Multi-organ assessment in mainly non-hospitalized individuals after SARS-CoV-2 infection: The Hamburg City Health Study COVID programme

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See the editorial comment for this article 'Post-COVID-19 illness trajectory in community patients: mostly reassuring results', by Colin Berry and Hannah K. Bayes, https://doi.org/10.1093/eurheartj/ehac057.

Aims	Long-term sequelae may occur after SARS-CoV-2 infection. We comprehensively assessed organ-specific functions in individuals after mild to moderate SARS-CoV-2 infection compared with controls from the general population.

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Methods and results	Four hundred and forty-three mainly non-hospitalized individuals were examined in median 9.6 months after the first positive SARS-CoV-2 test and matched for age, sex, and education with 1328 controls from a population-based German cohort. We assessed pulmonary, cardiac, vascular, renal, and neurological status, as well as patient-related outcomes. Bodyplethysmography documented mildly lower total lung volume (regression coefficient −3.24, adjusted $P = 0.014$) and higher specific airway resistance (regression coefficient 8.11, adjusted $P = 0.001$) after SARS-CoV-2 infection. Cardiac assessment revealed slightly lower measures of left (regression coefficient for left ventricular ejection fraction on transthoracic echocardiography −0.93, adjusted $P = 0.015$) and right ventricular function and higher concentrations of cardiac biomarkers (factor 1.14 for high-sensitivity troponin, 1.41 for N-terminal pro-B-type natriuretic peptide, adjusted $P \leq 0.01$) in post-SARS-CoV-2 patients compared with matched controls, but no significant differences in cardiac magnetic resonance imaging findings. Sonographically non-compressible femoral veins, suggesting deep vein thrombosis, were substantially more frequent after SARS-CoV-2 infection (odds ratio 2.68, adjusted $P < 0.001$). Glomerular filtration rate (regression coefficient −2.35, adjusted $P = 0.019$) was lower in post-SARS-CoV-2 cases. Relative brain volume, prevalence of cerebral microbleeds, and infarct residuals were similar, while the mean cortical thickness was higher in post-SARS-CoV-2 cases. Cognitive function was not impaired. Similarly, patient-related outcomes did not differ.
Conclusion	Subjects who apparently recovered from mild to moderate SARS-CoV-2 infection show signs of subclinical multi-organ affection related to pulmonary, cardiac, thrombotic, and renal function without signs of structural brain damage, neurocognitive, or quality-of-life impairment. Respective screening may guide further patient management.
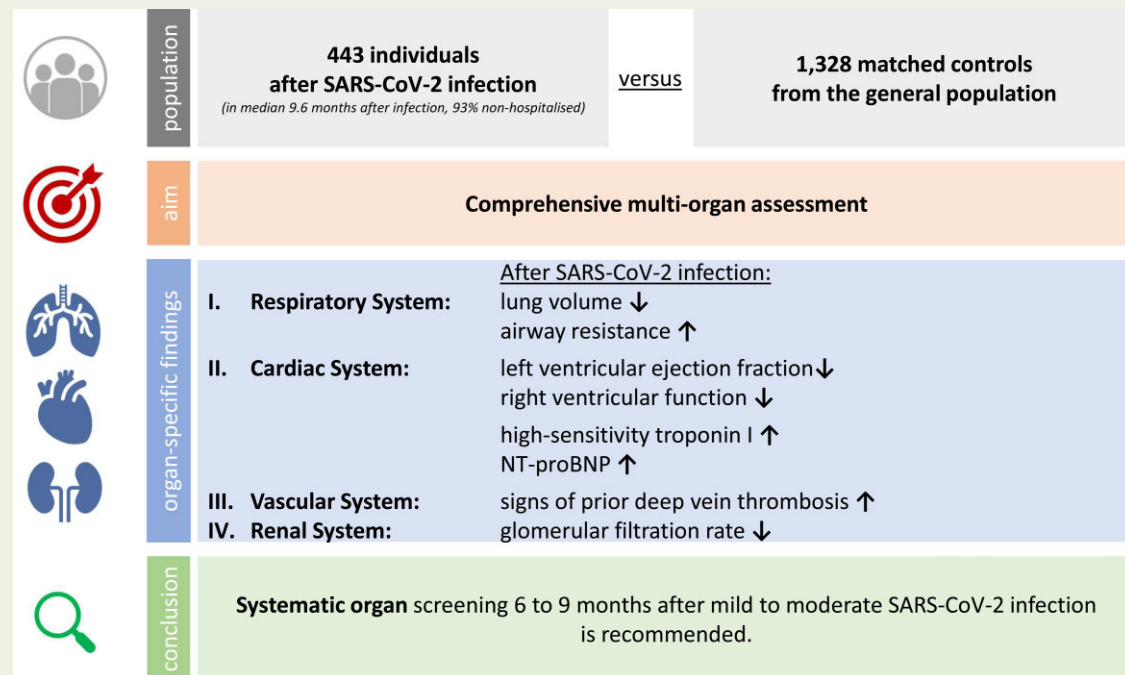
Graphical Abstract The key question is: How does a mild to moderate course of SARS-CoV-2 infection in mainly non-hospitalized individuals impact intermediate-term organ-specific functions in comparison to the general population? The key findings are (i) a mild to moderate course of SARS-CoV-2 infection is associated with subsequent signs of subclinical multi-organ affection; (ii) associations mainly affect the pulmonary, cardiac, coagulation, and renal system; and (iii) no systematic associations with structural brain damage, neurocognition, or quality of life were observed. The take-home message is systematic screening of multi-organ function even after mild to moderate SARS-CoV-2 infection is recommended to identify individuals at risk and initiate appropriate preventive therapies.

Introduction

As of December 2021, the coronavirus disease 2019 (COVID-19) pandemic has resulted in over 260 million confirmed positive cases of severe acute respiratory syndrome coronavirus type 2 (SARS-CoV-2) globally.[1] In Europe, 3.6% of infected patients required intensive care, 90% recovered in an ambulatory setting.[2,3]

Severe COVID-19 may affect multiple organ systems during the acute phase of the disease. The impairment of pulmonary, cardiac, and renal function as well as thromboembolism have been described in severely ill and deceased patients.[4–6] Furthermore, serious neurological complications, including cerebrovascular events, encephalopathy, and encephalitis, have been reported.[7]

While single reports describe persisting symptoms of pulmonary, renal, cardiac, or vascular dysfunction as well as fatigue or depression mostly derived from hospitalized cohorts,[8] a systematic investigation especially of subclinical changes of multi-organ structure and function is of particular interest in the current pandemic context.

With this study, we aimed to comprehensively determine the intermediate-term impact of a mild to moderate course of COVID-19 on organ-specific function. A COVID-19 disease course was classified as mild to moderate when not requiring intensive care unit treatment. We assessed multi-organ function by deep phenotyping in patients ~9 months after COVID-19 recovery in direct comparison with age-, sex-, and education-matched subjects from a population-based cohort study.

Methods

Study design

We report a cross-sectional study to compare the organ-specific functions and structures assessed in a cohort of subjects with prior mild to moderate SARS-CoV-2 infection with matched subjects from an ongoing, prospective, population-based cohort study.

Study population

The post-SARS-CoV-2 cohort consists of subjects living in the metropolitan area of Hamburg, Germany, with a laboratory-confirmed positive polymerase chain reaction (PCR) test for SARS-CoV-2, obtained between 1 March and 31 December 2020 at least 4 months prior to study enrollment. In correspondence to the age structure of the population-based Hamburg City Health Study (HCHS), which served as the control cohort, participants had to be 45–74 years old at the time of recruitment.[9]

Eligible participants after SARS-CoV-2 infection were invited to participate after they were automatically identified through the clinical information system of the University Medical Center Hamburg-Eppendorf (UKE), which performed SARS-CoV-2 diagnostics for the public health service. In addition, eligible inhabitants living in the metropolitan area were invited to participate via public announcement. At the time of study enrollment, all participants provided written informed consent. The local ethics committee (State of Hamburg Chamber of Medical Practitioners, PV5131) did not have objections against the recruitment of post-SARS-CoV-2 individuals and the study protocol as an extension of the HCHS, and the study was conducted in compliance with the Declaration of Helsinki. Categorization of disease severity was based on retrospectively documented, self-reported symptom severity at the time of SARS-CoV-2 infection and consisted of four groups: asymptomatic, mild symptoms, moderate symptoms without the need for hospitalization, and moderate symptoms with the need for hospitalization.

Matched controls were selected from the HCHS (ClinicalTrials.gov identifier NCT03934957), a large, ongoing, prospective, population-based cohort study conducted at the University Medical Center Hamburg-Eppendorf (UKE), Germany, since 2016 to gain knowledge about major chronic diseases.[9] As of July 2021, data from ~15 000 participants were available.

Hamburg City Health Study participants were only eligible as controls for this work if they had been enrolled prior to being at risk of infection with SARS-CoV-2 and during the same seasonal period as the patients after SARS-CoV-2 infection. The inclusion periods were November–April 2016–19.

Outcomes and data collection

Both patients after SARS-CoV-2 infection and matched controls underwent the same standardized 7 h assessment at the Epidemiological Study Center of the Population Health Research Department of the University Medical Center Hamburg-Eppendorf (UKE), which has been described elsewhere.[9]

Pulmonary phenotyping

Lung function was assessed by trained technicians performing body-plethysmography including forced spirometry (MasterScreen with SentrySuite, Vyaire Medical GmbH, Hoechberg, Germany) according to international guidelines.[10,11] Forced spirometry manoeuvres were performed to obtain forced expiratory volume in 1 s (FEV_1) and forced vital capacity (FVC). Resistance manoeuvres were performed a minimum of five times, all other manoeuvres were performed a minimum of three times. Bodyplethysmography was conducted to obtain the specific airway resistance (sRaw) and static lung volumes. Lung volumes were obtained by means of functional residual capacity (FRC) measured during gentle breathing against a shutter as well as expiratory reserve volume and vital capacity (VC). Residual volume was calculated by subtracting expiratory reserve volume from FRC, and total lung capacity (TLC) was calculated by the addition of residual volume and VC. For quality control, a multidimensional quality assurance concept was established, including regular calibration of the equipment, regular training and certification of the technicians, and a combination of algorithm-based data processing and visual control by a trained examiner. We used FEV_1, the ratio of FEV_1 and FVC, and sRaw to assess airway function and FVC, VC, TLC, and FRC to assess lung volumes.

Cardiac phenotyping

Transthoracic echocardiography was performed and analysed by cardiologists and sonographers (technicians) (Siemens Acuson SC2000 Prime, Siemens Healthineers, Erlangen, Germany). Technical details on transthoracic echocardiography data collection and quality assurance have been described elsewhere.[12] Left ventricular ejection fraction was assessed using dicks summation in two-dimensional (2D) and 3D loops. Mitral inflow pattern was assessed in the apical four-chamber view by placing pulsed-wave (PW) Doppler sample volume between mitral leaflet tips. Pulsed-wave tissue Doppler imaging (TDI) e′ velocity was measured in the apical four-chamber view by placing the sample volume at the lateral and septal basal regions of the mitral annulus. To assess the left ventricular diastolic function, E/e' was computed as the mean ratio of E-wave velocity divided by e′ lateral and septal velocity. The right ventricular longitudinal function was quantified by the systolic excursion of the tricuspid annulus. The tricuspid annular plane systolic excursion (TAPSE) was obtained by M-mode echocardiography in the apical four-chamber view, reflecting right ventricular ejection fraction. The peak tricuspid regurgitation velocity, representing right ventricular filling pressure, was measured in the apical four-chamber view by continuous-wave Doppler.

Cardiovascular magnetic resonance imaging (MRI) was performed on a 3 T scanner (MAGNETOM™ Skyra, Siemens Healthineers,

Erlangen, Germany). The protocol included standard electrocardiogram (ECG)-triggered steady-state free-precession cine series in short axis for biventricular estimation of ejection fraction using the commercially available software 'CVi42' (Circle Cardiovascular Imaging Inc., Calgary, Alberta, Canada). Endo- and epicardial borders of the left and right ventricles were manually delineated on short-axis cine images as recommended.[13,14] Pre-contrast (native) T1 mapping was performed using a commercially available modified Look-Locker inversion recovery (MOLLI) sequence with a 5b(3b)3b scheme, T2 mapping using a T2-prepared single-shot fast low-angle shot (FLASH) sequence. The myocardium was automatically divided into 16 segments according to the American Heart Association (AHA) segment model and the mid-anteroseptal myocardial AHA segment VIII providing the most robust measurements of myocardial T1 and T2 and predominantly used to assess diffuse myocardial tissue alterations in clinical routine was selected for the analyses.[15] Late gadolinium enhancement (LGE) imaging was performed (phase-sensitive inversion recovery, PSIR) in short-axis orientation covering the entire heart and in two-, three-, and four-chamber views. Late gadolinium enhancement was evaluated qualitatively.[14]

Electrocardiogram at rest was recorded with a standard digital 12-lead ECG (Schiller, Baar, Switzerland) and included a 2 min rhythm strip.

Vascular phenotyping

Study participants underwent a compression ultrasound of the left and right common femoral vein using a Siemens SC2000 with a 7.5 MHz linear probe (Siemens Healthcare GmbH, Erlangen, Germany). Trained technicians performed the ultrasound examination following a standard protocol, and a regular quality assurance of research data was conducted via statistical measures as well as random and risk-based controls by specialized physicians. A static grey-scale image (brightness mode) of the common femoral vein was captured at the level of the saphenofemoral junction both with and without external mechanical compression.

A vascular specialist reviewed images from all post-SARS-CoV-2 participants ($n = 443$) and from a random sample of the matched controls ($n = 400$). If the common femoral vein was not fully compressible, the corresponding image was independently reviewed by a second vascular specialist. Only in the case of agreement between both investigators, veins were declared non-compressible.

Carotid atherosclerosis and flow velocities were evaluated by ultrasound using a Siemens SC2000® with a 7.5 MHz linear array transducer. Measurements of ultrasound parameters were performed as reported previously.[16] In brief, carotid intima-media thickness (CIMT) was measured three times in the B-mode in a longitudinal view of the left and right common carotid artery (CCA) >1 cm proximal to the carotid bulbus, and mean values were calculated. The presence of atherosclerotic plaques was recorded. The peak systolic flow velocity (PSV) of the CCA and internal carotid artery (ICA) were measured by PW Doppler.

Neuroimaging phenotyping

Images were acquired using a 3 T scanner (MAGNETOM™Skyra, Siemens Healthineers, Erlangen, Germany). For 3D T1-weighted anatomical images, rapid acquisition gradient-echo sequence (MPRAGE) was used with the following sequence parameters: repetition time (TR) = 2500 ms, echo time (TE) = 2.12 ms, 256 axial slices, slice thickness (ST) = 0.94 mm, and in-plane resolution (IPR) = 0.83 × 0.83 mm^2. Three-dimensional T2-weighted fluid attenuated inversion recovery (FLAIR) images were acquired with the following sequence parameters: TR = 4700 ms, TE = 392 ms, 192 axial slices, ST = 0.9 mm, and IPR = 0.75 × 0.75 mm. Time of flight (TOF) magnetic resonance angiography was performed using TR = 25 ms, TE = 3.6 ms,

64 axial slices, ST = 0.5 mm, IPR = 0.7 × 0.7 mm, GRAPPA 2, flip angle (FA) = 25°, and bandwith 133 Hz/Px positioned transversal over the circle of Willis.

Cortical thickness, brain volume, and intracranial volumes were quantified based on T1-weighted imaging data using the standardized FreeSurfer processing pipeline (version 6.0.1).[17] Cortical thickness was averaged over both hemispheres using standard outputs of the FreeSurfer pipeline as described before.[18] Relative brain volume was calculated by normalizing brain volume for estimated intracranial volume to account for variation in head size.[19]

Laboratory parameters

Routine laboratory parameters such as leucocytes, creatinine, sodium, potassium, glucose, glycated haemoglobin, and high-sensitivity C-reactive protein (hs-CRP) were immediately measured in fresh samples. Glomerular filtration rate (GFR) was calculated using the Chronic Kidney Disease Epidemiology Collaboration (CKD-EPI) formula for creatinine.[20]

The following biomarkers were assayed in a blinded fashion using serum samples stored at −80°C in a dedicated biobank: (i) cardiac troponin I was measured using a high-sensitivity cardiac troponin I assay (ARCHITECT High Sensitive STAT Troponin I, Abbott Laboratories, Abbott Park, IL, USA) with a limit of detection of 1.9 ng/L;[21] (ii) N-terminal pro-B-type natriuretic peptide (NT-proBNP) was measured using an immunoassay (Alere NT-proBNP for ARCHITECT, Abbott Diagnostics) with measurement ranges between 8.2 and 35 000 ng/L; and (iii) cystatin C was determined by the Abbott Cystatin C ARCHITECT c8000 assay with measurement ranges between 0.05 and 8.33 mg/L and a limit of detection of 0.05 mg/L. Coagulation parameters, including D-dimers, fibrinogen, prothrombin time according to Quick's method, as well as activated partial thromboplastin time, were measured in sodium citrated plasma samples stored at −80°C on a Atellica COAG 360 System analyzer (Siemens Healthineers, Erlangen, Germany) in a blinded fashion.

SARS-CoV-2 serology

For the detection of past SARS-CoV-2 infection and the qualitative serologic assay targeting the viral nucleocapsid protein, the Elecsys® Anti-SARS-CoV2 (Roche, Mannheim, Germany; analyzer: cobas e411) assay was used. To analyse neutralizing antibody responses, the quantitative assay Liason® SARS-CoV-2 S1/S2 IgG (Diasorine, Saluggia, Italy; analyzer: Liaison XL) or the quantitative assay Elecsys® Anti-SARS-CoV-2 S (Roche; analyzer: cobas e411) were used. Quantitative values were normalized to the first World Health Organization standard for anti-spike antibodies (NIBSC code: 20/136; version 2.0, as of 17 December 2020) and presented as binding antibody unit (BAU)/mL [conversion factor 4.961 (Diasorin) and 0.9750 (Roche)]. All serological tests were performed according to the manufacturer's recommendations in an accredited laboratory (UKE Hamburg).

Cognitive function assessment

Mini-Mental State Examination (MMSE) was applied as a screening instrument for cognitive impairment and dementia.

Patient-related outcomes and quality-of-life assessment

Patient Health Questionnaire-9 and -15 (PHQ-9, PHQ-15), Generalized Anxiety Disorder Assessment (GAD-7), European Quality of Life 5 Dimensions (EQ-5D), as well as visual analogue scale (EQ-5D VAS) were assessed as central parameters of patient-reported outcomes (PRO).[9]

Main outcomes

Main outcomes per organ system were discussed and pre-defined among experts in a focus group to examine pre-defined hypotheses (Table 1).

Subgroup analyses

Subgroup analyses were performed to analyse potential differences between (i) severity of COVID-19 (asymptomatic or mild symptoms vs. moderate symptoms) and (ii) recruitment route of subjects with SARS-CoV-2 infection with respect to organ-specific main

Table 1 Main outcomes per organ system

Biological parameter	Corresponding surrogate
Respiratory system	
Lung volume	TLC in bodyplethysmography, % of predicted value
Airway function	sRaw in bodyplethysmography, % of predicted value
Cardiac system	
LV function	LVEF in TTE, %
RV function	TAPSE in TTE, mm
Focal myocardial fibrosis	Presence of late gadolinium enhancement on CMR
Myocardial damage	High-sensitivity cardiac troponin I, ng/L
Haemodynamic stress	NT-proBNP, ng/L
Vascular system	
Deep vein thrombosis	At least one non-compressible common femoral vein on venous ultrasound
Atherosclerosis	Presence of carotid plaques on carotid ultrasound
Renal system	
Renal function	GFR, mL/min/1.73 m2
	Potassium, mmol/L
Metabolic and inflammatory system	
Metabolics	Glycated haemoglobin, %
Inflammation	High-sensitivity CRP, mg/L
Neurological system	
Vascular brain damage	Presence of cerebral micro bleeds on brain MRI
	Presence of infarct residuals on brain MRI
Brain atrophy	Relative brain volume, %
	Cortical thickness, mm
Neurocognition	
Screening for cognitive impairment	Mini-Mental State Examination
Psychosocial outcomes	
Severity of depression	PHQ-9
Generalized anxiety	GAD-7

Main outcomes selected by clinical relevance.
CMR, cardiac magnetic resonance; CRP, C-reactive protein; GAD-7, Generalized Anxiety Disorder Assessment; GFR, glomerular filtration rate; LV, left ventricular; LVEF, left ventricular ejection fraction, MRI, magnetic resonance imaging; PHQ-9, Patient Health Questionnaire-9; RV, right ventricular; sRaw, specific airway resistance; TAPSE, tricuspid annular plane systolic excursion; TLC, total lung capacity; TTE, transthoracic echocardiogram.

outcomes. These analyses are explorative, given the resulting small sample sizes.

Statistics

Matching

The pool of possible controls was drawn from the cohort of the first 10 000 HCHS participants omitting seasonal effects and undetected COVID-19 infections. Matching was conducted using *k*-nearest neighbours without the re-use of controls and a variable ratio of cases and controls. Controls were matched for age at inclusion using a caliper of ± 2 years, dichotomized sex, the three categorical educational status according to the International Standard Classification of Education, and the availability of MRI examinations of the heart and the brain.[22] More information on the setup and quality of the matching can be found in the Supplementary material online. Matching groups were clustered by age, sex, and educational status of patients after SARS-CoV-2 infection leading to an *n:m* matching and more variability between clusters.

Imputation of missing data

Multivariate predictive mean imputation by chained equations with 100 imputations and 10 iterations was used to impute missing data.[23] Variables used for imputation included the matching cluster, age, sex, educational status as well as relevant comorbidities, and all variables used to analyse the hypotheses relevant to the medical speciality. Imputations were performed separately for patients after SARS-CoV-2 infection and controls as well as for each medical speciality. Information on specific handling of PRO can be found in the Supplementary material online.

Statistical analysis

In the descriptive analysis, continuous data are presented as the median and interquartile range (IQR), and categorical data as absolute numbers and percentage.

Continuous outcomes were analysed using multiple linear mixed models, binary outcomes using multiple logistic mixed regression, both with matching cluster as random intercept. Regression results are presented as a forest plot depicting the comparison of patients after SARS-CoV-2 infection and controls. Regression estimates are provided, i.e. betas for continuous variables and odds ratios (OR) for binary outcomes. Parameters with skewed distributions were log-transformed before being entered into regression analyses, then re-transformed to enable interpretation. These parameters were left ventricular end-diastolic volume, high-sensitivity cardiac troponin I, NT-proBNP, leucocytes, and anti-spike antibodies.

P-values and 95% confidence intervals (CI) for pre-defined main outcomes were corrected for multiple testing within each organ system using the Bonferroni method.

All analyses were performed in R version 4.0.3.

Results

Baseline characteristics

The baseline characteristics of the 1771 participants are outlined in Table 2. The median age was 55 (IQR 51, 60) years in 443 subjects after SARS-CoV-2 infection vs. 57 (IQR 52, 62) years in 1328 matched controls. Female sex was present in 52.6 vs. 54.1%, respectively. Participants after SARS-CoV-2 infection and matched controls were comparable regarding socio-demographic characteristics, vital signs, anthropometry, patient history, and risk

Table 2 Baseline characteristics

	Subjects after SARS-CoV-2 infection (n = 443)	Matched controls (n = 1328)
Demographics		
Age, years	55 (51, 60)	57 (52, 62)
Female sex	233 (52.6)	719 (54.1)
Education[a]		
Low	90 (20.3)	329 (24.8)
Medium	88 (19.9)	289 (21.8)
High	265 (59.8)	710 (53.5)
Employment		
Full time	246 (56.4)	661 (51.8)
Part time	129 (29.6)	303 (23.7)
Unemployed	61 (14.0)	312 (24.5)
COVID-19-specfic characteristics		
Self-reported disease severity at the time of infection*		
Asymptomatic	14 (3.2)	
Mild	253 (58.4)	
Moderate, no need for hospitalization	135 (31.2)	
Moderate, need for hospitalization, without ICU	31 (7.2)	
Detectable SARS-CoV-2 nucleocapsid antibody	411 (94.5)	
Neutralizing anti-spike antibodies, BAU/mL	189.83 (52.36, 536.21)	
Months between first positive SARS-CoV2 PCR test and study enrollment	9.58 (5.20, 10.60)	
Risk factors		
Hypertension	267 (61.5)	750 (58.6)
Diabetes mellitus	27 (6.4)	78 (6.3)
Dyslipidaemia	89 (20.9)	210 (16.8)
Smoking status		
Current	33 (7.5)	280 (21.2)
Former	163 (37.1)	556 (42.1)
Never	243 (55.4)	485 (36.7)
ESC score[b]	1.20 (0.60, 2.40)	1.20 (0.60, 2.40)
Vital signs and anthropometry		
Heart rate, b.p.m.	69 (61, 77)	68 (62, 76)
Systolic blood pressure, mmHg	137 (126, 150)	136 (125, 149)
Diastolic blood pressure, mmHg	87 (80, 94)	83 (76, 90)
BMI, kg/m^2	25.9 (23, 29)	25.7 (23.3, 29.0)
Medical history[c]		
Coronary artery disease	15 (3.4)	49 (3.8)
Myocardial infarction	7 (1.6)	24 (1.8)
Peripheral artery disease	13 (3.0)	24 (1.9)
Chronic kidney disease	49 (11.1)	126 (9.6)
Chronic lung disease	53 (12.0)	154 (11.8)
Neoplasia	43 (9.7)	151 (11.5)

Continuous variables are presented as median and interquartile range, and categorical variables are presented as absolute numbers and percentages.
Individuals after SARS-CoV-2 infection were mainly community patients as 93% were not hospitalized.
BAU, binding antibody unit; BMI, body mass index; ESC, European Society of Cardiology; ICU, intensive care unit; PCR, polymerase chain reaction.
[a]Educational status defined according to the International Standard Classification of Education (ISCED).
[b]ESC score calculated for a 6-year risk of fatal cardiovascular disease.
[c]Self-reported.
*P-value for association of disease severity with anti-spike antibody concentration in linear regression <0.001.

factors. In total, 92.8% of participants after SARS-CoV-2 infection were managed as outpatients, reporting none (3.2%), mild (58.4%), or moderate (31.2%) symptoms at the time of SARS-CoV-2 infection. Hospitalization without need for intensive care support was required in only 7.2% of all participants in the post-SARS-CoV-2 cohort. Participants after SARS-CoV-2

infection were enrolled at a median of 9.6 months after the first positive PCR test. In 94.5% of patients after SARS-CoV-2 infection, SARS-CoV-2 nucleocapsid antibodies were detectable. In median, these patients had a neutralizing anti-spike antibody concentration of 190 BAU/mL (IQR 52, 536). Anti-spike antibody concentrations were associated with perceived symptom severity ($P < 0.001$, Table 2) as well as age ($P < 0.001$) and body mass index ($P = 0.033$) but none of the other baseline characteristics (see Supplementary material online, Table S4). Baseline characteristics stratified by disease severity and recruitment route can be found in Supplementary material online Tables S5 and S6.

Organ-specific outcome measures

Respiratory system

Total lung capacity was lower in post-SARS-CoV-2 cases compared with controls [99.1% of the predicted value vs. 102.4% of the predicted value, regression coefficient −3.24 (95% CI −5.57, −0.91), adjusted $P = 0.014$]. At the same time, specific airway resistance, as assessed by sRaw, was significantly higher in patients after SARS-CoV-2 infection than matched controls [77.3% of the predicted value vs. 69.8% of the predicted value, regression coefficient 8.11 (95% CI 3.56, 12.65), adjusted $P = 0.001$, Table 3 and Figure 1].

Cardiac system

Left ventricular ejection fraction determined by both transthoracic echocardiography and cardiac MRI was numerically slightly decreased in patients after SARS-CoV-2 infection when compared with controls, while only the difference in left ventricular ejection fraction derived from transthoracic echocardiography remained statistically significant after adjustment for multiple testing [left ventricular ejection fraction on transthoracic echocardiography, 57.9 vs. 59.1%, regression coefficient −0.93 (95% CI −1.54, −0.32), adjusted $P = 0.015$]. Measures of left ventricular diastolic function, assessed by E/e' and left atrial volume, were not increased in subjects after SARS-CoV-2 infection. Right ventricular systolic function, quantified by TAPSE using transthoracic echocardiography, was significantly reduced in patients after SARS-CoV-2 infection [TAPSE, 23.0 vs. 23.9 mm, regression coefficient −0.72 (95% CI −1.24, −0.21), adjusted $P = 0.031$]. Additional right ventricular measures of systolic function such as right ventricular ejection fraction, assessed by cardiac MRI, and diastolic function such as peak tricuspid regurgitation velocity, showed no relevant intergroup changes.

There was a trend of more focal myocardial fibrosis, assessed by cardiac magnetic resonance LGE imaging, in participants after SARS-CoV-2 infection; however, LGE and myocardial T2 mapping (indicating myocardial oedema) did not differ significantly between patients after SARS-CoV-2 infection and matched controls. Furthermore, diffuse myocardial fibrosis was assessed by native myocardial T1 mapping and revealed comparable results in both groups, in particular participants after SARS-CoV-2 did not show any evidence for accelerated remodelling of the myocardium.

Both cardiac biomarkers, NT-proBNP [87.84 vs. 62.76 ng/L, multiplicative regression coefficient 1.41 (95% CI 1.29, 1.55), adjusted $P < 0.001$] and high-sensitivity cardiac troponin I [2.07 vs.

1.90 ng/L, multiplicative regression coefficient 1.14 (95% CI 1.05, 1.24), adjusted $P = 0.010$] were higher in patients after SARS-CoV-2 infection compared with matched controls. NT-proBNP concentrations ≥125 ng/L were more frequently observed in patients after SARS-CoV-2 infection than matched controls [33.2 vs. 18.2%, OR 2.39 (95% CI 1.82, 3.12), adjusted $P < 0.001$].

Whereas the corrected QTc interval was longer in patients after SARS-CoV-2 infection when compared with matched controls, no significant intergroup differences could be observed in further conduction abnormalities like PQ interval, QRS interval, atrioventricular blocks, or bundle branch blocks.

Vascular system

Sonographically non-compressible common femoral veins were found more frequently in participants after SARS-CoV-2 infection than in matched controls [43.2 vs. 22.2%, OR 2.68 (95% CI 1.77, 4.05), adjusted $P < 0.001$], independent of the side of measurement. Coagulation parameters did not differ between groups.

The mean CIMT assessed by carotid sonography was comparable between patients after SARS-CoV-2 infection and matched controls, but participants after SARS-CoV-2 infection more often presented atherosclerotic plaques [36.9 vs. 23.4%, OR 2.27 (95% CI 1.76, 2.93), adjusted $P < 0.001$]. In contrast, PSV was lower in participants after SARS-CoV-2 infection both in the ICA and CCA.

Renal system

Both markers of renal function, creatinine and cystatin C, were slightly elevated in patients after SARS-CoV-2 infection compared with matched controls. Ultimately, estimated GFR based on creatinine measurements was reduced in patients after SARS-CoV-2 infection by 2.35 mL/min/1.73 m2 (95% CI −4.04, −0.67, adjusted $P = 0.019$). Similarly, serum levels of sodium and potassium were slightly decreased in post-SARS-CoV-2 cases [for potassium, 3.8 vs. 3.9 mmol/L, regression coefficient −0.12 (95% CI −0.15, −0.09), adjusted $P < 0.001$].

Metabolic and inflammatory system

Whereas glucose concentrations were higher in patients after SARS-CoV-2 infection, glycated haemoglobin was similar in both groups [5.49 vs. 5.50%, regression coefficient −0.03 (95% CI −0.09, 0.03), adjusted $P = 0.60$]. Despite only minimal numerical differences, hs-CRP concentrations [0.09 vs. 0.11 mg/L, regression coefficient −0.05 (95% CI −0.09, −0.01), adjusted $P = 0.044$] were slightly lower in patients after SARS-CoV-2 infection.

Neurological system

Brain MRI was available in 188 post-SARS-CoV-2 cases and 483 matched controls. Numbers of cerebral microbleeds (3.7 vs. 4.5%, adjusted $P = 1.00$), infarct residuals (2.7 vs. 6.5%, adjusted $P = 0.27$), and stenoses of the intracerebral vessels (1.1 vs. 0.6%, $P = 0.34$) were comparable in patients after SARS-CoV-2 infection and matched controls. In quantitative analysis of brain structure, relative brain volume was comparable between groups [81.4 vs. 81.7%, regression coefficient −0.52 (95% CI −1.1, 0.06), adjusted $P = 0.31$], while patients after SARS-CoV-2 infection had higher mean cortical thickness [2.65 vs. 2.63 mm, regression coefficient

Table 3 Descriptive distributions and regression outcomes by organ systems

Biological parameter	Corresponding surrogate	Subjects after SARS-CoV-2 infection (n = 443)	Matched controls (n = 1328)	Regression estimate[a]	Nominal P-value for regression estimates	Adjusted P-values
Respiratory system						
Lung volume	TLC, % of predicted value	99.14 (90.37, 107.66)	102.48 (93.66, 112.03)	−3.24 (−5.57, −0.91)	0.0071	0.014
	VC, % of predicted value	100.54 (90.88, 110.86)	104.67 (95.08, 115.16)	−4.43 (−6.97, −1.9)	<0.001	
	FVC, % of predicted value	99.48 (91.53, 107.13)	101.16 (90.79, 110.71)	−1.21 (−3.42, 1.01)	0.29	
	FRC, % of predicted value	105.39 (93.15, 119.83)	107.69 (94.87, 122.44)	−1.67 (−5.03, 1.69)	0.33	
Airway function	sRaw, % of predicted value	77.26 (58.89, 99.49)	69.76 (51.89, 91.85)	8.11 (3.56, 12.65)	<0.001	0.001
	FEV_1, % of predicted value	98.35 (88.69, 107.17)	97.84 (87.55, 108.46)	0.05 (−2.26, 2.37)	0.97	
	FEV_1/ FVC ratio	0.78 (0.74, 0.81)	0.77 (0.73, 0.80)	0.01 (0, 0.02)	0.083	
	Raw, % of predicted value	74.15 (58.39, 95.56)	69.26 (51.09, 93.67)	4.57 (−0.63, 9.76)	0.087	
Cardiac system						
TTE						
LV function	LVEF TTE, %	57.88 (55.76, 60.27)	59.05 (55.46, 62.63)	−0.93 (−1.54, −0.32)	0.003	0.015
	E/e′ mean	6.66 (5.75, 7.74)	7.05 (6.03, 8.26)	−0.37 (−0.56, −0.18)	<0.001	
RV function	TAPSE, mm	22.95 (20.37, 26.20)	23.88 (21.29, 26.79)	−0.72 (−1.24, −0.21)	0.006	0.031
	TR Vmax, m/s	2.33 (2.12, 2.49)	2.29 (2.19, 2.49)	−0.04 (−0.1, 0.01)	0.14	
CMR						
Focal myocardial fibrosis	Focal late gadolinium enhancement	40 (36.0)	67 (29.0)	1.39 (0.86, 2.27)	0.18	0.92
Diffuse myocardial fibrosis	Native myocardial T1, ms	1172 (1145, 1196)	1182 (1158, 1202)	−10.83 (−20.65, −1.02)	0.032	
Myocardial oedema	Myocardial T2, ms	40 (38, 42)	40 (38, 42)	0.31 (−0.29, 0.92)	0.31	
LV function	LVEF CMR, %	68.11 (62.56, 72.97)	69.84 (64.93, 74.13)	−1.3 (−2.86, 0.26)	0.10	
LV hypertrophy	LV mass index, g/m^2	64.92 (57.96, 71.59)	64.26 (55.18, 73.66)	0.91 (−1.65, 3.47)	0.49	
LV volume	LVEDV, mL	125.227 (106.51, 142.75) N	124.38 (106.78, 148.64)	0.99 (0.95, 1.04)[b]	0.82	
Left atrium size	LAVI, mL/ml^2	20.20 (15.95, 25.45)	18.76 (15.07, 24.19)	0.76 (−0.33, 1.85)	0.17	
RV function	RVEF, %	57.23 (51.27, 63.97)	57.79 (51.94, 63.94)	−0.72 (−2.44, 1.0)	0.41	
Laboratory						
Myocardial damage	High-sensitivity troponin I, ng/L	2.07 (1.30, 3.23)	1.90 (1.25, 2.88)	1.14 (1.05, 1.24)[b]	0.002	0.010
Haemodynamic stress	NT-proBNP, ng/L	87.84 (55.65, 145.76)	62.76 (38.18, 106.39)	1.41 (1.29, 1.55)[b]	<0.001	<0.001
ECG						
Arrhythmia	Atrial fibrillation	13 (3.0)	45 (3.4)	0.94 (0.5, 1.77)	0.84	
Conduction disturbances	PQ interval, ms	160.84 (146.00, 179.33)	158.91 (145.74, 176.39)	2.68 (−0.22, 5.57)	0.070	
	QRS interval, ms	93.00 (85.61, 101.00)	92.00 (86.00, 99.69)	0.31 (−1.12, 1.74)	0.67	
	QTc time, ms	433.90 (414.39, 454.88)	429.64 (413.16, 444.91)	6.47 (3.57, 9.37)	<0.001	
	AV blocks	33 (7.5)	67 (5.1)	1.51 (0.98, 2.33)	0.064	
	Bundle branch blocks	41 (9.3)	104 (7.8)	1.24 (0.84, 1.83)	0.28	
Symptoms						
Dyspnoea	NYHA class					
	I	392 (88.5)	1211 (91.2)	Reference[c]		
	II	27 (6.1)	77 (5.8)	1.1 (0.68, 1.79)	0.69	
	III or IV	24 (5.4)	41 (3.1)	1.87 (1.06, 3.23)	0.032	

Continued

Table 3 Continued

Biological parameter		Corresponding surrogate	Subjects after SARS-CoV-2 infection ($n = 443$)	Matched controls ($n = 1328$)	Regression estimate[a]	Nominal *P*-value for regression estimates	Adjusted *P*-values
Vascular system							
Veins	Deep vein thrombosis	At least one non-compressible common femoral vein	99 (43.2)	49 (22.2)	2.68 (1.77, 4.05)	<0.001	<0.001
		Non-compressible common femoral vein right	70 (30.6)	28 (12.7)	3.06 (1.87, 4.99)	<0.001	
		Non-compressible deep femoral vein left	69 (30.1)	29 (13.1)	2.86 (1.76, 4.62)	<0.001	
Arteries	Atherosclerosis	Carotid intima-media thickness, mm	0.70 (0.62, 0.79)	0.71 (0.63, 0.81)	−0.01 (−0.02, 0.01)	0.35	
		Carotid plaques	164 (36.9)	311 (23.4)	2.27 (1.76, 2.93)	<0.001	<0.001
		Peak systolic velocity CCA, m/s	0.80 (0.71, 0.90)	0.86 (0.75, 0.99)	−0.07 (−0.09, −0.06)	<0.001	
		Peak systolic velocity ICA, m/s	0.64 (0.56, 0.75)	0.72 (0.62, 0.83)	−0.07 (−0.09, −0.05)	<0.001	
Laboratory	Coagulation	D-dimer, μg/L	0.24 (0.19, 0.37)	0.29 (0.20, 0.42)	−0.08 (−0.3, 0.14)	0.460	
		Fibrinogen, g/L	2.47 (2.06, 2.87)	2.41 (1.99, 2.78)	0.06 (−0.23, 0.35)	0.700	
		Quick prothrombin time, %	85.59 (75.59, 94.20)	79.89 (71.45, 87.31)	3.75 (−5.11, 12.62)	0.410	
		Activated partial thromboplastin time, s	34.59 (31.51, 38.12)	35.70 (33.08, 38.62)	−0.49 (−4.04, 3.05)	0.790	
Renal system							
	Renal function	GFR, mL/min/1.73 m^2	108.89 (86.22, 120.70)	109.14 (89.21, 120.44)	−2.35 (−4.04, −0.67)	0.006	0.019
		Creatinine, mg/dL	0.80 (0.73, 0.88)	0.78 (0.72, 0.86)	0.02 (0.01, 0.04)	0.002	
		Cystatin C, mg/L	0.97 (0.87, 1.07)	0.96 (0.86, 1.05)	0.02 (0, 0.03)	0.033	
		Potassium, nmol/L	3.8 (3.6, 3.9)	3.9 (3.7, 4.0)	−0.12 (−0.15, −0.09)	<0.001	<0.001
		Sodium, mmol/L	139 (138, 140)	139 (138, 141)	−0.28 (−0.5, −0.05)	0.016	
Metabolic and inflammatory system							
	Metabolics	HbA1c, %	5.49 (5.30, 5.61)	5.50 (5.30, 5.70)	−0.03 (−0.09, 0.03)	0.30	0.60
		Glucose, mg/dL	94.45 (88.02, 102.00)	90.98 (85.00, 98.00)	3.89 (1.9, 5.87)	<0.001	
	Inflammation	High-sensitivity CRP, mg/L	0.09 (0.05, 0.19)	0.11 (0.06, 0.26)	−0.05 (−0.09, −0.01)	0.022	0.044
		Leucocyte, count per μL	5.57 (4.71, 6.41)	5.90 (4.89, 7.10)	0.94 (0.91, 0.97)[b]	<0.001	<0.001
Neurological system							
	Vascular brain damage	Cerebral microbleeds	7 (3.72)	21 (4.45)	0.86 (0.35, 2.1)	0.74	1.00
		Infarct residuals	5 (2.66)	31 (6.49)	0.4 (0.15, 1.07)	0.068	0.27
		Intracerebral stenoses	2 (1.1)	3 (0.63)	2.72 (0.35, 21.14)	0.34	
	Brain atrophy	Relative brain volume, %	81.37 (78.88, 83.70)	81.69 (78.90, 83.85)	−0.52 (−1.1, 0.06)	0.077	0.31
		Cortical thickness, mm	2.65 (2.60, 2.72)	2.63 (2.57, 2.69)	0.03 (0.01, 0.05)	<0.001	0.002
Neurocognition							
	Screening for cognitive impairment	Mini-Mental State Examination	29 (28, 29)	28 (27, 29)	0.26 (0.1, 0.42)	0.002	0.002
Psychosocial outcomes							

Continued

Table 3 Continued

Biological parameter	Corresponding surrogate	Subjects after SARS-CoV-2 infection (n = 443)	Matched controls (n = 1328)	Regression estimate[a]	Nominal P-value for regression estimates	Adjusted P-values
Severity of depression	PHQ-9	3 (1, 6)	3 (1, 6)	0.34 (−0.06, 0.74)	0.10	0.20
Severity of somatic symptoms	PHQ-15	4 (2, 7)	4.00 (2, 7)	0.21 (−0.23, 0.64)	0.35	
Generalized anxiety	GAD-7	2 (0, 5.00)	2 (1, 4)	0 (−0.35, 0.35)	0.99	
Quality of life	EQ-5D: index	0.91 (0.87, 1.00)	0.91 (0.89, 1.00)	0 (−0.01, 0.01)	0.75	1.00
	EQ-5D: VAS	81 (75, 90)	82 (75, 90)	−1.5 (−3.23, 0.23)	0.090	

Continuous variables are presented as median and interquartile range, and categorical variables are presented as absolute numbers and percentages.
AV, atrioventricular; CCA, common carotid artery; CMR, cardiac magnetic resonance; CRP, C-reactive protein; ECG, electrocardiogram; EQ-5D, European Quality of Life 5 Dimensions; EQ-5D: VAS, European Quality of Life 5 Dimensions visual analogue scale; FEV₁, forced expiratory volume in 1 s; FRC, functional residual capacity; FVC, forced vital capacity; GAD-7, Generalized Anxiety Disorder Assessment; GFR, glomerular filtration rate; HbA1c, glycated haemoglobin; ICA, internal carotid artery; LAVI, left atrial volume indexed to body surface area; LV, left ventricular; LVEDV, left ventricular end-diastolic volume; LVEF, left ventricular ejection fraction; NT-proBNP, N-terminal pro-B-type natriuretic peptide; NYHA, New York Heart Association;PHQ-9, Patient Health Questionnaire-9; PHQ-15, Patient Health Questionnaire-15; Raw, airway resistance; RV, right ventricular; RVEF, right ventricular ejection fraction; sRaw, specific airway resistance; TAPSE, tricuspid annular plane systolic excursion; TLC, total lung capacity; TR Vmax, peak tricuspid regurgitation velocity; TTE, transthoracic echocardiogram; VC, vital capacity.
[a]Regression estimates for patients after SARS-CoV-2 infection vs. matched controls. Regression estimates are presented as beta and 95% confidence interval for continuous variables and odd ratios and 95% confidence interval for categorical variables.
[b]Regression estimates are presented as beta for retransformed logarithmic outcomes. Betas are multiplicative instead of additive.
[c]NYHA Class II, III, and IV compared with NYHA Class I in binary logistic regression.

0.03 (95% CI 0.01, 0.05), adjusted $P = 0.002$] when compared with matched controls.

Neurocognitive testing
The median score in MMSE was higher in patients after SARS-CoV-2 infection than in matched controls [29 vs. 28, regression coefficient 0.26 (95% CI 0.1, 0.42), adjusted $P = 0.002$].

Quality of life and psychosocial outcomes
There were no significant differences between patients after SARS-CoV-2 infection and matched controls in terms of quality of life, severity of depression, severity of somatic symptom, or generalized anxiety.

Associations of antibody concentration with main outcomes
Associations of anti-spike antibody concentration with main outcomes across the various organ systems can be found in Supplementary material online, Table S7. In brief, no systematic associations could be observed except for a direct association between antibody concentrations and infarct residuals on brain MRI.

Subgroup analyses
Within the subjects with prior SARS-CoV-2 infection, no systematic differences in organ-specific outcomes could be observed when comparing mild vs. moderate course of COVID-19 as well as recruitment route, except for higher levels of anxiety and depression in subjects with the moderate course of COVID-19 compared with subjects with a milder course (see Supplementary material online, Tables S8 and S9).

Discussion

In more than 1700 individuals, we extensively phenotyped multi-organ-specific structure and function and explored neurocognitive and PRO to comprehensively assess the intermediate to long-term effects of mild and moderate COVID-19 disease. We consistently observed organ-specific subclinical involvement (Graphical Abstract).

Autopsy studies indicate that SARS-CoV-2 affects multiple organs beyond the respiratory tract, including the heart, brain, and kidneys.[6,24] Some patients continue to suffer from heterogeneous symptoms after the acute phase of critical illness. These conditions are described as 'post-COVID-19 syndrome' or—if symptoms continue longer than 6 months—as 'long COVID-19 syndrome'. Clinical, imaging, or laboratory findings should accompany the diagnosis of post- or long COVID-19.[25] To date, longer term effects after a mild to moderate COVID-19 disease course remain largely unknown. A systematic and comprehensive exploration of potential multi-organ impairment is important to plan surveillance and potential diagnostic testing after recovery.

While there is evidence that post-COVID-19 patients with mild to moderate disease have preserved lung volumes,[26] we found a significantly lower TLC in subjects after mild to moderate infection compared with controls. This is in line with previous findings of a reduction in TLC beginning after mild COVID-19.[27] Residual
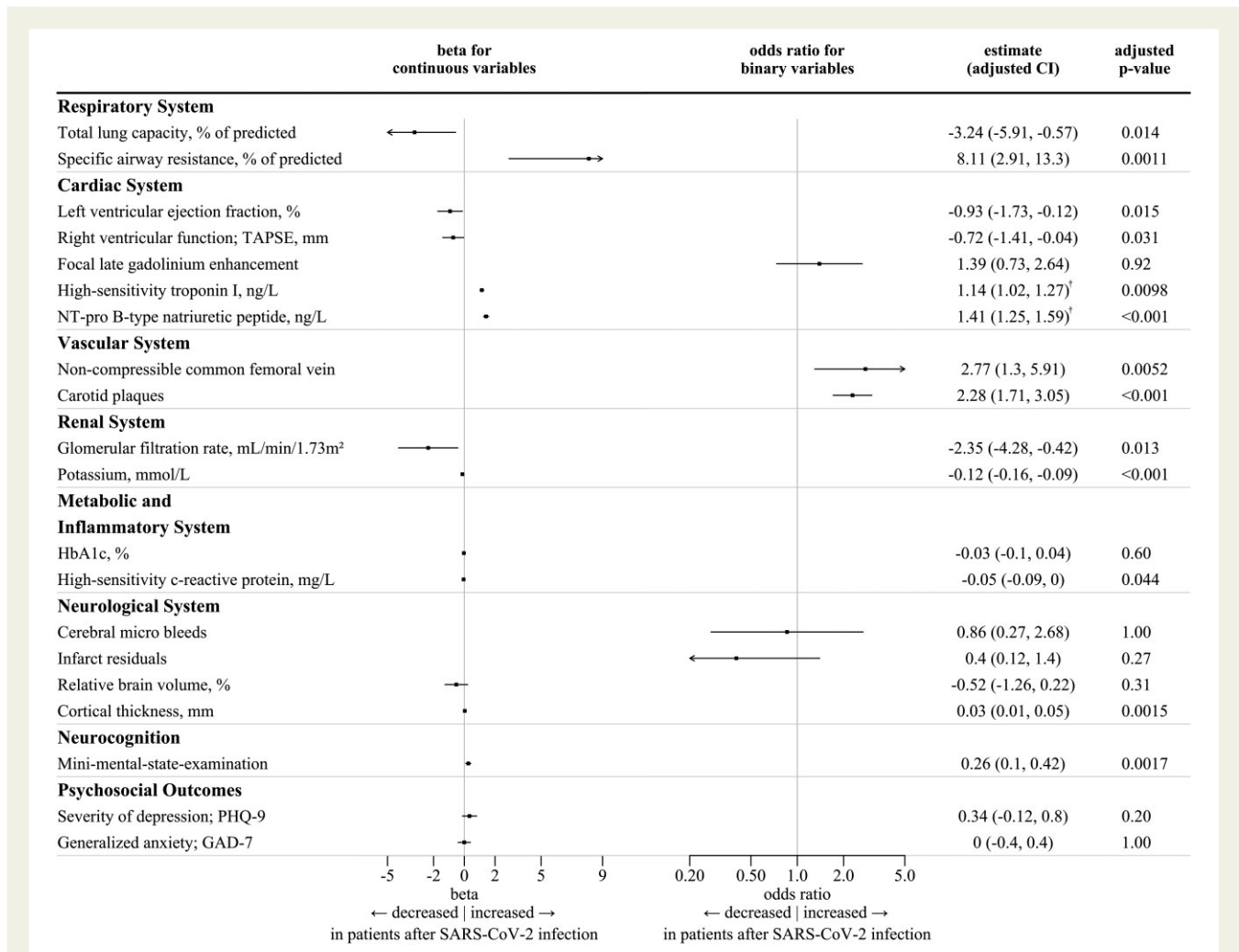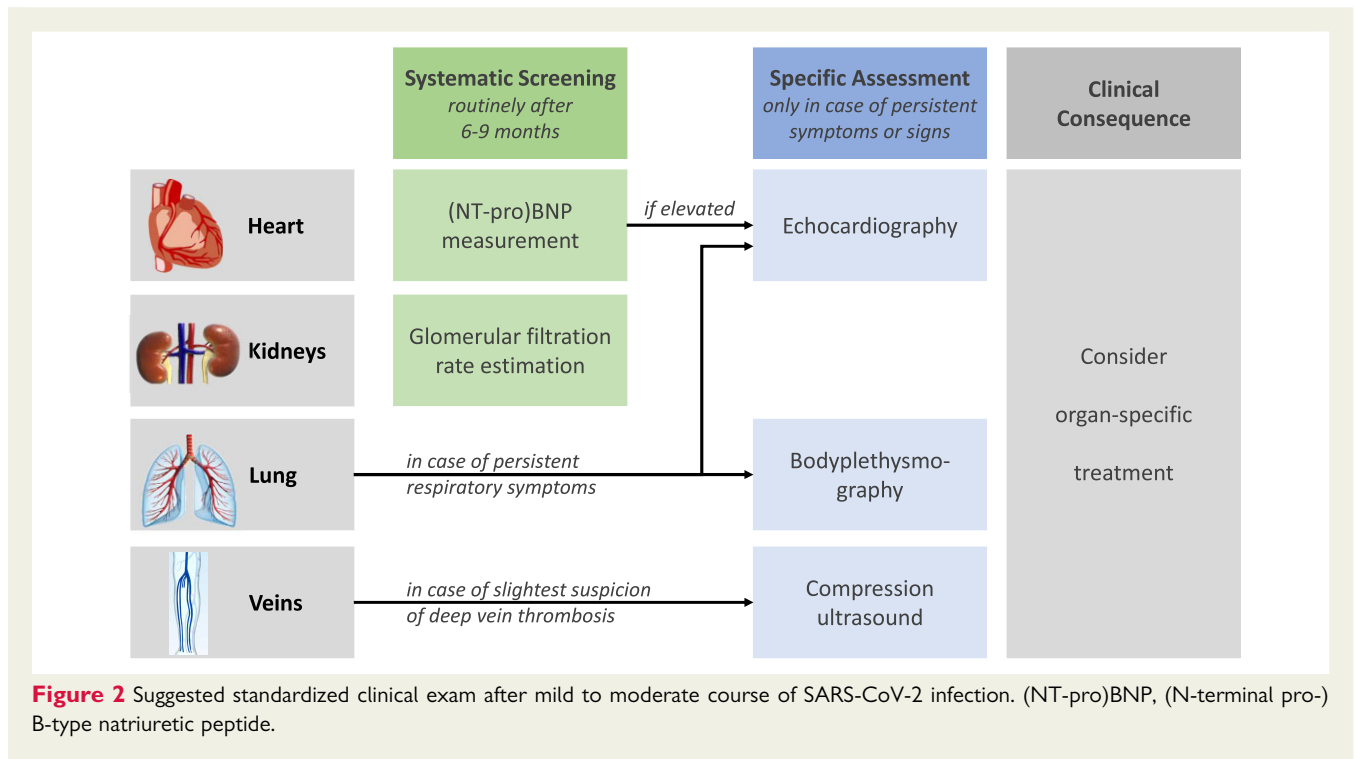
Figure 1 Forest plot depicting the association of prior SARS-CoV-2 infection with organ-specific main outcomes. Regression estimates for patients after SARS-CoV-2 infection vs. matched controls. Regression estimates are presented as beta and 95% confidence interval for continuous variables and odd ratios and 95% confidence interval for categorical variables. *P*-values are adjusted using the Bonferroni correction for main outcomes within each organ system. [†]Regression estimates are presented as beta for retransformed logarithmic outcomes. Betas are multiplicative instead of additive. GAD-7, Generalized Anxiety Disorder Assessment; HbA1c, glycated haemoglobin; NT-proBNP, N-terminal pro-B-type natriuretic peptide; PHQ-9, Patient Health Questionnaire-9; TAPSE, tricuspid annular plane systolic excursion.

inflammatory processes or beginning subclinical fibrotic remodelling might explain the slightly lower lung volumes observed in the post-SARS-CoV-2 cohort.[28] Congruent with histological findings, we found higher specific airway resistance in participants after SARS-CoV-2 infection.[29] Assessment of lung function should be considered after recovery from COVID-19 at slightest suspicion even in apparently healthy individuals (*Figure 2*).

From a cardiovascular perspective, a numerically small reduction of left ventricular ejection fraction with 1–2% difference was observed in participants after SARS-CoV-2 infection accompanied by higher concentration of cardiac biomarkers reflecting modest myocardial involvement. In a very long-term perspective, even a small reduction of left ventricular function and a slight increase in NT-proBNP concentration translate into an increased risk of mortality in the general population.[30,31] Therefore, the determination of NT-proBNP, followed by echocardiography control in the case of elevated concentrations, may be recommended after COVID-19 recovery to avoid untreated dysfunction of the heart.

Most importantly, our data suggest a significantly higher prevalence of deep venous thrombosis in participants after SARS-CoV-2 infection. Although affected by certain examiner bias, compression ultrasound is commonly accepted and widely used as the reference standard in everyday clinical practice.[32] The present findings extend the rapidly increasing evidence for an association between COVID-19 and venous thromboembolism,[4,33] while adding a prospectively enrolled cohort with mild or moderate disease. Of note, levels of coagulation parameters assessed at a median of 9.6 months after mild to moderate SARS-CoV-2 infection did not differ when compared with matched controls. Therefore, we may assume that acute thrombotic events themselves, associated with activation of the coagulation

Figure 2 Suggested standardized clinical exam after mild to moderate course of SARS-CoV-2 infection. (NT-pro)BNP, (N-terminal pro-) B-type natriuretic peptide.

and fibrinolytic cascade, occurred much earlier during the course of COVID-19. In this context, the non-compressible veins observed in our study are to be interpreted as remnants of the fibrotic processes replacing prior thrombotic material. Considering the existing evidence, the current study results suggest that a guideline-based surveillance with active screening for deep vein thrombosis in the case of minimal clinical suspicion early during COVID-19 infection should be considered.[32]

The assessment of the arterial system indicated more frequent carotid plaques in post-SARS-CoV-2 participants compared with matched controls, while CIMT was comparable between groups. Whether this finding may be fully explained by the higher susceptibility of patients with the prevalent atherosclerotic disease for COVID-19[34,35] or to some extent also by common immunological pathways involved in SARS-CoV-2 infections and atherosclerosis[36] needs to be addressed in future longitudinal studies. Lower peak systolic flow velocities in the carotid arteries in patients after SARS-CoV-2 infection are in line with the trend towards a lower ejection fraction and presence of carotid plaques, as these parameters were reported to be associated recently.[16]

Individuals after SARS-CoV-2 recovery were also found to have a subtle decrease in kidney function compared with the matched controls, which also does not appear clinically relevant at the time of investigation. However, autopsy studies have shown a distinct renal tropism related to SARS-CoV-2 and early urine abnormalities associated with mortality as well as multi-organ failure in hospitalized COVID-19.[6] The differences observed in our study might thus reflect a specific SARS-CoV-2-related injury with a starting trajectory to early chronic kidney disease, being an important risk factor for mortality and cardiovascular events.[37]

Therefore, we argue for a follow-up assessment of a renal function marker 6–9 months even after mild SARS-CoV-2 infections.

Brain MRI was available in a large subgroup of patients and did not reveal any signs of increased vascular brain damage. The presence and number of cerebral microbleeds and white matter hyperintensities as the most common hallmark of cerebral small vessel disease were comparable between participants recovered from mild to moderate COVID-19 and controls. This contrasts with findings in cohorts of severely ill COVID-19 patients, for whom both white matter lesions and microhaemorrhages have been reported.[38] The presence of vascular brain damage may thus be a phenomenon that is only observed in the severe course of COVID-19. Quantitative analysis of structural brain MRI did not show signs of overall brain atrophy, as relative brain volume was comparable between groups, and even showed a higher mean cortical thickness in patients after SARS-CoV-2 infection. This, again, is in contrast to a recent longitudinal analysis of brain imaging from participants of the UK Biobank who recovered from COVID-19, which showed brain atrophy and a regional reduction in grey matter thickness.[39] We also did not find any sign of cognitive impairment in patients after mild to moderate COVID-19. This adds to recent reports of persistent impaired cognitive function in few individuals who recovered from severe COVID-19.[40]

Central parameters of PRO like depression, anxiety, and quality of life were tested. In none of the five scales, significant differences were observed in the main analysis. However, in an explorative subgroup analysis, higher degrees of depression and anxiety were observed in subjects with moderate course of COVID-19 when compared with subjects with none or only mild symptoms. These findings extend and corroborate previous work describing serious long-COVID-19 symptoms after intensive care courses

of COVID-19, which also affect the psychosocial domain.[41] Individuals after mild to moderate disease might have normalized their previously elevated levels of anxiety and depression by the time of the survey.

The following limitations merit consideration. In this study, cases were only assessed after exposure to SARS-CoV-2. As assessment was not available prior to SARS-CoV-2 infection, multi-organ functions were compared with matched controls from the general population without exposure. Accordingly, as this is a cross-sectional study, we cannot infer on future developments and causality, but suggest that these findings represent signs of subtle medium to long-term multi-organ affection persisting after recovery from mild to moderate COVID-19, where most cases were not hospitalized.

Also, a selection bias might have occurred in the post-SARS-CoV-2 group of this study as the public invitation resulted in the participation of highly motivated post-SARS-CoV-2 study individuals, which particularly might have led to better performance in neurocognitive tests and quality-of-life questionnaires. However, the versatile multi-organ assessment consistently describes multi-organ affection even in the case of selection bias towards healthier or more motivated post-SARS-CoV-2 study individuals. As pharmacological COVID-19 therapies were not assessed in this study, we cannot quantify their impact on multi-organ function. However, given the high proportion of individuals with only mild symptoms and outpatient management in this data set, no relevant impact on reported outcomes is assumed. In addition, hs-CRP levels were very low and comparable in both groups. As levels were below the limit of detection at 0.1 mg/dL in about half of the populations, we may not comment on hypothetical very small differences in the extremely low end. Lastly, data on outcome variables were not completely available in all subjects. Multiple imputation was used to at least partly correct for this aspect.

Particular study strengths exist. The entire study population had been evaluated by one of the most comprehensive study protocols ever applied to evaluate multi-organ function, neurocognition, and PRO in patients recovered from mild to moderate COVID-19 within the frame of the HCHS. Participants after mild to moderate COVID-19 qualified with at least one positive PCR test. Further, antibody status was obtained. Finally, matched controls had been randomly recruited from the same geographic region via registration office within the frame of the HCHS.

Given that moderate SARS-CoV-2 infection may impact multi-organ function in the longer term, comprehensive preventive strategies such as hygiene measures, social distancing, and vaccination seem crucial.

In conclusion, in the HCHS COVID project, we demonstrate that patients who apparently recovered from mild to moderate COVID-19 suffer from modest subclinical multi-organ affection related to thrombotic, pulmonary, cardiac, and renal function without signs of structural brain damage, obvious impairment in cognitive function, or quality of life. Whereas the impact on very long-term outcome remains unclear, a standardized clinical exam of these conditions after recovery is recommended.

Data availability

The data underlying this article will be shared on reasonable request to the corresponding author.

Supplementary material

Supplementary material is available at *European Heart Journal* online.

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