Brief Summary of Findings on the Association Between Alpha-1 Antitrypsin Deficiency and Severe COVID-19 Outcomes Prepared and reviewed by:

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Six studies were included for analysis. Three cohort studies, two ecological studies, and one case report were retrieved that reported data on underlying Alpha-1 Antitrypsin Deficiency (A1AT) and severe COVID-19 outcomes.

• The evidence is inconsistent and inconclusive on the association between A1AT and mortality¹⁻⁵, ICU admission^{1, 5} intubation⁵, ventilation^{5, 6}, and hospitalization^{1, 5, 6}. Evidence was also insufficient to determine if the relationship between A1AT and mortality was influenced by severity or confounding with other underlying conditions.

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A. Methods

The aim of this review was to identify and synthesize the best available evidence on the association between alpha-1 antitrypsin (A1AT) deficiency and severe COVID-19 outcomes to update the Centers for Disease Control and Prevention (CDC) website on underlying conditions for a consumer and a provider-specific website with more rigorous information.

A.1. Literature Search

A list of search terms was developed to identify the literature most relevant to the population, exposure, comparator, and outcomes (PECO) question. Clinical experts and library scientists were consulted to develop a robust list of search terms. These terms were then incorporated into search strategies, and these searches were performed in OVID using the COVID-19 filter for all articles from the beginning of each database until July 19, 2021. The detailed search strategies for identifying primary literature and the search results are provided in *Part B*. References were included if retrieved by the literature search and reported exposures and outcomes relevant to this review.

A.2. Study Selection

Titles and abstracts from references were screened by dual review (A.H., M.M., D.O.S., or E.C.S.). Full-text articles were retrieved if they were:

- 1. relevant to the PECO question;
- 2. primary research; and
- 3. written in English.

Part B presents the full list of exclusion criteria. The full texts of selected articles were then screened by two independent reviewers, and disagreements were resolved by discussion (M.M, A.H., D.O.S., or E.C.S.). After the full-text screening was complete, a bibliography of the articles selected for inclusion was vetted with subject matter experts. Additional studies suggested by the subject matter experts were screened for inclusion as described above. The results of the study selection process are depicted in Figure 1.

Figure 1. Results of the Study Selection Process



A.4. Data Extraction and Synthesis

Methodologic data and results of relevant outcomes from the studies meeting inclusion criteria were extracted into standardized evidence tables. Data and analyses were extracted as presented in the studies. For the purposes of this review, statistical significance was defined as $p \le 0.05$.

A.5. Aggregation of the Evidence

The internal validity associated with each study was assessed using scales developed by the Division of Healthcare Quality Promotion and scores were recorded in the evidence tables. The *Part B* includes the questions used to assess the quality of each study design. The strength, magnitude, precision, consistency, and applicability of results were assessed for all comparators. The overall confidence in the evidence base is reported in the aggregation tables in the *Part B*.

A.6. Reviewing and Finalizing the Systematic Review

Draft findings, aggregation tables, and evidence tables, were presented to CDC subject matter experts for review and input. Following further revisions, the summary will be published on the CDC website.

B. Systematic Literature Review Results

B.1. Search Strategies and Results

Table 1 Alpha-1 Antitrypsin Deficiency Search Conducted July 19, 2021.

Database	Strategy	Records 07/19/2021
Medline	Alpha 1 antitrypsin OR a1 antitrypsin OR a-1 antitrypsin OR A1AT deficiency OR A1AD	32
(OVID)	AND	
1946-	Limit COVID-19 (validated filter)	
	2020- ;	
Embase	Alpha 1 antitrypsin OR a1 antitrypsin OR a-1 antitrypsin OR A1AT deficiency OR A1AD	62
(OVID)	AND	
1988-	Limit COVID-19 (validated filter)	-29
		duplicates
	2020- ; NOT pubmed/medline	
		=33
		unique items
Global Health	(novel coronavir* OR novel corona virus* OR 2019 coronavirus OR coronavirus disease OR	11
(OVID)	coronavirus 2019 OR betacoronavir* OR covid19 OR covid OR nCoV OR novel CoV OR CoV 2 OR CoV2	
	OR sarscov2 OR sars-cov OR sarscov OR 2019nCoV OR 2019-nCoV OR wuhan virus*)	-9
	AND	duplicates
	Alpha 1 antitrypsin OR a1 antitrypsin OR a-1 antitrypsin OR A1AT deficiency OR A1AD	
	2020-	=2
		unique items
CAB Abstracts	(novel coronavir* OR novel corona virus* OR 2019 coronavirus OR coronavirus disease OR	3
(OVID)	coronavirus 2019 OR betacoronavir* OR covid19 OR covid OR nCoV OR novel CoV OR CoV 2 OR CoV2	
	OR sarscov2 OR sars-cov OR sarscov OR 2019nCoV OR 2019-nCoV OR wuhan virus*)	-3
	AND	duplicates
	Alpha 1 antitrypsin OR a1 antitrypsin OR a-1 antitrypsin OR A1AT deficiency OR A1AD	
	2020-	=0
		unique items
PsycInfo	(novel coronavir* OR novel corona virus* OR 2019 coronavirus OR coronavirus disease OR	0
(OVID)	coronavirus 2019 OR betacoronavir* OR covid19 OR covid OR nCoV OR novel CoV OR CoV 2 OR CoV2	
1987-	OR sarscov2 OR sars-cov OR sarscov OR 2019nCoV OR 2019-nCoV OR wuhan virus*)	

Database	Strategy	Records 07/19/2021
	AND	
	Alpha 1 antitrypsin OR a1 antitrypsin OR a-1 antitrypsin OR A1AT deficiency OR A1AD	
	2020-	
CINAHL	("novel coronavir*" OR "novel corona virus*" OR "2019 coronavirus" OR betacoronavir* OR covid19	2
(EbscoHost)	OR covid OR nCoV OR "novel CoV" OR "CoV 2" OR CoV2 OR sarscov2 OR sars-cov OR sarscov OR	
	2019nCoV OR 2019-nCoV OR "wuhan virus*")	-2
	AND	duplicates
	"Alpha 1 antitrypsin" OR "a1 antitrypsin" OR "a-1 antitrypsin" OR A1AT deficiency OR A1AD	
		=0
	2020- ;	unique items
Academic Search	("novel coronavir*" OR "novel corona virus*" OR "2019 coronavirus" OR betacoronavir* OR covid19	7
Complete	OR covid OR nCoV OR "novel CoV" OR "CoV 2" OR CoV2 OR sarscov2 OR sars-cov OR sarscov OR	
	2019nCoV OR 2019-nCoV OR "wuhan virus*")	-2
	AND	duplicates
	"Alpha 1 antitrypsin" OR "a1 antitrypsin" OR "a-1 antitrypsin" OR A1AT deficiency OR A1AD	
		=5
	2020- ;	unique items
Scopus	TITLE-ABS("novel coronavir*" OR "novel corona virus*" OR "2019 coronavirus" OR betacoronavir*	27
	OR covid19 OR covid OR ncov OR "CoV 2" OR cov2 OR sarscov2 OR sars-cov OR sarscov OR 2019ncov	
	OR 2019-nCoV OR "novel CoV" OR "wuhan virus") AND TITLE-ABS("Alpha 1 antitrypsin" OR "a1	-25
	antitrypsin" OR "a-1 antitrypsin" OR A1AT deficiency OR A1AD)	duplicates
		=2
		unique items
WHO Global	"Alpha 1 antitrypsin" OR "a1 antitrypsin" OR "a-1 antitrypsin" OR A1AT deficiency OR A1AD	24
COVID Literature		24
Database		-24
		auplicates
		-0
		=U
		unique items

Database	Strategy	Records 07/19/2021
Coronavirus	TI,AB("Alpha 1 antitrypsin" OR "a1 antitrypsin" OR "a-1 antitrypsin" OR A1AT deficiency OR A1AD)	13
Research		
Database		-9
		duplicates
		=4
		unique items
Cochrane Library	"Alpha 1 antitrypsin" OR "a1 antitrypsin" OR "a-1 antitrypsin" OR A1AT deficiency OR A1AD	8
	AND	
	"novel coronavir*" OR "novel corona virus*" OR "2019 coronavirus" OR betacoronavir* OR covid19	-2
	OR covid OR ncov OR "CoV 2" OR cov2 OR sarscov2 OR sars-cov OR sarscov OR 2019ncov OR 2019-	duplicates
	nCoV OR "novel CoV" OR "wuhan virus"	
		=6
		unique items
Clinicaltrials.gov	"Alpha 1 antitrypsin" OR "a1 antitrypsin" OR "a-1 antitrypsin" OR A1AT DEFICIENCY OR A1AD	8
	"novel coronavir*" OR "novel corona virus*" OR "2019 coronavirus" OR betacoronavir* OR covid19	
	OR covid OR ncov OR "CoV 2" OR cov2 OR sarscov2 OR sars-cov OR sarscov OR 2019ncov OR 2019-	
	nCoV OR "novel CoV" OR "wuhan virus"	

B.2. Study Inclusion and Exclusion Criteria

Inclusion Criteria: Studies were included at the title and abstract screen if they:

- were relevant to the key question "What is the association between alpha-1 antitrypsin deficiency and severe COVID-19?";
 - exposures: Alpha-1 Antitrypsin (A1AT) Deficiency.
 - o outcomes: mortality, ICU admission, intubation, ventilation, and hospitalization
- were primary research;
- were written in English (can be seen as [language] in title); and
- examined humans only.

Exclusion Criteria: Studies were excluded at full-text review if they:

- were not available as full-text;
- were a conference abstract, poster, or reply letter;

- reported autopsy results; and
- reported only composite outcome measures for "severe covid-19".

B.3. Evidence Review: Alpha-1 Antitrypsin (A1AT) Deficiency and Severe COVID-19

B.3.a. Strength & Direction of Evidence

Table 2 The Association Between A1AT Deficiency and Severe COVID-19 Outcomes

Outcome	Results
Mortality	Overall, data from five studies ¹⁻⁵ (N = 14,402) is inconsistent and inconclusive on an association between underlying α -1
	antitrypsin deficiency (A1AT deficiency) and mortality. Two studies ^{2, 3} were found to have a high threat to internal validity, while
	one study ⁴ was found to have a moderate threat to internal validity. Internal validity assessments are not completed for studies ^{1,}
	⁵ with less than 10 people with A1AT deficiency.
	• Strength of Association: One small study reported a measure of association suggesting no difference. ¹
	 Precision of Association: No studies reported confidence intervals.
	Consistency of Association: Overall, the evidence is inconsistent.
	• Applicability of Association: The population and setting were directly applicable to the question. Two studies ^{1, 2} were
	conducted in European countries, two studies ^{3, 4} were conducted internationally, and one study ⁵ was conducted in the
	U.S. Studies were conducted among adults of all age groups including population-level, community, and hospital settings.
	Summary of Evidence:
	 Two ecological studies^{3, 4} (N = NR) reported correlation coefficients suggesting a positive correlation between mortality and A1AT deficiency in people with COVID-19.
	 One cross-sectional ecological study³ (N = NR) reported a moderate positive correlation suggests an association
	of national rates of A1AT deficiency with the national rates of COVID-19 fatalities [$R = 0.54$, $p < 0.01$]. This
	correlation remained significant even after adjusting for the human development index.
	 One ecological study⁴ (N = NR) reported data suggesting a strong correlation between global COVID-19 mortality
	and A1AT deficiency [R = 0.86, p = NR]. This correlation remained strong in a sub-analysis of data from European,
	and North and South American countries $[R = 0.89, p = NR]$ but did not persist in a sub-analysis of data from
	Asian and African countries [R = 0.025, p = NR].
	 Three studies^{1,2,3} (N = 14,402) reported proportions suggesting no association between mortality and A1AT deficiency
	among people with COVID-19.
	 One telephone survey¹ (N = 8) of Italians with severe A1AT deficiency reported eight cases of self-reported
	COVID-19+ status, with a mortality rate [12.5% (1/8)] similar to the mortality rate in the national population
	[13.9% (RR: 0.90)]. The survey only reached 35% of the Italian cohort with severe A1AT deficiency.

Outcome	Results
	 One cohort study² (N = 14,393) of people tested for genotype variants, suggested similar proportions of mortality
	among people with A1AT deficiency and COVID-19 compared to those with only COVID-19 [3.17% (53/1,670) vs.
	2.77% (353/12,723)].
	 One case report⁵ (N = 1) reported an immunosuppressed 67-year-old female with homozygous Z-allele mutation
	A1AT deficiency and COVID-19 who did not die. The patient had a history of several comorbidities, including liver
	transplant, chronic kidney disease stage IIIa, cirrhosis, COPD, hypertension, deep venous thrombosis, and
	uncontrolled insulin-dependent type 2 diabetes mellitus. Internal validity is not assessed for case reports.
ICU Admission	Overall, limited data from two studies ^{1, 5} (N = 9) is inconclusive on the association between underlying A1AT deficiency and ICU
	admission. Internal validity assessments are not completed for studies ^{1, 5} with less than 10 people with A1AT deficiency.
	 Strength of Association: No studies reported measures of association.
	Precision of Association: No studies reported confidence intervals.
	Consistency of Association: Overall, the number of events, and sample sizes were too small to determine consistency.
	 Applicability of Association: The population and setting were directly applicable to the question. One study¹ was
	conducted in a European country, and one ⁵ was conducted in the U.S. Studies were conducted among adults in
	community and hospital settings.
	Summary of Evidence:
	 Two studies^{1, 5} reported proportions or cases on ICU admission and A1AT deficiency in COVID-19 patients.
	 One cohort study¹ (N = 8) of Italians with severe A1AT deficiency and COVID-19 reported an ICU admission rate
	of 0% (0/8) among patients with COVID-19 and underlying A1AT deficiency.
	 One case report⁵ (N = 1) of an immunosuppressed 67-year-old female with homozygous Z-allele mutation A1AT
	deficiency in Pennsylvania, US reported that the patient was upgraded to the Medical intensive care unit (MICU)
	after presenting with COVID-19 mediated hypoxic respiratory failure. The patient had a history of several
	comorbidities, including liver transplant, chronic kidney disease stage IIIa, cirrhosis, COPD, hypertension, deep
	venous thrombosis, and uncontrolled insulin-dependent type 2 diabetes mellitus. Internal validity is not assessed
	for case reports.
Intubation	Overall, limited evidence from one study is insufficient to determine an association between A1AT deficiency and intubation.
	Aggregation indices are not evaluated for outcomes reported by only one study.

Outcome	Results
	 One case report⁵ (N = 1) reported intubation in an immunosuppressed 67-year-old female with homozygous Z-allele
	mutation A1AT deficiency after presenting with COVID-19 mediated hypoxic respiratory failure. Internal validity is not
	assessed for case reports.
Ventilation	Overall, the evidence from two studies ^{5, 6} (N = 10) is inconsistent and inconclusive on an association between underlying A1AT
	deficiency and ventilation. Internal validity assessments are not completed for studies ^{5, 6} with less than 10 people with A1AT
	deficiency.
	Strength of Association: No studies reported measures of association.
	Precision of Association: No studies reported confidence intervals.
	• Consistency of Association: Overall, the number of events, and sample sizes were too small to determine consistency.
	• Applicability of Association: The population and setting were directly applicable to the question. One study ⁵ was
	conducted in the U.S., and one study was conducted in a European country. Studies were conducted among adults in
	hospital settings.
	Summary of Evidence:
	• Two studies ^{5, 6} (N = 10) reported data on ventilation and A1AT deficiency in COVID-19 patients.
	 One cohort⁶ (N = 9) reported the proportion of COVID-19 patients with A1AT deficiency that required ventilation
	[0% (0/77)]. This study had a small sample size with a low number of events and no comparison group,
	decreasing confidence in the finding.
	 One case report⁵ (N = 1) of a 67-year-old female with A1AT deficiency in Pennsylvania, US reported that the
	patient was mechanically ventilated. Internal validity is not assessed for case reports.
Hospitalization	Overall, the evidence from three studies ^{1, 5, 6} (N = 18) is consistent, but inconclusive on an association between underlying A1AT
	deficiency and hospitalization. Internal validity assessments are not completed for studies ^{1, 5, 6} with less than 10 people with
	A1AT deficiency.
	• Strength of Association: No studies reported measures of association.
	Precision of Association: No studies reported confidence intervals
	 Consistency of Association: Descriptive statistics are consistent across studies.

Outcome	Results
	 Applicability of Association: The population and setting were directly applicable to the question. Two studies^{1, 6} were conducted in European countries, and one study⁵ was conducted in the U.S. Studies were conducted among adults in hospital settings.
	Summary of Evidence:
	 Three studies^{1, 5, 6} reported data on hospitalization and A1AT deficiency in COVID-19 patients. One cohort study¹ (N = 8) using telephone survey data reported the proportions of participants with underlying A1AT deficiency that were hospitalized among people with COVID-19 [37.5% (3/8)]. The study did not have a comparison group for this outcome and had a small sample size with a low number of events, decreasing confidence in this finding. One cohort study⁶ (N = 9) reported the proportion of participants with underlying A1AT deficiency that were hospitalized [4% (3/77)]. This study had a small sample size with a low number of events and no comparison group, decreasing confidence in the finding. One case report⁵ (N = 1) of a 67-year-old female with A1AT deficiency in Pennsylvania, US reported that the patient was admitted to the hospital. Internal validity is not assessed for case reports.

Table 3 Increasing Severity of Underlying A1AT Deficiency and Severe COVID-19 Outcomes

Outcome	Results
Mortality	Overall, the evidence is inconsistent and inconclusive on an association between the severity markers of A1AT deficiency and
	mortality. One study was found to have moderate ⁴ and the other study ² was found to have high threat to internal validity.
	• Strength of Association: The correlation coefficient ranged from 0.025 - 0.950.
	 Precision of Association: Confidence intervals were not reported.
	 Consistency of Association: Evidence is inconsistent
	 Applicability of Association: The population and setting were directly applicable to the question. One study² was conducted in Europe and one⁴ was conducted internationally.
	Summary of Evidence:
	• Two studies ^{2, 4} reported data on mortality and A1AT deficiency genotypes in COVID-19 patients.

One ecological study ⁴ (N = 68 countries) reported a stronger correlation between COVID-19 and mortality in
populations with the underlying PiZ genotype than in populations with the PiS genotype.
 One cohort study² (N = 14,393) of Biobank and national death registries data in the United Kingdom reported on people with different genotypes representing severity of A1AT deficiency. While proportions of mortality were
consistent across the genotypes, no conclusions can be drawn from this study because of a small number of events and lack of statistical significance

Table 4 Risk Markers of Underlying A1AT Deficiency Examined for Association With Severe COVID-19 Outcomes

Risk Marker	Results
Smoking	Overall, limited evidence from one study is insufficient to determine an association between hospitalization in people with COVID-
(median	19 and underlying A1AT deficiency who smoked. Aggregation indices are not evaluated for outcomes reported by only one study.
pack/year)	
	• One cohort study ⁶ reported hospitalization in smokers with A1AT deficiency among people with COVID-19.
	 One cohort study⁶ (N = 9) of a national COVID-19 status database and pulmonology records reported [33.3% (3/9)]
	people with A1AT deficiency and COVID-19 were hospitalized. These people were more likely to be heavier
	smokers than those who were not hospitalized [66.6% (6/9)].
Baseline	Overall, limited evidence is inconclusive to determine an association between hospitalization and low baseline diffusion capacity
diffusion	[DLCO] in people with COVID-19 with underlying A1AT deficiency. Aggregation indices are not evaluated for outcomes reported by
capacity (mean	only one study.
%)	
	• One cohort study ⁶ reported hospitalization in people with a lower baseline DLCO with A1AT deficiency among people with
	COVID-19.
	 One cohort study⁶ (N = 9) of national COVID-19 status database and pulmonology records reported that with
	underlying A1AT deficiency and COVID-19 and acute respiratory failure (ARF) who were hospitalized [33.3% (3/9)]
	were more likely to have a lower baseline DLCO than those who were not hospitalized [66.6% (6/9)].

Table 5 Comorbid Conditions and Underlying A1AT Deficiency Examined for Association With Mortality Due to COVID-19

Comorbid	Results
condition	

Severe lung	Overall, limited evidence is inconclusive to determine an association between severe lung impairment and mortality in people
impairment	with COVID-19 and underlying A1AT deficiency. Aggregation indices are not evaluated for outcomes reported by only one study.
	 One cohort study¹ reported hospitalization in an individual with A1AT deficiency and severe lung impairment prior to COVID-19 illness. One cohort study¹ (N = 8) of patients with severe A1AT deficiency reported on eight patients with COVID-19 and the one patient that died was already hospitalized because of severe lung impairment prior to COVID-19 illness. This study had a small sample size with a low number of events, decreasing confidence in this finding.

B.3.b. Extracted Evidence

Study	Population and Setting	Exposure	Definitions	Results
Author: De Souza ⁵	Population: N = 1	Medical Condition, n/N	Medical Condition(s):	Severe COVID-19:
		(%):	A1AT deficiency: hereditary co-	Mortality: No
Year: 2020	Setting: Hospital	Alpha-1 antitrypsin (A1AT)	dominant disorder resulting from	ICU admission: Yes, MICU
		deficiency: 1/1 (100%)	the replacement of a single	Intubation (or Invasive Ventilation): Yes, Intubation
Data Extractor: AH	Location: PA, US		amino acid in the serine protease	Ventilation (mechanical, or non-invasive ventilation): Yes,
			inhibitor clade A-member-1	Mechanical
Reviewer: DOS	Study dates: April 4 –		(SERPINA1) gene on the long arm	Hospitalization: Yes
Study design: Case	May 11, 2020		of chromosome 14;	
report			characterized by either the	General Progression
	Inclusion criteria: NR		complete absence of the ATT-1	• Case 1: An immunosuppressed 67-year-old female with
Study Objective:			enzyme (homozygote genotype	homozygous Z-allele mutation A1AT deficiency admitted to
NR	Exclusion criteria: NR		PiZZ) or a misfolded A1AT	hospital due to dyspnea and cough and tested positive for
			enzyme (heterozygote genotype	COVID-19 via real-time RT-PCR. Home medications included
IVA Score: Not			PiMZ), which can promote early-	tacrolimus, mycophenolate, and prednisone. Patient was
completed			onset emphysema due to "loss of	upgraded to COVID-19 designated MICU, requiring sedation,
			function mutation" and/or the	intubation, and medical ventilation. Given a low dose of
			retainment of misfolded protein	norepinephrine by infusion and started treatment with
			within hepatocytes leading to	ceftriaxone, doxycycline, and hydroxychloroquine.
			cirrhosis	Hydroxychloroquine was stopped prematurely due to QTc
				prolongation. Patient's vasopressor requirement continued to
			Severity Measure(s):	increase, and she was placed on thiamine, ascorbic acid, and
			Homozygous Z-allele mutation:	stress dose steroids. The patient was placed on continuous
			ND	renal replacement therapy on hospital day 6. During the second
				week of hospitalization, patient's clinical status declined. Chest
			Clinical marker: NR	X-ray showed increased opacities in the right lung base and
				ultrasound showed a hyperdynamic ejection fraction. She

Table 6 Extracted Studies Reporting the Association Between A1AT Deficiency and Severe COVID-19 Outcomes

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			Outcome Definitions: Mortality: ND ICU admission: MICU Intubation: ND Ventilation: mechanical ventilation Hospitalization: ND Non-elective readmissions: NR Comments: None	 received empiric treatment for ventilation-acquired pneumonia and a 7-day course of piperacillin-tazobactam. On hospital day 9, she was weaned off norepinephrine, cisatracurium, and hydrocortisone. Patient remained on continuous veno-veno- hemodiafiltration for 10 days and was extubated and transitioned to intermittent hemodialysis on hospital day 16. The patient had a repeat positive COVID-19 test, but her mental status showed improvement. She was downgraded from the MICU to the intermediate medical care unit where she completed a 10-day course of oral vancomycin after testing positive for <i>Clostridium difficile</i>. She continued to improve and following 2 negative COVID-19 tests and the removal of her dialysis catheter, she was discharged to a local rehabilitation center and then discharged home. Severity of Condition: <i>Case 1:</i> homozygous Z-allele mutation A1AT deficiency Duration of Condition: NR <i>Case 1:</i> Liver transplant patient with chronic kidney disease stage IIIa, cirrhosis, COPD, hypertension, deep venous thrombosis, uncontrolled insulin-dependent type 2 diabetes mellitus Risk Markers: NR Long-term Sequelae: Non-elective readmissions: Not applicable for this study type
Author: Earia6	Population: N = 77	Modical Condition n/N	Modical Condition(s):	Sovere COVID 19:
AUGIOL Falla	COVID 10 + N = 0		A1AT deficiency: ND	Jevere Covid-13.
Voor: 2021	COMD-13+ IN = 3	(70):	ALAT DEJICIENCY: ND	
redr: 2021	Catting Tartian	(100%)	Soucrity Managera(a), ND	Hospitalization: 3/ / (4%)
Data Estra ata m	secting: rertiary	(100%)	Severity weasure(s): NK	 No hospitalization: 74/77 (96%)
	nospital			All patients hospitalized had acute respiratory failure.
IVIIVI	Data Causa di LO	Control/Comparison Grou	Clinical Marker: NK	
P · POC	Data Source: national C	p, n/N (%): NA		Ventilation, n/N (%):
Reviewer: DOS	UVID-19		Outcome Definitions:	 Ventilation: 0/77 (0%)
	status database and		Mortality: NR	• No ventilation: 77/77 (100%)
Study Design: Retr	pulmonology records		ICU admission: NR	
ospective cohort			Intubation: NR	

	Location: Portugal		Ventilation: Non-invasive	Severity of Condition: NR
Study Objective: ⊤			ventilation or high flow nasal	
o address the risk	Study Dates: NR - Janua		cannula	Duration of Condition: NR
of COVID-19	ry 2021		Hospitalization: hospitalized due	
infection in a			to acute respiratory failure	Comorbid Conditions: NR
cohort of Alpha-	Inclusion Criteria: All A1		Non-elective readmissions: NR	
1 Antitrypsin	AT deficient			Risk Markers:
(A1AT) Deficiency	patients followed at pul		Comments: None	Smoking, pack/year, Mean ± std dev:
patients through a	monology			 Hospitalized: 30 ±25.2 pack/year
comparison of	consultation at			 Not hospitalized: 5.2 ±2.9 pack/year
A1AT patients with	study hospital.			• p=0.09
and without				P
COVID-19.	Exclusion Criteria: NR			Baseline Diffusion Capacity, Mean % ± std dev:
11/A				 Hospitalized: 53.0% ±18.8%
Score: 16 (High)				 Not hospitalized: 87.8% ±4.6%
Score. 10 (mgn)				• p=0.042
				Long-term Sequelae:
				Non-elective readmissions: NR
Author: Ferrarotti ¹	Population: N = 209	Medical Condition, n/N	Medical Condition(s):	Severe COVID-19
			incultur contaition(s):	
	COVID-19+ N = 8	(%):	A1AT deficiency: the most	RR: Relative Risk
Year: 2021	COVID-19+ N = 8	(%): A1AT deficiency: 8/8	A1AT deficiency: the most abundant serum proteinase,	RR: Relative Risk
Year: 2021	COVID-19+ N = 8 Setting: Community	(%): A1AT deficiency: 8/8 (100%)	A1AT deficiency: the most abundant serum proteinase, whose deficiency caused by	RR: Relative Risk Mortality, n/N (%):
Year: 2021 Data Extractor:	COVID-19+ N = 8 Setting: Community	(%): A1AT deficiency: 8/8 (100%)	A1AT deficiency: the most abundant serum proteinase, whose deficiency caused by pathological mutation(s) in the	RR: Relative Risk Mortality, n/N (%): • RR: 0.9
Year: 2021 Data Extractor: MM	COVID-19+ N = 8 Setting: Community Data Source: Telephone	(%): A1AT deficiency: 8/8 (100%) Control/Comparison	A1AT deficiency: the most abundant serum proteinase, whose deficiency caused by pathological mutation(s) in the SERPINA1 gene results in an	RR: Relative Risk Mortality, n/N (%): • RR: 0.9 • A1AT deficiency: 1/8 (12.5%)
Year: 2021 Data Extractor: MM	COVID-19+ N = 8 Setting: Community Data Source: Telephone survey and Istituto	(%): A1AT deficiency: 8/8 (100%) Control/Comparison Group, n/N (%):	A1AT deficiency: the most abundant serum proteinase, whose deficiency caused by pathological mutation(s) in the SERPINA1 gene results in an imbalance in proteinase activity	RR: Relative Risk Mortality, n/N (%): • RR: 0.9 • A1AT deficiency: 1/8 (12.5%) • General Italian population: N = NR/N = NR (13.9%)
Year: 2021 Data Extractor: MM Reviewer: DOS	COVID-19+ N = 8 Setting: Community Data Source: Telephone survey and Istituto Superiore di Sanita	(%): A1AT deficiency: 8/8 (100%) Control/Comparison Group, n/N (%): General population in	A1AT deficiency: the most abundant serum proteinase, whose deficiency caused by pathological mutation(s) in the SERPINA1 gene results in an imbalance in proteinase activity which may lead to proteolytic	RR: Relative Risk Mortality, n/N (%): • RR: 0.9 • A1AT deficiency: 1/8 (12.5%) • General Italian population: N = NR/N = NR (13.9%)
Year: 2021 Data Extractor: MM Reviewer: DOS	COVID-19+ N = 8 Setting: Community Data Source: Telephone survey and Istituto Superiore di Sanita	(%): A1AT deficiency: 8/8 (100%) Control/Comparison Group, n/N (%): General population in Italy: NR	A1AT deficiency: the most abundant serum proteinase, whose deficiency caused by pathological mutation(s) in the SERPINA1 gene results in an imbalance in proteinase activity which may lead to proteolytic tissue damage and premature	RR: Relative Risk Mortality, n/N (%): • RR: 0.9 • A1AT deficiency: 1/8 (12.5%) • General Italian population: N = NR/N = NR (13.9%) ICU admission, n/N (%):
Year: 2021 Data Extractor: MM Reviewer: DOS Study Design:	COVID-19+ N = 8 Setting: Community Data Source: Telephone survey and Istituto Superiore di Sanita Location: Italy	(%): A1AT deficiency: 8/8 (100%) Control/Comparison Group, n/N (%): General population in Italy: NR	A1AT deficiency: the most abundant serum proteinase, whose deficiency caused by pathological mutation(s) in the SERPINA1 gene results in an imbalance in proteinase activity which may lead to proteolytic tissue damage and premature emphysema and COPD	RR: Relative Risk Mortality, n/N (%): • RR: 0.9 • A1AT deficiency: 1/8 (12.5%) • General Italian population: N = NR/N = NR (13.9%) ICU admission, n/N (%): • Admitted to ICU: 0/8 (0%)
Year: 2021 Data Extractor: MM Reviewer: DOS Study Design: Cohort	COVID-19+ N = 8 Setting: Community Data Source: Telephone survey and Istituto Superiore di Sanita Location: Italy	(%): A1AT deficiency: 8/8 (100%) Control/Comparison Group, n/N (%): General population in Italy: NR	A1AT deficiency: the most abundant serum proteinase, whose deficiency caused by pathological mutation(s) in the SERPINA1 gene results in an imbalance in proteinase activity which may lead to proteolytic tissue damage and premature emphysema and COPD	RR: Relative Risk Mortality, n/N (%): • RR: 0.9 • A1AT deficiency: 1/8 (12.5%) • General Italian population: N = NR/N = NR (13.9%) ICU admission, n/N (%): • Admitted to ICU: 0/8 (0%) • Not admitted to ICU: 8/8 (100%)
Year: 2021 Data Extractor: MM Reviewer: DOS Study Design: Cohort	COVID-19+ N = 8 Setting: Community Data Source: Telephone survey and Istituto Superiore di Sanita Location: Italy Study Dates: May 2020	(%): A1AT deficiency: 8/8 (100%) Control/Comparison Group, n/N (%): General population in Italy: NR	AIAT deficiency: the most abundant serum proteinase, whose deficiency caused by pathological mutation(s) in the SERPINA1 gene results in an imbalance in proteinase activity which may lead to proteolytic tissue damage and premature emphysema and COPD Severity Measure(s): NR	RR: Relative Risk Mortality, n/N (%): • RR: 0.9 • A1AT deficiency: 1/8 (12.5%) • General Italian population: N = NR/N = NR (13.9%) ICU admission, n/N (%): • Admitted to ICU: 0/8 (0%) • Not admitted to ICU: 8/8 (100%)
Year: 2021 Data Extractor: MM Reviewer: DOS Study Design: Cohort Study Objective: To investigate	COVID-19+ N = 8 Setting: Community Data Source: Telephone survey and Istituto Superiore di Sanita Location: Italy Study Dates: May 2020	(%): A1AT deficiency: 8/8 (100%) Control/Comparison Group, n/N (%): General population in Italy: NR	AIAT deficiency: the most abundant serum proteinase, whose deficiency caused by pathological mutation(s) in the SERPINA1 gene results in an imbalance in proteinase activity which may lead to proteolytic tissue damage and premature emphysema and COPD Severity Measure(s): NR	RR: Relative Risk Mortality, n/N (%): • RR: 0.9 • A1AT deficiency: 1/8 (12.5%) • General Italian population: N = NR/N = NR (13.9%) ICU admission, n/N (%): • Admitted to ICU: 0/8 (0%) • Not admitted to ICU: 8/8 (100%) Hospitalization, n/N (%):
Year: 2021 Data Extractor: MM Reviewer: DOS Study Design: Cohort Study Objective: To investigate whether people	COVID-19+ N = 8 Setting: Community Data Source: Telephone survey and Istituto Superiore di Sanita Location: Italy Study Dates: May 2020 Inclusion Criteria: Subjects aged over 18	(%): A1AT deficiency: 8/8 (100%) Control/Comparison Group, n/N (%): General population in Italy: NR	A1AT deficiency: the most abundant serum proteinase, whose deficiency caused by pathological mutation(s) in the SERPINA1 gene results in an imbalance in proteinase activity which may lead to proteolytic tissue damage and premature emphysema and COPD Severity Measure(s): NR Clinical Marker: NR	RR: Relative Risk Mortality, n/N (%): • RR: 0.9 • A1AT deficiency: 1/8 (12.5%) • General Italian population: N = NR/N = NR (13.9%) ICU admission, n/N (%): • Admitted to ICU: 0/8 (0%) • Not admitted to ICU: 8/8 (100%) Hospitalization, n/N (%): • Hospitalized 3/8 (37.5%)
Year: 2021 Data Extractor: MM Reviewer: DOS Study Design: Cohort Study Objective: To investigate whether people with severe AAT	COVID-19+ N = 8 Setting: Community Data Source: Telephone survey and Istituto Superiore di Sanita Location: Italy Study Dates: May 2020 Inclusion Criteria: Subjects aged over 18 years old with severely	(%): A1AT deficiency: 8/8 (100%) Control/Comparison Group, n/N (%): General population in Italy: NR	A1AT deficiency: the most abundant serum proteinase, whose deficiency caused by pathological mutation(s) in the SERPINA1 gene results in an imbalance in proteinase activity which may lead to proteolytic tissue damage and premature emphysema and COPD Severity Measure(s): NR Clinical Marker: NR	RR: Relative Risk Mortality, n/N (%): • RR: 0.9 • A1AT deficiency: 1/8 (12.5%) • General Italian population: N = NR/N = NR (13.9%) ICU admission, n/N (%): • Admitted to ICU: 0/8 (0%) • Not admitted to ICU: 8/8 (100%) Hospitalization, n/N (%): • Hospitalized 3/8 (37.5%) • Not Hospitalized 5/8 (62.5%)
Year: 2021 Data Extractor: MM Reviewer: DOS Study Design: Cohort Study Objective: To investigate whether people with severe AAT deficiency (A1AT	COVID-19+ N = 8 Setting: Community Data Source: Telephone survey and Istituto Superiore di Sanita Location: Italy Study Dates: May 2020 Inclusion Criteria: Subjects aged over 18 years old with severely reduced serum AAT	(%): A1AT deficiency: 8/8 (100%) Control/Comparison Group, n/N (%): General population in Italy: NR	A1AT deficiency: the most abundant serum proteinase, whose deficiency caused by pathological mutation(s) in the SERPINA1 gene results in an imbalance in proteinase activity which may lead to proteolytic tissue damage and premature emphysema and COPD Severity Measure(s): NR Clinical Marker: NR Outcome Definitions: Mortality: ND	RR: Relative Risk Mortality, n/N (%): • RR: 0.9 • A1AT deficiency: 1/8 (12.5%) • General Italian population: N = NR/N = NR (13.9%) ICU admission, n/N (%): • Admitted to ICU: 0/8 (0%) • Not admitted to ICU: 8/8 (100%) Hospitalization, n/N (%): • Hospitalized 3/8 (37.5%) • Not Hospitalized 5/8 (62.5%)
Year: 2021 Data Extractor: MM Reviewer: DOS Study Design: Cohort Study Objective: To investigate whether people with severe AAT deficiency (A1AT deficiency) have an	COVID-19+ N = 8 Setting: Community Data Source: Telephone survey and Istituto Superiore di Sanita Location: Italy Study Dates: May 2020 Inclusion Criteria: Subjects aged over 18 years old with severely reduced serum AAT levels due to two	(%): A1AT deficiency: 8/8 (100%) Control/Comparison Group, n/N (%): General population in Italy: NR	A1AT deficiency: the most abundant serum proteinase, whose deficiency caused by pathological mutation(s) in the SERPINA1 gene results in an imbalance in proteinase activity which may lead to proteolytic tissue damage and premature emphysema and COPD Severity Measure(s): NR Clinical Marker: NR Outcome Definitions: Mortality: ND ICU admission: intensive care	RR: Relative Risk Mortality, n/N (%): • RR: 0.9 • A1AT deficiency: 1/8 (12.5%) • General Italian population: N = NR/N = NR (13.9%) ICU admission, n/N (%): • Admitted to ICU: 0/8 (0%) • Not admitted to ICU: 8/8 (100%) Hospitalization, n/N (%): • Hospitalized 3/8 (37.5%) • Not Hospitalized 5/8 (62.5%) Severity of Condition: NR
Year: 2021 Data Extractor: MM Reviewer: DOS Study Design: Cohort Study Objective: To investigate whether people with severe AAT deficiency (A1AT deficiency) have an increased risk of	COVID-19+ N = 8 Setting: Community Data Source: Telephone survey and Istituto Superiore di Sanita Location: Italy Study Dates: May 2020 Inclusion Criteria: Subjects aged over 18 years old with severely reduced serum AAT levels due to two inherited nathological	(%): A1AT deficiency: 8/8 (100%) Control/Comparison Group, n/N (%): General population in Italy: NR	A1AT deficiency: the most abundant serum proteinase, whose deficiency caused by pathological mutation(s) in the SERPINA1 gene results in an imbalance in proteinase activity which may lead to proteolytic tissue damage and premature emphysema and COPD Severity Measure(s): NR Clinical Marker: NR Outcome Definitions: Mortality: ND ICU admission: intensive care Intubation: NR	RR: Relative Risk Mortality, n/N (%): • RR: 0.9 • A1AT deficiency: 1/8 (12.5%) • General Italian population: N = NR/N = NR (13.9%) ICU admission, n/N (%): • Admitted to ICU: 0/8 (0%) • Not admitted to ICU: 8/8 (100%) Hospitalization, n/N (%): • Hospitalized 3/8 (37.5%) • Not Hospitalized 5/8 (62.5%) Severity of Condition: NR
Year: 2021 Data Extractor: MM Reviewer: DOS Study Design: Cohort Study Objective: To investigate whether people with severe AAT deficiency (A1AT deficiency) have an increased risk of severe COVID-19	COVID-19+ N = 8 Setting: Community Data Source: Telephone survey and Istituto Superiore di Sanita Location: Italy Study Dates: May 2020 Inclusion Criteria: Subjects aged over 18 years old with severely reduced serum AAT levels due to two inherited pathological alleles in the SERPINA1	(%): A1AT deficiency: 8/8 (100%) Control/Comparison Group, n/N (%): General population in Italy: NR	A1AT deficiency: the most abundant serum proteinase, whose deficiency caused by pathological mutation(s) in the SERPINA1 gene results in an imbalance in proteinase activity which may lead to proteolytic tissue damage and premature emphysema and COPD Severity Measure(s): NR Clinical Marker: NR Outcome Definitions: Mortality: ND ICU admission: intensive care Intubation: NR Ventilation: NR	RR: Relative Risk Mortality, n/N (%): • RR: 0.9 • A1AT deficiency: 1/8 (12.5%) • General Italian population: N = NR/N = NR (13.9%) ICU admission, n/N (%): • Admitted to ICU: 0/8 (0%) • Not admitted to ICU: 8/8 (100%) Hospitalization, n/N (%): • Hospitalized 3/8 (37.5%) • Not Hospitalized 5/8 (62.5%) Severity of Condition: NR Duration of Condition: NR
Year: 2021 Data Extractor: MM Reviewer: DOS Study Design: Cohort Study Objective: To investigate whether people with severe AAT deficiency (A1AT deficiency) have an increased risk of severe COVID-19.	COVID-19+ N = 8 Setting: Community Data Source: Telephone survey and Istituto Superiore di Sanita Location: Italy Study Dates: May 2020 Inclusion Criteria: Subjects aged over 18 years old with severely reduced serum AAT levels due to two inherited pathological alleles in the SERPINA1 gene were surveved via	(%): A1AT deficiency: 8/8 (100%) Control/Comparison Group, n/N (%): General population in Italy: NR	A1AT deficiency: the most abundant serum proteinase, whose deficiency caused by pathological mutation(s) in the SERPINA1 gene results in an imbalance in proteinase activity which may lead to proteolytic tissue damage and premature emphysema and COPD Severity Measure(s): NR Clinical Marker: NR Outcome Definitions: Mortality: ND ICU admission: intensive care Intubation: NR Ventilation: NR Hospitalization: ND	RR: Relative Risk Mortality, n/N (%): • RR: 0.9 • A1AT deficiency: 1/8 (12.5%) • General Italian population: N = NR/N = NR (13.9%) ICU admission, n/N (%): • Admitted to ICU: 0/8 (0%) • Not admitted to ICU: 8/8 (100%) Hospitalization, n/N (%): • Hospitalized 3/8 (37.5%) • Not Hospitalized 5/8 (62.5%) Severity of Condition: NR Duration of Condition: NR

IVA Score: 17 (High)	telephone. Subjects were derived from Italian Registry of Severe A1AT deficiency. COVID-19 was diagnosed via laboratory tests for SARS-CoV-2 on nasal swabs or blood samples. Exclusion Criteria: NR		Non-elective readmissions: NR Comments: Author's note: The survey only reached 35% of the Italian cohort with severe A1AT deficiency.	The one patient who died was already long-term hospitalized because of severe lung impairment before COVID-19 infection. Risk Markers: NR Long-term Sequelae: Non-elective readmissions: NR
Author: Schneider ²	Population:	Medical Condition, n/N	Medical Condition(s):	Severe COVID-19:
Year: 2021 Data Extractor: AH	N = 14,393 Setting: Community	(%): A1AT deficiency: 1,1670/14,393 (11.60%)	A1AT deficiency: included the most clinically relevant AAT variants Pi*Z (rs28929474) and Pi*S (rs17580)	Mortality, n/N (%): • A1AT deficiency: 53/1,670 (3.20%) • No A1AT deficiency: 353/12,723 (3.00%)
Reviewer: DOS	Biobank and national	Group, n/N (%):	Severity Measure(s):	Severity of Condition:
Study Design:	death registries	No A1AT deficiency:	Pi*MS: ND Pi*SS: ND	 Pi*MS: 37/1,156 (3.20%) Pi*SS: 1/28 (4.00%)
Cohort Study Objective: To study the association between alph-1- antitrypsin deficiency (A1AT deficiency) and SARS-CoV-2 infection. IVA Score: 15 (High)	Location: United Kingdom Study Dates: NR – February 18, 2021 Inclusion Criteria: Patients recruited to a community-based cohort from 22 centers who had genotyping available on the most clinically relevant AAT variants Pi*Z (rs28929474) and Pi*S (rs17580). Exclusion Criteria: NR		 Pi*MZ: heterozygous Pi*Z genotype results in mild A1AT deficiency Pi*SZ: combined presence of Pi*S and Pi*Z variants Pi*ZZ: homozygous Pi*Z genotype, the predominant cause of severe A1AT deficiency Clinical Marker: NR Outcome Definitions: Mortality: Death by COVID-19 ICU admission: NR Intubation: NR Ventilation: NR Hospitalization: NR Non-elective readmissions: NR 	 P1'SS: 1/28 (4.00%) Pi*MZ: 14/460 (3.04%) Pi*SZ: 1/23 (4.35%) Pi*ZZ: 0/3 (0.00%) Duration of Condition: NR Comorbid Conditions: NR Risk Markers: NR Long-term Sequelae: NR
			Comments: None	

countries(%):Alpha 1 antitrypsin (A1AT)R: Pearson Correlation (r); ANCOVA (Analysis	of Covariance)
Year: 2020 A1AT deficiency: NR deficiency: presence of any of	
Setting: Population- the following alleles: - PiS Mortality, n/N (%):	
Data Extractor: AH level Control/Comparison (SERPINA1 rs17580) and PiZ Correlation of national rates of population ad	ljusted COVID-19
Group, n/N (%): (SERPINA1 rs28929474,) mortality with A1AT deficiency allele frequence	cies. Adjusted for
Reviewer: ES Data Source: NA urbanization, age distribution, etc.	
Relevant literature: Severity Measure(s): ND Pearson R=0.56, p = 0.00000087	
Study Design: 2017 National PiS &	
Cross-sectional PiZ allele frequencies Clinical Marker: ND Correlation of national rates of COVID-19 mol	rtality with A1AT
ecological • United Nations deficiency allele frequencies	
database: 2018 date Outcome Definitions: • Pearson R=0.54, p = 0.00000198	
Study Objective: for all indicators Mortality: ND	
To examine a except for 2017 ICU admission: NR Severity of Condition: NR	
possible inbound tourism Intubation: NR	
association • The World Bank Open Ventilation: NR Duration of Condition: NR	
between the Data: 2018 Population Hospitalization: NR	
distributions of size, density, male Non-elective readmissions: NR Comorbid Conditions: NR	
common SERPINA1 percentage, urban	
single nucleotide percentage, and age Comments: None Risk Markers: NR	
polymorphisms composition	
(SNPs) underlying • Johns Hopkins	
A1AT deficiency University: 2020 Non-elective readmissions: NR	
and between National COVID-19	
COVID-19 infection rates	
epidemiology on a	
global scale.	
IVA Score: 17 Study Dates:	
(High) Up to September 7.	
2020	
Inclusion Criteria:	
Countries with a	
population >1million	
Exclusion Criteria: NR	
Author: Yoshikura ⁴ Population: N = 68 Medical Condition, n/N Medical Condition(s): Severe COVID-19: NR	
countries (%): A1AT deficiency: PI*S, PI*Z, and	
Year: 2021 A1AT deficiency: NR PI*SZ Severity of Condition:	
Setting: NR Mortality correlation coefficient:	

Data Extractor:		Control/Comparison	Severity Measure(s):	PI*S
MM	Data Source: WHO	Group, n/N (%):	A1AT deficiency PI*S: frequent	• All: 0.6326
	COVID-19 situation	No A1AT deficiency: NR	mutant allele and milder variant	Europe & America: 0.8244
Reviewer: DOS	reports, A1AT deficiency		A1AT deficiency PI*SZ: serum	Other regions: 0.4360
	prevalence data tables		level 75-150 mg/dL	PI*SZ
Study Design:	published in relevant		A1AT deficiency PI*Z: frequent	• All: 0.8585
Ecological	literature,		mutant allele and most severe	• Europe & America: 0.8864
	Worldometer, and		variant	Other regions: 0.0253
Study Objective:	World Bank data			PI*Z:
To examine the			Clinical Marker: NR	• All: 0.8713
epidemiological	Location: International			• Europe & America: 0.9503
correlation			Outcome Definitions:	Other regions: 0.4360
between the	Study Dates: January 21		Mortality: ND	
COVID-19	– June 18, 2020		ICU admission: NR	Duration of Condition: NR
epidemic and A1AT			Intubation: NR	
deficiency.	Inclusion Criteria:		Ventilation: NR	Comorbid Conditions: NR
	Countries for which		Hospitalization: NR	
IVA Score: 18	mortality data from		Non-elective readmissions: NR	Risk Markers: NR
(Moderate)	WHO COVID-19			
	situation reports and		comments: None	Long-term Sequelae:
	ATAT deficiency			Non-elective readmissions: NR
	Plance et al. (2017) is			
	Exclusion Criteria: NR			

B.3.c. Internal Validity Assessments of Extracted Studies

Table 7 Internal Validity Assessments of Extracted Studies Reporting the Association Between A1AT Deficiency and Severe COVID-19 Outcomes

		Schneider	Shapira	Yoshikura
	Author Year	2021 ²	2020 ³	20214
	Outcome	Mortality	Mortality	Mortality
Domain	Signaling question			
Study Elements	Design appropriate to research question	1	0	0
	Well described population	1	0	1
	Well described setting	1	0	0
	Well described intervention/ exposure	1	1	0
	Well described control/ comparator	1	0	0
	Well described outcome	1	1	1
	Clear timeline of exposures/ interventions and outcomes	0	1	1
Selection Bias: Sampling	Randomization appropriately performed	0	0	0
	Allocation adequately concealed	0	0	0
	Population sampling appropriate to study design	1	1	0
Selection Bias: Attrition	Attrition not significantly different between groups	0	0	1
	Attrition <10-15% of population	0	0	1
	Attrition appropriately analyzed	0	0	1
Information Bias:	Measure of intervention/ exposure is valid	1	1	0
Measurement	Measure of outcome is valid	1	1	1
and	Fidelity to intervention is measured	0	0	0
Wisclassification	Fidelity to intervention is valid	0	0	0
	Prospective study	1	1	1

	Adequately powered to detect result	0	1	0
Information	Outcome assessor blinded	0	0	0
Bias:	Study participant blinded	0	0	0
Performance &	Investigator/ data analyst blinded	0	0	0
Detection	Data collection methods described in sufficient detail	1	0	1
	Data collection methods appropriate	1	1	1
	Sufficient follow up to detect outcome	0	1	1
Information Bias: Analytic	Appropriate statistical analyses for collected data	0	1	1
	Appropriate statistical analyses are conducted correctly	0	1	1
	Confidence interval is narrow	0	0	0
Confounding	Potential confounders identified	0	1	1
	Adjustment for confounders in study design phase	0	0	0
	Adjustment for confounders in data analysis phase	0	1	1
Reporting Bias	All pre-specified outcomes are adequately reported	1	1	1
Other Bias	No other sources of bias	1	1	1
COI	Funding sources disclosed and no obvious conflict of interest	1	1	1
SCORE	Threat to internal validity	15	17	18
	Low, Moderate, High	High	High	Moderate

C. References

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D. Abbreviations

Acronym	Full
95% CI	95% confidence interval
A1AT	Alpha-1 Antitrypsin deficiency
aHR	adjusted hazard ratio
aOR	adjusted odds ratio
ARF	acute respiratory failure
BMI	body mass index
BPD	bronchopulmonary dysplasia
CF	cystic fibrosis
CFR	case fatality ratio
COI	conflict of interest
COPD	chronic obstructive pulmonary disease
CRD	chronic respiratory disease
ECMO	extracorporeal membrane oxygenation
EHR	electronic health record

EMR	electronic medical record
ERT	evidence review team
IQR	interquartile range
GLM	generalized linear model
HR	hazard ratio
ICD10	International Classification of Diseases 10
ICS	inhaled corticosteroids
ICU	intensive care unit
ILD	interstitial lung disease
IPF	idiopathic pulmonary fibrosis
IVA	Internal validity assessments
MICU	medical intensive care unit
MR	mortality rate
ND	not defined
NR	not reviewed
OR	odds ratio
PCR	polymerase chain reaction
PECO	population, exposure, comparator, and outcomes
RR	rate ratio
RT-PCR	real time polymerase chain reaction
SNP	single nucleotide polymorphisms