

Multinational Observational Cohort Study of COVID-19–Associated Pulmonary Aspergillosis¹

Nico A.F. Janssen, Rémy Nyga, Lore Vanderbeke, Cato Jacobs, Mehmet Ergün, Jochem B. Buil, Karin van Dijk, Josje Altenburg, Catherine S.C. Bouman, Hans I. van der Spoel, Bart J.A. Rijnders, Albert Dunbar, Jeroen A. Schouten, Katrien Lagrou, Marc Bourgeois, Marijke Reynders, Niels van Regenmortel, Lynn Rutsaert, Piet Lormans, Simon Feys, Yves Debavaye, Fabienne Tamion, Damien Costa, Julien Maizel, Hervé Dupont, Taieb Chouaki, Saad Nseir, Boualem Sendid, Roger J.M. Brüggemann, Frank L. van de Veerdonk, Joost Wauters,² Paul E. Verweij²

We performed an observational study to investigate intensive care unit incidence, risk factors, and outcomes of coronavirus disease–associated pulmonary aspergillosis (CAPA). We found 10%–15% CAPA incidence among 823 patients in 2 cohorts. Several factors were independently associated with CAPA in 1 cohort and mortality rates were 43%–52%.

Incidence of coronavirus disease (COVID-19)–associated pulmonary aspergillosis (CAPA) in hospital intensive care units (ICUs) is 3.8%–33.3% (1–9). Variations might be explained by differences in patient populations and CAPA definitions used, complicating direct comparisons between studies.

Diagnosing CAPA is complex because cases frequently lack typical radiologic features and European Organization for Research and Treatment of Cancer and the Mycoses Study Group Education and Research Consortium (EORTC/MSGERC) host factors (10) and because mycologic evidence is difficult to obtain. Serum galactomannan (GM) detection has low sensitivity in CAPA (7,10).

The European Confederation of Medical Mycology and International Society for Human and Animal Mycology (ECMM/ISHAM) published consensus criteria for a CAPA definition (11). We used these criteria to perform an observational cohort study to assess CAPA incidence in patients with COVID-19 admitted to ICUs during the first wave of the COVID-19 pandemic.

Author affiliations: Radboud University Medical Center, Nijmegen, the Netherlands (N.A.F. Janssen, M. Ergün, J.B. Buil, J.A. Schouten, R.J.M. Brüggemann, F.L. van de Veerdonk, P.E. Verweij); Amiens University Hospital, Amiens, France (R. Nyga, J. Maizel, H. Dupont, T. Chouaki); University Hospitals Leuven, Leuven, Belgium (L. Vanderbeke, C. Jacobs, K. Lagrou, Y. Debavaye, J. Wauters); Katholieke Universiteit Leuven, Leuven (L. Vanderbeke, K. Lagrou, J. Wauters); Amsterdam University Medical Centers, Amsterdam, the Netherlands (K. van Dijk, J. Altenburg, C.S.C. Bouman, H. van der Spoel); Erasmus Medical Center, Rotterdam, the Netherlands (B.J.A. Rijnders, A. Dunbar); AZ Sint-Jan Brugge-Oostende, Brugge, Belgium (M. Bourgeois, M. Reynders); ZNA Campus Stuivenberg, Antwerpen, Belgium (N. van Regenmortel, L. Rutsaert); AZ Delta Hospital, Roeselare, Belgium (P. Lormans, S. Feys); Rouen University Hospital, Rouen, France (F. Tamion, D. Costa); Lille University Hospital, Lille, France (S. Nseir, B. Sendid); University of Lille, Lille (S. Nseir, B. Sendid)

The Study

We collected partially prospective and partially retrospective data for 823 patients in 2 cohorts. The discovery cohort comprised patients with PCR-confirmed or clinically presumed COVID-19 admitted to 4 ICUs in the Netherlands and 4 ICUs in Belgium during February 28–May 27, 2020. The validation cohort comprised patients with PCR-confirmed COVID-19 admitted because of respiratory insufficiency to 3 participating ICUs in France during April 7–May 31, 2020 (Appendix Methods, Table 1, <https://wwwnc.cdc.gov/EID/article/27/11/21-1174-App1.pdf>).

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²These authors were co–principal investigators.

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Table 1. Demographic, clinical, and mycologic characteristics of the discovery cohort in a multinational observational study of COVID-19–associated pulmonary aspergillosis in 3 countries in Europe, 2020*

Characteristics	Total population, n = 519	CAPA, n = 42	CAPA excluded, n = 237	p value
Age, y	64 (55–72)	68 (61–73)	65 (57–71)	0.12
Sex				
F	141 (27)	8 (19)	58 (24)	
M	378 (73)	34 (81)	179 (76)	0.56
BMI, kg/m ²	27.2 (24.4–31.0); n = 507	27.4 (23.6–30.2); n = 40	26.9 (24.4–30.9); n = 231	0.72
Underlying conditions				
Cardiovascular disease†	291 (56)	25 (60)	130 (55)	0.62
Diabetes mellitus	139 (27)	9 (21)	61 (26)	0.70
Asthma	37 (7)	1 (2)	19 (8)	0.33
COPD	44 (9)	8 (19)	19 (8)	0.042
Liver cirrhosis	6 (1)	0	2 (0.8)	1.00
Rheumatological disease	31 (6)	5 (12)	14 (6)	0.18
HIV/AIDS	6 (1)	3 (7)	1 (0.4)	0.011
Solid organ malignancy	28 (5)	3 (7)	11 (5)	0.45
EORTC/MSGERC host factors				
Any‡	70 (16); n = 426	13 (33); n = 39	31 (19); n = 166	0.053
Recent neutropenia§	7 (2); n = 413	1 (3); n = 38	5 (3); n = 156	1.00
Hematologic malignancy	18 (4)	4 (10)	9 (4)	0.11
Receipt of allogeneic SCT	4 (0.8); n = 516	0	3 (1); n = 236	1.00
Receipt of SOT	6 (1)	1 (2)	2 (0.8)	0.39
Systemic corticosteroids ≤30 d before ICU admission, any dose	38 (9); n = 430	7 (18); n = 39	14 (9); n = 160	0.14
T or B cell immunosuppressants other than corticosteroids ≤90 d before ICU admission	31 (6); n = 514	7 (17)	12 (5); n = 233	0.014
Inherited severe immunodeficiency	0; n = 517	0	0; n = 236	NA
ICU treatment data				
Invasive mechanical ventilation	423 (82); n = 517	40 (98); n = 41	225 (95)	0.70
No. invasive ventilation days¶	14 (9–24); n = 395	16 (13–27); n = 37	18 (11–30); n = 212	0.98
RRT during ICU admission	93 (18); n = 516	17 (41)	44 (19); n = 236	0.004
Systemic corticosteroids during ICU admission	216 (42); n = 516	20 (48)	131 (56); n = 236	0.40
Outcome data				
ICU death	154 (30); n = 518	22 (52)	81 (34)	0.036
ICU LOS, d#	14 (8–24); n = 491	18 (12–27); n = 39	20 (12–32); n = 222	0.84
Mycologic diagnostic tests				
Serum GM OD >0.5, no. positive (%); no. values reported/no. performed	3 (2); 134/176	3 (11); 28/28	0; 106/148	NA
Serum GM OD**	0.10 (0.10–0.10); n = 134	0.10 (0.06–0.14); n = 28	0.10 (0.10–0.10); n = 106	0.95
Positive BALF/BL culture	17 (10); n = 166	17 (42); n = 41	0; n = 125	NA
BALF/BL GM OD ≥1.0, no. positive (%); no. OD values reported/no. BL/BALF performed	32 (19); 90/166	32 (78); 34/41	0; 55/125	NA
BALF/BL GM OD**	0.20 (0.10–1.50); n = 90	1.80 (1.00–3.90); n = 35	0.10 (0.10–0.20); n = 55	<0.001
Positive BALF/BL PCR, any C _t , no. positive (%); no. reported/no. tested	9 (5); 11/166	7 (17); 7/41	2 (2); 4/125††	NA
Days between ICU admission and first positive mycologic test‡‡	NA	6 (3–9); n = 41	NA	NA

*Data are presented as no. (%) or median (IQR) unless otherwise indicated. Continuous variables were compared by Mann-Whitney U test, categorical variables by Fisher exact test with omission of missing data, unless stated otherwise. Total percentages might not equal 100% because of rounding. Bold text indicates statistical significance. BAL, bronchoalveolar lavage; BALF, BAL fluid; BL, bronchial lavage; BMI, body mass index; CAPA, COVID-19–associated pulmonary aspergillosis; COPD, chronic obstructive pulmonary disease; COVID-19, coronavirus disease; C_t, cycle threshold; CT, computed tomography; ECMO, extracorporeal membrane oxygenation; EORTC/MSGERC, European Organization for Research and Treatment of Cancer and the Mycoses Study Group Education and Research Consortium; GM, galactomannan; ICU, intensive care unit; IQR, interquartile range; LOS, length of stay; NA, not applicable; NBL, nonbronchoscopic lavage; OD, optical density; RRT, renal replacement therapy; SCT, stem cell transplantation; SOT, solid organ transplant.

†Includes hypertension

‡Includes any use of systemic corticosteroids before ICU admission; If data on one or more EORTC host factors were missing, then data were regarded as missing for this variable.

§Neutropenia includes absolute neutrophil count of <0.5 × 10⁹ cells/L for >10 d.

¶If transferred to another hospital from ICU and still on ventilatory support of any kind, duration of invasive mechanical ventilatory support was regarded as missing data and not included in the analyses. The same holds true for those who received a tracheostomy for a prolonged weaning trajectory.

#Data on ICU LOS were regarded as missing if transfer to another hospital was the reason for ICU discharge because exact ICU LOS was unknown.

**When multiple values were reported for 1 patient, the median of these values was used for further calculations.

††Positive PCR with C_t values ≥36 as only positive mycologic criterion.

‡‡Mycologic test considered a criterion for proven, probable, or possible CAPA according to the 2020 European Confederation for Medical Mycology/International Society for Human and Animal Mycology classification (11).

We applied ECMM/ISHAM classification criteria for CAPA (11). We considered bronchial lavage (BL) equivalent to bronchoalveolar lavage (BAL). We assumed all CAPA classified patients demonstrated clinical factors and radiographic abnormalities. We defined 3 patient groups: CAPA, CAPA-excluded, and CAPA not classifiable (Figure 1; Appendix).

We included 519 patients in the discovery cohort; median age was 64 years, 73% were male, and 82% required invasive mechanical ventilation during ICU admission (Table 1; Appendix Table 2, 3, 4). Among patients in the discovery cohort, 279 (54%) were classifiable: 6 (2%) as CAPA proven, 32 (12%) as probable CAPA, and 4 (1%) as possible CAPA (Figure 1, panel A; Appendix Results, Tables 5, 6). CAPA incidence among classifiable patients was 15% (42/279); 85% were CAPA-excluded. Among patients in the discovery cohort, 46% (240/519) were not classifiable, including 3 who did not fulfill the criteria for possible CAPA (Figure 1, panel A). In patients with any EORTC/MSGERC host factor, CAPA incidence was 30% (13/44), compared with 16% (26/161) in patients with no host factors ($p = 0.053$).

Chronic obstructive pulmonary disease (COPD; $p = 0.04$) and HIV/AIDS ($p = 0.01$) were more

prevalent in CAPA patients (Table 1; Appendix Table 2). Among CAPA patients, 33% had ≥ 1 EORTC/MSGERC host factor, compared with 19% of CAPA-excluded patients ($p = 0.053$). Corticosteroid use was not more prevalent in the CAPA group ($p = 0.14$), in contrast to other immunosuppressant drugs ($p = 0.01$). In logistic regression analysis, corticosteroid use at any dose before or during ICU admission was not independently associated with CAPA development. However, COPD, HIV/AIDS, and use of other immunosuppressant drugs before ICU admission were associated with CAPA (Appendix Figure 1, panel A).

Among CAPA patients who underwent BAL or BL, *Aspergillus* culture was positive in 42%, GM was positive (optical density [OD] ≥ 1.0) in 78%, and *Aspergillus* PCR was positive in 17%. Among CAPA patients who underwent nonbronchoscopic lavage, 67% had positive cultures. Serum GM was positive in 11% of tested CAPA patients. Median time between ICU admission and first positive mycologic test was 6 (interquartile range [IQR] 3–9) days (Table 1; Appendix Table 7).

The proportion of patients receiving systematic corticosteroid treatment in ICUs was not

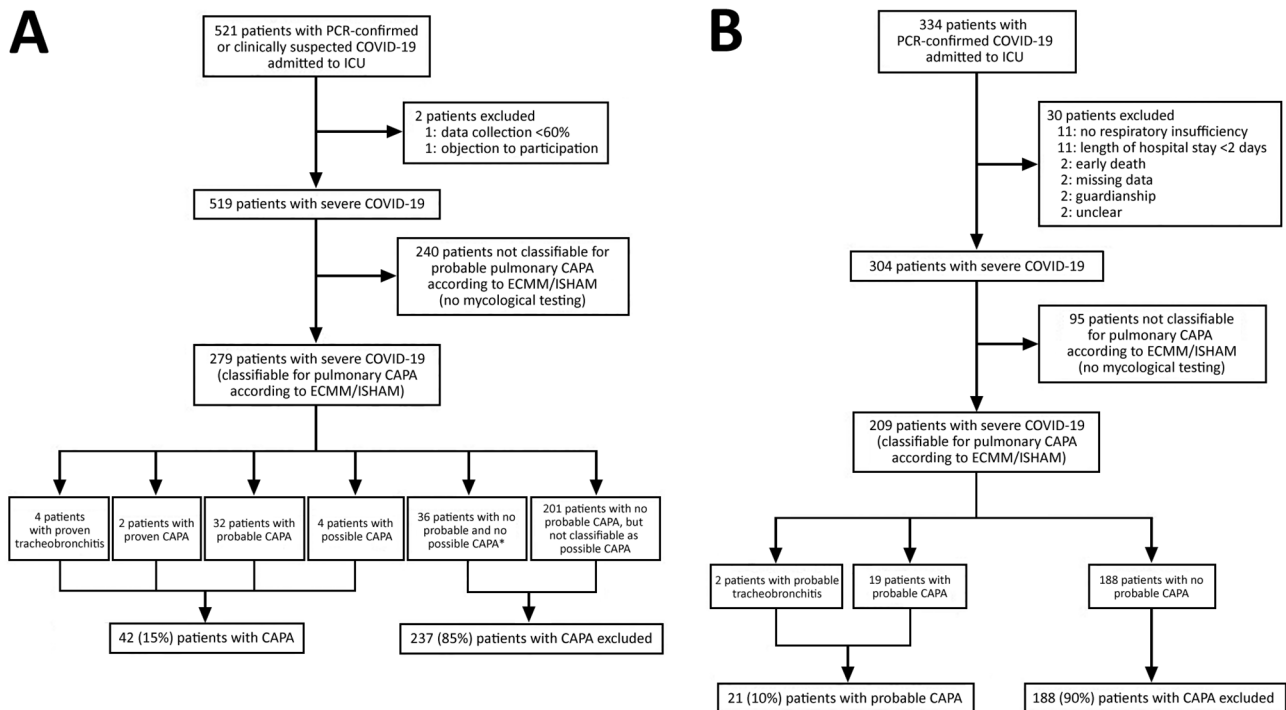


Figure 1. Flowchart of the study inclusion process for a multinational observational study of CAPA in 3 countries in Europe, 2020.

A) Discovery cohort; B) validation cohort. For further analyses, patients with proven, probable, and possible CAPA were designated to the CAPA group. Patients were classified to the CAPA excluded group when they had ≥ 1 negative mycological test according to 2020 ECMM/ISHAM classification consensus criteria (11). Patients who did not undergo any of the mycological tests were designated to the CAPA not classifiable group. *Value includes 6 patients in whom CAPA was excluded at the time of autopsy. CAPA, COVID-19–associated pulmonary aspergillosis; COVID-19, coronavirus disease; ECMM/ISHAM, European Confederation for Medical Mycology/International Society for Human and Animal Mycology; ICU, intensive care unit.

significantly different between CAPA and CAPA-excluded groups ($p = 0.40$), nor was corticosteroid dose ($p = 0.88$) (Table 1; Appendix Table 4). Antifungal treatment was administered to 16% (83/519) of patients, 88% of CAPA patients, and 15% of CAPA-excluded patients (Appendix Table 8). ICU mortality rates were significantly higher in CAPA patients (52%) than in CAPA-excluded patients (34%) ($p = 0.04$; Table 1; Appendix Table 4); mortality rates were 67% for patients with positive serum GM. CAPA patients demonstrated reduced survival ($p = 0.02$) (Figure 2, panel A); estimated median survival was 42 days after ICU admission. When correcting for covariates, CAPA was not independently associated with ICU mortality rates, but older age and acute kidney injury (AKI) during ICU stay were (Appendix Figure 1, panel B).

We included 304 patients in the validation cohort (Figure 1, panel B); median age was 63 years, 25% were male, and 76% required invasive mechanical ventilation (Table 2; Appendix Tables 9, 10). Ultimately, 209/304 (69%) patients were classifiable for CAPA: 21 (10%) probable CAPA and 188 (90%) CAPA excluded (Figure 1, panel B; Appendix Results, Tables 5, 11). Among patients with EORTC/MSGERC host factors, CAPA incidence was 13% (3/23), compared with 10% (18/186) among patients without host factors ($p = 0.71$).

All 21 probable CAPA patients were female; cardiovascular disease, excluding hypertension ($p = 0.02$), and bronchiectasis ($p = 0.03$) were more prevalent in this group (Table 2; Appendix Table 9). Use of corticosteroids before or during ICU admission or other immunosuppressant drugs before ICU admission were not independently associated with CAPA (Appendix Figure 1, panel C). In the validation cohort, 19% received antifungal treatment;

57% of the CAPA group received antifungal treatment (Appendix Table 8).

Corticosteroid use during ICU stay was not significantly different between the CAPA and CAPA-excluded groups ($p = 0.82$) in the validation cohort. ICU mortality rates were higher in the CAPA group than the CAPA-excluded group (43% vs. 25%; $p = 0.12$) (Table 2; Figure 2, panel B; Appendix Table 10). The ICU mortality rate was 50% in patients with positive serum GM. CAPA was not independently associated with ICU death, but older age and AKI during ICU admission were (Appendix Table 10, Figure 1, panel D).

Conclusions

We found CAPA incidence was 10%–15%, corresponding to the 14%–19% reported in other studies (8,9). Discovery cohort CAPA incidence was similar to influenza-associated pulmonary aspergillosis (IAPA) incidence in ICUs (12,13). CAPA seems to develop later after ICU admission than IAPA. Median time to first positive mycologic test in our study was 6 days after ICU admission, similar to other studies reporting 4–8 days (7–9) but in contrast to the median 3 days reported for IAPA (12,14).

Corticosteroids were not associated with CAPA in our study, consistent with previous reports (7–9), but contrasting associations seen with invasive pulmonary aspergillosis (IPA) and IAPA (12). This finding might be explained by possible dual effects of corticosteroids in COVID-19, impairing anti-*Aspergillus* immunity while simultaneously ameliorating the hyperinflammatory immune dysregulation and associated tissue damage conducive to IPA.

We found CAPA ICU mortality rates were 43%–52%, in line with previous reports (7–9) and

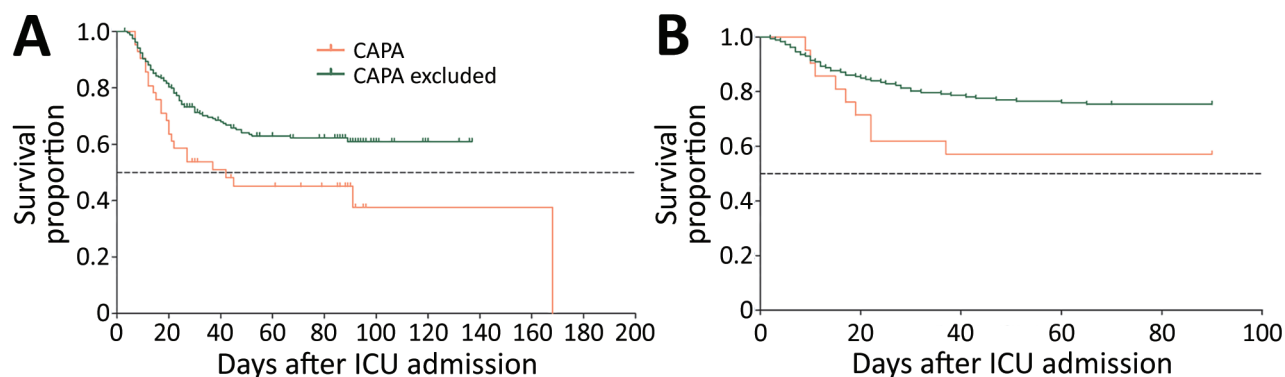


Figure 2. Kaplan-Meier survival curves comparing patients with CAPA and those classified as CAPA excluded in a multinational observational study. A) Discovery cohort; B) validation cohort. Survival analysis performed by using Mantel-Cox log rank test. Survival over time differs significantly in the discovery cohort ($n = 279$); median estimated survival in the CAPA group is 42.0 days ($p = 0.015$ by log rank test). In the validation cohort ($n = 209$), survival over time is not significantly different between the 2 groups ($p = 0.065$ by log rank test). CAPA, COVID-19–associated pulmonary aspergillosis; COVID-19, coronavirus disease; ICU, intensive care unit.

comparable to those for IAPA (12). We could not assess antifungal treatment effects on mortality rates, but CAPA patients in the validation cohort who received antifungal treatment demonstrated a trend toward improved survival (Appendix Figure 2).

Table 2. Demographic, clinical, and mycologic characteristics of the validation cohort in a multinational observational study of COVID-19-associated pulmonary aspergillosis in 3 countries in Europe, 2020*

Characteristics	Total population, n = 304	CAPA, n = 21	CAPA excluded, n = 188	p value
Age, y	63 (55–71)	67 (59–75)	62 (53–69)	0.06
Sex				
F	227 (75)	21 (100)	141 (75)	
M	77 (25)	0	47 (25)	0.005
BMI, kg/m ²	30.0 (26.0–34.4); n = 296	30.2 (26.1–32.8); n = 20	30.0 (26.4–34.5); n = 185	0.84
Underlying conditions				
Active hematologic malignancy	10 (3)	0	6 (3)	1.00
Cardiovascular disease†	185 (61)	17 (81)	112 (60)	0.06
Diabetes mellitus	92 (30)	9 (43)	62 (33)	0.47
Asthma	22 (7)	2 (10)	12 (6)	0.64
COPD	20 (7)	2 (10)	12 (6)	0.64
Liver cirrhosis‡	5 (2)	2 (10)	2 (1)	0.051
Autoimmune disease	16 (5)	2 (10)	11 (6)	0.63
HIV/AIDS	3 (1)	0	1 (0.5)	1.00
Active solid organ malignancy	4 (1)	1 (5)	3 (2)	0.35
Bronchiectasis	5 (2)	2 (10)	1 (0.5)	0.027
EORTC/MSGERC host factors				
Any§	35 (12)	3 (14)	20 (11)	0.71
Recent neutropenia¶	0; n = 303	0	0; n = 187	NA
Hematological malignancy	10 (3)	0	6 (3)	1.00
Receipt of SOT	9 (3)	1 (5)	5 (3)	0.48
Corticosteroids ≥0.3 mg/kg for ≥3 wks within previous 60 d	17 (6)	2 (10)	10 (5)	0.34
Other immunosuppressants <90 d before ICU admission	23 (8)	2 (10)	16 (9)	0.70
ICU treatment data				
Invasive mechanical ventilation	228 (76); n = 302	19 (95); n = 20	168 (89)	0.70
No. invasive ventilation days	15 (9–25); n = 212	18 (13–25); n = 17	15 (9–25); n = 157	0.21
RRT	64 (21); n = 303	11 (55); n = 20	47 (25)	0.008
Systemic corticosteroids during ICU admission	147 (49); n = 303	11 (52)	106 (57); n = 187	0.82
Outcome data				
ICU death	69 (23); n = 299	9 (43)	46 (25); n = 185	0.12
ICU LOS, d#	14 (8–26); n = 295	22 (12–35); n = 20	18 (10–28); n = 183	0.27
Mycologic diagnostic tests				
Serum GM OD >0.5	4 (2); n = 172**	4 (22); n = 18	0; n = 154††	NA
Serum GM OD	0.07 (0.04–0.12); n = 172**	0.10 (0.06–0.34); n = 18	0.06 (0.04–0.11); n = 154††	0.008
Positive BALF culture	11 (8); n = 135	11 (52) n = 21	0; n = 114	NA
BALF GM OD ≥1.0	13 (11); n = 123	13 (62) n = 21	0; n = 102	NA
BALF GM OD‡‡	0.12 (0.05–0.32); n = 123	1.10 (0.12–3.06); n = 21	0.11 (0.05–0.18); n = 102	<0.001
Positive BALF PCR, any Ct	8 (13); n = 64	8 (53); n = 15	0; n = 49	NA
Serum β-D-glucan value ≥80 pg/mL	37 (20); n = 184	8 (42); n = 19	29 (18); n = 160	0.030
Serum β-D-glucan value§§	31 (13–60); n = 184	34 (31–156); n = 19	31 (10–59); n = 160	0.055

*Data are presented as no. (%) or median (IQR), unless stated otherwise. Continuous variables were compared by Mann-Whitney U test, categorical variables by Fisher exact test with omission of missing data, unless stated otherwise. Total percentages might not equal 100% because of rounding. Bold text indicates statistical significance. BAL, bronchoalveolar lavage; BALF, BAL fluid; BL, bronchial lavage; BMI, body mass index; CAPA, COVID-19-associated pulmonary aspergillosis; COPD, chronic obstructive pulmonary disease; COVID-19, coronavirus disease; Ct, cycle threshold; CT, computed tomography; ECMO, extracorporeal membrane oxygenation; EORTC/MSGERC, European Organization for Research and Treatment of Cancer and the Mycoses Study Group Education and Research Consortium; ICU, intensive care unit; IQR, interquartile range; GM, galactomannan; LOS, length of stay; NA, not applicable; OD, optical density; RRT, renal replacement therapy; SAPS, simplified acute physiology score; SOT, solid organ transplant; TBA, tracheobronchial aspirate.

†Includes Factor V Leiden mutation and hypertension.

‡Includes hemochromatosis.

§Includes use of any systemic corticosteroids. We did not assess receipt of an allogeneic stem cell transplant, presence of an inherited severe immunodeficiency, and presence of acute graft-versus-host disease.

¶Neutropenia includes absolute neutrophil count of $<0.5 \times 10^9/L$ for >10 d.

#Data on ICU LOS were regarded as missing if still admitted at the time of data entry or if transfer to another hospital was the reason for ICU discharge.

**Serum GM performed in 173 patients, including 1 patient with an unknown result.

††Serum GM values known for 154 patients, unknown value in 1 patient.

‡‡One value of >6.0 entered as 6.0.

§§One value of >500 pg/mL entered as 500 pg/mL.

The first limitation of our study is that assuming clinical and imaging factors were available for all patients classified with CAPA possibly led to overreporting of CAPA. Excluding CAPA based on 1 negative mycologic test might have led to underreporting. Another limitation was that patients undergoing mycologic workup were likely more severely ill, which becomes apparent when comparing baseline and outcome data of the CAPA not classifiable group to the other 2 groups (Appendix Tables 5–12). Several classifications have been published or updated after we initiated this study; therefore, not all diagnostic modalities were evaluated, and we used some terms, such as BAL and BL, interchangeably (11,15).

In conclusion, we report CAPA incidence of 10%–15% in COVID-19 patients admitted to ICUs, CAPA ICU mortality rates of 43%–52%, and decreased survival over time. Clinicians should be aware of CAPA and that underlying factors, including COPD, immunosuppressant drugs other than corticosteroids, and HIV/AIDS, can increase the risk for CAPA.

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About the Author

Dr. Janssen is an infectious disease clinician and PhD candidate at the Radboud University Medical Center, Nijmegen, the Netherlands. His primary research interests are fungal diseases, including viral pneumonitis-associated invasive aspergillosis and chronic pulmonary aspergillosis.

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Address for correspondence: P.E. Verweij, Department of Medical Microbiology, Radboud University Medical Center, Geert Grooteplein-Zuid 10,6525 GA Nijmegen, the Netherlands; email: paul.verweij@radboudumc.nl

Multinational Observational Cohort Study of COVID-19–Associated Pulmonary Aspergillosis

Appendix

Additional Methods

Study Design and Participants

Data on the discovery cohort were largely collected prospectively and partially retrospectively; patients were admitted during February 28–May 27, 2020. Data on the validation cohort were collected retrospectively; patients in this cohort were admitted during April 7–May 31, 2020. The 2 cohorts were analyzed separately because of differences in inclusion criteria and data variables collected. The number of patients included per center varied (Appendix Table 1). Patients were followed for 90 days after ICU admission or until ICU discharge, whichever occurred last.

Ethics Statement

This study was performed in accordance with the latest version of the Declaration of Helsinki (<https://www.wma.net>), the International Conference on Harmonisation–Good Clinical Practice (ICH-GCP) guidelines (<https://ichgcp.net>) and local legislation and regulations. For all centers in the Netherlands, ethical approval was granted by the Ethics Board region Arnhem-Nijmegen; based on the observational nature of this study, informed consent was waived (approval no. CMO 2020–6339).

For centers in Belgium, ethical approval was granted by the University Hospitals Leuven Ethics Board (approval no. S64071) and local ethics boards of participating centers. Ethical approval for the separate study protocol for the sites in France was granted by the institutional committee of the Amiens University Hospital (registration no. PI2020_843_0028.). The study protocol was registered with the Commission Nationale de l’Informatique et des Libertés, France.

Data Collection

For the discovery cohort, local investigators collected pseudonymized patient data from medical files into Castor Electronic Data Capture (Castor EDC, <https://www.castoredc.com>) electronic case report form (eCRF). Pseudonymized data for the validation cohort were entered into a SharePoint eCRF (Department of Clinical Research, Amiens University Hospital, Amiens, France).

Definitions

Clinically presumed coronavirus disease 2019 (COVID-19) was based on clinical signs and symptoms and exposure to severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), as assessed by treating physicians or multidisciplinary COVID-19 team. COVID-19–associated pulmonary aspergillosis (CAPA) diagnosis was based on the 2020 European Confederation for Medical Mycology/International Society for Human and Animal Mycology (ECMM/ISHAM) consensus classification (1). We applied several necessary modifications needed because of the eCRF's design and differences in interpretation between participating centers regarding performance of bronchial lavage (BL) or bronchoalveolar lavage (BAL). Although a BL was intended to signify a nonbronchoscopic, nondirected bronchial lavage (NBL) and a BAL a directed, bronchoscopic lavage, these terms were used differently by several centers. To avoid missing any directed BALs performed, we regarded BLs as equivalent to BALs during data analysis. Due to missing data on clinical factors and radiological results in many patients, all patients classified with CAPA were considered to demonstrate these clinical characteristics and (infiltrative) abnormalities on thoracic imaging during ICU stay. Patients who underwent none of the mycological tests required for classification as proven or probable CAPA, including autopsy, were designated CAPA not classifiable. Patients who underwent mycological tests were further evaluated for the presence of CAPA. In the validation cohort, data were not available to classify patients as possible CAPA.

According to the 2020 ECMM/ISHAM classification (1), BL or BALF galactomannan (GM) results that were qualitatively positive without a known quantitative result were not regarded as a positive mycological result. Patients were classified into 3 defined groups for further analyses. The CAPA group comprised patients with proven, probable, or possible *Aspergillus* tracheobronchitis, pulmonary CAPA, or both. The CAPA excluded group comprised patients who underwent diagnostic workup, but had no evidence for proven, probable, or

possible CAPA, including no CAPA at autopsy, or patients without probable CAPA, but who were not classifiable as possible CAPA. The CAPA not classifiable group included patients who did not undergo any required mycological testing for proven or probable CAPA or those who were not classifiable for probable CAPA but had possible CAPA excluded.

We defined acute kidney injury (AKI) according to the Kidney Disease: Improving Global Outcomes (KDIGO; <https://kdigo.org>) criteria. AKI criteria include increase in serum creatinine of ≥ 26.5 $\mu\text{mol/L}$ (≥ 0.3 mg/dL) within 48 hours or increase in serum creatinine ≥ 1.5 times baseline values, which is known or presumed to have occurred within the prior 7 days or urine volume < 0.5 mL/kg/h for 6 hours (2). Prior to classification, correction of volume status and obstructive causes of AKI are allowed.

We also defined stages of AKI according to the KDIGO criteria. Patients were classified according to the criteria that resulted in the highest, that is the most severe, stage of injury. Stage 1 is increase in serum creatinine to 1.5–1.9 times baseline values or increase in serum creatinine by ≥ 26.5 $\mu\text{mol/L}$ (≥ 0.3 mg/dL) or reduction in urine output to < 0.5 mL/kg/hour for 6–12 hours. Stage 2 is increase in serum creatinine to 2.0–2.9 times baseline values, or reduction in urine output to < 0.5 mL/kg/hour for ≥ 12 hours. Stage 3 is increase in serum creatinine to 3.0 times baseline values, or increase in serum creatinine to ≥ 353.6 $\mu\text{mol/L}$ (≥ 4.0 mg/dL), or reduction in urine output to < 0.3 mL/kg/h for ≥ 24 hours, or anuria for ≥ 12 hours, or the initiation of renal replacement therapy.

Comparisons between CAPA Classification Criteria

CAPA diagnosis according to the 2020 ECMM/ISHAM classification was compared to the IAPA expert opinion case definition (3), the modified *Asp*ICU (*mAsp*ICU) classification (4,5), and the revised European Organization for Research and Treatment of Cancer and the Mycoses Study Group Education and Research Consortium (EORTC/MSGERC) classification (6). As we did for the ECMM/ISHAM classification, we presumed all patients classified with IPA had clinical factors, signs, symptoms, and pulmonary infiltrates on thoracic imaging present for all other classifications, if applicable. As such, only subset A of the IAPA expert opinion case definition criteria for probable IPA could be evaluated in our cohorts. For the *mAsp*ICU classification, BAL fluid (BALF) GM optical density (OD) ≥ 1.0 and serum GM OD ≥ 0.5 were added as mycological criteria for the diagnosis of putative IPA and positive *Aspergillus* BALF

culture was regarded as an entry criterion. For EORTC/MSGERC definitions, the eCRF was not designed to assess the presence of acute graft-versus-host-disease; therefore, we could not take this host factor into account. Due to frequent missing data regarding details on use of corticosteroids before admission to the ICU, all systemic corticosteroid use was considered a risk factor for invasive aspergillosis according to *mAspICU* and EORTC/MSGERC criteria. We compared the 2020 ECMM/ISHAM CAPA classification with the clinically reported occurrence of CAPA in patients' medical files, that is physician reported CAPA, regardless of fulfilment of any formal classification criteria.

Statistical Analysis

All data are expressed as no. (%) or median (interquartile range [IQR]). We compared variables by using Fisher exact test, Mann-Whitney U test, or Kruskal-Wallis test, as appropriate. We analyzed survival differences by using the Kaplan-Meier method and log rank test. We performed binary logistic regression analysis to detect independent predictors of CAPA occurrence and ICU death. In all cases, we considered $p < 0.05$ statistically significant. We did not apply corrections for multiple statistical testing during these analyses; readers should keep this in mind when interpreting results.

For binary logistic regression analysis, we used different independent variables as potential predictors for CAPA occurrence and ICU death. For CAPA occurrence, we used underlying conditions significantly more prevalent in the CAPA group in univariate analysis and corticosteroid use before and during ICU admission as independent variables. For ICU death, age, sex, AKI, renal replacement therapy (RRT), mechanical ventilation, use of vasopressors and/or inotropes and corticosteroids during ICU admission and presence of CAPA were used as independent variables or covariates. We performed statistical analyses by using SPSS Statistics for Windows version 25.0 (IBM Corp., <https://www.ibm.com>) or GraphPad Prism 5.03 for Windows (GraphPad Software Inc., <https://www.graphpad.com>).

Results

Discovery Cohort

Of the 521 patients admitted to the participating ICUs in the Netherlands and Belgium during the study period, 1 patient was excluded from further analysis because <60% of data were

collected due to an ICU stay of only several hours, at which point a non-ICU policy was instated and patient was discharged from the ICU. Another patient was excluded because of objection to participation (Figure 1, panel A).

Of the 519 included patients, 4 (0.8%) did not have a positive SARS-CoV-2 PCR result, 2 of whom later were shown to be SARS-CoV-2 IgG positive. Three of the SARS-CoV-2 PCR-negative patients had radiographic findings on thoracic computed tomography (CT) and 1 on thoracic x-ray suggestive of COVID-19. One patient (0.2%) had no known SARS-CoV-2 PCR results but had radiographic findings on thoracic CT suggestive of COVID-19.

ECMM/ISHAM CAPA Classification

In the discovery cohort, tracheobronchitis could be evaluated in 187 patients who underwent bronchoscopy with or without BAL or BL, underwent autopsy, or both. Diagnostic tests to classify patients as proven tracheobronchitis or pulmonary CAPA were performed in 41/519 (8%) patients. Tests for classification as probable (pulmonary) CAPA were performed in 273/519 (53%) patients, and tests for classification as possible CAPA in 43/519 (8%). Because positive culture, GM and PCR results were reported more readily than negative ones, we could not give exact total numbers of fungal culture, GM testing, and *Aspergillus* PCR performed in BAL, BL, and NBL samples obtained (Tables 1, 2; Appendix Tables 7, 12).

In the validation cohort, 127/304 (42%) of patients could be evaluated for invasive *Aspergillus* tracheobronchitis, and 209/304 (69%) could be evaluated for probable CAPA. Data to classify patients as possible CAPA were not collected in this cohort.

CAPA Patients with COPD or Bronchiectasis

The discovery cohort contained 8 patients with CAPA and underlying COPD, whereas the validation cohort contained 2 CAPA patients with COPD. Furthermore, the validation cohort included 2 patients with CAPA and bronchiectasis. Data on bronchiectasis were not collected in the discovery cohort. The 2020 ECMM/ISHAM classification requires a positive GM test result as confirmation of a positive *Aspergillus* culture or PCR result in patients with COPD or another chronic respiratory disease to rule out colonization or chronic aspergillosis (1). Of the 12 patients with COPD or bronchiectasis in both cohorts, 11 (92%) had a BAL or BL GM OD ≥ 1.0 , and 1 had a GM OD > 6.0 in an NBL sample, indicating high fungal load with hyphal formation in the respiratory tract and probably not mere colonization.

Discovery Cohort Microbiological Results

Among the 17 patients with positive BL or BAL cultures, *A. fumigatus* was found in 15/17 (88%), *A. nidulans* in 1 (6%) patient, and both *A. fumigatus* and *A. flavus* in 1 (6%) patient. None of the 5 patients for whom susceptibility data were available demonstrated voriconazole resistance. Among 9 patients in whom positive BAL or BL PCR results were reported, *A. fumigatus* was reported in 3 (33%); speciation was not possible or provided in the other 6 (67%). Azole resistance PCR test results were reported in 1 patient in whom wild-type *A. fumigatus* was found. *A. fumigatus* was found in all 6 patients with positive NBL cultures; 1 patient had a positive BALF culture and a positive NBL culture on different dates and both demonstrated *A. fumigatus*. In 5 of these patients, an *Aspergillus* PCR also was performed on the same NBL sample: in 3 patients *A. fumigatus* was found and no species were identified in the other 2.

Logistic Regression Analysis

In the discovery cohort, RRT was more prevalent in the CAPA group than in the CAPA excluded group. To account for any possible effects of this difference on ICU death, we explored an interaction term between RRT and CAPA in the logistic regression model, which demonstrated no interaction between the 2 variables (adjusted odds ratio [aOR] for the interaction term for ICU death 1.42, 95% CI 0.30–6.87, $p = 0.66$). In the validation cohort, we explored an interaction term for AKI during ICU admission and CAPA for the occurrence of ICU death, which also demonstrated no interaction between the 2 variables for ICU death (aOR 0.16, 95% CI 0.01–2.71, $p = 0.20$).

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Appendix Table 1. Number of included patients by participating center in a multinational observational study of COVID-19–associated pulmonary aspergillosis*

Participating center	No. patients
Discovery cohort	
Amsterdam University Medical Centers, Academisch Medisch Centrum (AMC), Amsterdam, the Netherlands	80
Amsterdam University Medical Centers, Vrije Universiteit Medisch Centrum (VUmc), Amsterdam	69
Erasmus Medical Center, Rotterdam, the Netherlands	85
Radboud University Medical Center, Nijmegen, the Netherlands	80
Ziekenhuis Netwerk Antwerpen (ZNA) Campus Stuivenberg, Antwerpen, Belgium	41
Algemeen Ziekenhuis (AZ) St-Jan Brugge-Oostende, Brugge, Belgium	14
University Hospitals Leuven, Leuven, Belgium	92
Algemeen Ziekenhuis (AZ) Delta Hospital, Roeselare, Belgium	58
Validation cohort	
Amiens University Hospital, Amiens, France	119
Lille University Hospital, Lille, France	128
Rouen University Hospital, Rouen, France	57

*COVID-19, coronavirus disease.

Appendix Table 2. Demographic and clinical characteristics of the discovery cohort in a multinational observational study of COVID-19–associated pulmonary aspergillosis*

Characteristics	Total population, n = 519	CAPA, n = 42	CAPA excluded, n = 237	CAPA not classifiable, n = 240	p value†	p value‡
Age, y	64 (55–72)	68 (61–73)	65 (57–71)	64 (53–73)	0.12	0.36
Sex						
F	141 (27)	8 (19)	58 (24)	75 (31)		
M	378 (73)	34 (81)	179 (76)	165 (69)	0.56	0.13
Ethnicity	N = 365	N = 37	N = 143	N = 185		
Caucasian	294 (81)	32 (87)	115 (80)	147 (80)	0.48	0.64
Northern African	20 (6)	0 (0)	10 (7)	10 (5)	0.22	0.28
Middle Eastern	6 (2)	0 (0)	3 (2)	3 (2)	1.00	1.00
Black or sub-Saharan African	18 (5)	1 (3)	5 (4)	12 (7)	1.00	0.49
Asian	18 (5)	2 (5)	6 (4)	10 (5)	0.67	0.83
Hispanic or Latino	6 (2)	2 (5)	1 (0.7)	3 (2)	0.11	0.15
Pacific Islander	3 (0.8)	0 (0)	3 (2)	0 (0)	1.00	0.13
Unknown	154 (30); n = 519	5 (12); n = 42	94 (40); n = 237	55 (23); n = 240	0.0004	<0.0001
Height, m	1.75 (1.68–1.80); n = 507	1.77 (1.69–1.81); n = 40	1.75 (1.70–1.80); n = 231	1.74 (1.65–1.80); n = 236	0.875	0.059
Weight, kg	84.3 (74.2–95.0); n = 515	83.3 (70.0–94.3); n = 42	84.1 (75.0–95.0); n = 234	85.0 (73.8–95.0); n = 239	0.321	0.58
BMI, kg/m ²	27.2 (24.4–31.0); n = 507	27.4 (23.6–30.2); n = 40	26.9 (24.4–30.9); n = 231	27.5 (24.3–31.5); n = 236	0.72	0.64
BMI >30 kg/m ²	158 (31); n = 507	10 (25); n = 40	69 (30); n = 231	79 (34); n = 236	0.58	0.51
Smoking status	N = 308	N = 23	N = 138	N = 147		
Yes, current smoker§	25 (8)	4 (17)	9 (7)	12 (8)	0.094	0.20
No, never	140 (46)	9 (39)	53 (38)	78 (53)	1.00	0.036
No, but former smoker	143 (46)	10 (44)	76 (55)	57 (39)	0.369	0.021
Unknown	211 (41); n = 519	19 (45); n = 42	99 (42); n = 237	93 (39); n = 240	0.736	0.67
Admission and disease course data						
Days between first signs/symptoms and hospital admission	7 (5–10); n = 467	7 (4–10); n = 36	7 (5–10); n = 221	7 (5–10); n = 210	0.72	0.40
Days between hospital admission and ICU admission	1 (0–3); n = 518	2 (0–5); n = 42	2 (0–4); n = 236	1 (0–3); n = 240	0.60	0.053

Characteristics	Total population, n = 519	CAPA, n = 42	CAPA excluded, n = 237	CAPA not classifiable, n = 240	p value†	p value‡
Days between first signs/symptoms and first positive SARS-CoV-2 PCR	7 (4–10); n = 459	9 (3–10); n = 37	7 (5–11); n = 213	7 (4–10); n = 209	0.48	0.13
Reason for ICU admission						
Respiratory insufficiency	507 (98)	42 (100)	234 (99)	231 (96)	1.00	0.16
Hemodynamic instability	12 (2)	1 (2)	4 (2)	7 (3)	0.56	0.60
Decreased consciousness	12 (2)	0 (0)	5 (2)	7 (3)	1.00	0.74
Other	11 (2)	1 (2)	3 (1)	7 (3)	0.48	0.44
Underlying conditions						
Any	447 (86)	39 (93)	201 (85)	207 (86)	0.23	0.41
Acute leukemia <90 d before ICU admission	1 (0.2)	0 (0)	1 (0.4)	0 (0)	1.00	0.54
Acute leukemia >90 d before ICU admission	3 (0.6)	1 (2)	2 (0.8)	0 (0)	0.39	0.07
SCT <90 d before ICU admission	0 (0)	0 (0)	0 (0)	0 (0)	NA	NA
SCT >90 d before ICU admission	6 (1)	0 (0)	5 (2)	1 (0.4)	1.00	0.27
Other hematological malignancy	20 (4)	3 (7)	11 (5)	6 (3)	0.45	0.18
Kidney transplantation	3 (0.6)	0 (0)	1 (0.4)	2 (0.8)	1.00	1.00
Lung transplantation	1 (0.2)	1 (2)	0 (0)	0 (0)	0.15	0.08
Heart transplantation	2 (0.4)	0 (0)	1 (0.4)	1 (0.4)	1.00	1.00
Liver or pancreas transplantation	0 (0)	0 (0)	0 (0)	0 (0)	NA	NA
Cardiovascular disease¶	291 (56)	25 (60)	130 (55)	136 (57)	0.62	0.83
Diabetes mellitus	139 (27)	9 (21)	61 (26)	69 (29)	0.70	0.58
Asthma	37 (7)	1 (2)	19 (8)	17 (7)	0.33	0.50
COPD	44 (9)	8 (19)	19 (8)	17 (7)	0.042	0.052
Cystic fibrosis	0 (0)	0 (0)	0 (0)	0 (0)	NA	NA
Pulmonary TB	0 (0)	0 (0)	0 (0)	0 (0)	NA	NA
Multiple sclerosis	4 (0.8)	0 (0)	2 (0.8)	2 (0.8)	1.00	1.00
Liver cirrhosis	6 (1)	0 (0)	2 (0.8)	4 (2)	1.00	0.81
Ulcerative colitis	3 (0.6)	0 (0)	0 (0)	3 (1)	NA	0.31
Crohn's disease	0 (0)	0 (0)	0 (0)	0 (0)	NA	NA
Rheumatological disease	31 (6)	5 (12)	14 (6)	12 (5)	0.18	0.22
Psoriasis	7 (1)	1 (2)	3 (1)	3 (1)	0.481	0.70
HIV/AIDS	6 (1)	3 (7)	1 (0.4)	2 (0.8)	0.011	0.008
Congenital immunodeficiency syndrome	0 (0)	0 (0)	0 (0)	0 (0)	NA	NA
Acquired immunodeficiency syndrome other than HIV/AIDS	3 (0.6)	1 (2)	2 (0.8)	0 (0)	0.388	0.07
Solid organ malignancy	28 (5)	3 (7)	11 (5)	14 (6)	0.45	0.64
Other malignancy	11 (2)	0 (0)	5 (2)	6 (3)	1.00	0.90
CKD requiring RRT	7 (1)	0 (0)	0 (0)	7 (3)	NA	0.023
CKD not requiring RRT	30 (6)	3 (7)	8 (3)	19 (8)	0.22	0.08
Thyroid disease	14 (3)	3 (7)	8 (3)	3 (1)	0.22	0.05
Other	215 (41)	13 (31)	110 (46)	92 (38)	0.066	0.07
None	72 (14)	3 (7)	36 (15)	33 (14)	0.23	0.41
EORTC/MSGERC host factors						
Any#	70 (16); n = 426	13 (33); n = 39	31 (19); n = 166	26 (12); n = 221	0.053	0.003
Recent neutropenia**	7 (2); n = 413	1 (3); n = 38	5 (3); n = 156	1 (0.5); n = 219	1.00	0.09
Hematological malignancy	18 (4)	4 (10)	9 (4)	5 (2)	0.11	0.041
Receipt of allogeneic SCT	4 (0.8); n = 516	0 (0)	3 (1); n = 236	1 (0.4); n = 238	1.00	0.55
Receipt of SOT	6 (1)	1 (2)	2 (0.8)	3 (1)	0.39	0.53
T or B cell	31 (6); n = 514	7 (17)	12 (5); n = 233	12 (5); n = 239	0.014	0.024
immunosuppressant drugs, other than corticosteroids ≤90 d before ICU admission						
Inherited severe immunodeficiency	0 (0); n = 517	0 (0)	0 (0); n = 236	0 (0); n = 239	NA	NA
Corticosteroid use						
Systemic corticosteroids ≤30 d before ICU admission	38 (9); n = 430	7 (18); n = 39	14 (9); n = 160	17 (7); n = 231	0.14	0.12
Cumulative corticosteroid dose ≤30 d before ICU admission, mg prednisone equivalent/kg bodyweight	1.82 (1.18–3.48); n = 36	1.74 (1.28–2.37); n = 7	2.03 (0.61–3.73); n = 13	1.82 (1.42–3.60); n = 16	0.76	0.87

Characteristics	Total population, n = 519	CAPA, n = 42	CAPA excluded, n = 237	CAPA not classifiable, n = 240	p value†	p value‡
Inhalational corticosteroids ≤30 d before ICU admission	39 (10); n = 406	5 (14); n = 37	16 (11); n = 150	18 (8); n = 219	0.57	0.46
COVID-19 diagnosis						
PCR-confirmed COVID-19 (any positive PCR)	514 (99); n = 518	42 (100)	232 (98); n = 236	240 (100)	1.00	0.14
Chest x-ray or CT suggestive of COVID-19	352 (79); n = 445††	22 (69); n = 32	160 (79); n = 202	170 (81); n = 211	0.32	0.58
CT severity score at admission	15 (11–18); n = 155	15 (13–17); n = 7	16 (11–19); n = 75	15 (10–17); n = 73	0.64	0.16
Clinical data at ICU admission						
APACHE II score	14 (11–19); n = 163	20 (13–27); n = 12	16 (12–21); n = 64	13 (9–17); n = 87	0.24	0.001
Microbiology at admission						
Positive <i>Aspergillus</i> culture respiratory sample ≤6 mo before ICU admission	1 (0.3); n = 368	1 (3); n = 34	0 (0); n = 126	0 (0); n = 208	0.21	0.09
Positive <i>Aspergillus</i> culture respiratory sample at the time of COVID-19 diagnosis	4 (2); n = 186	2 (9); n = 22	2 (2); n = 89	0 (0); n = 75	0.18	0.037

*Data are presented as no. (%) or median (IQR), unless otherwise stated. Continuous variables were compared by Mann-Whitney U test, categorical variables by Fisher exact test with omission of missing data, unless stated otherwise. Percentages might not accumulate to 100% because of rounding off of numbers. Bold text indicates statistical significance. AKI, acute kidney injury APACHE, acute physiology and chronic health evaluation; BAL, broncho-alveolar lavage; BALF, broncho-alveolar lavage fluid BL, bronchial lavage; BMI, body mass index; CAPA, COVID-19-associated pulmonary aspergillosis; CKD, chronic kidney disease; COPD, chronic obstructive pulmonary disease; COVID-19, coronavirus disease; C_t, cycle threshold; CT, computed tomography; ECMO, extracorporeal membrane oxygenation; EORTC/MSGERC, European Organization for Research and Treatment of Cancer and the Mycoses Study Group Education and Research Consortium (6); GM, galactomannan; ICU, intensive care unit; IQR, interquartile range; LOS, length of stay; NA, not applicable; NBL, nonbronchoscopic lavage; NO, nitric oxide; RRT, renal replacement therapy; SARS-CoV-2, severe acute respiratory syndrome coronavirus 2; SCT, stem cell transplantation; SOT, solid organ transplant; TB, tuberculosis.

†CAPA versus CAPA excluded; comparisons made by Mann-Whitney U test.

‡CAPA versus CAPA excluded versus CAPA not classifiable; comparisons made by Kruskal-Wallis test.

§A patient who smoked ≤1 month before ICU admission was considered a current smoker.

¶Includes hypertension.

#Includes any use of systemic corticosteroids before ICU admission; If data ≥1 EORTC host factors were missing, then data were regarded as missing for this variable.

**Neutropenia <0.5 × 10⁹/L for >10 d.

††At least one thoracic X-ray or CT scan result at admission known for 445 patients.

Appendix Table 3. Laboratory results at ICU admission in the discovery cohort in a multinational observational study of COVID-19-associated pulmonary aspergillosis*

Laboratory tests	Total population, n = 519	CAPA, n = 42	CAPA excluded, n = 237	CAPA not classifiable, n = 240	p value†	p value‡
CRP, mg/L	175 (101–263); n = 447	177 (100–248); n = 34	197 (114–278); n = 200	161 (96–247); n = 213	0.327	0.048
Haemoglobin, mmol/L	7.6 (6.8–8.4); n = 457	7.7 (6.6–8.8); n = 34	7.5 (6.9–8.4) (n = 207)	7.7 (6.8–8.4); n = 216	0.666	0.87
Leukocytes, cells × 10 ⁹ /L	8.2 (5.9–11.3); n = 457	6.9 (5.2–10.2); n = 33	8.7 (6.4–11.6); n = 207	7.7 (5.6–11.1); n = 217	0.061	0.035
Neutrophils, cells × 10 ⁹ /L	6.50 (4.48–9.24); n = 347	5.84 (4.10–7.40); n = 31	6.97 (5.00–9.39); n = 154	6.10 (4.20–9.21); n = 162	0.100	0.16
Lymphocytes, cells × 10 ⁹ /L	0.75 (0.50–1.02); n = 350	0.58 (0.45–1.02); n = 31	0.79 (0.50–1.10); n = 156	0.78 (0.51–1.00); n = 163	0.211	0.36
NLR	9.0 (5.1–14.5); n = 345	9.1 (5.1–15.1); n = 31	9.09 (5.61–14.50); n = 153	8.7 (4.5–14.3); n = 161	0.900	0.67
Platelets, cells × 10 ⁹ /L	223 (163–294); n = 456	218 (174–299); n = 33	222 (156–303); n = 207	228 (172–288); n = 216	0.989	0.89
BUN, mmol/L	7.9 (5.1–14.1); n = 442	11.7 (7.4–21.3); n = 34	7.8 (5.4–11.8); n = 199	7.5 (4.8–15.0); n = 209	0.002	0.012
Creatinine, µmol/L	81 (61–108); n = 460	84 (60–152); n = 34	81 (63–105); n = 210	81 (60–111); n = 216	0.397	0.75
Total bilirubin, µmol/L	8.2 (6.0–12.0); n = 429	8.6 (6.8–10.9); n = 33	8.8 (6.8–12.0); n = 188	8.0 (5.5–12.0); n = 208	0.533	0.23
Direct bilirubin, µmol/L	5.1 (3.5–7.4); n = 114	5.6 (4.0–9.1); n = 10	5.3 (3.8–8.6); n = 42	5.0 (3.1–6.5); n = 62	0.880	0.25
ALAT, U/L	37 (24–59); n = 432	34 (21–52); n = 33	38 (25–59); n = 193	38 (23–64); n = 206	0.143	0.32
GGT, U/L	61 (35–125); n = 407	64 (34–112); n = 31	65 (39–136); n = 178	60 (32–118); n = 198	0.591	0.29
Ferritin, µg/L	1,390 (718–2,146); n = 245	1,597 (960–3,761); n = 18	1,471 (884–2,424); n = 117	1,154 (544–1,672); n = 110	0.194	0.002
Albumin, g/L	29 (24–32); n = 402	28.5 (26.0–33.0); n = 31	27 (22–32); n = 190	30 (26–33); n = 181	0.061	<0.001
Total protein level, g/L	63 (59–68); n = 126	63 (57–66); n = 18	61 (59–65); n = 39	64 (59–70); n = 69	0.757	0.09
INR	1.2 (1.1–1.3); n = 312	1.1 (1.1–1.2); n = 27	1.2 (1.1–1.3); n = 113	1.1 (1.1–1.3); n = 172	0.724	0.90
D-dimer ≥500 ng/mL	277 (93); n = 298	25 (96); n = 26	128 (94); n = 136	124 (91); n = 136	1.000	0.66
D-dimer, ng/mL; if >500 ng/mL	1,720 (1,128–3,965); n = 277	2,170 (1,245–5,959); n = 25	2,135 (1,265–5,878); n = 128	1,398 (989–2,555); n = 124	0.927	0.001

*Data are presented as no. (%) or median (IQR), unless otherwise stated. The most strongly abnormal laboratory results obtained during the first 24 h of ICU admission are shown. Continuous variables were compared by Mann-Whitney U test or Kruskal-Wallis test, categorical variables by Fisher exact test with omission of missing data, unless stated otherwise. Percentages might not accumulate to 100% because of rounding off of numbers. Bold text indicates statistical significance. ALAT, alanine aminotransferase; BUN, blood urea nitrogen; CAPA, COVID-19-associated pulmonary aspergillosis; COVID-19, coronavirus disease; CRP, C-reactive protein; GGT, gamma glutamyl transpeptidase; ICU, intensive care unit; INR, international normalized ratio; IQR, interquartile range; NLR, neutrophil/lymphocyte ratio.

†CAPA versus CAPA excluded; comparisons made by Mann-Whitney U test.

‡CAPA versus CAPA excluded versus CAPA not classifiable; comparisons made by Kruskal-Wallis test.

Appendix Table 4. Treatment and outcome data in the discovery cohort in a multinational observational study of COVID-19-associated pulmonary aspergillosis*

Characteristics	Total population, n = 519	CAPA, n = 42	CAPA excluded, n = 237	CAPA not classifiable, n = 240	p value†	p value‡
ICU treatment data						
Invasive mechanical ventilation	423 (82); n = 517	40 (98); n = 41	225 (95)	158 (66); n = 239	0.70	<0.0001
Ventilation in prone position	323 (63); n = 517	26 (63); n = 41	195 (82)	102 (43); n = 239	0.011	<0.0001
Ventilation with NO	17 (3); n = 518	5 (12)	8 (3)	4 (2); n = 239	0.031	0.008
ECMO§	16 (3); n = 518	3 (7)	10 (4)	3 (1); n = 239	0.42	0.029
No. invasive ventilation days¶	14 (9–24); n = 395	16 (13–27); n = 37	18 (11–30); n = 212	10 (6–14); n = 146	0.98	<0.001
Any vasopressors and/or inotropes during ICU admission	413 (80); n = 516	41 (98)	223 (95); n = 235	149 (62); n = 239	0.70	<0.0001
AKI at any time during ICU admission#	264 (54); n = 486	29 (71); n = 41	137 (63); n = 218	98 (43); n = 227	0.38	<0.0001
AKI stage 1	89 (35); n = 258	7 (24); n = 29	44 (33); n = 135	38 (40); n = 94	0.51	0.23
AKI stage 2	47 (18); n = 258	4 (14); n = 29	31 (23); n = 135	12 (13); n = 94	0.33	0.13
AKI stage 3	122 (47); n = 258	18 (62); n = 29	60 (44); n = 135	44 (47); n = 94	0.10	0.23
RRT during ICU admission	93 (18); n = 516	17 (41)	44 (19); n = 236	32 (13); n = 238	0.004	0.0004
Systemic corticosteroids during ICU admission	216 (42); n = 516	20 (48)	131 (56); n = 236	65 (27); n = 238	0.40	<0.0001
Cumulative corticosteroid dose during ICU admission, mg prednisone equivalent/kg bodyweight	5.37 (1.14–13.99); n = 210	5.30 (2.36–14.30); n = 20	6.63 (1.03–16.73); n = 126	4.21 (1.22–10.80); n = 64	0.88	0.13
COVID-19 treatment data during hospital admission						
Chloroquine	162 (31)	7 (17)	114 (48)	41 (17)	0.0001	<0.0001
Hydroxychloroquine	182 (35)	21 (50)	61 (26)	100 (42)	0.003	0.0001
Remdesivir	21 (4)	1 (2)	18 (8)	2 (0.8)	0.33	0.0004
Lopinavir/ritonavir	20 (4)	1 (2)	6 (3)	13 (5)	1.00	0.24
Anakinra	26 (5)	0 (0)	25 (11)	1 (0.4)	0.019	<0.0001
Tocilizumab	5 (1)	1 (2)	2 (0.8)	2 (0.8)	0.39	0.59
Other	101 (20)	9 (21)	39 (17)	53 (22)	0.51	0.28
None of the above	155 (30)	14 (33)	50 (21)	91 (38)	0.11	0.0003
Outcome data						
ICU death	154 (30); n = 518	22 (52)	81 (34)	51 (21); n = 239	0.036	<0.0001
ICU LOS, d**	14 (8–24); n = 491	18 (12–27); n = 39	20 (12–32); n = 222	10 (5–15); n = 230	0.84	<0.001
ICU LOS, if alive at ICU discharge, d**	15 (9–25); n = 337	21 (16–41); n = 17	23 (13–34); n = 141	11 (6–17); n = 179	0.61	<0.001
Hospital LOS, d††	22 (13–35); n = 428	25 (15–38); n = 33	28 (17–45); n = 195	18 (10–27); n = 200	0.32	<0.001
Hospital LOS, if alive at ICU discharge, d††	26 (17–39); n = 274	34 (25–40); n = 11	36 (23–51); n = 114	20 (14–29); n = 149	0.60	<0.001

*Data are presented as no. (%) or median (IQR), unless otherwise stated. Continuous variables were compared by Mann-Whitney U test or Kruskal-Wallis test, categorical variables by Fisher exact test with omission of missing data, unless stated otherwise. Percentages might not accumulate to 100% because of rounding off of numbers. Bold text indicates statistical significance. AKI, acute kidney injury; CAPA, COVID-19-associated pulmonary aspergillosis; COVID-19, coronavirus disease; ECMO, extracorporeal membrane oxygenation; ECCO2R, extracorporeal CO₂ removal; ICU, intensive care unit; IQR, interquartile range; LOS, length of stay; NO, nitric oxide; RRT, renal replacement therapy.

†CAPA versus CAPA excluded; comparisons made by Mann-Whitney U test.

‡CAPA versus CAPA excluded versus CAPA not classifiable; comparisons made by Kruskal-Wallis test.

§Including one patient with Extracorporeal CO₂ removal (ECCO2R).

¶If transferred to another hospital from ICU and still on ventilatory support of any kind, duration of invasive mechanical ventilatory support was regarded as missing data and not taken into account in the analyses. The same holds true for those who received a tracheostomy for a prolonged weaning trajectory.

#Acute kidney injury definition and staging according to Kidney Disease: Improving Global Outcomes (KDIGO; <https://kdigo.org>) criteria.

**Data on ICU LOS were regarded as missing if transfer to another hospital was the reason for ICU discharge.

††Data on hospital LOS were regarded as missing if patients deceased after ICU discharge because no data were available whether or not death occurred in hospital.

Appendix Table 5. CAPA diagnosis among participating centers in a multinational observational study of COVID-19–associated pulmonary aspergillosis*

Participating center	CAPA	CAPA excluded	CAPA not classifiable	CAPA in classifiable patients
Discovery cohort				
Amsterdam University Medical Centers, Academisch Medisch Centrum (AMC), Amsterdam, the Netherlands	7/80 (9)	32/80 (40)	41/80 (51)	7/39 (18)
Amsterdam University Medical Centers, Vrije Universiteit Medisch Centrum (VUmc), Amsterdam, the Netherlands	7/69 (10)	39/69 (57)	23/69 (33)	7/46 (15)
Erasmus Medical Center, Rotterdam, the Netherlands	3/85 (4)	22/85 (26)	60/85 (71)	3/25 (12)
Radboud University Medical Center, Nijmegen, the Netherlands	3/80 (4)	77/80 (96)	0/80 (0)	3/80 (4)
Ziekenhuis Netwerk Antwerpen (ZNA) Campus Stuivenberg, Antwerpen, Belgium	10/41 (24)	15/41 (37)	16/41 (39)	10/25 (40)
Belgium				
Algemeen Ziekenhuis (AZ) St-Jan Brugge_Oostende, Brugge, Belgium	1/14 (7)	7/14 (50)	6/14 (43)	1/8 (13)
Validation cohort				
Amiens University Hospital, Amiens, France	13/119 (11)	75/119 (63)	31/119 (26)	13/88 (15)
Lille University Hospital, Lille, France	8/128 (6)	101/128 (79)	19/128 (15)	8/109 (7)
Rouen University Hospital, Rouen, France	0/57 (0)	8/57 (14)	49/57 (86)	0/8 (0)

*Results are reported as no. patients/no. patients included per center (%). CAPA classification according to the European Confederation for Medical Mycology/International Society for Human and Animal Mycology (ECMM/ISHAM) classification (1). CAPA, COVID-19–associated pulmonary aspergillosis; COVID-19, coronavirus disease.

Appendix Table 6. Application of different classification systems in the discovery cohort in a multinational observational study of COVID-19–associated pulmonary aspergillosis*

Classification	Physician reported, n = 45	ECMM/ISHAM proven or probable CAPA, n = 38†,‡	IAPA expert case definition, set A, proven or probable IPA, n = 37‡	mAspICU proven or putative IPA, n = 31‡,§	EORTC/MSGERC proven or probable IPA, n = 20‡,§
Physician reported, n = 45		32	31	24	14
ECMM/ISHAM proven or probable CAPA, n = 38†,‡	32		37	26	16
IAPA expert case definition, set A, proven or probable IPA, n = 37‡	31	37		26	16
mAspICU proven or putative IPA, n = 31‡,§	24	26	26		19
EORTC/MSGERC proven or probable IPA, n = 20‡,§	14	16	16	19	

*CAPA, COVID-19–associated pulmonary aspergillosis; COVID-19, coronavirus disease; ECMM/ISHAM, European Confederation for Medical Mycology/International Society for Human and Animal Mycology; EORTC/MSGERC, European Organization for Research and Treatment of Cancer and the Mycoses Study Group Education and Research Consortium; IAPA, influenza-associated pulmonary aspergillosis; IPA, invasive pulmonary aspergillosis; mAspICU, modified *Aspergillus* intensive care unit classification.

†European Confederation for Medical Mycology/International Society for Human and Animal Mycology (ECMM/ISHAM) classification (1).

‡Radiological findings were not considered for any of the classification systems; therefore, for the IAPA expert case definition classification, only subset A for probable pulmonary IPA, which does not require cavitating infiltrates specifically on chest radiography, could be assessed.

§For the mAspICU (4,5) and EORTC/MSGERC classifications (6), clinical factors were not taken into account. For the mAspICU classification, all lower respiratory tract cultures positive for *Aspergillus*, including bronchial lavage (BL), broncho-alveolar lavage (BAL) and mini-BAL fluid cultures, were considered as a positive entry criterion.

Appendix Table 7. Mycological diagnostic test results according to diagnostic group in the discovery cohort in a multinational observational study of COVID-19–associated pulmonary aspergillosis*

Characteristics	Total population, n = 519	CAPA, n = 42	CAPA excluded, n = 237	CAPA not classifiable, n = 240	p value†	p value‡
Proven CAPA	6 (2); n = 279	6 (14)	NA	NA	NA	NA
Histological examination of biopsy	5 (83); n = 6	5 (83); n = 6	NA	NA	NA	NA
Autopsy indicative of CAPA	2 (33); n = 6	2 (33); n = 6	NA	NA	NA	NA
Serum GM						
Serum GM performed	176 (34)	28 (67)	148 (62)	0 (0)	0.73	<0.0001
>1 Serum GM performed on separate days	95 (71); n = 134	14 (50); n = 28	81 (76); n = 106	NA	0.010	0.010
Serum GM OD >0.5	3 (2); n = 176§	3 (11); n = 28	0 (0); n = 148	NA	NA	NA
Serum GM OD >0.5 on multiple, separate days	0 (0); n = 3	0 (0); n = 3	NA	NA	NA	NA
Serum GM OD						
All¶	0.10 (0.10–0.10); n = 134	0.10 (0.06–0.14); n = 28	0.10 (0.10–0.10); n = 106	NA	0.95	0.95
If OD >0.5¶	1.00 (0.80–.); n = 3	1.00 (0.80–.); n = 3	NA	NA	NA	NA
Days between ICU admission and first reported serum GM#	5 (3–8); n = 133	7 (5–14); n = 27	5 (3–7); n = 106	NA	0.005	0.005
Days between ICU admission and first reported serum GM OD >0.5#	15 (4–.); n = 3	15 (4–.); n = 3	NA	NA	NA	NA
BAL/BL						
Any performed/reported	166 (32)	41 (98)	125 (53)	NA	<0.0001	<0.0001
Positive BALF/BL culture	17 (10); n = 166	17 (42); n = 41	0 (0); n = 125	NA	NA	NA
BALF/BL GM obtained on multiple, separate days	33 (37); n = 90	16 (46); n = 35	17 (31); n = 55	NA	0.18	0.18
BALF/BL GM OD ≥1.0	32 (19); n = 166**	32 (78); n = 41††	0 (0); n = 125‡‡	NA	NA	NA
BALF/BL GM OD ≥1.0 on multiple, separate days	8 (25); n = 32	8 (25); n = 32	NA	NA	NA	NA
BALF/BL GM OD value						
All¶	0.20 (0.10–1.50); n = 90	1.80 (1.00–3.90); n = 35	0.10 (0.10–0.20); n = 55	NA	<0.001	<0.001
If OD ≥1.0¶	2.72 (1.75–4.00); n = 32	2.72 (1.75–4.00); n = 32	NA	NA	NA	NA
Days between ICU admission and first reported BALF/BL GM#	9 (5–14); n = 88	7 (4–14); n = 33	10 (5–14); n = 55	NA	0.21	0.21
Days between ICU admission and first reported BALF/BL GM OD ≥1.0#	7 (5–13); n = 30	7 (5–13); n = 30	NA	NA	NA	NA
Positive BALF/BL PCR, any C _t value	9 (5); n = 166§§	7 (17); n = 41¶¶	2 (2); n = 125##	NA	NA	NA
NBL						
Any performed/reported	42 (8)	9 (21)	30 (13)	3 (1)	0.15	<0.0001
Positive NBL culture	6 (14); n = 42	6 (67); n = 9	0 (0); n = 30	0 (0); n = 3	NA	NA
Positive plasma, serum, or whole blood PCR	ND	ND	ND	ND	NA	NA
1 NBL GM OD >4.5	5 (12); n = 42	5 (56); n = 9	0 (0); n = 30	0 (0); n = 3	NA	NA
≥2 NBL GM OD >1.2	0 (0); n = 7***	0 (0); n = 3	0 (0); n = 4	0 (0); n = 0	NA	NA
NBL GM OD >1.2 and positive PCR	5 (56); n = 9†††	5 (83); n = 6	0 (0); n = 3	0 (0); n = 0	NA	NA
Time to first positive mycological test						
Days between ICU admission and first positive mycological test†††	NA	6 (3–9); n = 41	NA	NA	NA	NA

*Data are presented as no. (%) or median (IQR), unless otherwise stated. Continuous variables were compared by Mann-Whitney U test or Kruskal-Wallis test, categorical variables by Fisher exact test with omission of missing data, unless stated otherwise. Percentages might not accumulate to 100% because of rounding off of numbers. Bold text indicates statistical significance. BAL, bronchoalveolar lavage; BALF, bronchoalveolar lavage fluid; BL, bronchial lavage; BMI, body mass index; CAPA, COVID-19–associated pulmonary aspergillosis; COVID-19, coronavirus disease; C_t, cycle threshold; GM, galactomannan; ICU, intensive care unit; IQR, interquartile range; ND, not done; NA, not applicable; OD, optical density.

†CAPA versus CAPA excluded; comparisons made by Mann-Whitney U test.

‡CAPA versus CAPA excluded versus CAPA not classifiable; comparisons made by Kruskal-Wallis test.

§Serum GM values available for 134 patients.

Characteristics	Total population, n = 519	CAPA, n = 42	CAPA excluded, n = 237	CAPA not classifiable, n = 240	p value†	p value‡
¶When multiple values were reported for one patient, the median of these values was used for further calculations.						
#Dates were regarded as missing if inconsistencies regarding date of obtaining test results existed.						
**BALF/BL GM OD values reported for 90 patients, but 166 patients underwent BALF/BL sampling.						
††BALF/BL GM OD values reported for 35 patients.						
‡‡BALF/BL GM OD values reported for 55 patients.						
§§BALF/BL PCR results reported for 11 patients.						
¶¶BALF/BL PCR results reported for 7 patients.						
## BALF/BL PCR results reported for 4 patients. Positive PCR with C _v values ≥36 as only positive mycological criterion.						
***Multiple NBL GM OD values reported for 7 patients.						
†††Results for both NBL GM and PCR reported in 9 patients.						
‡‡‡ Mycological test considered a criterion for proven, probable, or possible CAPA according to the 2020 ECMM/ISHAM classification.						

Appendix Table 8. Antifungal treatment in the discovery and validation cohorts in a multinational observational study of COVID-19–associated pulmonary aspergillosis*

Cohort and antifungal treatment	Total population	CAPA	CAPA excluded	CAPA not classifiable
Discovery cohort	N = 519	N = 42	N = 237	N = 240
Any antifungal treatment during ICU admission	83 (16)	37 (88)	36 (15)	10 (4)
Initial antifungal treatment	N = 83	N = 37	N = 36	N = 10
Voriconazole	24 (29)	20 (54)	3 (8)	1 (10)
Other azole monotherapy	13 (16)	2 (5)	3 (8)	8 (80)
Echinocandin monotherapy	8 (10)	1 (3)	7 (19)	0 (0)
Azole + echinocandin combination therapy	33 (40)	13 (35)	19 (53)	1 (10)
Liposomal amphotericin B monotherapy	2 (2)	1 (3)	1 (3)	0 (0)
Other combination therapy	1 (1)	0 (0)	1 (3)	0 (0)
Unspecified	2 (2)	0 (0)	2 (6)	0 (0)
Validation cohort	N = 304	N = 21	N = 188	N = 95
Any antifungal treatment used	57 (19)	12 (57)	37 (20)	8 (8)
Specified antifungal treatment	N = 57	N = 12	N = 37	N = 8
Voriconazole	14 (25)	7 (58)	7 (19)	0 (0)
Isavuconazole	3 (5)	3 (25)	0 (0)	0 (0)
Fluconazole	11 (19)	0 (0)	8 (22)	3 (38)
Caspofungin	21 (37)	1 (8)	18 (49)	2 (25)
Amphotericin B	3 (5)	2 (17)	1 (3)	0 (0)
Combination of isavuconazole and liposomal amphotericin B	1 (2)	1 (8)	0 (0)	0 (0)
Unspecified antifungal treatment	8 (14)	0 (0)	5 (14)	3 (38)

*Data are presented as no. (%) and for the discovery cohort include antifungal treatment administered in ICU, including treatment initiated before ICU admission, but continued in ICU. CAPA, COVID-19–associated pulmonary aspergillosis; COVID-19, coronavirus disease; ICU, intensive care unit.

Appendix Table 9. Demographic and clinical characteristics of the validation cohort in a multinational observational study of COVID-19–associated pulmonary aspergillosis*

Characteristics	Total population, n = 304	CAPA, n = 21	CAPA excluded, n = 188	Not classifiable, n = 95	p value†	p value‡
Age, y	63 (55–71)	67 (59–75)	62 (53–69)	64 (56–72)	0.06	0.1
Sex						
F	227 (75)	21 (100)	141 (75)	65 (68)		
M	77 (25)	0 (0)	47 (25)	30 (32)	0.005	0.004
BMI, kg/m ²	30.0 (26.0–34.4); n = 296	30.2 (26.1–32.8); n = 20	30.0 (26.4–34.5); n = 185	29.4 (25.4–36.1); n = 91	0.84	0.77
BMI >30 kg/m ²	142 (48); n = 296	10 (50); n = 20	90 (49); n = 185	42 (46); n = 91	1.00	0.89
Smoking status						
Current smoker	10 (3); n = 295	0 (0)	7 (4); n = 180	3 (3); n = 94	1.00	1.00
Not current smoker	285 (97); n = 295	21 (100)	173 (96); n = 180	91 (97); n = 94	1.00	1.00
Unknown	9 (3)	0 (0)	8 (4)	1 (1)	1.00	0.37
Underlying conditions						
Active hematological malignancy	10 (3)	0 (0)	6 (3)	4 (4)	1.00	0.87
Hypertension	165 (54)	15 (71)	96 (51)	54 (57)	0.106	0.18
Cardiovascular disease§	64 (21)	9 (43)	35 (19)	20 (21)	0.020	0.046
Cardiovascular disease, including hypertension§	185 (61)	17 (81)	112 (60)	56 (59)	0.06	0.15
Diabetes mellitus	92 (30)	9 (43)	62 (33)	21 (22)	0.47	0.07
Asthma	22 (7)	2 (10)	12 (6)	8 (8)	0.64	0.67
COPD	20 (7)	2 (10)	12 (6)	6 (6)	0.64	0.80
Liver cirrhosis¶	5 (2)	2 (10)	2 (1)	1 (1)	0.051	0.046
Autoimmune disease	16 (5)	2 (10)	11 (6)	3 (3)	0.63	0.32
HIV/AIDS	3 (1)	0 (0)	1 (0.5)	2 (2)	1.00	0.41

Characteristics	Total population, n = 304	CAPA, n = 21	CAPA excluded, n = 188	Not classifiable, n = 95	p value†	p value‡
Active solid organ malignancy	4 (1)	1 (5)	3 (2)	0 (0)	0.35	0.16
CKD	18 (6)	3 (14)	10 (5)	5 (5)	0.13	0.25
Sleep apnea syndrome	36 (12)	2 (10)	21 (11)	13 (14)	1.00	0.81
Bronchiectasis	5 (2)	2 (10)	1 (0.5)	2 (2)	0.027	0.018
EORTC/MSGERC host factors						
Any#	35 (12)	3 (14)	20 (11)	12 (13)	0.71	0.74
Recent neutropenia**	0 (0); n = 303	0 (0)	0 (0); n = 187	0 (0)	NA	NA
Hematological malignancy	10 (3)	0 (0)	6 (3)	4 (4)	1.00	0.87
Receipt of SOT	9 (3)	1 (5)	5 (3)	3 (3)	0.48	0.63
Corticosteroids ≥0.3 mg/kg for ≥3 wks within the past 60 d	17 (6)	2 (10)	10 (5)	5 (5)	0.34	0.61
Other immunosuppressant drugs ≤90 d before ICU admission	23 (8)	2 (10)	16 (9)	5 (5)	0.70	0.60
Clinical data at ICU admission						
SAPS II score	41 (30–58); n = 285	48 (36–68); n = 17	44 (32–63); n = 185	35 (24–46); n = 83	0.38	<0.001
ARDS††	212 (70); n = 303	17 (85); n = 20	150 (80)	45 (47)	0.77	<0.001

*Data are presented as no. (%) or median (IQR), unless otherwise stated. Continuous variables were compared by Mann-Whitney U test or Kruskal-Wallis test, categorical variables by Fisher exact test with omission of missing data, unless stated otherwise. Percentages might not accumulate to 100% because of rounding off of numbers. Bold text indicates statistical significance. APACHE, acute physiology and chronic health evaluation; ARDS, acute respiratory distress syndrome; BMI, body mass index; CAPA, COVID-19-associated pulmonary aspergillosis; CKD, chronic kidney disease; COPD, chronic obstructive pulmonary disease; COVID-19, coronavirus disease; EORTC/MSGERC, European Organization for Research and Treatment of Cancer and the Mycoses Study Group Education and Research Consortium (6); ICU, intensive care unit; IQR, interquartile range, NA, not applicable; SAPS, simplified acute physiology score; SOT, solid organ transplant.

†CAPA versus CAPA excluded; comparisons made by Mann-Whitney U test.

‡CAPA versus CAPA excluded versus CAPA not classifiable; comparisons made by Kruskal-Wallis test.

§Includes Factor V Leiden mutation.

#Includes any use of systemic corticosteroids We did not assess receipt of an allogeneic stem cell transplant, presence of an inherited severe immunodeficiency, and presence of acute graft-versus-host disease.

**Neutropenia $<0.5 \times 10^9/L$ for >10 d.

††ARDS according to the Berlin definition.

Appendix Table 10. Treatment and outcome data in the validation cohort in a multinational observational study of COVID-19–associated pulmonary aspergillosis*

Characteristics	Total population; n = 304	CAPA; n = 21	CAPA excluded; n = 188	Not classifiable; n = 95	p value†	p value‡
ICU treatment data						
Invasive mechanical ventilation	228 (76); n = 302	19 (95); n = 20	168 (89)	41 (44); n = 94	0.70	<0.0001
Ventilation in prone position	159 (53); n = 302	15 (75); n = 20	116 (62)	28 (30); n = 94	0.33	<0.0001
ECMO	41 (14); n = 303	4 (20); n = 20	34 (18)	3 (3)	0.77	0.0004
No. invasive ventilation days	15 (9–25); n = 212	18 (13–25); n = 17	15 (9–25); n = 157	14 (10–25); n = 38	0.21	0.44
Any vasopressors and/or inotropes	189 (63); n = 302	16 (80); n = 20	138 (74); n = 187	35 (37)	0.79	<0.0001
AKI at any time during ICU admission§	126 (42); n = 303	16 (80); n = 20	94 (50)	16 (17)	0.017	<0.0001
RRT	64 (21); n = 303	11 (55); n = 20	47 (25)	6 (6)	0.008	<0.0001
Systemic corticosteroids during ICU admission	147 (49); n = 303	11 (52)	106 (57); n = 187	30 (32)	0.82	0.0003
COVID-19 treatment data						
Hydroxychloroquine	56 (18)	2 (10)	35 (19)	19 (20)	0.38	0.59
Azithromycin	63 (21)	3 (14)	43 (23)	17 (18)	0.58	0.55
Remdesivir	10 (3)	0 (0)	9 (5)	1 (1)	0.60	0.29
Lopinavir/ritonavir	115 (38)	6 (29)	68 (36)	41 (43)	0.63	0.37
Tocilizumab	13 (4)	2 (10)	6 (3)	5 (5)	0.19	0.24
Interferon β	12 (4)	0 (0)	11 (6)	1 (1)	0.61	0.11
Oseltamivir	20 (7)	3 (14)	15 (8)	2 (2)	0.40	0.037
Outcome data						
Candidemia	4 (1)	0 (0)	4 (2)	0 (0)	1.00	0.48
ICU mortality rate	69 (23); n = 299	9 (43)	46 (25); n = 185	14 (15); n = 93	0.12	0.017
30-Day mortality rate	56 (19); n = 301	8 (38)	37 (20); n = 186	11 (12); n = 94	0.09	0.015
90-Day mortality rate	69 (23); n = 301	9 (43)	46 (25); n = 186	14 (15); n = 94	0.11	0.016
ICU LOS, d¶	14 (8–26); n = 295	22 (12–35); n = 20	18 (10–28); n = 183	9 (5–15); n = 92	0.27	<0.001
ICU LOS if alive at ICU discharge, d¶	14 (8–26); n = 230	30 (14–40); n = 12	18 (10–28); n = 139	8 (5–15); n = 79	0.13	<0.001

*Data are presented as no. (%) or median (IQR), unless otherwise stated. Continuous variables were compared by Mann-Whitney U test or Kruskal-Wallis test, categorical variables by Fisher exact test with omission of missing data, unless stated otherwise. Percentages might not accumulate to 100% because of rounding off of numbers. Bold text indicates statistical significance. AKI, acute kidney injury; CAPA, COVID-19–associated pulmonary aspergillosis; COVID-19, coronavirus disease; ECMO, extracorporeal membrane oxygenation; ICU, intensive care unit; IQR, interquartile range; LOS, length of stay; RRT, renal replacement therapy.

†CAPA versus CAPA excluded; comparisons made by Mann-Whitney U test.

‡CAPA versus CAPA excluded versus CAPA not classifiable; comparisons made by Kruskal-Wallis test.

§Acute kidney injury definition and staging according to Kidney Disease: Improving Global Outcomes (KDIGO; <https://kdigo.org>) criteria.

¶Data on ICU LOS were regarded as missing if still admitted at the time of data entry or if transfer to another hospital was the reason for ICU discharge.

Appendix Table 11. Application of different classification systems in the validation cohort in a multinational observational study of COVID-19–associated pulmonary aspergillosis*

Classification	Physician reported, n = 20	ECMM/ISHAM probable CAPA, n = 21†,‡	IAPA expert case definition, set A, probable IA, n = 19‡	mAsplCU putative IPA, n = 13‡,§	EORTC/MSGERC probable IPA, n = 3‡,§
Physician reported, n = 20	20	19	19	11	2
ECMM/ISHAM probable CAPA, n = 21†,‡	19	21	19	11	2
IAPA expert case definition, set A, proven or probable IPA, n = 19‡	19	19	19	11	2
mAsplCU putative IPA, n = 13‡,§	11	11	11	13	3
EORTC/MSGERC probable IPA, n = 3‡,§	2	2	2	3	3

*CAPA, COVID-19–associated pulmonary aspergillosis; COVID-19, coronavirus disease; ECMM/ISHAM, European Confederation for Medical Mycology/International Society for Human and Animal Mycology; EORTC/MSGERC, European Organization for Research and Treatment of Cancer and the Mycoses Study Group Education and Research Consortium; IAPA, influenza-associated pulmonary aspergillosis; IPA, invasive pulmonary aspergillosis; mAsplCU, modified *Aspergillus* intensive care unit classification.

†European Confederation for Medical Mycology/International Society for Human and Animal Mycology (ECMM/ISHAM) classification (1).

‡Radiological findings were not considered for any of the classification systems; therefore, for the IAPA expert case definition classification, only subset A for probable pulmonary IPA, which does not require cavitating infiltrates specifically on chest radiography, could be assessed.

§For the mAsplCU (4,5) and EORTC/MSGERC classifications (6), clinical factors were not taken into account. For the mAsplCU classification, all lower respiratory tract cultures positive for *Aspergillus*, including broncho-alveolar lavage (BAL) fluid cultures, were considered as a positive entry criterion.

Appendix Table 12. Mycological diagnostic test results according to diagnostic group in the validation cohort in a multinational observational study of COVID-19–associated pulmonary aspergillosis*

Characteristics	Total population, n = 304	CAPA, n = 21	CAPA excluded, n = 188	Not classifiable, n = 95	p value†	p value‡
Serum GM						
Serum GM performed	173 (57)	18 (86)	155 (82)	0 (0)	NA	NA
Serum GM OD >0.5	4 (2); n = 172§	4 (22); n = 18	0 (0); n = 154¶	NA	NA	NA
Serum GM OD						
All	0.07 (0.04–0.12); n = 172§	0.10 (0.06–0.34); n = 18	0.06 (0.04–0.11); n = 154¶	NA	0.008	0.008
If OD >0.5	0.66 (0.54–3.84); n = 4	0.66 (0.54–3.84); n = 4	NA	NA	NA	NA
BAL						
BAL performed/reported	135 (44)	21 (100)	114 (61)	0 (0)	NA	NA
Positive BALF culture	11 (8); n = 135	11 (52)	0 (0); n = 114	NA	NA	NA
BALF GM performed	123 (91); n = 135	21 (100)	102 (89); n = 114	NA	NA	NA
BALF GM OD ≥1.0	13 (11); n = 123	13 (62)	0 (0); n = 102	NA	NA	NA
BALF GM OD value						
All#	0.12 (0.05–0.32); n = 123	1.10 (0.12–3.06)	0.11 (0.05–0.18); n = 102	NA	<0.001	<0.001
If OD ≥1.0#	1.70 (1.24–5.38); n = 13	1.70 (1.24–5.38); n = 13	NA	NA	NA	NA
Positive BALF PCR, any Ct value	8 (13); n = 64	8 (53); n = 15	0 (0); n = 49	NA	NA	NA
Serum β-D-glucan						
Serum β-D-glucan performed	184 (61)	19 (91)	160 (85)	5 (5)	0.75	<0.0001
Serum β-D-glucan ≥80 pg/mL	37 (20); n = 184	8 (42); n = 19	29 (18); n = 160	0 (0); n = 5	0.030	0.038
Serum β-D-glucan, pg/mL**						
All	31 (13–60); n = 184	34 (31–156); n = 19	31 (10–59); n = 160	31 (16–46); n = 5	0.055	0.15
If value ≥80 pg/mL	171 (120–288); n = 37	180 (116–316); n = 8	171 (130–283); n = 29	NA	1.00	0.99
Other mycological tests						
TBA performed	131 (43)	8 (38)	94 (50)	29 (31)	0.36	0.006
Positive TBA culture	7 (5); n = 131	2 (25); n = 8	5 (5); n = 94	0 (0); n = 29	0.09	0.05
Positive TBA PCR	3 (18); n = 17	0 (0); n = 1	3 (43); n = 7	0 (0); n = 9	1.00	0.08
Sputum obtained	34 (11)	3 (14)	21 (11)	10 (11)	0.72	0.84
Positive sputum culture	0 (0); n = 34	0 (0); n = 3	0 (0); n = 21	0 (0); n = 10	NA	NA
Positive sputum PCR	2 (29); n = 7	0 (0); n = 0	2 (67); n = 3	0 (0); n = 4	NA	0.14
Positive plasma, serum, or whole blood PCR	ND	ND	ND	ND	NA	NA

*Data are presented as no. (%) or median (IQR), unless stated otherwise. Continuous variables were compared by Mann-Whitney U test or Kruskal-Wallis test, categorical variables by Fisher exact test with omission of missing data, unless stated otherwise. Percentages might not accumulate to 100% because of rounding off of numbers. Bold text indicates statistical significance. BAL, broncho-alveolar lavage; BALF, BAL fluid; CAPA, COVID-19–associated pulmonary aspergillosis; COVID-19, coronavirus disease; Ct, cycle threshold; GM, galactomannan; ICU, intensive care unit; IQR, interquartile range; NA, not applicable; ND, not done; OD, optical density; TBA, tracheobronchial aspirate.

†CAPA versus CAPA excluded; comparisons made by Mann-Whitney U test.

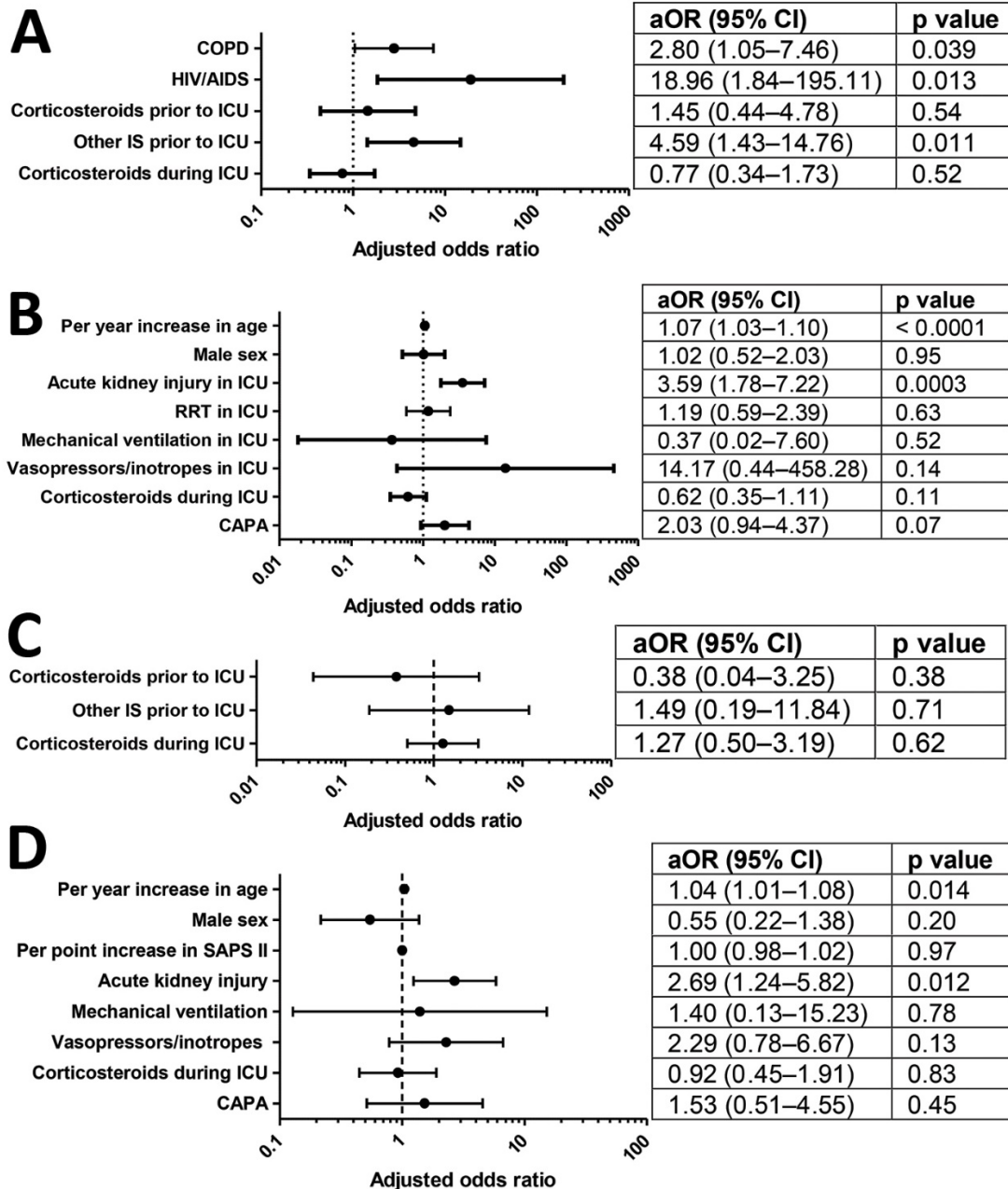
‡CAPA versus CAPA excluded versus CAPA not classifiable; comparisons made by Kruskal-Wallis test.

§Serum GM performed in 173 patients, including 1 patient with an unknown result.

¶Serum GM OD values known for 154 patients, unknown value in 1 patient.

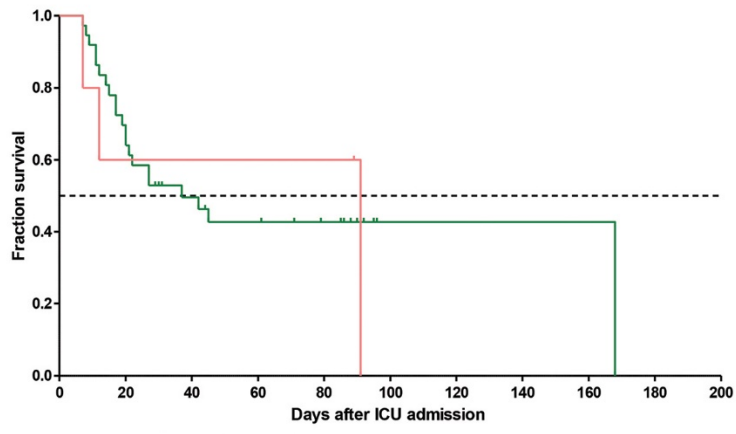
#One value of >6.0 entered as 6.0.

**One value of >500 pg/mL entered as 500 pg/mL.



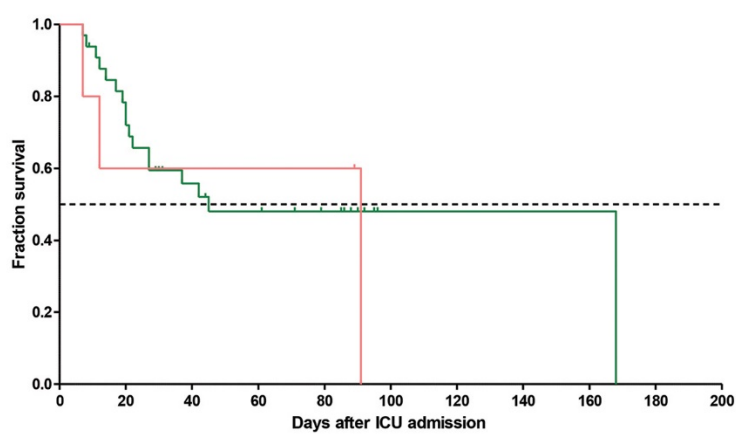
Appendix Figure 1. Forest plots of risk factors for the development of COVID-19–associated pulmonary aspergillosis (CAPA) and intensive care unit (ICU) death in a multinational observational study of CAPA. A) Binary logistic regression analysis of risk factors for the development of CAPA in the discovery cohort. B) Binary logistic regression analysis of risk factors for ICU death in the discovery cohort. C) Binary logistic regression analysis of risk factors for the development of CAPA in the validation cohort. D) Binary logistic regression analysis of risk factors for ICU death in the validation cohort. Error bars indicate 95% CI; dots indicate aOR. Dotted vertical lines indicate an aOR of 1. aOR, adjusted odds ratio; 95% CI COVID-19, coronavirus disease; COPD, Chronic obstructive pulmonary disease; IS, immunosuppressant drugs; RRT, renal replacement therapy; SAPS, simplified acute physiology score.

A



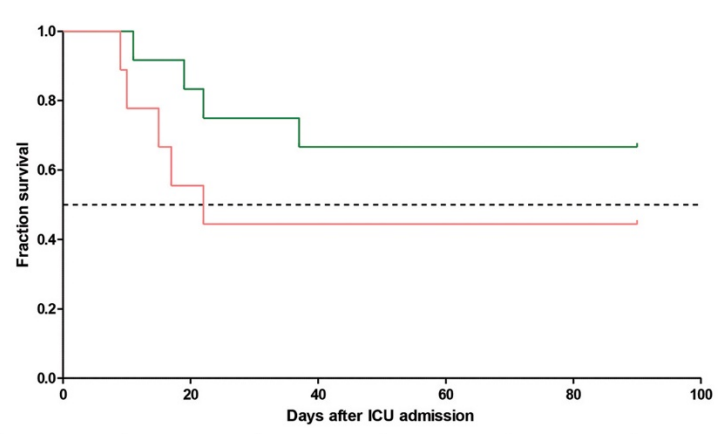
Number at risk										
CAPA, AF	37	25	16	13	10	2	2	2	2	2
CAPA, no AF	5	4	4	4	4	1	1	1	1	1

B



Number at risk										
CAPA, AF	33	25	16	13	10	2	2	2	2	1
CAPA, no AF	5	4	4	4	4	1	1	1	1	1

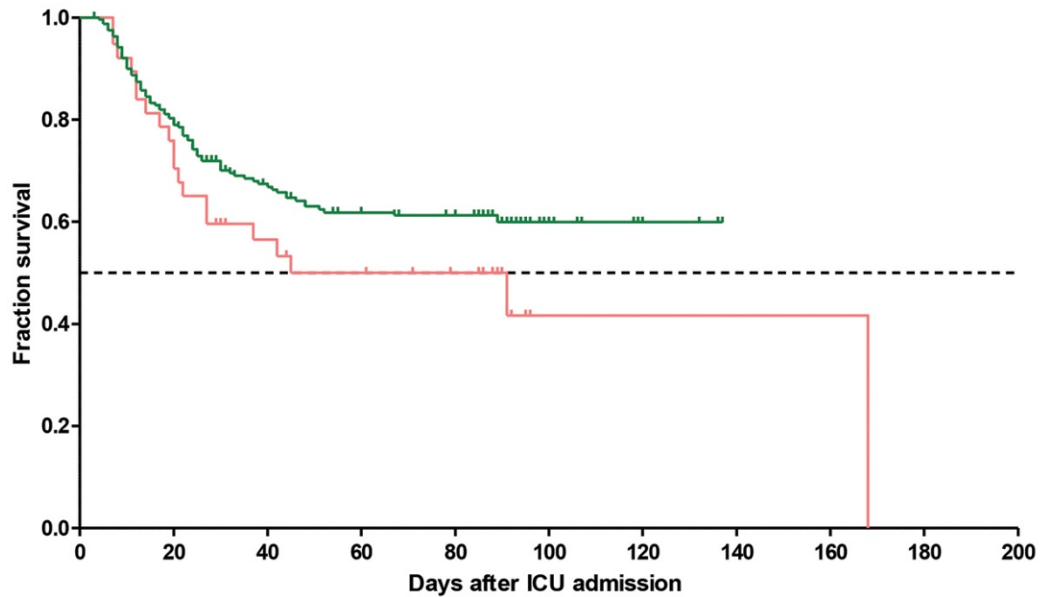
C



Number at risk						
CAPA, AF	12	11	9	9	9	8
CAPA, no AF	9	6	5	5	5	4

Appendix Figure 2. Kaplan-Meier survival curves comparing CAPA patients who did and did not receive any antifungal (AF) treatment during intensive care unit (ICU) admission in a multinational observational study of COVID-19–associated pulmonary aspergillosis (CAPA). Green indicates CAPA patients receiving

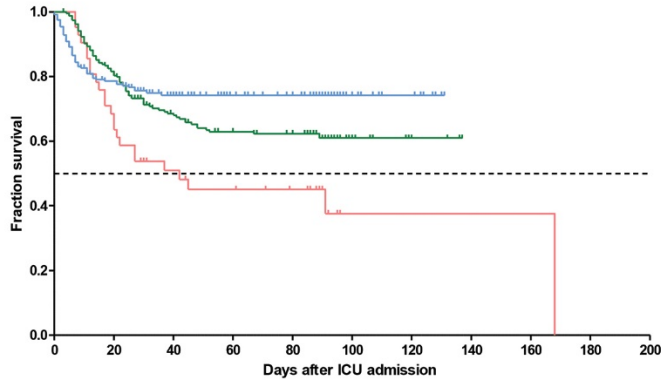
AF; salmon indicates patients not receiving AF. Tables show the number of patients at risk for ICU death per time point after ICU admission. Dashed horizontal lines indicate a survival fraction of 0.5. A) Discovery cohort (n = 42); (p = 0.869); B) discovery cohort after designating patients with possible CAPA to the CAPA excluded group (n = 38) (p = 0.683); C) validation cohort (n = 21) (p = 0.212). Survival analysis was performed by Mantel-Cox log rank test. COVID-19, coronavirus disease.



Number at risk										
CAPA	38	28	19	16	13	2	2	2	2	1
CAPA excluded	241	187	124	107	101	10	4	1	1	1

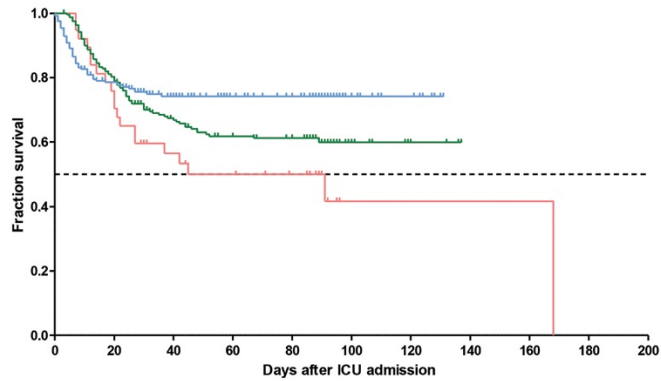
Appendix Figure 3. Kaplan-Meier survival curves comparing patients in a multinational observational study COVID-19–associated pulmonary aspergillosis (CAPA). The graph compares patients classified with CAPA (salmon) with CAPA excluded patients (green) when possible CAPA was classified as CAPA excluded (n = 279). Survival analysis was calculated by Mantel-Cox log rank test and shows survival over time is no longer significantly different within the discovery cohort when patients with possible CAPA are designated to the CAPA excluded group (p = 0.134). Median estimated survival in the CAPA group is 45 days. Tables show the number of patients at risk for ICU death per time point after ICU admission. Dashed horizontal lines indicate a survival fraction of 0.5. COVID-19, coronavirus disease.

A



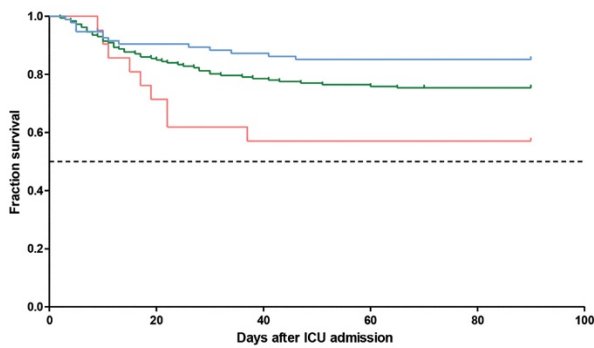
Number at risk										
CAPA	42	28	19	16	13	2	2	2	2	1
CAPA excluded	237	187	124	107	101	10	4	1	1	1
CAPA not classifiable	240	167	94	77	65	17	11	2	2	2

B



Number at risk										
CAPA	38	28	19	16	13	2	2	2	2	1
CAPA excluded	241	187	124	107	101	10	4	1	1	1
CAPA not classifiable	240	167	94	77	65	17	11	2	2	2

C



Number at risk						
CAPA	21	16	13	13	13	12
CAPA excluded	188	160	148	143	141	140
CAPA not classifiable	95	86	83	81	81	80

Appendix Figure 4. Kaplan-Meier survival curves comparing patients in a multinational observational study of COVID-19–associated pulmonary aspergillosis (CAPA). The graph compares patients classified with CAPA (salmon), CAPA excluded (green) and CAPA not classifiable (blue). Survival analysis was performed by Mantel-Cox log rank test. A) Discovery cohort (n = 519); survival over time is significantly different between the 3 groups in this cohort (p = 0.007). B) Discovery cohort after designating possible

CAPA patients to the CAPA excluded group (n = 519), survival over time remained significantly different between the 3 groups (p = 0.040). C) Validation cohort (n = 304); survival over time is significantly different between the 3 patient groups (p = 0.014). Tables show the number of patients at risk for ICU death per time point after ICU admission. Dashed horizontal line indicates a survival fraction of 0.5. COVID-19, coronavirus disease.