

# A Multifaceted and Multi-Institutional Analysis of the COVID19-Associated Mucormycosis **Outbreak in the Delhi Area Indicates the Simultaneous Convergence of Multiple Risk Factors**

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#### Background

A major outbreak of COVID-19-associated mucormycosis (CAM) in India in spring 2021 aggravated the death toll of COVID-19. As the causes of that CAM outbreak remain unclear, we performed a multifaceted study of environmental, host-, pathogen-, and healthcarerelated factors in CAM patients in the metropolitan New Delhi area.

## Methods

<u>Case/control design</u>: We reviewed all adult patients (≥18 years of age, n = 50) who were diagnosed with culture- or biopsy-proven mucormycosis at 7 participating public and private hospitals in the New Delhi area between April 1 and June 30, 2021 and within 60 days of a prior COVID-19 infection. Prior Covid-19 infection was defined as COVID-19 symptoms and at least one positive SARS-CoV-2 PCR. Sixty-nine contemporary adult patients admitted to the same hospitals for treatment of PCR-confirmed COVID-19 infection served as the control cohort. We performed two distinct case/control analyses, as summarized in Table 1.

#### Table 1: Summary of case/control analyses performed.

	Analysis 1	Analysis 2
Inclusion criteria for the CAM cohort	Patients hospitalized for CAM, regardless of their hospitalization status for COVID-19 (n = 50)	Patients developing CAM who had already been hospitalized for COVID-19 (n = 31)
CAM cohort includes patients with	Severe COVID-19 Moderate COVID-19 Mild COVID-19	Severe COVID-19 Moderate COVID-19
Control cohort for both analyses	Contemporary patients (r participating hospitals fo	n = 69) hospitalized at the r treatment of COVID-19
Control cohort includes patients with	Severe C Moderate	OVID-19 COVID-19

Statistical analyses: Univariate comparisons of continuous variables were performed using the Wilcoxon rank-sum test. Categorial variables were compared using Chi-square or Fisher's exact test, as appropriate. A logistic regression model with backward elimination was used to identify independent predictors of CAM development. All tests were 2-sided with a significance level of p < 0.05.

Collection of meteorological data: Temperature, relative humidity, and evaporation at the Agrometeorological Observatory New Delhi (28°38'23"N, 77°09'27"E, altitude: 229 m above sea level) were recorded daily by the Indian Agricultural Research Institute.

Quantification of fungal spore concentrations in outdoor air: Fungal spore concentrations in environmental air were determined by the Vallabhbhai Patel Chest Institute using a 24-hour volumetric trap air sampler (Burkard Manufacturing Co Ltd) with an air flow of 10 L/min. Total mold particle recovery per day was determined by microscopic examination of slides fitted in the lid assembly of the air sampler.

Microbiological analyses: Tissue specimens were processed for direct microscopic examination by 10% KOH-Blankophor staining. Additionally, tissue samples and hospital fomite swabs were cultured on Sabouraud dextrose agar. Patient isolates were identified by MALDI-TOF mass spectrometry and ITS sequencing.

Whole genome sequencing (WGS): Fungal DNA from selected patient isolates was sequenced with a NOVASEQ6000 sequencer (Illumina). Paired-end reads were aligned to reference isolates and additional species-typed Sequence Read Archive isolates (NCBI).

ble 2: Univariate comparison of pat						
		Analysis 1			Analysis 2	
nless indicated otherwise, number of	CAM cases	Controls	P-value	CAM cases	Controls	P-value
atients and percentages are given	n = 50	n = 69	0.07	n = 31	n = 69	0.00
ge (years), median (range)	57 (34 – 77)	50 (19 – 86)	0.07	58 (34 – 77)	50 (19 – 86)	0.09
ender, male	35 (70)	47 (68)	0.83	21 (68)	47 (68)	0.97
ving conditions	7/40 (47)		0.04	<b>E</b> (00, (40)		0.04
Rural	7/42 (17)	3/68 (4)		5/28 (18)	3/68 (4)	
Urban	35/42 (83)	65/68 (96)		23/28 (82)	65/68 (96)	
Unknown	8	1		3	1	
nderlying conditions						
Arterial Hypertension	15 (30)	27 (39)	0.30	9 (29)	27 (39)	0.33
Chronic kidney disease	1 (2)	2 (3)	> 0.99	0 (0)	2 (3)	> 0.99
Chronic liver disease	2 (4)	0 (0)	0.17	2 (6)	0 (0)	0.09
Heart failure or coronary artery disease	1 (2)	4 (6)	0.40	1 (3)	4 (6)	> 0.99
Cancer (any malignancy)	5 (10)	0 (0)	0.01	4 (13)	0 (0)	< 0.01
Hematological malignancy	3 (6)	0 (0)	0.07	3 (10)	0 (0)	0.03
Solid tumor	2 (4)	0 (0)	0.17	1 (3)	0 (0)	0.31
Chronic lung disease	2 (4)	7 (10)	0.30	1 (3)	7 (10)	0.43
Asthma	2 (4)	5 (7)	0.70	1 (3)	5 (7)	0.66
COPD	0 (0)	2 (3)	0.51	0 (0)	2 (3)	> 0.99
Surgery within last 14 days	1 (2)	0 (0)	0.42	1 (3)	0 (0)	0.31
nmunosuppressive therapy <sup>a</sup>	4 (8)	0 (0)	0.03	3 (10)	0 (0)	0.03
ytopenia	- (-)					
Neutropenia (ANC < 1000)	1/42 (2)	0/68 (0)	0.38	1/27 (4)	0/68 (0)	0.28
Lymphopenia (ALC < 1000)	20/43 (47)	20/68 (29)	0.07	13/27 (48)	20/68 (29)	0.08
OVID-19 severity	20/40 (47)	20/00 (23)	< 0.001		20/00 (23)	< 0.01
Mild	19 (38)	N/A	< 0.001	N/A	N/A	< 0.01
Moderate	18 (36)			18 (58)		
	· · · · ·	57 (83)			57 (83)	
Severe	13 (26)	12 (17)	< 0.001		12 (17)	ΝΙ/Δ
ospitalization for COVID-19	31 (62)	69 (100)	< 0.001	50 (100)	69 (100)	N/A
U admission for COVID-19	14 (28)	34 (49)	0.02	14 (45)	34 (49)	0.70
owest SO <sub>2</sub> at room air, median (range)	90% (60% – 98%)	85% (65% – 98%)	< 0.01		85% (65% – 98%)	0.24
upplemental oxygen for COVID-19	30 (60)	66 (96)	< 0.001	22 (71)	66 (96)	< 0.01
ighest level of oxygen support			0.01			< 0.01
Nasal cannula	12/30 (40)	23/66 (35)		5/22 (23)	23/66 (35)	
Face mask/non-rebreathing mask	5/30 (17)	31/66 (47)		4/22 (18)	31/66 (47)	
BiPap/non-invasive ventilation	11/30 (37)	9/66 (14)		11/22 (50)	9/66 (14)	
Invasive ventilation	2/30 (7)	3/66 (5)		2/22 (9)	3/66 (5)	
Iucocorticosteroids for COVID-19	40 (80)	67 (97)	< 0.01	26 (84)	67 (97)	0.03
igh dose glucocorticosteroids	24/34 (55)	58/68 (85)	< 0.001	19/30 (63)	58/68 (85)	0.02
Unknown	6	1		1	1	
ays of steroids, median (range)	11 (5 – 20)	15 (3 – 45)	0.05	11 (5 – 20)	15 (3 – 45)	0.12
onoclonal antibodies	2 (4)	4 (6)	> 0.99	2 (6)	4 (6)	> 0.99
Tocilizumab	2 (4)	0 (0)		2 (4)	0 (0)	
Itolizumab	0 (0)	1 (1)		0 (0)	1 (1)	
Bevacizumab	0 (0)	3 (4)		0 (0)	3 (4)	
ther COVID-19 therapeutics	(-/					
Remdesivir	10 (20)	30 (43)	< 0.01	10 (32)	30 (43)	0.29
Intravenous immunoglobulins	2 (4)	0 (0)	0.17	2 (6)	0 (0)	0.09
Baricitinib	0 (0)	8 (12)	0.02	0 (0)	8 (12)	0.06
abetes mellitus (DM)	0 (0)	0 (12)	0.02			0.00
Previously diagnosed DM	33 (66)	19 (28)	< 0.001	21 (69)	19 (28)	< 0.001
	· · · · ·	<b>x 7</b>		21 (68)		
Newly diagnosed DM	12/17 (71)	7/50 (14)	< 0.001	6/10 (60)	7/50 (14)	< 0.01
Diabetic ketoacidosis	4/45 (9)	0/25 (0)	0.29		0/25 (0)	0.49
Latest HbA1c, median (range)	8.0 (4.2 – 16.6)	6.7 (3.2 – 12.4)	< 0.001	7.6 (4.2 – 14.8)	6.7 (3.2 – 12.4)	< 0.001
Latest HbA1c ≥ 8.0	25 (50)	7/65 (11)	< 0.001	12 (39)	7/65 (11)	< 0.01
Highest glucose level <sup>b</sup>	385 (180 – 600)	303 (110 – 588)	0.04	373 (180 – 570)	303 (110 – 588)	0.16

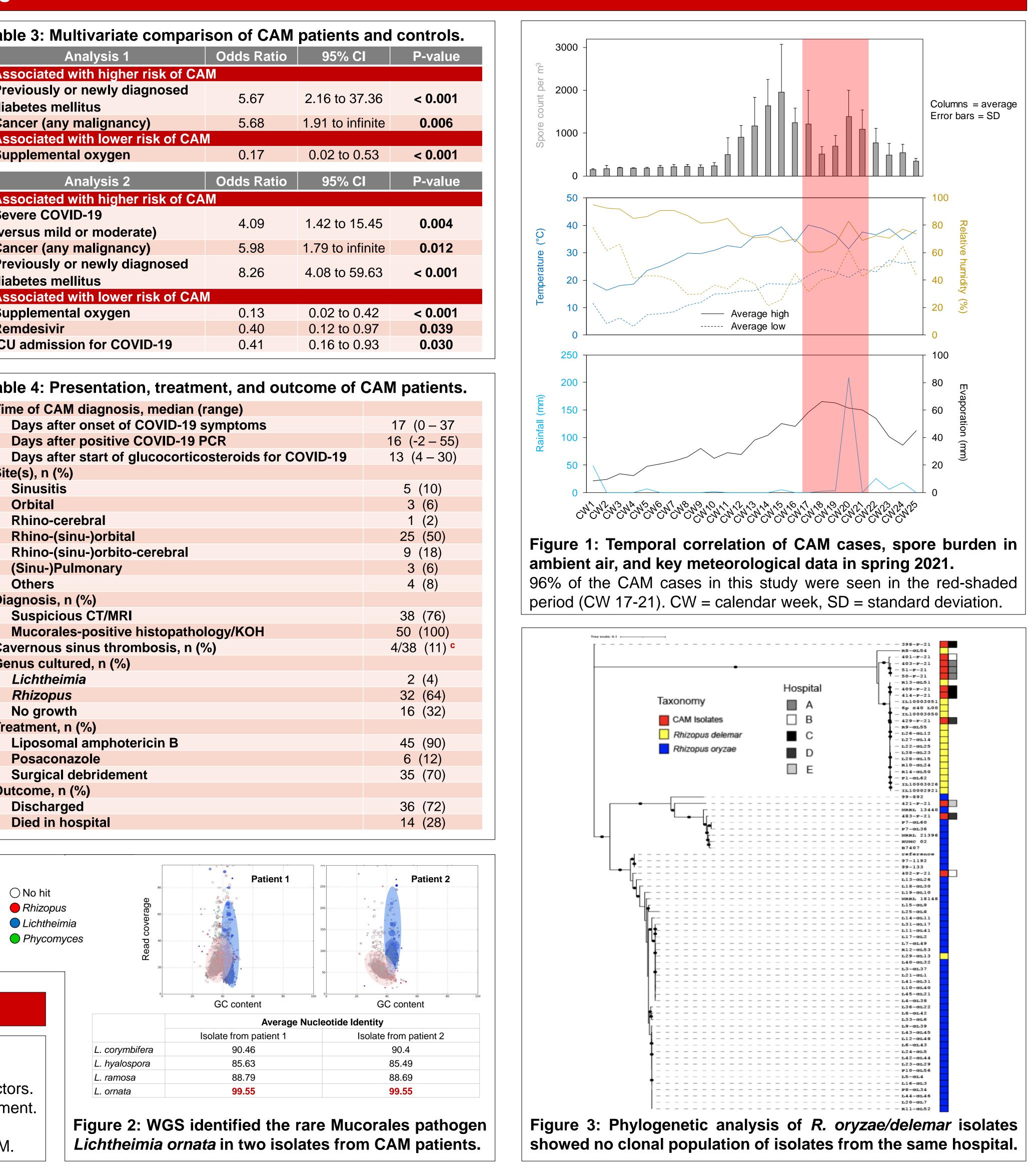
> Consistent with prior studies,<sup>1,2</sup> previously or newly diagnosed diabetes mellitus was a key predictor of CAM risk, especially when poorly controlled. > Surrogates of access to advanced treatment of COVID-19 (ICU admission, remdesivir, supplemental oxygen) were associated with lower CAM risk > The CAM incidence peak was preceded by a significant uptick in environmental spore concentrations but was not linked to specific meteorological factors.  $\succ$  Fomite cultures were negative (data not shown) and WGS showed no clonal population of patient isolates  $\rightarrow$  no link of CAM cases to hospital environment. > Rhizopus was the predominant causative genus (64%), but two cases of the rare pathogen Lichtheimia ornata were detected by WGS. > Altogether, our data suggest that an intersection of host, environmental, pathogen and healthcare-related factors contributed to the emergence of CAM.

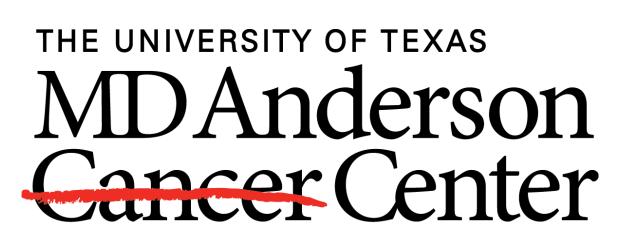
#### Conclusions

Footnotes: <sup>a</sup> any (systemic) immunosuppressive agent received prior to the diagnosis of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glucose level recorded during the treatment of COVID-19; <sup>b</sup> highest glu

### Results

Table 3: Multivariate comparAnalysis 1	Odds Ratio	95% CI
Analysis 1 Associated with higher risk of C		
Previously or newly diagnosed		
diabetes mellitus	5.67	2.16 to 37.3
Cancer (any malignancy)	5.68	1.91 to infini
Associated with lower risk of CA		1.91 (0 11111
Supplemental oxygen	0.17	0.02 to 0.5
Analysis 2	Odds Ratio	95% CI
Associated with higher risk of C	AM	
Severe COVID-19	4.09	1.42 to 15.4
(versus mild or moderate)		
Cancer (any malignancy)	5.98	1.79 to infini
Previously or newly diagnosed	8.26	4.08 to 59.6
diabetes mellitus		
Associated with lower risk of CA		
Supplemental oxygen	0.13	0.02 to 0.42
Remdesivir	0.40	0.12 to 0.9
ICU admission for COVID-19	0.41	0.16 to 0.9
Table 4: Presentation, treatTime of CAM diagnosis, medianDays after onset of COVID-19	(range)	come of C
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Time of CAM diagnosis, median Days after onset of COVID-19 Days after positive COVID-19 Days after start of glucocortic Site(s), n (%) Sinusitis	(range) symptoms PCR	
Time of CAM diagnosis, median Days after onset of COVID-19 Days after positive COVID-19 Days after start of glucocortic Site(s), n (%) Sinusitis Orbital	(range) symptoms PCR	
Time of CAM diagnosis, median Days after onset of COVID-19 Days after positive COVID-19 Days after start of glucocortic Site(s), n (%) Sinusitis Orbital Rhino-cerebral	(range) symptoms PCR	
Time of CAM diagnosis, median Days after onset of COVID-19 Days after positive COVID-19 Days after start of glucocortic Site(s), n (%) Sinusitis Orbital Rhino-cerebral Rhino-(sinu-)orbital	(range) symptoms PCR	
Time of CAM diagnosis, median Days after onset of COVID-19 Days after positive COVID-19 Days after start of glucocortic Site(s), n (%) Sinusitis Orbital Rhino-cerebral Rhino-cerebral Rhino-(sinu-)orbital Rhino-(sinu-)orbito-cerebral	(range) symptoms PCR	
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