The relationship between cardiac injury, inflammation and coagulation in predicting COVID-19 outcome

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Research Article
Abstract

Introduction: High sensitivity troponin T (hsTnT) is a strong predictor of adverse outcome during SARS-CoV-2 infection. However, its determinants remain partially unknown. We aimed to assess the relationship between severity of inflammatory response/coagulation abnormalities and hsTnT in Coronavirus Disease 2019 (COVID-19). We then explored the relevance of these pathways in defining mortality and complications risk and the potential effects of the treatments to attenuate such risk.

Methods: In this single-center, prospective, observational study we enrolled 266 consecutive patients hospitalized for SARS-CoV-2 pneumonia. Primary endpoint was in-hospital COVID-19 mortality.

Results: hsTnT, even after adjustment for confounders, was associated with mortality. D-dimer and CRP presented stronger associations with hsTnT than PaO$_2$. Changes of hsTnT, D-dimer and CRP were related but only D-dimer was associated with mortality. Moreover, low molecular weight heparin showed attenuation of the mortality in the whole population, particularly in subjects with higher hsTnT.

Conclusions: D-dimer possessed a strong relationship with hsTnT and mortality. Anticoagulation treatment showed greater benefits with regard to mortality. These findings suggest a major role of SARS-CoV-2 coagulopathy in hsTnT elevation and its related mortality in COVID-19. A better understanding of the mechanisms related to COVID-19 might pave the way to therapy tailoring in these high-risk individuals.

Introduction

Elevated high sensitivity troponin T (hsTnT) at admission, a well-established marker of cardiac injury, has been shown to be common and highly predictive of adverse outcome in patients with Coronavirus Disease 2019 (COVID-19) [1, 2, 3]. The mechanisms underlying to troponin elevation in COVID-19 are thought to be multifactorial. Hypoxia related to interstitial pneumonia potentially leading to non-ischemic myocardial injury, as well as the exaggerated immune-inflammatory response, marked by an elevation of the C-reactive protein (CRP), are amongst the two most commonly cited mechanisms potentially leading to hsTnT elevation in these patients [4, 5].

More recently, activation of coagulation pathways has been described in a substantial proportion of patients with severe COVID-19 and their presence has been related to the development of multi-organ dysfunction, resulting in an increased risk of death [6]. An hyperactivation of the coagulation system, marked by an elevation of the D-dimer, has been associated to an increased risk of mortality in COVID-19 [3, 7]. Post-mortem studies have confirmed the presence of a COVID-19-related coagulopathy characterized by the development of thrombotic microangiopathy in multiple organs, including the heart vessels [8]. Such anatomical alterations can contribute to the elevation of hsTnT independently from the severity of the inflammatory response, as they can further aggravate the delivery of an adequate oxygen supply to the myocardium. The importance of control the coagulatory homeostasis in COVID-19 is recently under as some studies have reported a beneficial effect of anticoagulation therapies in these patients[7, 9]. However, no studies have accurately explored the relative contribution of these three pathways to the increase in hsTnT and their importance in defining the risk of mortality in COVID-19. A better understanding of the plethora of ischemic and non-ischemic causes of myocardial injury related to COVID-19 might inform on the most appropriate treatment to be used in these high-risk patients.
In this study, we evaluated the relative importance of the hypoxia, inflammatory response and activated coagulation in the elevation of the hsTnT levels in patients with a diagnosis of Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2). We also assessed the complex interaction between hsTnT, D-dimer and C-reactive protein (CRP) in defining the risk of an adverse outcome or hospital complications in the same population. Finally, given the potential contribution of inflammation and coagulation abnormalities to the myocardial injury in COVID-19, we assessed the capacity of immunomodulating drugs or low molecular weight heparin to modify the relationship between high hsTnT levels and increased risk of mortality.

Methods

Population and data collection

This single-center, prospective, observational study included 266 consecutive patients hospitalized for SARS-CoV-2 pneumonia at the tertiary care University hospital of Pisa, Tuscany, Italy, between 4th March and 31st March. All patients had a diagnosis of SARS-CoV-2 infection confirmed by positive results of PCR testing of a nasopharyngeal swab. Pneumonia was established with a plain chest CT reviewed by trained radiologists. Demographic, clinical, laboratory, instrumental, treatment, and outcome data were collected. Venous and arterial blood samples for standard biochemistry and arterial blood gas analysis were collected at the time of hospitalization and repeated according to clinical practice and depending on the patient clinical conditions. The study complied with the edicts of the 1975 Declaration of Helsinki and. All experimental protocols were approved by the institutional ethic committee Comitato Etico Area Vasta Nord Ovest (CEAVNO).

Outcomes

The primary endpoint of the study was considered the in-hospital mortality. Secondary endpoints were considered the admission to Intensive Care Unit (ICU), the development of acute respiratory distress syndrome (ARDS, according to the Berlin Definition)[10] and the need for respiratory support by invasive or non-invasive mechanical ventilation.

Statistics

Continuous variables are presented as mean±standard deviation or median (interquartile range) for variables not following normal distribution. Nominal variables are summarized as absolute counts and percentage values (%). Normal distribution of all continuous variables was graphically assessed by histograms and quantile-quantile plots. Differences in variables of interest between pre-defined groups (i.e. according to sex, history of CVD, increased admission levels of hsTnT) were examined with the independent’s samples Student’s T Test or the non-parametric Mann-Whitney test and the chi-squared test for continuous and nominal variables, respectively. To identify independent determinants of admission hsTnT concentrations, we used multivariable linear regression models. Subsequently, we employed Cox proportional-hazards models to evaluate the association between admission variables and the occurrence of the primary endpoint (all-cause mortality). We used the Schoenfeld residuals to test the proportional hazards assumption. Kaplan Meier curves were plotted to illustrate the difference in survival probabilities (by implementing the Log rank test) according to admission hsTnT levels. Respectively, we used multivariable logistic regression analysis to appraise the association of hsTnT with
secondary outcomes, including incidence of ARDS, invasive and/or non-invasive mechanical ventilation and transfer to ICU. Finally, we utilized two-level linear mixed models with random effects (random intercept and random coefficient) and unstructured variance-covariance matrix to assess whether longitudinal changes in hsTnT levels were associated with fluctuations in levels of CRP and D-dimer across the hospitalization. In all multivariable regression models (Cox regression, logistic regression and linear mixed models), we controlled for a pre-specified set of biologically plausible covariates, including age, gender, history of CVD and creatinine levels; thus, no selection technique was implemented. In linear regression and linear mixed models, we ensured the normal distribution of the dependent outcome by transforming continuous hsTnT with the natural logarithm. Associations are presented as Hazard Ratio (HR) or Odds Ratio (OR) or coefficients with 95% confidence intervals (CI).

In this study, we included all consecutive patients with COVID-19 admitted to the Hospital of Pisa, Italy, until the 31st March 2020. In this context, no formal power analysis was performed. Still, we sought to retain a ratio of 5 to 10 events or observations per covariate used in Cox or logistic regression and linear regression models. In Cox regression models with a large number of covariates (i.e. predictors of mortality among different treatments after adjustment for hsTnT, age, gender, history of CVD and creatinine levels), we performed bootstrapping with 200 replicates and derived corrected confidence intervals.

Statistical analysis was conducted with STATA package, version 12.1 (StataCorp, College Station, Texas USA). All tests were 2-tailed. We deemed statistical significance at p<0.05.

**Results**

**Prognostic role of hsTnT**

The baseline clinical characteristics of the population stratified by mortality status at follow up are reported in Table 1. Mortality rate was high (32%) at the end of the follow-up. People who died were older and presented a higher blood pressure, prevalence of male sex, CVD comorbidities and chronic liver and kidney diseases. Their lymphocyte and platelet counts were lower, while they showed an elevated AST, creatinine, CRP, D-dimer and creatine kinase. Lower PaO$_2$ and PaO$_2$/FiO$_2$ on admission were also detected in patients who died during hospitalization. Remarkably, hsTnT was the parameter with the greatest difference between the group of patients who died during admission compared to subjects who survived (5-fold difference). In Cox regression models, hsTnT levels were strongly associated with the risk of in-hospital mortality, even after adjustment for age, sex, previous history of CVD, creatinine levels (Table 2 and Figure 1). Weaker or no associations were observed between hsTnT and risk of ARDS, ICU admission and ventilation. The association between hsTnT and mortality was significant in both sexes, as well as in people with and without previous history of CVD. Only age modified the relationship between hsTnT and risk of mortality, so that in subjects >75 years old this relationship was stronger than in those <75 years old (HR=1.012, p=0.005 vs HR=1.001, p=0.051).

Similar to hsTnT, D-dimer was associated with the risk of in-hospital mortality at Cox regression analysis, while weaker or no associations were identified with the other outcomes (Table 3). The association between D-dimer and mortality risk remained significant even after adjustment for age, sex, previous history of CVD and levels of creatinine (HR=1.11, 95% CI 1.03-1.185, p=0.005). Conversely, CRP was not associated with an increased
mortality risk, although it was related with the risk of ARDS, ICU admission and requirement of respiratory ventilation (Table 3).

Determinants of hsTnT levels

The Table 4 shows the characteristics of the population stratified by the median value of hsTnT. Comorbidities including diabetes mellitus, chronic obstructive pulmonary disease, arterial hypertension, prior history of CVD and cerebrovascular disease, dementia, chronic kidney disease were more prevalent in patients with higher hsTnT levels. They also had increased levels of D-dimer (P=0.021) and CRP (P<0.001) and lower PaO$_2$ (P<0.001) and PaO$_2$/FiO$_2$ ratio (P=0.001).

In linear regression models, including PaO$_2$, D-dimer and CRP as independent variables further adjusted for history of CVD, creatinine, sex and age, D-dimer and CRP presented a stronger association with the levels of hsTnT than PaO2 (Table 5). Linear mixed model analysis including the subgroup of patients with available serial measurements of hsTnT, D-dimer and CRP, changes in D-Dimer followed the fluctuations in hsTnT concentrations subjects after controlling for age, gender, creatinine and history of CVD (coefficient=0.014, 95% CI 0.01-0.024, p=0.003). A weaker association was also detected between changes in CRP and changes of hsTnT concentrations after adjustment (coefficient=0.053, 95% CI 0.0173-0.089, p= 0.004).

Attenuation of the risk related to high hsTnT levels by COVID-19 treatment

The prognostic role of hsTnT at admission remained significant also after adjustment for the main drugs used for the treatment of COVID-19, including low molecular weight heparin (LMWH), hydroxychloroquine, antibiotic, immunomodulating drugs (baricitinib and tocilizumab) and corticosteroids. In multivariable Cox regression models including the same drugs and controlling for age, sex, history of CVD, creatinine and hsTnT, only LMWH remained significantly associated with mortality after bootstrapping (HR=0.271, 95% C.I. 0.140-0.525, p<0.001). Thus, we tested whether the risk of mortality related to high hsTnT level was somewhat attenuated by the treatment with heparin. When added to the model testing the relationship between hsTnT levels and mortality, the interaction between hsTnT and use/non-use of heparin during hospitalization was highly significant (HR=0.424, 95% C.I. 0.239-0.752, p=0.003), suggesting that heparin could modify the relationship between hsTnT levels and mortality risk. Indeed, in the group of subjects with levels of hsTnT above the median values the treatment with heparin significantly lowered the risk of mortality compared to untreated subjects (HR=0.294, 95% C.I. 0.152-0.569, p<0.001), while the same difference was not detected in subjects with hsTnT lower than the median (HR=0.06, 95% CI 0.002-2.100, p=0.121) (Figure 2).

Discussion

In this single-centre, prospective, observational study we report that: hsTnT five-fold elevation is the greatest difference observed among a large plethora of clinical and biochemical parameters characterizing baseline phenotype of 266 patients hospitalized for COVID-19 and stratified for mortality; baseline hsTnT and D-dimer, but not CRP, are associated with in-hospital COVID-19 mortality; D-dimer accounts for a substantial proportion of the hsTnT variability; anticoagulant treatment during hospitalization substantially reduced the mortality risk, a benefit particularly evident in subjects with elevated hsTnT.
Our study aimed at exploiting, from a clinical in-hospital point of view, the complex interplay between inflammation, coagulation and cardiac injury. We showed that the activation of inflammatory/coagulation pathways might explain a large proportion of the increase in hsTnT levels commonly observed in the COVID-19, probably higher than the severity of hypoxia. Although changes in hsTnT levels during hospitalization were significantly related to the fluctuations of the CRP and D-dimer concentrations, only D-dimer presented an association with mortality similar to hsTnT. These findings suggest that cardiac injury and coagulation pathways might be relevant in defining the risk of mortality related to elevated hsTnT levels in patients with SARS-CoV-2-related pneumonia.

In our observation we found hsTnT, among a large numbers of serum biomarkers, the parameter with the major difference at baseline between survived vs deceased patients hospitalized for COVID-19. Cardiac injury, marked by troponin elevation, is frequent in patients with COVID-19 and related with adverse outcomes[11]. In fact, while the most common clinical presentation of the SARS-CoV-2 infection is bilateral interstitial pneumonia often associated to respiratory failure, several reports have supported the capacity of cardiac injury, indirectly assessed by cardiac troponin due to the in-hospital real-life setting, to inform on the mortality risk in COVID-19 [1, 2, 3, 11]. Troponin elevation has been related with a large set of inflammatory serum markers, such as CRP, ferritin, IL-6, tumour necrosis factor-α and also thrombotic markers such as D-dimer. No previous studies[11], however, have accurately explored the potential and complex interaction between these three biologic systems and thus defined the putative determinants of hsTnT elevation levels in COVID-19.

In accordance with current literature, we document that hsTnT is associated to an increased risk of in-hospital mortality in subjects with and without previous history of CVD[11], suggesting that the exclusive presence of coronary atherosclerosis is unlikely to completely explain the high mortality risk of these patients[12]. One plausible hypothesis is that cytokine storm induced by the SARS-CoV-2-related pneumonia can promote the activation of the coagulation system which, in turn, can result in a systemic microangiopathy, multi-organ failure and increased mortality[13]. However, the relevance of the inflammation in defining the mortality outcome has recently been debated, as observed by Sinha et al[14]. Indeed, similar to hsTnT, an increased D-dimer concentrations have been related to the mortality risk of infected patients [3, 7].

In our observation, also, we found that CRP, beyond its association with hsTnT, was related to the risk of ICU admission and requirement of mechanical ventilation, both outcomes defining the severity of the pneumonia. However, CRP was not predictive of mortality. Instead, D-dimer and hsTnT presented a strong relationship with mortality, but they were not informative on the risk of respiratory complications eventually leading to ICU admission. It’s plausible that while CRP might reflect more closely the pulmonary stage of the disease and the severity of cytokine storm related to pneumonia, the elevation of D-dimer and hsTnT might sign the evolution towards a systemic involvement of the infection. This is in line with the fact that, in our study, we observe a larger variability of hsTnT explained through D-dimer levels instead that through the severity of the hypoxia related to pneumonia. The development of disseminated intravascular coagulation (DIC) in patients with more severe interstitial pneumonia could partially explain the stronger relationship of D-dimer and hsTnT with the mortality risk compared to CRP. However, in ours and other studies, the pattern of coagulopathy observed during SARS-CoV-2 infection was substantially different to the DIC commonly seen in sepsis [15]. Indeed, our population did not show severe thrombocytopenia which usually accompanies the sepsis-related DIC, and the D-dimer concentrations often reached higher levels than those expected in DIC. It has been repeatedly emphasized that most patients with COVID-19 related coagulopathy would not be classified as having DIC according to the
recommendations of the International Society on Thrombosis and Haemostasis [7, 15]. However, a large involvement of the coagulation system in COVID-19 is undeniable: in post-mortem studies, a microangiopathy characterized by typical microvascular platelet-rich thrombotic depositions has been described not only in small vessels of the lung parenchyma but also in other tissues, such as the myocardium [8]. Such microangiopathy could account more directly for the multi-organ failure and death in COVID-19 patients. Following this interpretation, the elevation of hsTnT might represent also an early marker of a systemic macroangiopathic involvement during COVID-19 infection, and its high prognostic capacity might depend on its higher sensitivity and specificity compared to other markers of tissue injury [4].

In our study only the treatment with low molecular weight heparin showed convincing evidence for an attenuation of the mortality risk. This is in line with the current literature: observations about anticoagulation therapy show promising results and expert opinions have recommended the use of prophylactic low molecular weight heparin in patients with COVID-19 in absence of clinical contraindications [6, 16]. Ayerbe et al reported an association between heparin and lower mortality in 2075 patients admitted with COVID-19 in 17 hospitals in Spain [9]. In a retrospective study including 449 patients with severe COVID-19 in China, 99 of which received heparin, 28-day mortality of heparin users was lower than that of non-users in patients with D-dimer levels more than six-fold the upper limit of normal [7]. An observational study including 2,773 patients hospitalized with laboratory-confirmed COVID-19 within the Mount Sinai Health System in New York City showed that systemic anticoagulation was associated with improved outcomes [6]. As for other systemic diseases with high prevalence in older population, the prescription of an anticoagulation treatment should be weighed against the risk of bleeding. Otherwise, this may lead to controversial results. Indeed, Pesavento et al showed that, in a cohort of 324 patients showed that (sub)therapeutic doses of anticoagulants, when compared to standard prophylactic ones, are related to a major risk of bleeding (HR 3.89; 95%CI, 1.90 to 7.97) despite no beneficial effect on overall mortality[17].

In our study, we report, for the first time, that the benefit from anticoagulation therapy was more evident in subjects with higher hsTnT elevation. The positive effects obtained from the prescription of low molecular weight heparin and particularly in patients with higher hsTnT levels confirm the contribution of systemic coagulopathy to the myocardial injury detected in COVID-19. Moreover, due to the bleeding risk implied with the LMWH treatment, the identification of parameters that enable selection of patients who could receive the greatest benefits from anticoagulation therapies might be crucial to optimize the risk-benefit balance related to this treatment. Our data suggest that a prophylactic dose of low molecular weight heparin during admission might be particularly indicated in patients presenting with mild to moderate elevation of hsTnT levels. If these patients might benefit from the use of higher dose thromboprophylaxis than is generally given remains unknown. However, ongoing multicentre, randomized, controlled trials will be able to address this question (NCT04372589, NCT04367831, NCT04345848, and NCT04366960).

Our study has some strengths and limitations. The availability of makers informing on the severity of hypoxia, inflammatory response and coagulation abnormalities provided us with the opportunity to explore their relative importance in causing an increase in the hsTnT level and its association with mortality. In a subset of patients, serial measures of hsTnT, CRP and D-dimer were also available, making it possible, for the first time, to relate changes in hsTnT levels with the fluctuation of the inflammatory and coagulation responses to the disease. The presence of several outcome measurements enabled the assessment of the relative importance of the different pathways explored in the study with multiple complications of patients with COVID-19 infection. However, our
study is monocentric, thus the results might be affected by local practice in the management of the COVID-19 infection and not be generalizable to other health care settings. Also, we used only D-dimer and CRP as markers of coagulation abnormalities and inflammation. The health care emergency imposed by the elevated number of SARS-CoV-2 positive patients admitted to the hospital in a short period of time did not provide us the opportunity to obtain a more accurate assessment of the COVID-19-related coagulopathy and inflammatory response. However, previous report have shown that the elevation in the circulating levels of D-dimer is amongst the most typical characteristics of the coagulopathy characterizing the SARS-CoV-2 infection and its assessment is commonly used in clinical practice as a screening tool for the presence of trombo-embolic events in other diseases[18]. Similarly, CRP is commonly used in clinical practice to monitor the severity of the inflammatory response resulting from acute infections and is also considered a good marker to assess the inflammatory-related risk of cardiovascular events [19]. The analysis on the beneficial effects of treatments should be interpreted cautiously, as it was not conducted on randomized groups and might be therefore affected by several measured and unmeasured confounding factors. Furthermore, we were not able to recover clear information on the indication for anticoagulation treatment.

In summary, our data support the hypothesis that the increase of hsTnT levels detected in COVID-19 infection is multifactorial, with inflammation and coagulation responses representing important factors contributing to this clinical manifestation. CRP can be used to monitor inflammation but might not represent the most accurate marker to discriminate between the severity of the respiratory and systemic inflammatory involvement during SARS-CoV-2-related pneumonia. Instead, an elevation of the D-dimer might sign the evolution towards a systemic involvement of the infection, with the development of a coagulopathy that could lead to multiorgan failure. Low molecular weight heparin is, indeed, a promising treatment to counteract the risk of mortality related to systemic COVID-19. Further studies are needed to clearly define the pathogenesis of COVID-19 and provide useful insight for tailored therapies. Troponin, in a context where risk stratification is of primary importance, due to its ability to predict mortality and the relevance of if prognostic capacity even in terms of identification of patients who would benefit more from specific treatments (such as anticoagulation), might play a fundamental role.

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**Declarations**

**Acknowledgments:** PISA Covid Study Group members and affiliations: Agostini o Degl’Innocenti Sabrina, Biancalana Martina, Borselli Matteo, Nencini Elia and Spinelli Stefano from the Emergency Medicine Unit, Azienda Ospedaliera Universitaria Pisana; Antognoli Rachele, Calsolorio Valeria and Paterni Simone from the...
Geriatrics Unit, Azienda Ospedaliera Universitaria Pisana; Baldassarri Rubia, Bertini Pietro, Brizzi Giulia, Corradi Francesco, Della Rocca Alessandra, Guarracino Fabio, Malacarne Paolo, Monfroni Marco and Piagnani Chiara from the Department of Anaesthesia and Intensive Care, Azienda Ospedaliera Universitaria Pisana; Celi Alessandro from the Respiratory Unit, Azienda Ospedaliera Universitaria Pisana; Cinotti Francesco, Park Naria and Cipriano Alessandro from the Emergency Department, Azienda Ospedaliera Universitaria Pisana; Colangelo Luciano, Sonato Chiara and Galfo Valentina from the Infectious Diseases Unit, Azienda Ospedaliera Universitaria Pisana; Monica Uliana from the Forth Medical Unit, Internal Medicine Department, Azienda Ospedaliera Universitaria Pisana; Ruberti Francesca, Forotti Giovanna and Sciuto Maria from the Fifth Medical Unit, Internal Medicine Department, Azienda Ospedaliera Universitaria Pisana; Pisa (Italy).

**Funding:** none.

**Conflict of interest/Competing interests:** The authors declare that they have no interest.

**Ethics approval:** The study complied with the edicts of the 1975 Declaration of Helsinki. All experimental protocols were approved by Comitato Etico Area Vasta Nord Ovest (CEAVNO).

**Consent to participate:** Informed consent was obtained from all individual participants or their legal guardians included in the study.

**Consent to publication:** Informed consent for data publications was obtained from all individual participants or their legal guardians included in the study.

**Availability of data and material:** Data are available upon reasonable request.

**Code availability:** not applicable

**Author contribution:** AM: study design, data analysis and interpretation, manuscript write-up; GG: data analysis and interpretation, manuscript write-up; MF, GT, NRP, MAD, LG, GB, FF, LC, MS, FM, SDM, FM: data collection, manuscript editing; AV and SM: study design and supervision, data collection, analysis and interpretation, manuscript editing and critical revision. All authors read and approved the final submitted version of the manuscript. AM, AV and SM are the guarantors of this work and, as such, have full access to all the data in the study and take responsibility for the integrity of the data and the accuracy of the data analysis.

**Abbreviations**

**COVID-19:** Coronavirus Disease 2019

**CRP:** C-reactive protein

**CVD:** Cardiovascular disease

**hsTnT:** high-sensitivity troponin T

**SARS-CoV-2:** Severe Acute Respiratory Syndrome - Coronavirus – 2
**LMWH: Light Molecular Weight Heparin**

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### Tables

**Table 1.** Differences in clinical characteristics at admission according to survival status

| Variable                           | N      | Survived (N = 202) | Died (N = 64) | P-Value |
|------------------------------------|--------|--------------------|---------------|---------|
| **In-hospital mortality**          |        |                    |               |         |
| Age, years                         | 266    | 63 (15)            | 79.3 (10)     | <0.001  |
| Male sex, n (%)                    | 266    | 130 (64.4)         | 50 (78.1)     | 0.040   |
| Current smoking, %                 | 243    | 9 (5)              | 3 (5)         | 0.925   |
| Systolic blood pressure, mmHg      | 240    | 125 (22)           | 133 (20)      | 0.011   |
| Diastolic blood pressure, mmHg     | 240    | 71 (15)            | 76 (13)       | 0.011   |
| Heart rate, bpm                    | 242    | 90 (77-110)        | 85 (75-95)    | 0.057   |
| **PaO**2*                          | 240    | 71 (60-86)         | 59 (41-73)    | <0.001  |
| **PaO**2/FiO**2*                   | 254    | 290 (233-362)      | 200 (150-276) | <0.001  |
| Glucose, mg/dL                     | 252    | 132 (59)           | 145 (51)      | 0.145   |
| Haemoglobin, g/dl*                 | 258    | 13.5 (12-14.7)     | 13 (11.6-14.2)| 0.467   |
| Leukocyte count, cells/μL*         | 259    | 6580 (4960-9140)   | 6640 (5150-8940)| 0.860  |
| Lymphocyte count, cells/μl*        | 257    | 1000 (680-1317)    | 750 (508-1110)| 0.001   |
| Neutrophils count, cells/μl*       | 256    | 4700 (3220-6940)   | 5700 (3790-7320)| 0.164  |
| Platelets count, cells/μl*         | 259    | 195 (149-247)      | 172 (125-215) | 0.021   |
| INR*                               | 248    | 1.22 (1.15-1.32)   | 1.23 (1.17-1.50)| 0.183  |
| Total Bilirubin, mg/dL*            | 249    | 0.50 (0.36-0.74)   | 0.55 (0.38-0.78)| 0.509  |
| Aspartate transaminase, U/L*       | 237    | 32 (23-46)         | 39 (26.5-54.5)| 0.025   |
| Alanine transaminase, U/L*         | 253    | 27 (19-48)         | 25 (16-45)    | 0.327   |
| Creatinine, mg/dL*                 | 255    | 0.94 (0.78-1.14)   | 1.3 (1.05-1.91)| <0.001  |
| Sodium, mEq/L*                     | 258    | 138 (136-140)      | 138 (135-142) | 0.565   |
| Potassium, mEq/L*                  | 253    | 3.98 (3.73-4.28)   | 4.15 (3.78-4.64)| 0.011  |
| D-dimer, mg/dL*                    | 84     | 0.45 (0.22-1.04)   | 0.74 (0.54-2.86)| 0.043  |
| Lactate Dehydrogenase, U/L*        | 139    | 304 (216-427)      | 385 (255-502) | 0.077   |
| Creatine phosphokinase, U/L*       | 119    | 85 (48-166)        | 218 (80-465)  | 0.012   |
| High sensitivity troponin T, ng/L* | 200    | 12 (7-23)          | 59 (33-119)   | <0.001  |
| Procalcitonin, ng/mL*              | 222    | 0.11 (0.06-0.26)   | 0.21 (0.12-0.69)| <0.001  |
| C-reactive protein, mg/dl*         | 247    | 6.26 (2.60-13.60)  | 8.68 (5.36-17.23)| 0.008  |

**COMORBIDITIES**

| Chronic obstructive pulmonary disease, n (%) | 266 | 17 (8.42) | 19 (29.69) | <0.001 |
| Asthma, n (%)                                | 265 | 16 (7.92) | 2 (3.17)  | 0.191  |
| Diabetes mellitus, n (%)                     | 266 | 36 (17.82) | 15 (23.44) | 0.320  |
| Arterial hypertension, n (%)                 | 266 | 81 (40.10) | 41 (64.06) | <0.001 |
| History of cardiovascular disease, n (%)     | 266 | 52 (25.74) | 34 (53.13) | <0.001 |
| History of cerebrovascular disease, n (%)    | 266 | 14 (7)    | 13 (20)   | 0.002  |
| Diagnosis of dementia, n (%)                 | 266 | 13 (6)    | 7 (11)    | 0.234  |
| Diagnosis of Cancer, n (%)                   | 266 | 28 (14)   | 11 (17)   | 0.512  |
| Haematological malignancy, n (%)             | 266 | 3 (1.49)  | 3 (4.69)  | 0.133  |
| Chronic kidney disease, n (%)                | 266 | 11 (5)    | 12 (19)   | 0.001  |
| Dialysis, n (%)                              | 266 | 3 (1)     | 2 (3)     | 0.400  |
| Chronic liver disease, n (%)                 | 266 | 4 (2)     | 7 (11)    | 0.002  |

P-values are derived from independent samples Student’s T Test or the Mann Whitney test for continuous variables and the chi squared test for categorical variables; *Median and interquartile range
Table 2. HRs for each outcome for 1 SD increase in hsTnT levels

| Outcome            | HR     | 95% CI     | P-value |
|--------------------|--------|------------|---------|
| All cause death    |        |            |         |
| Unadjusted         | 1.2    | 1.07-1.34  | 0.002   |
| Adjusted           | 2.28   | 1.41-3.68  | 0.001   |
| ARDS               |        |            |         |
| Unadjusted         | 2.12   | 0.824-5.45 | 0.119   |
| Adjusted           | 1.58   | 0.659-3.8  | 0.305   |
| ICU admission      |        |            |         |
| Unadjusted         | 1.11   | 0.838-1.48 | 0.462   |
| Adjusted           | 1.19   | 0.886-1.6  | 0.219   |
| Ventilatory support* |        |            |         |
| Unadjusted         | 1.19   | 0.863-1.64 | 0.289   |
| Adjusted           | 1.01   | 0.532-1.91 | 0.983   |

Adjusted associations are controlled for the effect of age, gender, cardiovascular disease and admission creatinine. *Ventilatory support = requirement for invasive and non-invasive ventilation

Table 3. Unadjusted and multi-adjusted associations of 1 SD increase of C-reactive protein and DD with the main study outcomes
| Outcome                      | CRP               | D-dimer          |
|------------------------------|-------------------|------------------|
|                              | Univariable       | Multivariable    | Univariable       | Multivariable    |
|                              | HR/OR 95% CI     | P-value          | HR/OR 95% CI     | P-value          |
|                              |                   |                  |                   |                  |
| Death*                       | 1.01(0.993-1.02)  | 0.337            | 1.026(0.996-1.057)| 0.095            |
|                              | 1.09 (1.03-1.15)  | 0.001            | 1.11(1.03-1.185)  | 0.005            |
| ARDS¥                        | 1.03(1.003-1.06)  | 0.032            | 1.026(0.995-1.06) | 0.107            |
|                              | 1.32(0.925-1.88)  | 0.127            | 1.28(0.896-1.832) | 0.174            |
| ICU transfer¥                | 1.03(1.002-1.06)  | 0.038            | 1.04(1.01-1.08)   | 0.02             |
|                              | 1.03(0.936-1.14)  | 0.531            | 1.02(0.925-1.13)  | 0.664            |
| Ventilatory support ¥        | 1.04(1.01-1.07)   | 0.012            | 1.05(1.02-1.085)  | 0.005            |
|                              | 1.04(0.939-1.15)  | 0.473            | 1.02(0.918-1.126) | 0.747            |

* Results obtained from Cox regression. ¥ results obtained from logistic regression. Adjusted associations are controlled for the effect of age, gender, cardiovascular disease and admission creatinine.

**Table 4.** Differences in the clinical characteristics of the population according to median levels of admission hsTnT
| Variable                      | N  | Admission hs Troponin | P-Value   |
|-------------------------------|----|------------------------|-----------|
|                              |    | Below the median       | Above the median |
| Age, years                   | 200| 59 (13)                | 78 (11)    | <0.001 |
| Male sex, n (%)              | 200| 39 (38)                | 30 (31)    | 0.302  |
| Current smoking, %           | 187| 4 (4)                  | 2 (2)      | 0.494  |
| PaO2*                        | 182| 77 (63-87)             | 61 (50-74) | <0.001 |
| PaO2/FiO2*                   | 194| 300 (240-371)          | 257 (195-324) | 0.001 |
| Glucose, mg/dL               | 194| 121 (34)               | 148 (72)   | <0.001 |
| Haemoglobin, g/dl*           | 195| 14 (12.7-14.9)         | 12.9 (11.4-14.6) | 0.006  |
| Leukocyte count, cells/μl*   | 197| 6380 (4780-8330)       | 6690 (5240-10060) | 0.079  |
| Lymphocyte count, cells/μl*  | 197| 1100 (690-1330)        | 835 (540-1116) | <0.001 |
| Neutrophils count, cells/μl* | 194| 4500 (3120-6240)       | 5600 (3447-7665) | 0.036  |
| Platelets count, cells/μl*   | 197| 196 (149-248)          | 175 (129-238) | 0.126  |
| INR*                         | 193| 1.21 (1.13-1.28)       | 1.25 (1.18-1.48) | 0.002  |
| Total Bilirubin, mg/dL*      | 194| 0.44 (0.32-0.63)       | 0.56 (0.38-0.76) | 0.038  |
| Aspartate transaminase, U/L* | 186| 32 (24-50)             | 34 (25-51)  | 0.737  |
| Alanine transaminase, U/L*   | 195| 33 (23-49)             | 22 (14-36)  | <0.001 |
| Creatinine, mg/dL*           | 195| 0.89 (0.78-1.03)       | 1.24 (0.96-1.83) | <0.001 |
| Sodium, mEq/L*               | 196| 138 (135-140)          | 138(134-141) | 0.467  |
| Potassium, mEq/L*            | 192| 3.98 (3.73-4.25)       | 4.13 (3.84-4.47) | 0.007  |
| D-dimer, mg/L*               | 75 | 0.32 (0.21-0.69)       | 0.67 (0.35-1.10) | 0.021  |
| Lactate Dehydrogenase, U/L*  | 109| 283 (216-405)          | 357 (251-467) | 0.039  |
| Creatine phosphokinase, U/L* | 100| 76 (50-213)            | 92 (56-207)  | 0.587  |
| High sensitivity troponin T, ng/L* | 200 | 9 (6-12)               | 41 (26-85)  | <0.001 |
| Procalcitonin, ng/mL*        | 177| 0.08 (0.05-0.16)       | 0.19 (0.10-0.63) | <0.001 |
| C-reactive protein, mg/dl*   | 193| 4.31 (1.79-10.67)      | 8.88 (4.87-18.49) | <0.001 |

P-values are derived from independent samples Student’s T Test or the Mann Whitney test for continuous variables and the chi squared test for categorical variables; *Median and interquartile range

| Variable                                | N  | Univariable          | Multivariable          |
|-----------------------------------------|----|----------------------|------------------------|
|                                         |    | *Standardized        | P-value                |
|                                         |    | Coefficient          |                        |
| Age                                     | 200| 0.607                | <0.001                 |
| Sex                                     |    | -0.048               | 0.500                  |
| History of Cardiovascular disease       | 200| 0.319                | <0.001                 |
| Admission creatinine                    | 200| 0.321                | <0.001                 |
| Admission CRP                           | 200| 0.338                | <0.001                 |
| Admission PaO2                          | 200| -0.179               | 0.016                  |
| Admission D-Dimer (n=84)                | 200| 0.303                | 0.008                  |

Table 5. Determinants of admission hsTnT levels by linear regression analysis

* coefficients correspond to 1-unit change in natural log transformed hs Troponin.