

## Common cardiovascular risk factors and in-hospital mortality in 3,894 patients with COVID-19: survival analysis and machine learning-based findings from the multicentre Italian CORIST Study

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## KEYWORDS

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**Abstract** *Background and aims:* There is poor knowledge on characteristics, comorbidities and laboratory measures associated with risk for adverse outcomes and in-hospital mortality in European Countries. We aimed at identifying baseline characteristics predisposing COVID-19 patients to in-hospital death.

*Methods and results:* Retrospective observational study on 3894 patients with SARS-CoV-2 infection hospitalized from February 19th to May 23rd, 2020 and recruited in 30 clinical centres distributed throughout Italy. Machine learning (random forest)-based and Cox survival analysis. 61.7% of participants were men (median age 67 years), followed up for a median of 13 days. In-hospital mortality exhibited a geographical gradient, Northern Italian regions featuring more than twofold higher death rates as compared to Central/Southern areas (15.6% vs 6.4%, respectively). Machine learning analysis revealed that the most important features in death classification were impaired renal function, elevated C reactive protein and advanced age. These findings were confirmed by multivariable Cox survival analysis (hazard ratio (HR): 8.2; 95% confidence interval (CI) 4.6–14.7 for age  $\geq 85$  vs 18–44 y); HR = 4.7; 2.9–7.7 for estimated glomerular filtration rate levels  $<15$  vs  $\geq 90$  mL/min/1.73 m<sup>2</sup>; HR = 2.3; 1.5–3.6 for C-reactive protein levels  $\geq 10$  vs  $\leq 3$  mg/L). No relation was found with obesity, tobacco use, cardiovascular disease and related-comorbidities. The associations between these variables and mortality were substantially homogenous across all sub-groups analyses.

*Conclusions:* Impaired renal function, elevated C-reactive protein and advanced age were major predictors of in-hospital death in a large cohort of unselected patients with COVID-19, admitted to 30 different clinical centres all over Italy.

## Introduction

As of July 10, 2020, there have been over 12 million of confirmed cases of COVID-19, with 549,247 deaths worldwide [1]. Robust knowledge of correlates and possible predictors of death among patients diagnosed with COVID-19 is crucial to target patients at highest risk through early and more intensive interventions. Since the COVID-19 pandemic outbreak, several studies have evaluated different factors that may predispose individuals to a higher risk of death from COVID-19, but evidence on this is still limited, especially from European countries. Recent meta-analyses [2,3] indicated comorbidities (hypertension, coronary heart disease and diabetes) as major predictors of higher mortality. Moreover, some laboratory parameters such as elevated levels of C-reactive protein (CRP), cardiac troponins and interleukin-6 resulted associated with a higher risk of death. However, those meta-analyses mainly relied on data from China and the United States. Subsequent studies have confirmed a role for age, male gender and the presence of comorbidities as major risk factors for mortality [4,5]; other investigators have also found higher death risk associated with smoking [6] and obesity [7–9].

Since the beginning of the pandemic, Italy was one of the most heavily hit countries, with 34,026 deaths recorded as of July 10, 2020, the largest part occurring in Northern regions [10]. Data from small studies conducted in Lombardy during the first wave (February–March 2020) of the Italian pandemic, showed high in-hospital mortality and higher rate of complications in cardiac patients [11], while older age, obesity and more advanced illness were major risk factors for 30-day mortality [9]. Data from 500 patients hospitalized in Milan, confirmed cardiovascular comorbidities as major predictors of death along with the chest X-ray quantitative radiographic assessment of lung oedema (RALE) score at admission [12]. No evidence to date is available from the Central/Southern regions of Italy, which were partially spared by the COVID-19-pandemic, with less than 20 percent of the total cases recorded nationwide [13]. Only recently, a larger study including also COVID-19 patients from Central-Southern regions of Italy provided insights into factors predisposing to in hospital death, however without a separate North-to-South analysis [14].

Low total number of cases was also documented in other countries bordering the Mediterranean Sea, with about 3000 confirmed cases and less than 200 deaths [15]. Given such broad geographic variability in the pandemic outbreak, risk factors for mortality might vary, in type and impact, across different geographical areas. Finally, most studies have mainly analysed data on patients hospitalized

during the first wave of the pandemic, at the earliest stage of the outbreak, February–March 2020 [6,9,11,12,16,17].

To take into account potential non-linear relations of and interaction effects among clinical and sociodemographic risk factors for severity of COVID-19, other studies attempted to estimate mortality risk in COVID-19 patients using supervised machine learning algorithms, as recently reviewed [18]. Although these models were not always precisely described and thoroughly investigated and often only exploited limited sample sizes ( $N < 1000$ ), these algorithms suggested a prominent effect of age, sex, comorbidities, like hypertension, cardiovascular and chronic respiratory disease, cancer and circulating biomarkers (CRP, lymphocyte count and lactate dehydrogenase) [19–25]. Moreover, most of these studies were based on Chinese patients and none of them included an Italian population of COVID-19 patients, which may be important to investigate whether there are differences in the prognostic values of the variables identified so far, based on the ethnicity of cases analysed.

Our report, therefore, aims at extending knowledge on factors predisposing to higher in-hospital death risk, based on data from the CORIST Collaboration [26], an observational multicentre study including 3894 patients with laboratory-confirmed SARS-CoV-2 infection, hospitalized in 30 Italian clinical centres, 41.5% of whom resident in Central/Southern Italian regions. This was accomplished through a composite approach involving the development of a machine learning algorithm to determine the predictive power of the features available and their relative importance in influencing mortality risk, followed by a classical statistical approach to analyse the directions of effects and the relation with time-to-events in a longitudinal setting. The CORIST cohort allowed to identify baseline factors associated with a higher risk of in-hospital death in COVID-19 patients and their efficacy as prognostic factors in a single machine learning algorithm. Moreover, the data analyzed here allowed to assess whether there was a geographic gradient in Italy in the association between predictors and in-hospital death risk and to evaluate whether there was a time-dependent pandemic wave-related difference in the type and/or strength of the associations of risk factors with in-hospital death.

## Methods

### Setting

This national retrospective observational study was conceived, coordinated and analysed within the CORIST Collaboration Project (ClinicalTrials.gov ID: NCT04318418). Initially focussed on the inhibitors of renin-angiotensin

system, the CORIST Collaboration is a set of multicentre observational investigations launched in March 2020 and aimed at testing the association of risk factors and/or therapies with disease severity and mortality in COVID-19 hospitalised patients [26]. The study was approved by the institutional Ethics Board of the Istituto di Ricovero e Cura a Carattere Scientifico (IRCCS) Neuromed, Pozzilli, and of all recruiting centres, including the IRCCS “Lazzaro Spallanzani”, Rome, which was the Coordinating Centre for clinical studies on COVID-19 in Italy. Data for the present analyses were provided by 30 hospitals distributed throughout Italy. Each hospital provided data from hospitalised adult ( $\geq 18$  years of age) patients who all had a positive test result for the SARS-CoV-2 virus at any time during their hospitalisation from February 19th to May 23rd, 2020. The follow-up continued through May 29th, 2020.

### Data sources

We developed a cohort comprising 3971 patients with laboratory-confirmed SARS-CoV-2 infection in an in-patient setting. The SARS-CoV-2 status was declared on the basis of laboratory results (polymerase chain reaction on a nasopharyngeal swab) from each participating hospital. Clinical data were abstracted at one-time point from electronic medical records or charts, and collected using either a centrally-designed electronic worksheet or a centralized web-based database. Collected data included patients' demographics, laboratory test results, medication administration, historical and current medication lists, historical and current diagnoses, and clinical notes. In addition, specific information on the most severe manifestations of COVID-19 that occurred during hospitalisation was retrospectively captured. Maximum clinical severity observed was classified as: either light-mild pneumonia; or severe pneumonia; or acute respiratory distress syndrome (ARDS) [27]. Specifically, we obtained the following information for each patient: hospital; date of admission and date of discharge or death; age; gender; the first recorded inpatient laboratory tests at entry (creatinine, CRP); past and current diagnoses of chronic degenerative disease or risk factors (myocardial infarction, heart failure, diabetes, hypertension, chronic pulmonary disease and cancer), and in-hospital drug therapies for COVID-19. Chronic kidney disease was classified as: stage 1: normal or increased glomerular filtration rate (eGFR) ( $\geq 90$  mL/min/1.73 m<sup>2</sup>); stage 2: kidney damage with mild reduction in eGFR (60–89 mL/min/1.73 m<sup>2</sup>); stage 3a: moderate reduction in eGFR (45–59 mL/min/1.73 m<sup>2</sup>); stage 3 b: moderate reduction in eGFR (30–44 mL/min/1.73 m<sup>2</sup>); stage 4: severe reduction in eGFR (15–29 mL/min/1.73 m<sup>2</sup>); stage 5: kidney failure (eGFR  $< 15$  mL/min/1.73 m<sup>2</sup> or dialysis). eGFR was calculated by the Chronic Kidney Disease Epidemiology Collaboration (CKD-EPI) equation. CRP was classified as  $\leq 3$ , 3–10 and  $\geq 10$  mg/L.

The study index date was defined as the date of hospital admission. Index dates ranged from February 19th, 2020 to May 23rd, 2020. The study end point was the time from

study index to death. The number of patients who either died or had been discharged alive, or were still admitted to hospital as of May 29th, 2020, were recorded, and hospital length of stay was determined. Patients alive had their data censored on the date of discharge or as the date of the respective clinical data collection.

Of the initial cohort of 3971 patients, 77 patients were excluded from the present analysis because of one or more missing data at baseline or during follow-up, including time to event ( $n = 59$ ), outcome (death/alive,  $n = 8$ ), COVID-19 severity ( $n = 4$ ), age ( $n = 4$  with missing data and  $n = 2$  with age  $< 18$  years) or gender ( $n = 2$ ).

At the end, the analysed cohort consisted of  $N = 3894$  patients. Distribution of missing values for covariates is shown in Table 1.

### Machine learning analysis

To take into account potential non-linear relations of and interaction effects among the investigated variables with the risk of death, we performed an exploratory machine learning analysis to compute the predictive power of a potential classification algorithm for the risk of death and to establish variable importance in more complex mortality prediction models. To this purpose, we trained a Random Forest (RF) algorithm in R [28], using age, gender, obesity and smoking status, chronic comorbidities (history of myocardial infarction, hypertension, diabetes, lung disease, heart failure and cancer), CRP and eGFR as input features. Random forest is an ensemble of decision trees which is often used in classification tasks, and represents one of the most used machine learning algorithms applied to risk prediction in COVID-19 patients [18]. Missing data were imputed through a k-nearest neighbour algorithm (*kNN()* function) of the *VIM* package [29], while continuous features (CRP, eGFR and age) underwent min–max normalization before analysis, with CRP transformed on the natural logarithm scale to attain normality. The resulting dataset ( $N = 3894$ ) was divided in a random training and a test set (2725 and 1169 patients, respectively, 70:30 ratio). We then performed hyperparameter tuning through the *train()* function of the *caret* package [30], in a 10-fold cross validation setting, to optimize the algorithm over two varying parameters: the number of variables randomly sampled as candidate predictors at each node split in the decision tree (*mtry*, varying between 1 and 10), and the number of trees to grow in the random forest (*ntree* alternative values: 100, 500, 1000). Finally, we trained the optimized model (*mtry* = 3, *ntree* = 1000) within the training set, predicted the label (death/no death) in the independent test set, and performed a permutation feature importance (PFI) analysis to identify those variables showing the largest influence on the prediction of death. This implies shuffling measures of one marker at a time and then comparing the loss function (cross-entropy of the classification) of the perturbed RF model with that of



**Table 1** Incidence rates and univariable hazard ratios for death in COVID-19 patients.

	Patient at risk N = 3894 (%) <sup>a</sup>	Death N = 712 (%) <sup>b</sup>	Person-days	Death Rate ( $\times 1,000$ person-days)	Univariable HR (95% CI)
<b>Gender</b>					
Female	1491 (38.3)	243 (16.3)	24,327	10.0	-1-
Male	2403 (61.7)	469 (19.5)	39,559	11.9	1.18 (1.01–1.38)
<b>Age, years</b>					
18–44	348 (8.9)	6 (1.7)	4030	1.5	-1-
45–64	1413 (36.3)	75 (5.3)	22,387	3.4	2.40 (1.04–5.51)
65–74	808 (20.8)	145 (18.0)	15,437	9.4	7.12 (3.14–16.14)
75–84	849 (21.8)	266 (31.3)	14,462	18.4	13.56 (6.03–30.52)
$\geq 85$	476 (12.2)	220 (30.9)	7570	29.1	21.65 (9.60–48.82)
<b>Hypertension</b>					
No	1899 (48.8)	219 (11.5)	29,184	7.5	-1-
Yes	1943 (49.9)	461 (23.7)	33,994	13.6	1.85 (1.58–2.18)
Missing data	52 (1.3)	32 (61.5)			
<b>Diabetes</b>					
No	3103 (79.7)	481 (15.5)	49,303	9.8	-1-
Yes	739 (19.0)	203 (27.5)	13,855	14.7	1.56 (1.32–1.84)
Missing data	52 (1.3)	28 (53.9)			
<b>Myocardial infarction</b>					
No	3421 (87.9)	527 (15.4)	55,963	9.4	-1-
Yes	388 (10.0)	140 (36.1)	6641	21.1	2.31 (1.92–2.79)
Missing data	85 (2.2)	45 (52.9)			
<b>Heart Failure</b>					
No	3375 (86.7)	500 (14.8)	54,728	9.1	-1-
Yes	431 (11.1)	165 (38.3)	7710	21.4	2.44 (2.04–2.91)
Missing data	88 (2.3)	47 (53.4)			
<b>Cancer</b>					
No	3437 (88.3)	552 (16.1)	55,960	9.9	-1-
Yes	399 (10.3)	128 (32.1)	7074	18.1	1.88 (1.55–2.28)
Missing data	58 (1.5)	32 (55.2)			
<b>Lung disease</b>					
No	3269 (84.0)	513 (14.6)	53,097	9.7	-1-
Yes	557 (14.3)	167 (29.6)	9681	17.3	1.83 (1.53–2.17)
Missing data	68 (1.8)	32 (47.1)			
<b>Obesity, BMI <math>\geq 30</math> kg/m<sup>2</sup></b>					
No	2173 (55.8)	410 (18.9)	36,159	11.3	-1-
Yes	376 (9.7)	69 (18.4)	6683	10.3	0.92 (0.71–1.19)
Missing data	1345 (34.5)	233 (17.3)			
<b>Smoking</b>					
Non-smoker	2140 (55.0)	420 (19.6)	33,670	12.5	-1-
Current Smoker	319 (8.2)	59 (18.5)	5995	9.8	0.83 (0.63–1.09)
Missing data	1435 (36.9)	233 (16.2)			
<b>CRP, mg/L</b>					
$\leq 3$	884 (22.7)	50 (5.7)	12,906	3.9	-1-
3–10	835 (21.4)	165 (19.8)	12,384	13.3	3.39 (2.47–4.65)
$\geq 10$	1962 (50.4)	457 (23.3)	34,644	13.2	3.51 (2.62–4.71)
Missing data	213 (5.5)	40 (18.8)			
<b>eGFR, CKD stage, mL/min/1.73 m<sup>2</sup></b>					
$\geq 90$	1368 (35.1)	67 (4.9)	21,427	3.2	-1-
60–89	1409 (36.2)	205 (14.6)	23,950	8.6	2.75 (2.09–3.62)
45–59	433 (11.1)	131 (30.3)	7806	16.8	5.57 (4.15–7.48)
30–44	310 (8.0)	131 (42.3)	5317	24.6	8.17 (6.08–10.96)
15–29	197 (5.1)	110 (55.8)	2999	36.7	12.00 (8.85–16.26)
$< 15$	87 (2.2)	47 (54.0)	1336	35.2	11.71 (8.06–17.00)
Missing data	90 (2.3)	21 (23.3)			

Abbreviations: BMI: body mass index; CKD: chronic kidney disease; CRP: C-reactive protein; eGFR: glomerular filtration rate.

<sup>a</sup> Column %.<sup>b</sup> Row %: Death/Patient at risk.

the full model (i.e. with no permuted feature). This analysis was carried out through the *explain()* and the *variable\_importance()* functions of the *DALEX* package [31].

### Other statistical analyses

“Classical” (non-machine learning) statistical analyses were performed in SAS software for Windows, version 9.4.

**Table 2** Subgroup analyses showing hazard ratio for mortality, according to gender. Multiple imputation analysis, N = 3894 patients and N = 712 deaths.

	Gender	
	Women	Men
<b>Patient at risk</b>	1491	2403
<b>Death</b>	243	469
<b>Person-days</b>	24,327	39,559
<b>Death Rate, x1,000 PD</b>	10.0	11.9
	HR (95% CI)	HR (95% CI)
<b>Age, years</b>		
18–44	-1-	-1-
45–64	1.77 (0.33–9.54)	1.77 (0.76–4.12)
65–74	3.37 (0.87–13.06)	4.06 (1.65–9.95)
75–84	6.24 (1.52–25.59)	6.18 (2.70–14.17)
≥85	8.82 (2.27–34.23)	8.06 (3.35–19.44)
<b>Hypertension</b>		
No	-1-	-1-
Yes	0.65 (0.51–0.82)	0.98 (0.79–1.22)
<b>Diabetes</b>		
No	-1-	-1-
Yes	1.05 (0.77–1.43)	0.99 (0.76–1.29)
<b>Myocardial infarction</b>		
No	-1-	-1-
Yes	1.27 (0.82–1.98)	1.16 (0.86–1.56)
<b>Heart Failure</b>		
No	-1-	-1-
Yes	1.02 (0.69–1.50)	1.09 (0.78–1.53)
<b>Cancer</b>		
No	-1-	-1-
Yes	1.33 (0.96–1.84)	1.41 (1.11–1.78)
<b>Lung disease</b>		
No	-1-	-1-
Yes	0.98 (0.64–1.52)	1.31 (1.01–1.69)
<b>Obesity, BMI ≥30 kg/m<sup>2</sup></b>		
No	-1-	-1-
Yes	1.39 (0.80–2.41)	1.15 (0.87–1.52)
<b>Smoking</b>		
Non-smoker	-1-	-1-
Current Smoker	0.77 (0.45–1.32)	0.98 (0.68–1.41)
<b>CRP, mg/L</b>		
≤3	-1-	-1-
3–10	2.53 (1.57–4.10)	2.77 (2.07–3.72)
≥10	2.27 (1.42–3.63)	2.31 (1.41–3.77)
<b>eGFR, CKD stage, mL/min/1.73 m<sup>2</sup></b>		
≥90	-1-	-1-
60–89	1.14 (0.60–2.20)	1.72 (1.20–2.49)
45–59	2.01 (1.04–3.88)	2.51 (1.64–3.85)
30–44	3.05 (1.57–5.96)	3.06 (1.82–5.16)
15–29	4.81 (2.69–8.61)	3.44 (2.37–4.99)
<15	7.93 (3.86–16.27)	3.47 (1.83–6.56)

Controlling for age, diabetes, hypertension, history of myocardial infarction, heart failure, chronic pulmonary disease, chronic kidney disease, CRP, obesity, smoking habit and anti-COVID19 drugs during hospitalization as fixed effects and repeated measures within hospital.

Abbreviations: BMI: body mass index; CKD: chronic kidney disease; CRP: C-reactive protein; eGFR: glomerular filtration rate; PD: person-days.

For the purpose of the survival analyses, we used a multiple imputation technique (SAS PROC MI, followed by PROC MIANALYZE) to maximize data availability for all variables and get robust results over different simulations (n = 10 imputed datasets).

Cox proportional-hazards regression models were used to estimate the association between characteristics of patients at hospital admission and in-hospital mortality. Since the multiple imputation technique was applied, the final standard error was obtained using the Rubin's rule based on the robust variance estimator in Cox regression [32].

Multiple imputed (for all the variables) multivariable analysis controlling for gender, age, diabetes, hypertension, history of myocardial infarction, heart failure, chronic pulmonary disease, chronic kidney disease, CRP, obesity (body mass index (BMI) ≥30 kg/m<sup>2</sup>), smoking habit and in-hospital anti-COVID-19 therapy as fixed effects; accounting for clustering within hospital by using the robust sandwich estimator was considered as the main analysis. As secondary analyses, we considered a) multiple imputation for all the variables with the exception of smoking and obesity (which were the variables with the largest number of missing data); b) case complete analysis for all the variables except smoking and obesity (N = 3454 patients); c) case complete analysis for all variables (N = 1632 patients) (see [Supplementary Table 1](#)); d) accounting for clustering within hospital by inclusion of the hospital index as random effect or strata variable ([Supplementary Table 2](#)); e) censoring of follow up at 35 days ([Supplementary Table 3](#)). Data from 4 hospitals that provided less than 25 patients each were combined in a unique group for clustering. Sensitivity analyses were conducted according to temporal pandemic waves, geographical location of the hospitals, age or gender of patients or number of risk factors at baseline ([Tables 2–5 and Supplementary Fig. 1](#)).

For the latter we included the following: previous myocardial infarction, heart failure, hypertension, diabetes, cancer, chronic pulmonary disease, obesity and smoking. The presence of each of these conditions was summed up to obtain a score of risk factors potentially ranging from 0 to 8 ([Supplementary Fig. 1](#)).

## Results

We included in the final analyses 3894 patients who were hospitalized with laboratory-confirmed SARS-CoV-2 infection at 30 clinical centres across Italy and either died or had been discharged or were still in hospital as of May 29, 2020. Baseline characteristics are shown in [Table 1](#).

Among COVID-19 patients, there was a higher prevalence of men (61.7%) and elderly (54.8% of patients aged ≥65 years; median age 67 years, interquartile range (IQR): 55–79 years) ([Table 1](#)). Half of the patient sample reported a diagnosis of hypertension and about a fifth had diabetes. The prevalence of chronic degenerative diseases at admission (previous myocardial infarction, heart failure, cancer and lung disease) ranged from 10.0% to 14.3%.

At hospital admission, 71.8% of COVID-19 patients had high levels of CRP (>3 mg/L) and only 35.1% had normal eGFR (≥90 mL/min/1.73 m<sup>2</sup>).

At the end of follow-up, out of 3894 patients, 712 died (18.3%), 2650 were discharged alive (68.0%) while 532

**Table 3** Subgroup analyses showing hazard ratio for mortality, according to age classes. Multiple imputation analysis, N = 3894 patients and N = 712 deaths.

	Age classes		
	18–64 years	65–74 years	≥75 years
<b>Patient at risk</b>	1761	808	1315
<b>Death</b>	81	145	486
<b>Person-days</b>	26,417	15,437	22,032
<b>Death Rate, x1,000 PD</b>	3.1	9.4	22.1
	HR (95% CI)	HR (95% CI)	HR (95% CI)
<b>Gender</b>			
Female	-1-	-1-	-1-
Male	1.43 (0.69–2.96)	1.87 (1.33–2.64)	1.31 (1.07–1.61)
<b>Hypertension</b>			
No	-1-	-1-	-1-
Yes	0.85 (0.54–1.33)	0.92 (0.61–1.39)	0.87 (0.67–1.14)
<b>Diabetes</b>			
No	-1-	-1-	-1-
Yes	2.00 (1.15–3.50)	0.86 (0.59–1.25)	0.96 (0.78–1.19)
<b>Myocardial infarction</b>			
No	-1-	-1-	-1-
Yes	1.19 (0.43–3.28)	1.28 (0.75–2.20)	1.06 (0.82–1.39)
<b>Heart Failure</b>			
No	-1-	-1-	-1-
Yes	1.38 (0.56–3.37)	1.12 (0.66–1.92)	0.95 (0.74–1.23)
<b>Cancer</b>			
No	-1-	-1-	-1-
Yes	4.76 (2.46–9.21)	1.65 (0.95–2.86)	1.10 (0.86–1.40)
<b>Lung disease</b>			
No	-1-	-1-	-1-
Yes	1.09 (0.41–2.88)	1.76 (1.16–2.67)	1.11 (0.89–1.39)
<b>Obesity, BMI ≥30 kg/m<sup>2</sup></b>			
No	-1-	-1-	-1-
Yes	1.36 (0.75–2.46)	1.50 (0.92–2.45)	1.05 (0.73–1.52)
<b>Smoking</b>			
Non-smoker	-1-	-1-	-1-
Current Smoker	1.09 (0.47–2.49)	0.66 (0.39–1.11)	1.02 (0.71–1.46)
<b>CRP, mg/L</b>			
≤3	-1-	-1-	-1-
3–10	3.87 (1.74–8.60)	1.50 (0.81–2.78)	3.04 (2.19–4.22)
≥10	3.75 (1.92–7.34)	1.19 (0.68–2.08)	2.66 (1.62–4.35)
<b>eGFR, CKD stage, mL/min/1.73 m<sup>2</sup></b>			
≥90	-1-	-1-	-1-
60–89	1.76 (0.94–3.32)	1.49 (0.86–2.59)	1.10 (0.61–2.01)
45–59	3.30 (1.46–7.46)	1.93 (1.11–3.37)	1.84 (1.02–3.32)
30–44	3.24 (1.41–7.45)	3.33 (1.78–6.23)	2.36 (1.38–4.02)
15–29	13.89 (3.09–62.5)	4.86 (2.18–10.82)	3.00 (1.62–5.54)
<15	6.65 (1.66–26.73)	4.77 (1.63–13.9)	3.65 (1.92–6.96)

Controlling for gender, diabetes, hypertension, history of myocardial infarction, heart failure, chronic pulmonary disease, chronic kidney disease, CRP, obesity, smoking habit and anti-COVID19 drugs during hospitalization as fixed effects and repeated measures within hospital.

Abbreviation. BMI: body mass index; CKD: chronic kidney disease; CRP: C-reactive protein; eGFR: glomerular filtration rate; PD: person-days.

(13.7%) were still hospitalised. The median follow-up was 13 days (IQR: 7 to 22). Death rate (per 1000 person-days) and univariable hazard ratios for in-hospital death are shown in [Table 1](#).

### Main analyses

The trained Random Forest model proved to be quite accurate and robust in predicting survival in the test set (sensitivity 95.2%, specificity 30.8%, classification accuracy 83.4%, F1 value 90.4%). PFI analysis revealed that the most

important features in classification were eGFR and CRP (average drop of loss function >5% in the perturbed models), followed by age (loss-drop >4%). All the other input features showed a lower influence within the model ([Fig. 1](#)).

Feature importance was in line with the multivariable hazard ratios for in-hospital mortality associated with demographic and clinical characteristics as obtained by survival analyses ([Fig. 2](#)). Indeed, men and the elderly COVID-19 patients had higher mortality as compared to their counterparts. In particular, patients aged ≥85 years had an 8-fold higher mortality as compared with the

**Table 4** Subgroup analyses showing hazard ratio for mortality, according to geographical location of the hospitals. Multiple imputation analysis, N = 3894 patients and N = 712 deaths.

	Geographical location of the hospitals	
	Northern regions	Central and Southern regions
<b>Patient at risk</b>	2278	1616
<b>Death</b>	513	199
<b>Person-days</b>	32,839	31,047
<b>Death Rate, x1,000 PD</b>	15.6	6.4
	HR (95% CI)	HR (95% CI)
<b>Gender</b>		
Female	-1-	-1-
Male	1.20 (0.95–1.52)	1.36 (0.97–1.91)
<b>Age, years</b>		
18–44	-1-	-1-
45–64	1.55 (0.67–3.57)	1.87 (0.45–7.71)
65–74	4.19 (1.87–9.36)	1.75 (0.67–4.61)
75–84	5.63 (2.48–12.78)	4.97 (1.90–13.04)
≥85	8.42 (3.87–18.31)	8.60 (3.54–20.94)
<b>Hypertension</b>		
No	-1-	-1-
Yes	1.00 (0.90–1.12)	0.63 (0.38–1.04)
<b>Diabetes</b>		
No	-1-	-1-
Yes	1.09 (0.85–1.38)	0.83 (0.62–1.12)
<b>Myocardial infarction</b>		
No	-1-	-1-
Yes	1.21 (0.96–1.51)	1.31 (0.81–2.12)
<b>Heart Failure</b>		
No	-1-	-1-
Yes	1.07 (0.83–1.38)	1.11 (0.77–1.60)
<b>Cancer</b>		
No	-1-	-1-
Yes	1.16 (0.92–1.48)	1.66 (1.21–2.27)
<b>Lung disease</b>		
No	-1-	-1-
Yes	1.13 (0.91–1.41)	1.70 (1.34–2.14)
<b>Obesity, BMI ≥30 kg/m<sup>2</sup></b>		
No	-1-	-1-
Yes	1.24 (0.94–1.62)	0.80 (0.45–1.41)
<b>Smoking</b>		
Non-smoker	-1-	-1-
Current Smoker	1.11 (0.82–1.50)	1.17 (0.76–1.81)
<b>CRP, mg/L</b>		
≤3	-1-	-1-
3–10	2.59 (1.77–3.80)	2.35 (1.40–3.93)
≥10	2.53 (1.71–3.74)	2.09 (0.90–4.86)
<b>eGFR, CKD stage, mL/min/1.73 m<sup>2</sup></b>		
≥90	-1-	-1-
60–89	1.45 (1.05–2.00)	1.42 (0.87–2.29)
45–59	2.06 (1.45–2.92)	2.87 (1.84–4.47)
30–44	3.11 (2.09–4.63)	2.99 (1.65–5.42)
15–29	3.29 (2.48–4.37)	4.89 (2.98–8.03)
<15	6.13 (3.62–10.38)	3.85 (2.00–7.41)

Controlling for gender, age, diabetes, hypertension, history of myocardial infarction, heart failure, chronic pulmonary disease, chronic kidney disease, CRP, obesity, smoking habit and anti-COVID19 drugs during hospitalization as fixed effects and repeated measures within hospital.

Abbreviation. BMI: body mass index; CKD: chronic kidney disease; CRP: C-reactive protein; eGFR: glomerular filtration rate; PD: person-days.

Northern regions: Novara, Milano, Cinisello Balsamo, Monza, Varese, Cremona, Padova, Forlì, Ravenna, Modena, Pavia.

Central-Southern regions: Firenze, Pisa, Roma, Chieti, Pescara, Napoli, Pozzilli, Acquaviva delle Fonti, San Giovanni Rotondo, Taranto, Catanzaro, Catania, Palermo.



**Table 5** Subgroup analyses showing hazard ratio for mortality, according to pandemic wave. Multiple imputation analysis, N = 3894 patients and N = 712 deaths.

	Pandemic wave	
	From 19/02/2020 to 19/03/2020	From 20/03/2020 to 23/05/2020
<b>Patient at risk</b>	1712	2182
<b>Death</b>	324	388
<b>Person-days</b>	29,149	34,737
<b>Death Rate, x1,000 PD</b>	11.1	11.2
	HR (95% CI)	HR (95% CI)
<b>Gender</b>		
Female	-1-	-1-
Male	1.53 (1.27–1.85)	1.32 (1.04–1.68)
<b>Age, years</b>		
18–44	-1-	-1-
45–64	1.73 (0.55–5.49)	1.74 (0.58–5.24)
65–74	4.15 (1.51–11.42)	3.40 (1.24–9.29)
75–84	6.46 (2.17–19.25)	5.74 (1.94–17.02)
≥85	6.61 (2.29–19.02)	9.08 (3.38–24.37)
<b>Hypertension</b>		
No	-1-	-1-
Yes	1.06 (0.80–1.41)	0.73 (0.55–0.97)
<b>Diabetes</b>		
No	-1-	-1-
Yes	0.96 (0.68–1.36)	1.09 (0.87–1.38)
<b>Myocardial infarction</b>		
No	-1-	-1-
Yes	1.07 (0.77–1.48)	1.36 (0.98–1.89)
<b>Heart Failure</b>		
No	-1-	-1-
Yes	1.07 (0.73–1.58)	1.03 (0.77–1.39)
<b>Cancer</b>		
No	-1-	-1-
Yes	1.38 (0.92–2.07)	1.31 (1.01–1.68)
<b>Lung disease</b>		
No	-1-	-1-
Yes	1.30 (0.99–1.71)	1.18 (0.88–1.58)
<b>Obesity, BMI ≥30 kg/m<sup>2</sup></b>		
No	-1-	-1-
Yes	1.27 (0.92–1.75)	1.07 (0.70–1.64)
<b>Smoking</b>		
Non-smoker	-1-	-1-
Current Smoker	0.89 (0.54–1.48)	1.04 (0.74–1.46)
<b>CRP, mg/L</b>		
≤3	-1-	-1-
3–10	2.62 (1.58–4.36)	2.59 (1.74–3.86)
≥10	2.31 (1.40–3.82)	2.26 (1.29–3.97)
<b>eGFR, CKD stage, mL/min/1.73 m<sup>2</sup></b>		
≥90	-1-	-1-
60–89	1.46 (0.97–2.20)	1.48 (0.97–2.25)
45–59	2.23 (1.34–3.72)	2.25 (1.44–3.53)
30–44	2.45 (1.43–4.19)	3.42 (2.13–5.47)
15–29	2.99 (1.86–4.83)	4.71 (2.86–7.77)
<15	7.50 (4.32–13.01)	3.85 (2.01–7.38)

\*Controlling for gender, age, diabetes, hypertension, history of myocardial infarction, heart failure, chronic pulmonary disease, chronic kidney disease, CRP, obesity, smoking habit and anti-COVID19 drugs during hospitalization as fixed effects and repeated measures within hospital.

Abbreviations: BMI: body mass index; CKD: chronic kidney disease; CRP: C-reactive protein; eGFR: glomerular filtration rate; PD: person-days.

youngest (18–64 years) patients. Reduced kidney function was associated with a proportionally higher increase of death risk. High levels of CRP, a well-known, sensitive marker of inflammation, were also associated with an increased mortality risk.

Among chronic degenerative diseases, history of or active cancer was associated with a high in-hospital mortality (HR: 1.33; 95% CI 1.09–1.63) and an upward trend of risk was found for previous myocardial infarction (HR: 1.20; 95% CI 0.93–1.55), chronic pulmonary disease (HR: 1.20; 95% CI 0.97–1.48) and obesity (HR: 1.21; 95% CI 0.91–1.61). On the contrary, current smoking, diabetes and hypertension were not associated with mortality in our COVID-19 patients. However, a score of risk factors ≥3 was associated with a 47% increase in mortality risk ([Supplementary Fig. 1](#)).

These results were confirmed in a number of secondary analyses that were run to test the robustness of the main findings by using both multiple imputation and complete-case analysis ([Supplementary Table 1](#)) or through accounting of clustering within hospitals ([Supplementary Table 2](#)) and censoring of follow up at 35 days ([Supplementary Table 3](#)).

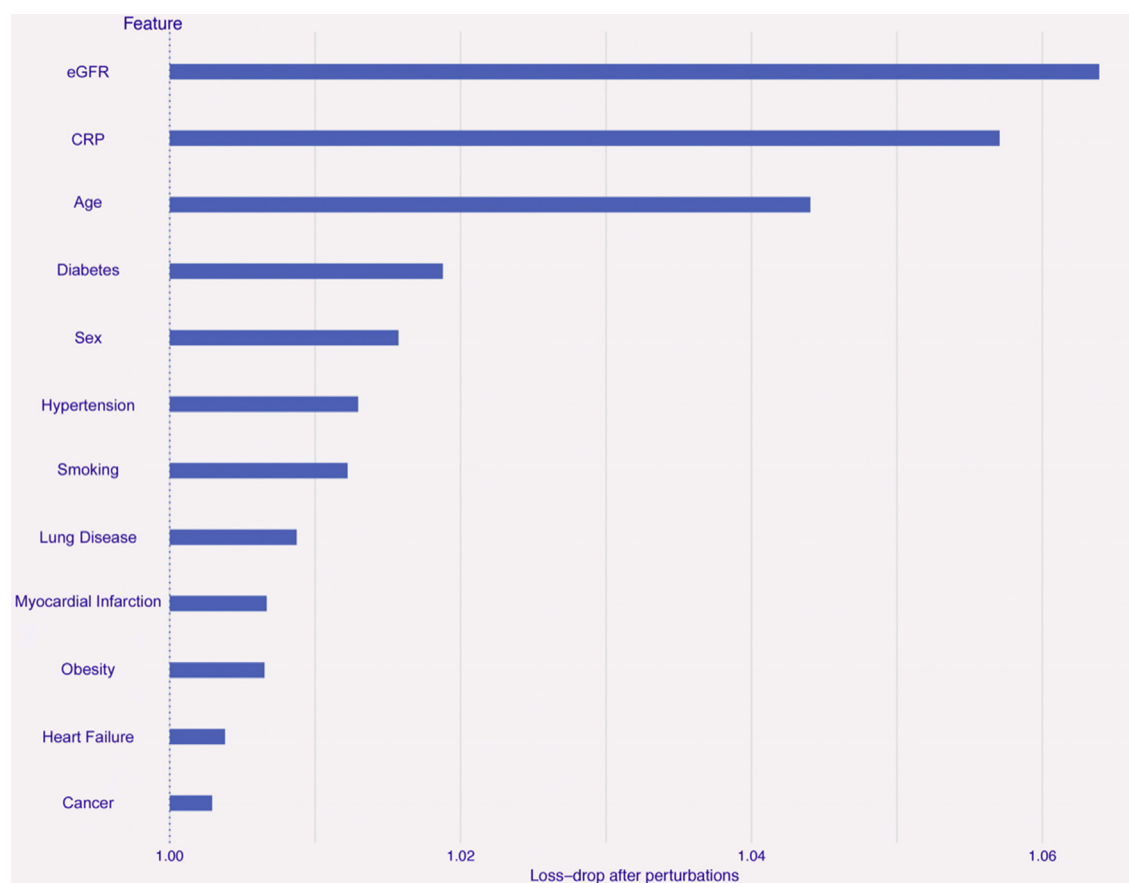
### Subgroup analyses

Subgroup analyses according to gender, age classes, geographical location of the hospitals (Northern, and Central/Southern regions) and temporal pandemic waves are presented in [Tables 2–5](#). Additional information on age- and gender-adjusted prevalence of the here-studied determinants of in-hospital mortality according to gender, age classes, geographical location of the hospitals and pandemic wave are reported in [Supplementary Table 4](#).

A higher death rate was observed in the elderly (age ≥ 75 years: 22.1 deaths x 1000 person-days) compared to younger patients (18–64 years: 3.1 deaths x 1000 person-days and 65–74 years: 9.4 deaths x 1000 person-days), as well as in patients admitted to hospitals of Northern regions (15.6 deaths x 1000 person-days) in comparison with those from Central/Southern regions (6.4 deaths x 1000 person-days) ([Tables 3 and 4](#)).

In general, subgroup analyses confirmed the findings reported in [Fig. 1](#). However, some differences could be observed. A hypertensive status was associated with a reduction of mortality among women and those patients hospitalised during the second wave of the pandemic emergency ([Tables 2 and 5](#)). However, major risk factors were also confirmed among those patients with hypertension ([Supplementary Table 5](#)).

Diabetes turned out to be associated with a higher risk of in-hospital mortality in younger COVID-19 patients (18–64 years) ([Table 3](#)). The magnitude of the association between history of cancer and in-hospital mortality was stronger in younger than in older patients ([Table 3](#)).



**Figure 1** Permutation Feature Importance analysis for the prediction of mortality. Bars indicate the importance of each feature used for the prediction of death in the Random Forest algorithm within the training set ( $N = 2,725$ , 70% of the total imputed sample available), based on the ratio between loss function in the perturbed model (i.e., after permutation of a given variable) and loss function in the full model (with no permuted variable). The higher the ratio, the more the perturbed model is altered and the more important is the permuted feature. Abbreviations. CRP: C-reactive protein; eGFR: glomerular filtration rate.

Chronic pulmonary disease was associated with in-hospital mortality both in men and in patients aged 65–74 years, as well as in those admitted to hospitals of Central-Southern regions (Tables 2–4). Additionally, non-hypertensive patients with chronic pulmonary diseases showed a 2-fold higher risk as compared with those without lung disease (Supplementary Table 5).

We found no association between obesity and in-hospital mortality, data confirmed by complete-case analysis restricted to 1517 patients without missing data on BMI and by distinguishing obesity as stage 1 ( $BMI = 30\text{--}34.9 \text{ kg/m}^2$ ) and stage 2 ( $BMI \geq 35 \text{ kg/m}^2$ ) (Supplementary Table 6).

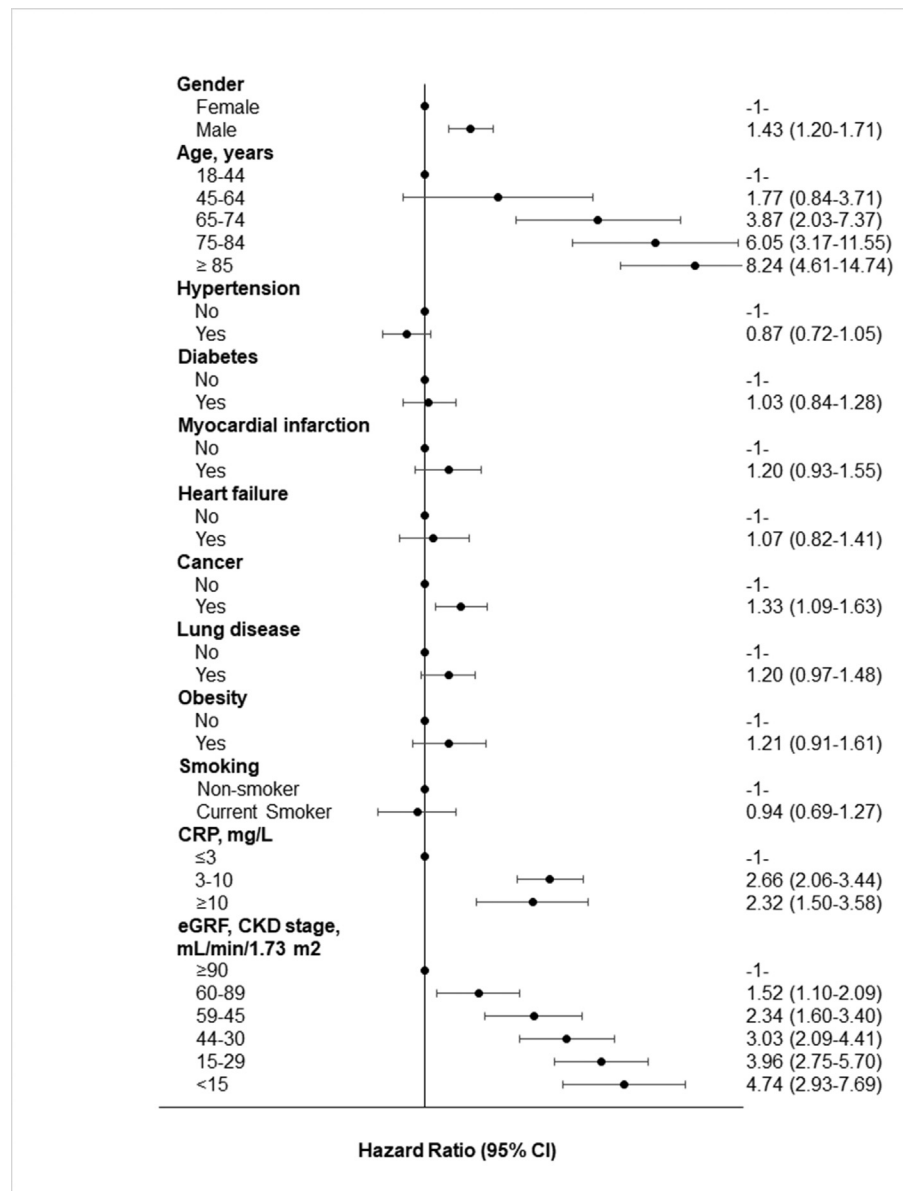
## Discussion

We analysed data from a large cohort of patients hospitalised between February 19 and May 23, 2020 with laboratory-confirmed SARS-CoV-2 infection, with the aim of defining an algorithm to classify death risk and identifying major predictors of in-hospital death.

Our sample had a higher prevalence of men (61.7%) and elderly aged  $\geq 65$  years (54.8%) as compared to a large

study on COVID-19-patients in New York City (49.5% of men and less than 30% of elderly individuals [4]), while in our cohort obese subjects were under-represented (9.7% vs 35% in the US cohort).

Through a supervised machine learning approach, we built a random forest algorithm to classify death risk in our cohorts, also identifying the most influential predictors of this risk and taking into account potential non-linear relations and interaction effects among them. This algorithm reached a good ( $>83\%$ ) accuracy, with a high ( $>90\%$ ) sensitivity for classification of survivors ( $>95\%$ ), probably due to the fact that the majority of our patients did not die, which made the model more prone and fitted to recognise these outcomes. Moreover, this approach revealed that age, renal function (eGFR) and circulating inflammation (hs-CRP) were the most important features predicting survival. While CRP and age have been already identified as top influential features in previous machine learning algorithms predicting mortality risk [21,22,25], we are not aware of any comparable approach identifying renal function among top features. This may be due to the huge heterogeneity of settings and of clinical variables available from COVID-19 patients in different studies, which make it hard to compare the studies among themselves. Further



**Figure 2** Multivariable hazard ratios for in-hospital mortality for different characteristics of patients. Multiple imputation analysis, N = 3894 patients and N = 712 deaths. Controlling for gender, age, diabetes, hypertension, history of myocardial infarction, heart failure, chronic pulmonary disease, chronic kidney disease, CRP, obesity, smoking habit and anti-COVID19 drugs during hospitalization as fixed effects and repeated measures within hospital. Abbreviations. BMI: body mass index; CKD: chronic kidney disease; CRP: C-reactive protein; eGFR: glomerular filtration rate.

investigations with more homogeneous settings and clinical variables are needed to further substantiate these findings. However, “classical” (non-machine learning based) statistical analyses (see below) revealed data in line with the results of the permutation feature importance analysis applied to the random forest algorithm, providing robustness to our findings. Indeed, Cox survival regressions revealed that advanced age, impaired renal function and elevated levels of CRP, as well as male gender and cancer, were independent predictors of in-hospital death.

These data are in line with prior evidence on a small sample of Italian patients admitted to hospital during the first wave of the COVID-19 pandemic, documenting

increased risk of death associated with older age, critical disease and high levels of CRP [9]. Accordingly, investigations from the U.S. and China also identified advanced age [2,4,5], cancer [4] and inflammation [2,4,5] as major predictors of death.

At variance from other studies [4,7–9], however, we failed to find any association between obesity and smoking on admission with the risk of in-hospital death, although findings on tobacco use were in accordance with what seen in a U.S. cohort [4]. Obesity was not a risk factor in a large U.S. cohort [5] and in another Italian series of COVID-19 patients [33].

Our findings do not support a pivotal role for mortality of previous known cardiovascular disease and other

comorbidities, data apparently contrasting from those of previous studies suggesting a major association of pre-existing cardiovascular conditions with higher mortality [5,12] and poor prognosis [34].

However, neither the study by Cummings [5] nor that by Ciceri [12] did account for smoking status in their multivariable analysis, and this may have biased their results; moreover, the Italian analysis [12] was conducted on a small sample of subjects from one single centre, as in the study by Colaneri and colleagues [35]. In another setting, the associations between cardiovascular disease and related comorbidities were mitigated after multivariable adjustments [36]. In addition, studies with a larger sample size, similar to ours, failed to find any association between cardiovascular comorbidities and in-hospital mortality after multivariable adjustments [4].

Age is one of the strongest predictors of mortality in COVID-19 patients, and the prevalence of cardiovascular comorbidities obviously increases with age. In univariable analysis, all cardiovascular risk factors are associated with a higher mortality. Adjustment for age strongly reduced almost all associations, and further adjustment for kidney impairment completely eliminated any residual association, suggesting that the latter were essentially driven by these two main factors (Supplementary Table 7), as also discussed previously [37,38].

The importance of renal function is reinforced by the observation that its deterioration was predictive of mortality across all sub-groups analysed in this study, consistent with data from 701 Chinese patients with COVID-19 showing that kidney disease was associated with a higher risk of in-hospital death [39]. We acknowledge that our analysis is based on one-time measurement of eGFR values at the hospital entry that does not allow to discern infection-induced acute kidney injury (AKI). It has been recently reported that patients with hospital-acquired AKI as opposed to those with community-acquired AKI had higher rates of in-hospital death [40].

Moreover, in a small sample of Italian patients, low levels of albumin were associated with higher risk [41]. It is known that SARS-CoV-2 can infect podocytes and tubular epithelial cells [42]. Thus, viral infection may be the cause of renal abnormalities, partially explaining the early involvement of kidney impairment in the severity of COVID-19. Alternatively – or possibly complementary to this – renal dysfunction may predispose affected individuals to a more rapid deterioration and death.

We carried out a number of subgroup analyses in order to test whether the association between variables found to predict in-hospital death was likely to vary according to age, gender, geographical area and temporal pandemic waves. Our results indicate a similar magnitude of the association between selected determinants and the outcome, although the strength of the relation appeared to differ for some baseline characteristics.

First, in-hospital mortality was likely to follow a geographical gradient, death rates being more than double in hospitals located in Northern regions as compared to Central-Southern centres. These inter-regional disparities

in death rates are likely linked to the timing of the COVID-19 outbreak that first hit Northern regions, forced to face an almost unknown and unexpected infectious agent that only later spread to Central-Southern areas of the country. Moreover, evidence-based management guidelines for patients diagnosed with COVID-19 were not available during the early stages of the pandemic. In addition, a viral load in Northern Italian coronavirus infections higher than in other parts of the country may have contributed to the different case-fatality ratio.

Analyses between temporal pandemic waves revealed similar death rates in the first and second wave, although the impact of hypertension on mortality was likely mitigated during the second wave (end of March–end of May), as compared to the first pandemic wave (mid-February–mid-March), while the other predictors were substantially homogenous. These data indicate that the risk factors analysed here were basically independent from the advancement in therapies against COVID-19 that occurred over time, and highlight the importance of prevention strategies targeting high risk individuals more aggressively. Finally, our data are unlikely to support the assumption of a reduced viral load over time.

Risk predictors slightly varied between genders. Gender-related differences are well established both for infectious and non-communicable diseases [43,44]; more recently, testosterone has been suspected as playing a critical role in driving the excess of risk observed among men tested positive for COVID-19 [45].

Among subjects under 65 years, major risk factors were kidney disease and inflammation, as well as cancer and diabetes, while for those aged  $\geq 75$ , male gender, kidney disease and inflammation were the only factors independently associated with fatality.

Overall, our data suggest that among young patients pre-existing diseases represent a major risk factor for a clinical outcome such as mortality, while they have a marginal role in the elderly.

A laboratory predictor, such as CRP levels, was consistently and independently associated with death risk, suggesting that pre-hospitalisation inflammation (possibly in response to viral infection) might have been even more important than baseline personal characteristics and comorbidities. Our data on excess of death risk associated with inflammation support the hypothesis of a 'cytokine storm' caused by SARS-CoV-2 being a main player in the high mortality of COVID-19 pneumonia and ARDS [46].

### Strengths and limitations

To the best of our knowledge, this is the largest study on COVID-19 patients actually available in Italy and one of the largest multicentre studies of patients with COVID-19 in Europe. Moreover, we are not aware of other larger studies attempting to classify death risk in COVID-19 patients through a machine learning approach, in a scenario where most studies show a sample size  $<1000$  [17,32]. Indeed, an unselected patient sample from 30 hospitals, covering the



entire Italian territory and all the overt epidemic period in Italy, was collected and analysed.

A second strength is the use of a multifaceted approach to estimate the risk and the most influential predictors of in-hospital death, which implied i) a supervised machine learning algorithm to classify death and survival and ii) multivariable, time-varying analyses to describe in detail the relationship of independent risk factors with in-hospital death. This allowed to also take into account potential non-linear and synergistic relationships among the risk factors. Moreover, several statistical approaches were used to test the robustness of the associations and overcome biases due to the observational nature of the study.

On the other hand, the results of this study should be interpreted in the light of several limitations. First, results may not be generalized to other populations with different geographical and socioeconomic conditions, differences in national health service or insurance-based health expenses, and in the natural history of COVID-19. Due to the retrospective nature of our study, some information (such as smoking and obesity) was not available in all patients. Lack of data on vital signs (e.g. respiratory rate or oxygen saturation) represents another limitation of this study although CRP levels and renal function were likely to be proxies of patients' severity upon admission.

Finally, the possibility of unmeasured residual confounders cannot be ruled-out due to the observational nature of our study.

## Conclusions

Impaired renal function, elevated levels of CRP and advanced age at hospital admission were powerful predictors of higher in-hospital death in a large cohort of unselected patients with COVID-19 admitted to 30 different clinical centres all over Italy.

Some of these risk factors are likely to vary somewhat according to gender, age and geographical location for reasons that are not fully understood, although being possibly associated, at least in part, with timing of the pandemic outbreak and the local management modalities of patients. Along with epidemiological, pathophysiologic and possibly epigenetic mechanisms underlying such differences need to be elucidated.

Our data also support the importance of measuring inflammatory markers other than CRP at admission and during hospital stay in patients with COVID-19.

Finally, our findings contribute to identify some potential effect modifiers in the association between risk factors and in-hospital mortality among COVID-19 diagnosed patients. Future comparison with other European and Mediterranean Countries, possibly including assessments of other factors such as socioeconomic status [47], will be useful to elucidate possible differences in risk

factors and outcomes observed across different geographical areas.

## Authors' contribution

ADC, LI, RDC designed the study. ADC and LI supervised research and led the data collection. All Authors contributed to collect, re-analyse, check and pool data, critically reviewed and approved the manuscript. SC and AG analysed the data and prepared the results. GV advised statistical analyses. MO managed the web-based database. MB and SC wrote the first draft of the manuscript with input from RDC, KdD, ADC, AG, LI, and the other Authors.

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## Additional information

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## Declaration of Competing Interest

All Authors declare no competing interests.

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The Authors alone are responsible for the views expressed in this article. They do not necessarily represent the views, decisions, or policies of the Institutions with which they are affiliated. AG was supported by Fondazione Umberto Veronesi.

## Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.numecd.2020.07.031>.

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