 7 6.2% 1.8% Prevotella sp. 6 5.3% 1.6% Stenotrophomonas maltophilia 6 5.3% 1.6% Unable to identify 5 4.4% 1.3% Pseudomonas fluorescens-putida group 4 3.5% 1.0% Pasteurella multocida 4 3.5% 1.0% Sphingomonas paucimobilis 3 2.7% 0.8% Morganella morganii 3 2.7% 0.8% Veillonella sp. 2 1.8% 0.5% Moreavella sp. 2 1.8% 0.5% <td></td> <td>1</td> <td>0.4%</td> <td>0.3%</td>		1	0.4%	0.3%
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Neisseria sicca group 1 0.9% 0.3% Anaerobiospirillum succiniciproducens 1 0.9% 0.3% Cardiobacterium hominis 1 0.9% 0.3% Alistipes sp. 1 0.9% 0.3% Chryseobacterium indologenes 1 0.9% 0.3% Brevundimonas vesicularis 1 0.9% 0.3% Proteus vulgaris 1 0.9% 0.3% Wautersiella falseni 1 0.9% 0.3% Eikenella corrodens 1 0.9% 0.3% Flavobacterium odoratum 1 0.9% 0.3%	Leptotrichia sp.	2	1.8%	0.5%
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Cardiobacterium hominis 1 0.9% 0.3% Alistipes sp. 1 0.9% 0.3% Chryseobacterium indologenes 1 0.9% 0.3% Brevundimonas vesicularis 1 0.9% 0.3% Proteus vulgaris 1 0.9% 0.3% Wautersiella falseni 1 0.9% 0.3% Eikenella corrodens 1 0.9% 0.3% Flavobacterium odoratum 1 0.9% 0.3%	Neisseria sicca group	1	0.9%	0.3%
Alistipes sp. 1 0.9% 0.3% Chryseobacterium indologenes 1 0.9% 0.3% Brevundimonas vesicularis 1 0.9% 0.3% Proteus vulgaris 1 0.9% 0.3% Wautersiella falseni 1 0.9% 0.3% Eikenella corrodens 1 0.9% 0.3% Flavobacterium odoratum 1 0.9% 0.3%	Anaerobiospirillum succiniciproducens	1	0.9%	0.3%
Chryseobacterium indologenes 1 0.9% 0.3% Brevundimonas vesicularis 1 0.9% 0.3% Proteus vulgaris 1 0.9% 0.3% Wautersiella falseni 1 0.9% 0.3% Eikenella corrodens 1 0.9% 0.3% Flavobacterium odoratum 1 0.9% 0.3%	Cardiobacterium hominis	1	0.9%	0.3%
Brevundimonas vesicularis 1 0.9% 0.3% Proteus vulgaris 1 0.9% 0.3% Wautersiella falseni 1 0.9% 0.3% Eikenella corrodens 1 0.9% 0.3% Flavobacterium odoratum 1 0.9% 0.3%	Alistipes sp.	1	0.9%	0.3%
Proteus vulgaris 1 0.9% 0.3% Wautersiella falseni 1 0.9% 0.3% Eikenella corrodens 1 0.9% 0.3% Flavobacterium odoratum 1 0.9% 0.3%	Chryseobacterium indologenes	1	0.9%	0.3%
Wautersiella falseni 1 0.9% 0.3% Eikenella corrodens 1 0.9% 0.3% Flavobacterium odoratum 1 0.9% 0.3%	Brevundimonas vesicularis	1	0.9%	0.3%
Eikenella corrodens10.9%0.3%Flavobacterium odoratum10.9%0.3%	Proteus vulgaris	1	0.9%	0.3%
Flavobacterium odoratum 1 0.9% 0.3%	Wautersiella falseni	1	0.9%	0.3%
	Eikenella corrodens	1	0.9%	0.3%
Grand Total 387 100.0% 100.0%	Flavobacterium odoratum	1	0.9%	0.3%
	Grand Total	387	100.0%	100.0%

Prepared By: Andrew Watkins, PharmD – PGY2 Infectious Diseases Pharmacy Resident

Reviewed By: Trevor Van Schooneveld, MD; Scott Bergman, Pharm.D; Paul Fey, PhD; Hannah Creager, PhD; Bryan

Alexander, PharmD; Jasmine Marcelin, MD; Erica Shohs, MD

Developed: Fall 2019