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Evaluation of Optimal Blood Culture Incubation Time To Maximize Clinically Relevant Results from a Contemporary Blood Culture Instrument and Media System

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ABSTRACT  Timely diagnosis of microorganisms in blood cultures is necessary to optimize therapy. Although blood culture media and systems have evolved for decades, the standard interval for incubation prior to being discarded as negative has remained 5 days. Here, we evaluated the optimal incubation time for the BacT/Alert Virtuo blood culture detection system (bioMérieux) using FA Plus (aerobic) and FN Plus (anaerobic) resin culture bottles in routine clinical use. Following institutional review board (IRB) approval, a retrospective review evaluated the outcomes of 158,710 bottles collected between November 2018 and October 2019. The number of positive blood bottles was 13,592 (8.6%); 99% of positive aerobic and anaerobic bottles flagged positive by 91.5 and 108 h, respectively. The mean (median) times to positivity were 18.4 h (15.6 h) for Staphylococcus aureus, 12.3 h (9.5 h) for Escherichia coli, 22.2 h (15.9 h) for Pseudomonas aeruginosa, and 48.9 h (42.9 h) for Candida spp. Only 175 bottles (0.1% of all bottles) flagged positive after 4 days of incubation; 89 (51%) of these bottles grew Cutibacterium (Propionibacterium) species. Chart review of blood cultures positive after 4 days (96 h) rarely had a clinical impact and sometimes had a negative impact on patient care. Finally, a seeded study of the HACEK group (i.e., Haemophilus, Aggregatibacter, Cardiobacterium, Eikenella, and Kingella), historically associated with delayed blood culture positivity, demonstrated no benefit to extended incubation beyond 4 days. Collectively, these findings demonstrated that a 4-day incubation time was sufficient for the Virtuo system and media. Implementation of the 4-day incubation time could enhance clinically relevant results by reducing recovery of contaminants and finalizing blood cultures 1 day earlier.

KEYWORDS  blood culture, Virtuo, time to positivity, BacT/Alert, FA Plus, FN Plus, Staphylococcus aureus, growth rate, Cutibacterium, Escherichia coli, Pseudomonas aeruginosa

Continuously monitored blood culture systems are used in clinical microbiology laboratories to test for life-threatening conditions like bacteremia, fungemia, and endocarditis. Historically, blood culture bottles were incubated on these systems for up to 7 days before being discarded and reported as “no growth.” Today, a 5-day minimum incubation is recommended for most commercial systems; however, there have been reports suggesting a 5-day incubation may not be necessary for modern blood culture systems (1–4). Optimization of the blood culture incubation period can support clinical care, with the goal of incubating long enough to recover clinically significant microorganisms while also minimizing the recovery of contaminants. The latter is important to minimize unnecessary antimicrobial therapy and can reduce waste in the...
clinical laboratory. In addition, a more rapid determination of a blood culture as negative has the potential to expedite patient discharges and more quickly deescalate broad-spectrum antibiotic usage to improve antimicrobial stewardship.

Our institution implemented the BacT/Alert Virtuo (Virtuo) blood culture detection system with FA Plus (aerobic) and FN Plus (anaerobic) resin culture bottles (bioMérieux) in mid-2018. This system has been shown to have a faster time to positivity compared to other systems using seeded cultures (5–7) and paired clinical cultures (8–10). These findings, along with anecdotal observations from the laboratory noting mostly growth of clinically insignificant microbes after 4 days of incubation, prompted a review of blood cultures from the Virtuo system. The goal of this work was to reassess optimal incubation time and microorganism recovery using a 1-year, retrospective analysis of Virtuo time-to-positivity data combined with medical chart review to determine clinical utility of a 5-day incubation. In addition, because they were encountered infrequently in the clinical specimens, a seeded study was included to evaluate HACEK microorganisms (Haemophilus, Aggregatibacter, Cardiobacterium, Eikenella, and Kingella).

**MATERIALS AND METHODS**

**Clinical setting.** The central microbiology laboratory at the Barnes-Jewish Institute of Health in St. Louis, MO, serves Barnes-Jewish Hospital (a 1,250-bed tertiary care academic medical center) as well as a stand-alone academic children’s hospital and three community hospitals in the St. Louis metropolitan area.

**Culture and identification.** Following approval from the Washington University in St. Louis Human Research Protection Office, all 158,710 blood culture bottles from November 2018 to October 2019 submitted to the laboratory were evaluated in the present study. The laboratory’s standard procedures recommend collecting blood cultures as a set consisting of two bottles: FA Plus (aerobic) and FN Plus (anaerobic) resin culture bottles. For adults, the recommended collection volume is 10 ml per bottle (i.e., 20 ml per set), and for children, approximately 1 ml of blood per year of patient age. No bottles were rejected by the laboratory or excluded from analyses based on blood volumes below the recommended volume. Bottles were continuously monitored until positive for up to 5 days at 37°C on the BacT/Alert Virtuo detector (bioMérieux). While an extended incubation was available by physician request, there were no requests during the time frame of this study. The Virtuo system screens bottles almost continuously during the first 74.4 h of incubation then every 144 min thereafter. Positive bottles underwent Gram staining and were processed according to laboratory standard operating procedures. If no organisms were seen by Gram stain, acridine orange staining was performed and the bottles were subcultured to solid medium.

All positive bottles were subcultured to agar media: sheep’s blood agar, chocolate agar, MacConkey’s agar, or colistin-nalidixic acid agar if Gram stain reveals mixed microbes, including Gram-positive organisms; Campylobacter selective agar if Gram stain reveals curved Gram-negative bacilli; CandiSelect and Sabouraud dextrose agar if Gram stain reveals fungal elements or yeast; TH11 agar if Gram stain reveals mycobacterium-like structures; and/or Brucella blood agar if an anaerobic bottle is flagged positive or if Gram stain is suggestive of anaerobic growth. Routine agars were incubated and imaged on the Kiestra Total Lab Automation system (Becton, Dickinson Microbiology Systems) (11). Anaerobic plates were incubated offline in conventional anaerobic chambers (either an Anaerobe Systems anaerobic workstation). Microbial identification was routinely performed by matrix-assisted laser desorption ionization–time of flight mass spectrometry (MALDI-TOF MS) with a Bruker BioTyp (Bruker, Billerica, MA).

**Data compilation.** A retrospective review evaluated the outcomes of 158,710 blood culture bottles, and, if positive, the microbial species/grouping. Bottles inoculated with non-blood specimens (e.g., pleural fluid) were excluded from analyses. Linking the recovered species to time to positivity required the merging of two large data sets. First, all time-to-positivity data were pulled from the Virtuo system’s MYLA middleware system using the culture accession number. Next, the reported species was pulled from the blood culture report logs using the Cerner Millennium laboratory information system. Using the accession number, cultures with a single positive bottle (with any number of reported species) or a single reported species (with any number of positive bottles) were linked. For cultures with multiple bottles with multiple species, these cultures required manual laboratory information system review to determine which species were recovered from which bottles.

**Medical chart review.** A medical chart review in EPIC (Madison, WI) was performed on patients with a bottle time to positivity of greater than 4 days (96 h) of incubation. Patients’ electronic health records were reviewed for clinical actions prior to and after reporting of the blood culture. This included admission status, antimicrobial treatments, comorbidities, diagnoses, deceased status, and any mention of the culture in the medical chart, especially from infectious disease physicians. Examples of “No Impact” included (i) bottles with no microorganisms observed/recovered on a subculture, (ii) bottles that flagged positive after patient expired, (iii) bottles from cultures in a two-bottle set and where the companion bottle had a time to positivity of less than 4 days and reported the same microbe, (iv) bottles that were specifically called out in medical record as a “contaminant” by the physician, and (v) bottles that were from patients lost to follow-up.
Seeded study (HACEK). Clinical isolates of HACEK microorganism recovered from frozen storage were first subcultured, and species were confirmed by MALDI-TOF MS. The species were *Aggregatibacter aphrophilus*, *Cardiobacterium hominis*, *Eikenella corrodens*, and *Kingella denitrificans*, identified using MALDI-TOF MS. For the seeded study, *Escherichia coli* strain ATCC 25922 was included as a positive control. After subculture, a few well-isolated colonies were used to make a 1.0 McFarland standard (3.0 × 10^8 CFU/ml). Subsequent dilutions were used to achieve a target inoculum of 30 CFU in 5 ml of human blood (packed red blood cell remnant units provided courtesy of the Barnes-Jewish Hospital transfusion service). This final concentration (6 CFU/ml) was determined by companion colony counts. If input colony counts were 10 or >80 CFU total on the agar plates using 0.1 ml of the prior 3 × 10^8 CFU/ml dilution, data were excluded from analysis and retested. A saline-only inoculum was used as a negative control. Aerobic and anaerobic bottles were inoculated in triplicate using standard practices and incubated on the Virtuo system per routine procedures for 5 days.

RESULTS

Cohort summary. A total of 158,710 blood culture bottles were tested at the Barnes-Jewish Hospital microbiology laboratory between November 2018 and October 2019. Of these bottles, 13,592 (8.6%) flagged positive by the 5-day incubation cutoff (Fig. 1). The breakdowns of bottles from adults and children were 12,769 and 823, respectively.

Time to positivity. The times for 50% and 90% of bottles to signal positive were 16.5 and 38.5 h, respectively (Table 1). The mean time to positivity was 21.1 h. After 24 h of incubation, 10,431 bottles, or 76.74%, flagged positive. An additional 2,286 bottles flagged positive between 24 and 48 h, increasing the positivity rate to 93.56%. By the end of 4 days, 98.71% of bottles flagged positive—all but 175 bottles. Similar trends were observed from children (defined as <18 years of age [Fig. 2]). For the cultures collected from children, 630 bottles, or 76.55%, flagged positive after 24 h of incubation; by the end of 4 days, 99.39% flagged positive—all but five bottles.

Time to positivity under aerobic and anaerobic growth conditions. To compare differences between aerobic and anaerobic growth conditions, the aforementioned data were parsed into aerobic (FA) and anaerobic (FN) bottle types. For aerobic bottles, the times until 50% and 90% of the bottles signaled positive were 15.5 and 36.5 h, respectively (Table 1). The mean time to positivity was 20.0 h. After 24 h of incubation, 5,565 bottles, or 80.19%, flagged positive (Fig. 3, top panel). An additional 957 bottles flagged positive between 24 and 48 h, increasing the positivity rate to 93.98%. By the end of 4 days, 99.27% of bottles flagged positive—all but 51 bottles.

For anaerobic bottles, a broader distribution was observed spanning the first 30 h. The times to reach 50% and 90% positivity were 17.5 and 36.5 h, respectively (Table 1). The mean time to flag positive was 22.3 h. After 24 h of incubation, 4,866 bottles, or
73.15%, flagged positive (Fig. 3, bottom panel). An additional 1,329 bottles flagged positive between 24 and 48 h, increasing the total positivity rate to 93.13%. By the end of 4 days, 98.14% of bottles flagged positive—all but 124 bottles.

**Species-specific analyses.** A total of 217 different species or microbial "groupings" (e.g., *Bacteroides fragilis* group or coagulase-negative *Staphylococcus* species) were reported from positive bottles. Of these, 52 different species or microbial groupings were found in ≥20 bottles (see Table S1 in the supplemental material); these included 31 Gram-positive species or groupings, 17 Gram-negative bacteria, and 4 species of yeast. Forty-six bottles (0.3% of positive bottles) flagged positive, but no organism was ever recovered or detected by acridine orange staining.

Histograms of select medically relevant Gram-positive microbes [*Staphylococcus aureus* (*n* = 3,893), *Staphylococcus epidermidis* (*n* = 1,248), *Streptococcus pneumoniae* (*n* = 192), and *Enterococcus faecium/Enterococcus faecalis* (*n* = 859)] are shown in Fig. 4. All four species reached 95% and 99% times to positivity more quickly than the overall

<table>
<thead>
<tr>
<th>Grouping</th>
<th>No. of bottles</th>
<th>Time to positivity (h)</th>
<th>Mean</th>
<th>T50</th>
<th>T90</th>
<th>T95</th>
<th>T99</th>
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</thead>
<tbody>
<tr>
<td>All</td>
<td>13,592</td>
<td>21.1</td>
<td>16.5</td>
<td>38.5</td>
<td>56.5</td>
<td>103.5</td>
<td></td>
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<tr>
<td>Aerobic bottles</td>
<td>6,940</td>
<td>20.0</td>
<td>15.5</td>
<td>36.5</td>
<td>54.0</td>
<td>91.5</td>
<td></td>
</tr>
<tr>
<td>Anaerobic bottles</td>
<td>6,652</td>
<td>22.3</td>
<td>17.5</td>
<td>39.5</td>
<td>59.5</td>
<td>108.0</td>
<td></td>
</tr>
<tr>
<td>Bottles from children</td>
<td>823</td>
<td>20.0</td>
<td>16.5</td>
<td>36.0</td>
<td>44.0</td>
<td>84.0</td>
<td></td>
</tr>
<tr>
<td><em>S. aureus</em></td>
<td>3,893</td>
<td>18.4</td>
<td>16.0</td>
<td>29.5</td>
<td>39.0</td>
<td>72.0</td>
<td></td>
</tr>
<tr>
<td>Coagulase-negative <em>Staphylococcus</em> spp.</td>
<td>2,208</td>
<td>23.5</td>
<td>20.5</td>
<td>35.5</td>
<td>46.0</td>
<td>86.5</td>
<td></td>
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<tr>
<td><em>E. coli</em></td>
<td>1,382</td>
<td>12.3</td>
<td>10.0</td>
<td>17.5</td>
<td>27.5</td>
<td>66.0</td>
<td></td>
</tr>
<tr>
<td><em>S. epidermidis</em></td>
<td>1,248</td>
<td>20.9</td>
<td>19.5</td>
<td>29.5</td>
<td>34.5</td>
<td>54.5</td>
<td></td>
</tr>
<tr>
<td><em>E. faecium/E. faecalis</em></td>
<td>859</td>
<td>15.7</td>
<td>13.0</td>
<td>25.5</td>
<td>34.5</td>
<td>64.5</td>
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</tr>
<tr>
<td><em>Candida</em> spp.</td>
<td>314</td>
<td>48.9</td>
<td>43.0</td>
<td>82.0</td>
<td>108.0</td>
<td>115.5</td>
<td></td>
</tr>
<tr>
<td><em>P. aeruginosa</em></td>
<td>298</td>
<td>22.2</td>
<td>16.0</td>
<td>45.0</td>
<td>57.5</td>
<td>96.0</td>
<td></td>
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<tr>
<td><em>S. pneumoniae</em></td>
<td>192</td>
<td>10.6</td>
<td>10.5</td>
<td>14.5</td>
<td>17.5</td>
<td>30.5</td>
<td></td>
</tr>
<tr>
<td><em>Cutibacterium</em> spp.</td>
<td>121</td>
<td>103.9</td>
<td>106.0</td>
<td>115.5</td>
<td>118.0</td>
<td>118.0</td>
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<tr>
<td><em>B. fragilis</em></td>
<td>76</td>
<td>31.8</td>
<td>28.0</td>
<td>52.0</td>
<td>57.5</td>
<td>60.0</td>
<td></td>
</tr>
<tr>
<td><em>S. maltophilia</em></td>
<td>25</td>
<td>25.9</td>
<td>21.5</td>
<td>56.0</td>
<td>57.5</td>
<td>79.5</td>
<td></td>
</tr>
<tr>
<td>No growth</td>
<td>46</td>
<td>33.0</td>
<td>14.5</td>
<td>94.0</td>
<td>106.0</td>
<td>118.0</td>
<td></td>
</tr>
</tbody>
</table>

*The total numbers of aerobic and anaerobic bottles were 7,113 and 6,804, respectively.*

*The "Coagulase-negative *Staphylococcus* spp." grouping includes isolates not otherwise specified by row in the table.*
average (Table 1). *S. pneumoniae* had the shortest time to positivity of any microbe, with 98.44% of bottles with this microorganism flagging positive within 24 h and 100% within 48 h. The grouping coagulase-negative *Staphylococcus* was commonly found in the data set and exhibited a histogram similar to those of other Gram-positive microbes. Key Gram-negative bacteria recovered from positive bottles included *Escherichia coli* (*n* = 1,382), *Pseudomonas aeruginosa* (*n* = 298), *Stenotrophomonas maltophilia* (*n* = 25), and *Bacteroides fragilis* group (*n* = 76) (Fig. 5). *E. coli* reached 50% positivity faster than any other microbe, at 10 h (Table 1). While 94.43% of bottles flagged positive in 24 h, a long histogram tail was observed, with 25 bottles flagging positive over days 3 to 5. *P. aeruginosa* had 97.99% of bottles flag positive after 3 days and only two bottles flag positive after 4 days. The data sets for *S. maltophilia* and *B. fragilis* were noteworthy because the overwhelming majority of bottles (all but 1) flagged positive within 3 days.

To determine the growth patterns associated with fungemia, all *Candida* species were evaluated (Fig. 6). Time to positivity was more evenly distributed across the 5-day incubation time than the bacterial histograms. By the fourth day, 92.68% of bottles flagged positive, leaving 23 remaining. The delayed time to positivity was also evident in the times to achieve 50, 90, 95, and 99% positivity (Table 1). *Cutibacterium* (formerly *Propionibacterium*) species are often considered contaminants, although this microbe has been implicated in causing serious disease (12–14). The histogram of *Cutibacterium* species (Fig. 4) was strikingly different from those of other microbial histograms. Only two bottles flagged positive after 3 days (1.65%). An additional 30 bottles flagged positive by day 4 (26.45%), and 89 (73.55%) bottles

![Graph of Aerobic Bottles](image1.png)

**FIG 3** Evaluation of time to positivity for aerobic and anaerobic blood culture bottles. Positive aerobic (*n* = 6,940) and anaerobic (*n* = 6,652) bottles were plotted in 30-min intervals. After 4 days of incubation, 99.27% of aerobic bottles and 98.14% of anaerobic bottles flagged positive.
flagged positive in the final 24 h. This delayed time to positivity was also notable in Table 1.

Clinical impact of positive bottles after 4 days. Only 175 bottles (0.1% of all bottles and 1.3% of positive bottles) flagged positive after 4 days of incubation. There were 160 bottles that had no recorded impact in the patient’s medical record. Of these bottles, 38 were specifically mentioned by medical staff as being “contaminated.” Twenty-two bottles had a companion bottle in the culture flagged positive with the same organism before 96 h. Three of the 175 bottles resulted in orders for additional tests.
that were never acted upon. Another bottle that grew *Cutibacterium* spp. delayed a patient’s discharge by 2 days. A different bottle that grew *Cutibacterium* spp. resulted in a visit to the Emergency Department, only to have repeat blood cultures show no growth. Another bottle grew *Lactobacillus* spp. that resulted in an unremarkable clinic visit. The remaining nine bottles caused a treatment adjustment. While it was difficult to determine true significance, there was documented reservation of true pathogenic-

**FIG 5** Evaluation of time to positivity for select Gram-negative bacteria. Positive bottles were plotted in 30-min intervals. For each grouping, greater than 99.3% of bottles flagged positive after 4 days.

**FIG 6** Evaluation of time to positivity for *Candida* species. Positive bottles (*n* = 314) were plotted in 30-min intervals. After 4 days of incubation, 92.68% of bottles flagged positive.
ity for at least three of the nine cultures, and three other cultures were from patients who expired in less than 96 h. Even by the most conservative of estimates, the last day of incubation was beneficial for <10 bottles (<0.006%).

**Seeded experiment.** Historically, microbes in the HACEK group have been considered slow growing, and an extended incubation might be necessary for recovery. HACEK organisms were infrequently encountered in the retrospective review (e.g., Eikenella species, n = 3), with the exception of Haemophilus species (n = 31). To enhance the evaluable instances of the other HACEK organisms, a seeded study was performed in triplicate using representative species: Aggregabacter aphrophilus, Cardiobacterium hominis, Eikenella corrodens, Kingella denitrificans, and Kingella kingae. The time to positivity of all isolates was less than 4 days for at least one bottle of the paired aerobic and anaerobic set (Fig. 7). Bottles that did not flag positive are shown at the top of the graph. Importantly, no bottles flagged positive between 4 and 5 days, suggesting no added benefit of incubation time past 4 days for the HACEK group.

*S. aureus* bacteremia. Prolonged positivity and time to culture clearance are important blood culture metrics when assessing a patient’s bacteremia. As shown in Fig. 4, S. aureus was routinely recovered in ≤4 days, but nine bottles flagged positive after 4 days. None of these nine bottles resulted in therapeutic implications. For three bottles, the companion bottle in the blood culture set had already flagged positive in ≤4 days. Another three bottles had a different culture collected the same day that had already flagged positive with *S. aureus*. Two bottles were from patients with intermittent *S. aureus*-positive blood cultures that had flanking cultures collected before and after flagging positive. Finally, one bottle was the last *S. aureus*-positive blood culture in a series of cultures, with all subsequent blood cultures being negative.

**Postimplementation.** Based on the data reported in this study, our institution implemented a 4-day incubation cutoff for blood cultures. Physicians still have the option to call the laboratory and request extended incubation. As a result, the laboratory is on track to report >60,000 blood culture tests 1 day earlier in just the first year of implementation. Another potential benefit has been a reduction in reporting of blood culture contaminants. While still preliminary, with only 2 months of data, we have seen a reduction in the recovery rate of *Cutibacterium* species relative to total positives (0.07%; predicted reduction based on shortened incubation, 0.23%; previous rate, 0.87%). Lastly, we were able to create additional instrument capacity without purchasing additional units in our Virtuo system by removing all the negative bottles 1 day earlier. This proved especially valuable in the era of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) and the corresponding massive increases in blood culture requests.
DISCUSSION

Bloodstream infections are a significant cause of morbidity and mortality in the United States, with an estimated burden of 72,000 to 85,000 deaths annually (15). Continuously monitored blood culture systems are a mainstay in clinical microbiology laboratories to test for these potentially life-threatening conditions. The goal of this study was to assess the optimal incubation time for blood cultures using the BacT/Alert Virtuo system paired with FA Plus (aerobic) and FN Plus (anaerobic) resin culture bottles. A retrospective review of 158,710 blood culture bottles collected over 1 year found only 175 (0.11%) flagged positive after 4 days of incubation. Chart review of these 175 bottles revealed that less than 10 bottles had a positive impact on clinical care, and some positives contributed to additional, unnecessary testing, delayed discharges, and unnecessary health care visits. These findings suggested that a routine incubation time of 4 days is sufficient for clinical laboratories using the Virtuo system. Optimizing culture incubation time to minimize the recovery of clinically insignificant cultures is appealing as blood culture contaminants can result in increased health care costs, prolonged length of stay, and laboratory waste (16–18).

There is a precedent for reducing the standard incubation time of blood cultures on automated systems. For years, the standard was 7 days. Publications from the 1990s demonstrated that commonly used systems like the BacT/Alert could have a shortened incubation period (19–22). The optimal incubation length varied by blood culture system. Hardy et al. reported that ≥99% of cultures from pathogenic species were positive by 5 days and noted a 5-day incubation would significantly reduce recovered Cutibacterium (Propionibacterium) species by 77% (19), strikingly similar to the work presented here. The Difco ESP system was reported to have sufficient results, with a 4-day incubation detecting 97.8% of all positives (3). The AccuMed ESP-384 system with a 3-day incubation was considered sufficient, with 94.2% of positives detected (23).

There are reports of the Virtuo system having an even faster time to positivity compared to other blood culture systems—BD Bactec FX (6, 24, 25) and the BacT/Alert 3D system (8–10, 24, 26–28)—although the vast majority of these investigations used seeded studies. It should be noted that the epidemiological prevalence of species throughout these studies was similar to that in the present study. The most commonly recovered microbes were coagulase-negative staphylococci, S. aureus, Enterococcus species, E. coli, and S. pneumoniae. As for the theoretically missed species with a reduced incubation, these were also similar: Cutibacterium (Propionibacterium) species, Candida glabrata, and microbes of questionable clinical significance. Taken together, the evidence within the present article supporting a 4-day incubation closely resembles that of the past that led to the now 5-day standard incubation time.

To our knowledge, this is the largest study of its kind evaluating the outcomes of blood cultures on the Virtuo system along with supplementary medical chart reviews. Even a study of this size will not include the rarest of slow-growing pathogens. To mitigate this limitation, a seeded experiment of the HACEK group was included and showed no difference between 4- and 5-day incubations. The HACEK group was historically often thought to require extended incubation for detection, but this has been refuted previously (29, 30). An additional limitation of this study is the findings may not be applicable to other blood culture systems or other laboratories with different local microbiological populations. Furthermore, a criticism of this work might be that the data come from a single microbiology laboratory, but cultures were received from five hospitals, including an academic hospital, community hospitals, and a children’s hospital. Lastly, there may be interest in establishing an even shorter incubation time in the future. If a 3-day incubation was proposed using our data set, the five most impacted species would be S. aureus, coagulase-negative Staphylococcus species, Cutibacterium species, Candida species, and Sphingomonas species. Future studies are needed to determine if shorter incubation periods are clinically feasible and could further optimize patient care and laboratory testing.

If an institution was to reduce the blood culture incubation time, national guidelines...
from pertinent organizations must be considered. The Clinical and Laboratory Standards Institute (CLSI) guidelines recommend a 5-day incubation time while also acknowledging 4 days, or even 3 days, might be adequate for some systems (31). The CLSI M47 document (published in 2007) states adequate data from a shorter incubation time may permit laboratories to adopt shorter protocols; this may be revised in updates to the M47 document. Other guidelines like those from the Infectious Diseases Society of America (IDSA) may be impacted by a shortened incubation period. For example, uncomplicated bacteremia is defined as patients with positive blood cultures and follow-up blood cultures performed on specimens obtained 2 to 4 days after the initial set that do not grow methicillin-resistant S. aureus (MRSA) (32). This 2- to 4-day time period may need to be adjusted in the setting of contemporary blood culture systems and media.

In conclusion, we found a 4-day incubation cutoff to detect the vast majority of microorganisms detected on the BacT/Alert Virtuo system. We project that a 4-day incubation on the Virtuo blood culture system not only will improve microbiological workflows but also will improve patient care.

SUPPLEMENTAL MATERIAL
Supplemental material is available online only.

SUPPLEMENTAL FILE 1, PDF file, 0.4 MB.

ACKNOWLEDGMENTS
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