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Value of β -D-glucan and Candida albicans germ tube antibody for discriminating between Candida colonization and invasive candidiasis in patients with severe abdominal conditions

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On behalf of the CAVA II Study Group. The members of the CAVA II Study Group are given in the Appendix.

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Abstract *Purpose:* To assess the value of $(1 \rightarrow 3)$ - β -D-glucan (BDG), Candida albicans germ tube antibody (CAGTA), C-reactive protein (CRP), and procalcitonin (PCT) levels for the diagnosis of invasive candidiasis (IC) and for differentiating Candida spp. colonization from infection in ICU patients with severe abdominal conditions (SAC). Methods: Prospective study of 176 nonneutropenic patients, with SAC at ICU admission, and expected to stay at least 7 days. Surveillance cultures and BDG, CAGTA, CRP, and PCT levels were performed on the third day of ICU stay and twice a week for four consecutive weeks. Patients were grouped into invasive candidiasis (IC), Candida colonization, and neither colonized/nor infected. The classification and regression tree (CART) analysis was used to predict IC in colonized patients. The discriminatory ability of the obtained prediction rule was assessed by the area under the ROC curve (AUC). Results: The probabilities of IC were 59.3 % for the terminal node of BDG greater than 259 pg/mL and 30.8 % for BDG less than 259 pg/mL and CAGTA positivity, whereas there was a 93.9 % probability in predicting the absence of IC for BDG less than 259 pg/mL and negative CAG-TA. Using a cutoff of 30 % for IC probability, the prediction rule showed 90.3 % sensitivity, 54.8 %

specificity, 42.4 % positive predictive not found. Conclusions: BDG with value, and 93.9 % negative predictive value with an AUC of 0.78 (95 % confidence interval 0.76–0.81). Significant differences in CRP (p = 0.411) and PCT (p = 0.179)among the studied groups were

a positive test for CAGTA accurately differentiated Candida colonization from IC in patients with SAC, whereas CRP and PCT did not.

Keywords $(1\rightarrow 3)$ - β -D-Glucan · Candida albicans germ tube antibody (CAGTA) · C-reactive protein · Procalcitonin · Critically ill patients · Abdominal conditions

Introduction

The diagnosis of invasive candidiasis (IC) in patients admitted to the ICU still poses a challenge [1]. Clinical prediction rules are relevant elements enabling proper diagnosis but some of them are not easy to fulfill, have not been previously validated, or may be eventually unhelpful in patients having abdominal surgery (e.g., the Candida score) [2–5]. Specific antigen components of the fungal cell wall have been exploited for the development of diagnostic assays that can detect the presence of these components in the serum. An important component of the cell wall of the majority of fungi is $(1\rightarrow 3)$ - β -D-glucan (BDG). Although the test is not Candida specific, it has been found to be a promising tool to diagnose invasive fungal infections, most of them being IC [6]. The accuracy of BDG for the diagnosis of invasive fungal infections has been discussed in a recent meta-analysis [7]. Also, a test based on the detection of antibodies against the surface of C. albicans germ tubes (CAGTA) has been commercialized [8]. Two newly published studies showed a significant decrease in mortality in ICU patients with a CAGTA-positive result especially in those with increasing CAGTA values who had been treated with antifungals [9, 10]. C-reactive protein (CRP) [11] and procalcitonin (PCT) [12, 13] have been also reported to be valuable markers.

The efficiency of diagnostic biomarkers in the field of fungal infections has become the object of intensive investigation. The aim of this study was to explore the accuracy of the following biomarkers: BDG, CAGTA, CRP, PCT, and some of them combined, for discriminating between Candida spp. colonization and IC in non-neutropenic critically ill patients with severe abdominal conditions (SAC) and to establish a model for the prediction of IC.

Patients and methods

Design and setting

This was a prospective, cohort, observational, and multicenter study conducted in the ICU setting in routine clinical practice. The study protocol was approved by the ethics committees of the participating centers, and

informed consent was obtained from the patients or their representatives.

Study population

All patients older than 18 years with SAC on ICU admission who were admitted to 18 medical-surgical ICUs of tertiary care hospitals in Spain between 1 April 2009 and 30 June 2010 were eligible. To be included in the study, an expected ICU stay of at least 7 days was required. At ICU entry, patients with neutropenia (total leukocyte count less than 1,000/mm³) were excluded as were those with other conditions (see Supplementary Material).

Screening, microbiological cultures, and Candida score

Surveillance cultures for *Candida* spp. were performed using samples obtained from feces (or rectal swabs), urine, skin (axillary surface), tracheal aspirates (or protected specimen brush or bronchoalveolar lavage), gastric or pharyngeal aspirates, and the peripheral blood. Other samples from the peripheral blood, vascular lines, wound/ drainage exudates, or infected foci were obtained at the discretion of the attending physician. (See Supplementary Material for details of microbiological cultures). Results were considered positive in the presence of Candida growth in the culture medium. The different Candida isolates were identified at species level. As soon as the results of surveillance cultures were available and at the time of starting antifungal treatment in patients with IC, the Candida score (CS) was calculated [4], with a cutoff point of at least 3 for discriminating between Candida spp. colonization and IC.

Serological biomarkers

Reference values for the serological tests were 80 pg/mL for the BDG assay, at least 1/160 for positive CAGTA, 0-5 mg/dL for CRP, and 0.5 ng/mL for PCT. Technical details of these assays are described in the Supplementary Material.

For each patient, maximum values recorded for *Candida* score and serologic biomarkers at or before the episode of IC were used in the analysis. When an episode of IC did not develop, the highest value of all observed values was used.

Definitions

A SAC was defined as the process that caused gastrointestinal dysfunction or failure in the context of a medical-surgical abdominal illness, including nonsurgical diseases (e.g., pancreatitis), emergency or elective surgical procedures, and related complications (e.g., gastrointestinal perforation, hepatobiliary and pancreatic disorders, peritonitis, intra-abdominal abscess, anastomotic leak), and prolonged postoperative stay after complicated abdominal surgery.

Candida colonization was considered unifocal when Candida spp. was isolated from one site and multifocal when Candida spp. was simultaneously isolated from various noncontiguous sites, even if two different Candida spp. were isolated [14, 15].

The diagnosis of IC required one of the following criteria: presence of candidemia, i.e., documentation of one or more blood culture(s) that yielded a *Candida* spp. in a patient with consistent clinical manifestations, isolation of *Candida* spp. from a normally sterile body fluids (e.g., pleural fluid, pericardial fluid) or candidal peritonitis, ophthalmic examination consistent with candidal endophthalmitis in a patient with clinical sepsis, or histologically documented candidiasis. IC was also considered if histopathological examination revealed typical patterns, such as pseudohyphae or true hyphae, in a relevant clinical context [16]. The definitions of candidal peritonitis [17], candidal endophthalmitis [18], catheter-related candidemia [19], and candiduria are described in the Supplementary Material.

Patients were classified into the groups of neither colonized nor infected, *Candida* spp. colonization without IC, and IC. Otherwise, the decision to treat a patient with antifungal drugs was left to the investigators' discretion.

Study protocol and collection of data

Once the patient was included in the study, the following data were recorded on the third day of ICU stay and twice a week thereafter for four consecutive weeks until ICU discharge or death: APACHE II score, SOFA score, surveillance cultures, *Candida* score, and presence or absence of sepsis, severe sepsis, or septic shock. Blood samples for the measurement of BDG, CAGTA, CRP, and PCT were drawn at the same time periods. Patients were followed until ICU and/or hospital discharge, or death. Other variables recorded are detailed in the Supplementary Material.

Sample size and statistical analysis

In a previous study [4], BDG was assessed in 65 of 217 non-neutropenic critically ill patients who underwent abdominal operations, yielding 11 patients classified into the neither colonized nor infected group, 45 into the *Candida* spp. colonization group, and 9 into the IC group. The area under the receiver operating characteristics (ROC) curve to assess the discriminatory power of BDG to distinguish *Candida* spp. colonization from IC was 0.731 [95 % confidence interval (CI) 0.620–0.841]. According to these results, 38 patients would be required in the IC group and 95 in the *Candida* spp. colonization group to estimate the area under the ROC curve (AUC) with an error bound of 10 % corresponding to the BDG.

Categorical variables are expressed as frequencies and percentages, and continuous variables as mean and standard deviation (SD) when data followed a normal distribution, or as median and interquartile (25th-75th percentile) range (IQR) when distribution departed from normality. The percentages were compared using the Chisquare (χ^2) test, the means by the F test, and the medians by the Kruskal-Wallis test. In patients with fungal colonization, a model for the prediction of the IC was obtained using the classification and regression trees (CART) procedure [20]. CART classifies data using a sequence of if-then rules. The basis of the decision tree algorithms is the binary recursive partitioning of the data. The most discriminative variable is first selected to partition the data set into child nodes. The splitting continues until some stopping criterion is reached. At each terminal node, the probability of IC was estimated as the proportion of patients belonging to that node that developed the event. The tree was constructed according to the following algorithm: in the first stage, the tree grows until all cases are correctly classified, and in the second stage, we used the tenfold cross-validation method of successive pruning [20]. Finally, the tree that minimized the error measurement (deviance) was chosen. For this predictor the corresponding ROC curve was obtained and the AUC was estimated by means of a 95 % CI. The predictive rule identified that a patient had an IC risk when the probability to develop the IC was 30 % or above. Sensitivity, specificity, and predictive values of the prediction rule were calculated. The data analysis was carried out using the R-package.

Results

Study population and salient findings

Of the initial 338 eligible patients, 162 were excluded because of lack of fulfillment of the inclusion criteria, incomplete clinical data collection, or blood samples to determine biologic biomarkers were not drawn (Fig. 1). The study population consisted of 176 patients, 65.9 % men, with a mean age (SD) of 64.1 (15.0) years. The mean (SD) APACHE II score and SOFA score on ICU admission was 18.7 (6.1) and 7.4 (3.6), respectively. The reason for ICU admission was medical in 18.8 % of patients, surgical in 76.1 %, and trauma in 5.1 %. The median (IQR) length of ICU and hospital stay was 15 (10–27) and 38 (24–57) days, respectively, which were significantly higher in patients with IC compared to the other two groups. ICU and hospital crude mortality rates were 26.7 and 32.4 %, respectively.

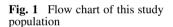
Abdominal conditions were gastrointestinal perforation (small and large intestine) 50 (28.4 %), prolonged postoperative state after complicated abdominal surgery (e.g., postoperative scheduled gastrointestinal neoplasia, intestinal ischemia, etc.) 47 (26.7 %), pancreatitis (no surgical) 32 (18.1 %), hepatobiliary-pancreatic pathology (cholecystitis/gallbladder perforation, liver abscess, pancreatic/peripancreatic abscesses) 25 (14.2 %), anastomotic leakage (esophagus, small and large intestine) 14 (7.9 %), and peritonitis with interbowel abscesses 8 (4.5 %).

There were 61 patients in the neither colonized nor infected group, 84 in the *Candida* spp. colonization group, and 31 in the IC group. Table 1 provides the

clinical characteristics of the participants. The rate of emergency surgery was significantly higher (p < 0.001) in the IC group than in the remaining two groups. Patients in the IC group showed significant differences as compared with the remaining groups in the maximum APACHE II and SOFA scores before or during the IC event, the *Candida* score, and the length of ICU stay and hospital stay. Data of the 31 patients with IC are summarized in Table 2. Most cases of IC were detected in the second week of ICU stay. Twenty-one of them had previous multifocal colonization, with a median of 3 days between colonization and IC (IOR 0–9).

Serological biomarkers and Candida score

A total of 766 blood samples (median 3, IQR 2–6) were drawn for the measurement of serological biomarkers. In the first study control on the third day after ICU admission, only the CS was significantly different in the IC group as compared with the other two groups. However, when the maximum values and rates were considered, BDG and CAGTA in addition to CS were also significantly different (Table 3). In relation to the influence of antifungal treatment on serological biomarkers, the median value of the variation between pre-treatment and



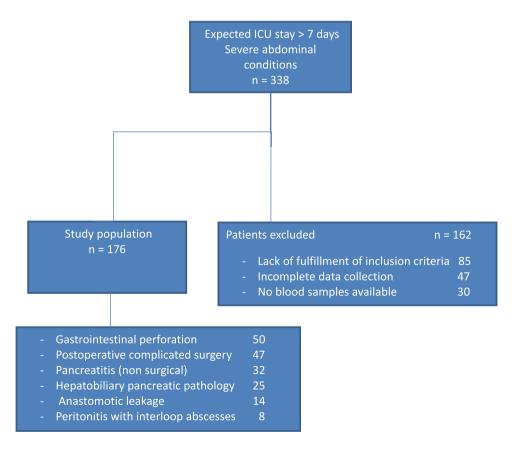


Table 1 Characteristics of the study patients according to colonization and infection status

	All patients $(n = 176)$	Study groups			P value
		Neither colonized nor infected (<i>n</i> = 61)	Candida spp. colonization $(n = 84)$	Invasive candidiasis $(n = 31)$	
Age, years, mean (±SD)	64.1 ± 15.0	60.2 ± 15.6	66.2 ± 15.1	65.8 ± 12.4	0.046
Male/female (%)	65.9/34.1	73.8/26.2	61.9/38.1	61.3/38.7	0.276
APACHE II score, mean (±SD) ICU admission	18.7 ± 6.1	17.9 ± 5.4	18.6 ± 6.1	20.4 ± 7.4	0.161
Maximum before/during development	17.7 ± 6.1	17.9 ± 3.4 16.1 ± 6.6	17.7 ± 6.5	20.4 ± 7.4 20.6 ± 5.7	0.101
SOFA score, mean (±SD)	17.7 ± 0.5	10.1 ± 0.0	17.7 ± 0.5	20.0 ± 3.7	0.007
ICU admission	7.4 ± 3.6	7.1 ± 3.7	7.3 ± 3.6	8.2 ± 3.7	0.428
Maximum before/during development	7.5 ± 4.0	6.8 ± 4.1	7.4 ± 4.0	9.2 ± 3.6	0.024
Maximum CS before/during development	4 (2–4)	2 (2–4)	4 (3–5)	5 (3–5)	< 0.001
IC, median (IQR)	15 (10, 25)	11 (0. 20)	15 (10, 22)	22 (21 45)	0.001
ICU length of stay, days, median (IQR)	15 (10–27)	11 (9–20)	15 (10–23)	32 (21–45)	< 0.001
Hospital length of stay, days, median (IQR) ICU mortality, no. (%)	38 (24–57) 47 (26.7)	31 (20–47) 15 (24.6)	38 (26–62) 19 (22.6)	52 (34–73) 13 (41.9)	0.005 0.104
Overall mortality, no. (%)	57 (32.4)	17 (27.9)	27 (32.1)	13 (41.9)	0.104
Types of patients on ICU admission, no. (%)	37 (32.4)	17 (27.9)	27 (32.1)	13 (41.9)	0.068
Medical	33 (18.8)	16 (26.2)	15 (17.9)	2 (6.5)	0.000
Surgical	134 (76.1)	44 (72.1)	62 (73.8)	28 (90.3)	
Trauma	9 (5.1)	1 (1.6)	7 (8.3)	1 (3.2)	
Total multifocality, no. (%)	111 (96.5)	_	83 (98.8)	28 (90.3)	0.059*
Clinical condition at the second week					0.258
of ICU stays, no. (%)	44 (25.0)	15 (24.6)	22 (26 2)	T (22.6)	
No sepsis	44 (25.0)	15 (24.6)	22 (26.2)	7 (22.6)	
Sepsis	49 (27.8)	21 (34.4)	24 (28.6)	4 (12.9)	
Severe sepsis Septic shock	39 (22.2) 44 (25.0)	9 (14.8) 16 (26.2)	20 (23.8) 18 (21.4)	10 (32.3) 10 (32.3)	
Diagnosis on admission, no. (%)	44 (23.0)	10 (20.2)	10 (21.4)	10 (32.3)	
Postoperative emergency gastrointestinal surgery	91 (51.7)	29 (47.5)	40 (47.6)	22 (71.0)	0.061
Postoperative elective gastrointestinal surgery	22 (12.5)	10 (16.4)	11 (13.1)	1 (3.2)	0.191
Mesenteric infarction	3 (1.7)	1 (1.6)	1 (1.2)	1 (3.2)	0.755
Peritonitis	48 (27.3)	17 (27.9)	21 (25.0)	10 (30.2)	0.734
Pancreatitis	39 (22.2)	17 (27.9)	17 (20.2)	5 (16.1)	0.370
Surgery (characteristics), no. (%)	22 (12.5)	12 (21 2)	0 (10.7)	0	0.001
No surgery	22 (12.5)	13 (21.3)	9 (10.7)	0	
Emergency	128 (72.7)	39 (63.9)	58 (69.0)	31 (100)	
Elective Abdominal surgery, no. (%)	26 (14.8) 144 (82.3)	9 (14.8) 48 (78.6)	17 (20.2) 68 (81.9)	0 28 (90.3)	0.382
Surgical procedures, no. (%)	144 (62.3)	40 (76.0)	00 (01.9)	26 (90.3)	0.004
None	22 (12.6)	13 (21.7)	9 (10.8)	0	0.001
One	77 (44.3)	25 (41.7)	42 (50.6)	10 (32.3)	
Two or more	75 (43.1)	22 (36.7)	32 (38.6)	21 67.7)	
Underlying illnesses, no. (%)					
Diabetes mellitus	41 (23.3)	14 (23.0)	23 (27.4)	4 (12.9)	0.264
Chronic liver failure	12 (6.8)	5 (8.2)	5 (6.0)	2 (6.5)	0.866
Hematologic malignancy	3 (1.7)	0	1 (1.2)	2 (6.5)	0.069
Chronic renal failure	17 (9.7)	4 (6.6)	9 (10.7)	4 (12.9)	0.562
Heart failure (NYHA, class III, IV) Solid tumor	6 (3.4) 35 (19.9)	2 (3.3) 9 (14.8)	3 (3.6) 23 (27.4)	1 (3.2) 3 (9.7)	0.994 0.050
Chronic obstructive pulmonary disease	30 (17.0)	12 (19.7)	13 (15.5)	5 (16.1)	0.030
Alcoholism	30 (17.0)	15 (24.6)	11 (13.1)	4 (12.9)	0.153
Risk factors, no. (%)	(-7.0)	(=)	()	. (/	
Urinary catheter	173 (98.3)	61 (100)	82 (97.6)	30 (96.8)	0.424
Central venous catheter	176 (100)	61 (100)	84 (100)	31 (100)	
Mechanical ventilation	151 (85.8)	51 (83.6)	71 (84.5)	29 (93.5)	0.391
Broad spectrum antibiotics	172 (97.7)	60 (98.4)	82 (97.6)	30 (96.8)	0.886
Arterial catheter	148 (84.1)	51 (83.9)	71 (84.5)	26 (83.9)	0.988
Enteral nutrition	76 (43.2)	20 (32.8)	39 (46.4)	17 (54.8)	0.092
Total parenteral nutrition	166 (94.3)	56 (91.8)	80 (95.2)	30 (96.8)	0.548

Table 1 continued

	All patients $(n = 176)$	Study groups			P value
		Neither colonized nor infected (n = 61)	Candida spp. colonization $(n = 84)$	Invasive candidiasis $(n = 31)$	
Corticosteroids Renal replacement therapy Selective digestive decontamination	74 (42.0) 45 (25.6) 37 (21.0)	21 (34.4) 13 (21.3) 11 (18.0)	38 (45.2) 20 (23.8) 19 (22.6)	15 (48.4) 12 (38.7) 7 (22.6)	0.314 0.171 0.778
(no antifungal drugs) Antifungal treatment	53 (30.1)	4 (6.6)	21 (25.0)	28 (90.3)	< 0.001

ICU intensive care unit, IC invasive candidiasis, APACHE II Acute Physiology and Chronic Health Evaluation, CS Candida score, SOFA Sequential Organ Failure Assessment, GI gastrointestinal, NYHA New York Heart Association

Table 2 Characteristics of the 31 patients with invasive candidi- CART predictive model asis (IC)

	No. patients
Abdominal conditions	
Gastroduodenal perforation	9
Pancreatitis	5
Intestinal perforation (large intestine)	9 5 5 3 3 3
Cholecystitis	3
Peritonitis with interloop abscesses	3
Anastomotic leakage	3
Postoperative complex complicated	2
abdominal surgery	
Liver abscess	1
Multifocal Candida spp. colonization	28
Candidemia	7 (22.6 %)
Catheter-related candidemia	2
Candidal peritonitis	23 (74.2 %)
Candidemia and candidal peritonitis	1 (3.8 %)
Causative <i>Candida</i> spp.	
C. albicans	15
C. tropicalis	6
C. glabrata	7
C. krusei	1
C. lusitanie	1
$C.\ albicans + C.\ glabrata$	1
Time between ICU admission and	9 (5–20)
IC diagnosis, days, median (IQR)	
Patients with candidemia	13 (10–22)
Patients with candidal peritonitis	7 (4–17)
Antifungal treatment	28
Azoles	10
Echinocandins	8
Combination of 2 or 3 antifungals	10
(second choice therapy)	

ICU intensive care unit, IQR interquartile (25th-75th percentile) range

post-treatment levels was zero for CAGTA and 0 (IQR -37; 31) for BDG. Also, serum levels of CRP and PCT were similar in the three study groups (Table 3).

The CART decision tree model showed a probability of IC of 59.3 % in the terminal node of BDG with levels greater than 259 pg/mL and a probability of 30.8 % for the combination of BDG less than 259 pg/mL and CAGTA positive results. In the presence of BDG levels less than 259 pg/mL and CAGTA negative results, the probability of having IC was only 6.1 % (Fig. 2). Using a cutoff of 30 % for the probability of IC, chosen by the CART algorithm to minimize the deviance, the resulting prediction rule showed a 90.3 % sensitivity, 54.8 % specificity, 42.4 % positive predictive value, and 93.9 % negative predictive value. This prediction rule was better than for the individual analysis of BDG, CAGTA, and CS (Table 4). The AUC of the ROC curve was 0.78 (95 % CI 0.76-0.81). However, a cutoff point of serum BDG of 80 pg/mL showed much worse predictive values with 67.7 % sensitivity, 54.7 % specificity, 35.5 % positive predictive value, and 82.1 % negative predictive value. The total number of patients with Candida spp. colonization was 115. The number of patients that would have received unnecessary antifungal treatment (false positives) was 38 (33.3 %). However, among the 31 patients with IC, there were only three false negatives. Table 5 shows the application of the CART-derived prediction rule.

Discussion

The combination of two biomarkers, BDG and CAGTA, allowed a novel structured diagnostic approach to IC in patients with SAC. This finding is clinically relevant not only for the population to which the prediction rule can be applied, but also because it is very easy to remember and may help intensivists to decide when to start antifungal treatment.

^{*} Colonized versus IC groups; median (25th-75th percentiles)

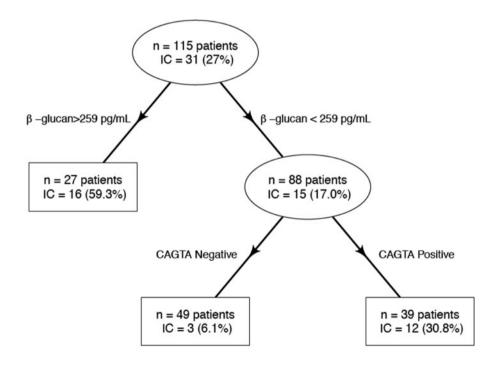
Table 3 Candida score and serological biomarkers in the first control and maximum values before or during the development of invasive candidiasis or the highest value when invasive candidiasis did not develop

	Assessment	Neither colonized nor infected $(n = 61)$	Candida spp. colonization $(n = 84)$	Invasive candidiasis $(n = 31)$	P value
Candida score	1st	2 (2–4)	3 (2–4)	4 (2–5)	< 0.001
	Max.	2 (2–4)	4 (3–5)	5 (3–5)	< 0.001
$(1\rightarrow 3)$ - β -D-glucan (pg/mL)	1st	9 (9–91)	45 (9–152)	54 (9–350)	0.110
	Max.	52 (9–145) ^a	66 (21–168) ^a	268 (50–444) ^b	0.003
CAGTA (%)	1st	25.0	25.0	43.3	0.129
	Max.	39.3 ^a	41.7 ^a	71.0 ^b	0.009
C-reactive protein (mg/L)	1st	201 (103–334)	207 (99–335)	172 (107–282)	0.911
	Max.	248 (142–373)	241 (125–383)	283 (177–426)	0.411
Procalcitonin (ng/mL)	1st	0.89 (0.2–3.21)	0.58 (0.23–5.48)	1.11 (0.29–6.14)	0.590
	Max.	1.25 (0.33–5.0)	0.59 (0.3–7.14)	3.33 (0.74–6.34	0.179

Values are medians and interquartile range (25th–75th percentile) CAGTA Candida albicans germ tube antibodies, 1st first control, Max maximum value before or during invasive candidiasis (when invasive candidiasis did not develop, the highest of all observed values was recorded)

 a,b Superscripts indicate significant differences (P < 0.05) among rates of BDG and positive CAGTA values among the three study

Fig. 2 Prediction rule for the diagnosis of invasive candidiasis (IC) in nonneutropenic adult critically ill patients with severe abdominal conditions. Each terminal node shows the probability of the predicted event



IC: invasive candidiasis.

BDG: $[1\rightarrow 3]$ - β -D-glucan;

CAGTA: Candida albicans germ tube antibodies.

n= number of patients with IC.

reported to be associated with other fungi, such as cular catheters, hemodialysis, and administration of

A few studies have investigated the predictive value of *Pneumocystis jirovecii* infections [26], Gram-positive and BDG on IC in non-neutropenic critically ill patients Gram-negative bloodstream infections, exposure to gauze [21-25]. Elevated concentrations of BDG have been or other materials that contain glucans, biofilms on vas-

Table 4 Diagnostic accuracy of CART-derived prediction rule, BDG (cutoff, >259 pg/mL), CAGTA (cutoff, any positive value), and CS for the diagnosis of invasive candidiasis

	Area under ROC	Sensitivity %	Specificity %	Predictive value		
	curve (95 % CI) ^a	(95 % CI)	(95 % CI)	Positive % (95 % CI)	Negative % (95 % CI)	
CART analysis BDG CAGTA CS	0.78 (0.76–0.81) 0.66 (0.59–0.74) 0.67 (0.64–0.71) 0.62 (0.58–0.66)	90.3 (75.1–96.6) 51.6 (34.8–68.0) 71.0 (53.4–83.9) 93.5 (79.2–98.2)	54.8 (44.1–65.0) 86.9 (78.0–92.5) 57.3 (46.5–67.5) 18.1 (11.3–27.7)	42.4 (31.2–54.4) 59.3 (40.7–75.5) 38.6 (27.1–51.6) 29.9 (21.7–39.6)	93.9 (83.5–97.9) 83.0 (73.8–89.4) 83.9 (72.2–91.3) 88.2 (65.7–96.7)	

Total number of patients with Candida spp. colonization = 115

CART classification and regression tree analysis, BDG beta-p-glucan, CAGTA Candida albicans germ tube antibody, CS Candida score
^a Original scale

Table 5 CART-derived prediction rule applied for all the study population

	Node BDG <259 and CAGTA negative	Node BDG <259 and CAGTA positive	BDG >259	Total
Neither colonized nor infected, n (%)	31 (50.8)	18 (29.5)	12 (19.7)	61
Candida spp. colonization, n (%)	46 (54.8)	27 (32.1)	11 (13.1)	84
Invasive candidiasis, n (%)	3 (9.7)	12 (38.7)	16 (51.6)	31
Total, <i>n</i> (%)	80 (45.5)	57 (32.4)	39 (22.2)	176

CART classification and regression tree analysis, BDG beta-D-glucan, CAGTA Candida albicans germ tube antibody

intravenous immunoglobulins, albumin, coagulation factors, and plasma protein fractions [27–29]. In addition, Pickering et al. [29] showed in a cross-contamination experiment that excess manipulation of a sample can result in its contamination with BDG. In our IC patients, 12 (38.7 %) had renal replacement therapy, 3 (9.6 %) had Gram-positive bacteremia, and 3 (9.6 %) received betalactam antibiotics. None of them presented Gram-negative bacteremia or received immunoglobulins, albumin, coagulation factor, or plasma protein fractions. However, Candida multifocal colonization was documented in 28 (90.3 %) of the IC patients and occult candidemia cannot be totally excluded. In a retrospective study performed in patients admitted for 7 or more days to nine ICUs, it has been shown that those patients with IC had mean concentrations of BDG, measured twice weekly, persistently high over time that decreased slowly after approximately 4 weeks [30]. Our results showed that CAGTA and BDG values are apparently not affected by antifungal treatment. We observed that CAGTA and BDG responses after antifungal treatment were quite diverse and this explains why the median value before and after treatment was zero. Although it seems that antifungal use did not impact the performance of BDG [31], biomarker kinetics in this scenario and particularly responses after starting antifungal treatment deserve further investigation.

There are only two previous reports of the same group [9, 10] assessing the predictive value of CAGTA for IC in a cohort of 53 critically ill non-neutropenic patients in which CAGTA was measured twice a week. Twenty-two patients (41.5 %) had CAGTA-positive results and none of them had a positive blood culture for Candida. The authors concluded that CAGTA detection may be important for the diagnosis of IC in ICU surgical patients. This test provides a rapid and quite simple laboratory IC diagnosis with a sensitivity of 84.4 % and a specificity of 94.7 % and it has been claimed to be independent of Candida colonization or antifungal treatment. Although CAGTA is designed for C. albicans germ tube antibody detection we also realized that it works for non-albicans species, which is an interesting finding when diagnosing an IC in the critical ill setting.

Although previous studies have shown the usefulness of BDG for the diagnosis of IC in surgical patients [6, 22, 25], our findings confirm that BDG is highly specific for a cutoff value of 259 pg/mL but with a very low sensitivity (51.6 %). CAGTA (negative/limit-positive) has a better sensitivity than BDG but its specificity is lower. Our predictive rule based on CART that includes both biomarkers demonstrates a sensitivity of 90.3 %, which is greater than the sensitivities found in both biomarkers, and an adequate specificity. When we analyzed the

dynamics of the most representative cases of both biomarkers, there were 13 cases in which BDG was lower than the cutoff value of 259 pg/mL and in 11 of them, CAGTA was positive, thus showing the complementariness of both biomarkers expressed by means of the CART analysis.

This study shows that BDG levels greater than 259 pg/ mL combined with CAGTA-positive results accurately discriminate Candida spp. colonization from IC in nonneutropenic critically ill patients with SAC. The predictive rule rests on BDG levels, which is the main node of the tree, and when these levels are greater than 259 pg/ mL, the patients have a nearly 60 % probability of having an IC. Moreover, using a cutoff of 30 % for the probability of IC, the CART-derived decision model had a very high negative predictive value, with a 93.9 % probability of not having IC for BDG less than 259 pg/mL combined with CAGTA-negative results. Other commonly used biomarkers, such as CRP and PCT, were not useful to distinguish between IC and *Candida* spp. colonization. In agreement with previous studies, the *Candida* score was significantly higher in patients with IC than in the remaining groups, but the usefulness of this prediction rule is limited in patients undergoing abdominal surgery because in these circumstances the score is consistently higher than the discriminating cutoff point of at least 3. The CART analysis showed a very high negative predictive value, which is in agreement with data obtained by Ostrosky-Zeichner et al. [3] with their prediction rule as well as with the CS in our two previous studies [2, 4]. However, the CART positive predictive value was 42.4 %, which is clearly higher than 9 % obtained by the prediction rule of Ostrosky-Zeichner et al. [3] and 13.8 % with CS in our previous study [4].

In patients admitted to an ICU with SAC if serum BDG levels are over 259 pg/mL or if BDG levels are lower than 259 pg/mL but CAGTA levels are positive, the IC risk is 59.3 and 30.8 %, respectively. Despite the 42.4 % positive predictive value of the CART model, we think that in this specific scenario clinicians should start AF therapy. This means that for every 100 treated patients, the expected number of overtreated would be 57.6. On the contrary, in similar patients with a BDG level lower than 259 pg/mL and negative CAGTA limits, no AF treatment should be started owing to the very high negative predictive value shown by the CART analysis even though for every 100 patients with IC, the expected number of untreated patients would be of 9.7.

Potential limitations are as follows: BDG testing was performed in batches or frozen samples and we cannot exclude that some negative results may have stemmed from sample instability. Also, the presence of intestinal mucositis may facilitate the translocation of *Candida* spp. through the gastrointestinal barrier and eventually might interfere with BDG determinations [24]. Although CAGTA was developed for *C. albicans* detection, it may

be also useful for the diagnosis of IC in cases of nonalbicans strains. However, our study population was not sufficiently large for a reliable analysis of the differences between *C. albicans* and non-albicans strains.

In summary, non-neutropenic critically ill patients with SAC at ICU entry and an expected stay of at least 7 days, BDG greater than 259 pg/mL with a CAGTA-positive result accurately differentiated *Candida* spp. colonization from IC. Further studies are needed to confirm the present data.

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Appendix

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