

**Individual factors including age, BMI and heritable factors underlie temperature variation in sickness and in health: an observational, multi-cohort study**

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## **Abstract:**

**Background:** Ageing affects immunity, potentially altering fever response to infection. We assess effects of biological variables on basal temperature, and during COVID-19 infection, proposing an updated temperature threshold for older adults  $\geq 65$  years.

## **Methods:**

Participants were from four cohorts: 1089 unaffected adult TwinsUK volunteers; 520 adults with emergency admission to a London hospital with RT-PCR confirmed SARS-CoV-2 infection; 757 adults with emergency admission to a Birmingham hospital with RT-PCR confirmed SARS-CoV-2 infection and 3972 adult community-based COVID Symptom Study participants self-reporting a positive RT-PCR test. Heritability was assessed using saturated and univariate ACE models; mixed-effect and multivariable linear regression examined associations between temperature, age, sex and BMI; multivariable logistic regression examined associations between fever ( $\geq 37.8^{\circ}\text{C}$ ) and age; receiver operating characteristic (ROC) analysis was used to identify temperature threshold for adults  $\geq 65$  years.

## **Results:**

Among unaffected volunteers, lower BMI ( $p=0.001$ ), and increasing age ( $p<0.001$ ) associated with lower basal temperature. Basal temperature showed a heritability of 47% (95% Confidence Interval 18-57%). In COVID-19+ participants, increasing age was associated with lower

temperatures in Birmingham and community-based cohorts ( $p < 0.001$ ). For each additional year of age, participants were 1% less likely to demonstrate a fever  $\geq 37.8^\circ\text{C}$  (OR 0.99;  $p < 0.001$ ). Combining healthy and COVID-19+ participants, a temperature of  $37.4^\circ\text{C}$  in adults  $\geq 65$  years had similar sensitivity and specificity to  $37.8^\circ\text{C}$  in adults  $< 65$  years for discriminating infection.

**Conclusions:** Ageing affects temperature in health and acute infection, with significant heritability, indicating genetic factors contribute to temperature regulation. Our observations suggest a lower threshold ( $37.4^\circ\text{C}/97.3^\circ\text{F}$ ) for identifying fever in older adults  $\geq 65$  years.

**Keywords:** thermoregulation; fever; immunosenescence; infection; COVID-19

## **Background**

Normal body temperature of older adults has been observed to be lower than that of younger people, with more limited tolerance of thermal extremes (1,2). Thermoregulation involves multiple systems, including cardiovascular, respiratory and musculoskeletal. Decreasing basal temperature with age may reflect natural changes in these systems and reduced ability to maintain homeostasis in response to ambient temperature changes.

Normal human body temperature is approximately 37°C (98.6°F); however, variation is observed, with daily variations as much as 0.25 to 0.5°C (3). Individual biological factors, such as genetics, may account for some of the observed variation in human body temperatures as well as metabolic determinants. Commercially important species, such as cattle and poultry, show heritability of thermo-regulation under different heat conditions (4-6). However, human heritability data are more limited. A small study of 53 adult female twin-pairs (mean age 52) used continuous wrist temperature monitoring to demonstrate Circadian system heritability (7). Monozygotic pairs showed higher intra-pair correlation than dizygotic twins for most parameters, including temperature, with genetic factors responsible for 46%-70% of variance.

Fever is a physiological response to infection and is clinically important in identifying infection. There is no widespread consensus on the definition of fever, although a number of thresholds have been proposed. Infectious Diseases Society of America guidelines define fever as a single

oral temperature  $>37.8^{\circ}\text{C}$  ( $>100.0^{\circ}\text{F}$ ); repeated oral temperatures  $>37.2^{\circ}\text{C}$  ( $>99.0^{\circ}\text{F}$ ) or rectal temperatures  $>37.5^{\circ}\text{C}$  ( $>99.5^{\circ}\text{F}$ ); or an increase in temperature of  $>1.1^{\circ}\text{C}$  ( $>2.0^{\circ}\text{F}$ ) above baseline (8). It has been observed that fever may be absent or blunted in older adults, as much as 20-30% of the time (9). Many have therefore questioned applicability of these thresholds in older adults (10,11); indeed, lowering the threshold to  $37.2^{\circ}\text{C}$  ( $99.0^{\circ}\text{F}$ ) has been shown to increase fever detection in older people (8,12). A previous study found that only 20-30% of patients aged  $\geq 60$  with infection present with a temperature of  $37.8^{\circ}\text{C}$  ( $100.0^{\circ}\text{F}$ ) or above (13). Many older adults display atypical presentations, which may be associated with worse outcomes (14,15). Age-related changes in immune function may affect pyrogen production and fever response. Altered immune function, or “immunesenescence”, is a well-established feature of physiological ageing (16). Immunesenescence affects both innate and adaptive arms of the immune system and is thought to be responsible, at least in part, for an increased incidence and severity of infections in older adults (16). Proposed age-related changes in adaptive immunity include limited diversity in B- and T-cell receptor repertoire, decreased numbers of naive T and B cells and reduced antigen-specific antibody production (17). Neutrophil function is also compromised by age and frailty and has been proposed as a possible therapeutic target to enhance immune responses in frail, older adults during infection (18).

Coronavirus Disease-19 (COVID-19) is an acute respiratory disease caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2). Fever has been cited as a “typical” symptom of COVID-19, with reported prevalence of between 64-98% in hospitalised patients across a number of international studies (19-23). Current United Kingdom (UK) public health guidance proposes a temperature  $\geq 37.8^{\circ}\text{C}$  to define a possible COVID-19 case (24). As for other

infections, atypical presentations of COVID-19 have been documented in older adults, with symptoms including weakness, headache and delirium, often without fever (20,25,26). In work done by our research group on point-of-care testing for COVID-19, hypothermia ( $T < 36.0^{\circ}\text{C}$ ) was noted to be an early clinical sign in a minority of patients (27).

Demonstrating relationships between age and temperature in a healthy cohort and cohorts with confirmed SARS-CoV-2 infection is important in understanding effects of ageing on temperature regulation and the fever response to infection. Identifying a more sensitive temperature threshold for older adults may help detect cases of infectious disease, including COVID-19. Case identification, especially in the case of COVID-19, is relevant for infection control.

In this study, we aim to:

1. Identify individual factors associated with baseline temperature in a sample of healthy volunteers, including age, sex, body mass index (BMI) and temperature heritability;
2. Determine if any identified associations are consistent with those observed during the acute temperature response to SARS-CoV-2 infection, in community-based and hospitalised cohorts;
3. Identify a more appropriate temperature threshold for discriminating SARS-CoV-2 infection in adults  $\geq 65$  years of age than  $37.8^{\circ}\text{C}$  ( $100.0^{\circ}\text{F}$ ) – the temperature proposed by UK public health guidance to screen for COVID-19 cases (24).

## **Methods**

### **Participants:**

This observational cohort study uses four distinct cohorts.

(a) Unaffected twin volunteers: Measurements were collected between January 2017-April 2018 on 1089 adult twin volunteers, a convenience sample of the TwinsUK registry enrolled at the Department of Twin Research (DTR), King's College London (KCL) (28). Demographic information recorded on twins in this cohort includes age, sex, zygosity and BMI.

Hospital COVID-19+ cohorts: Data was collected as part of the international COVIDCollab study, led by the Geriatric Medicine Research Collaborative (29,30). All data is routinely collected and includes age, sex, BMI, and vital signs, including temperature. Full data collection information has been published previously (29). Only cases of COVID-19 infection confirmed by RT-PCR of nasopharyngeal swab were included. For patients readmitted during study period, data from index admission was used.

(b) London: First documented point-of-care temperature measurements were collected on all adult patients with COVID-19 infection confirmed by RT-PCR of nasopharyngeal swab, with unscheduled admission to St Thomas' Hospital, London, UK between March 1st-May 4th, 2020.



(c) Birmingham: First documented point-of-care temperature measurements were collected on all adult patients with confirmed COVID-19 infection, with unscheduled admission to Queen Elizabeth Hospital, Birmingham, UK between March 4th 2020-April 13th, 2020.

(d) Community-based “app” COVID + cohort: Data was obtained from UK-based participants enrolled in the COVID Symptom Study (CSS). CSS participants log using a free smartphone application “app”, which was developed by Zoe Limited (London, UK) with scientific input from researchers and clinicians at King’s College London (London, UK). The app was launched in the UK on 24th March 2020 and captures information related to COVID-19 symptoms, self-reported or reported-by-proxy. On first accessing the app, users record baseline demographic factors e.g. location, age, ethnicity and health risk factors (31). Consent to sharing and analysis of data is obtained during signup for the app. Data was extracted for all UK participants self-reported to have tested positive for COVID-19 by RT-PCR between 24th March-9th June 2020 who self-reported “fever or chills” and logged at least one recorded temperature using the app during the study period.

Measurements:

(a) Unaffected volunteers: Three infrared tympanic temperature measurements were taken for each twin participant during a single routine research visit to the Guy’s and St Thomas’ Clinical Research Facility. A mean temperature for each participant was used for analysis.

(b) & (c) Hospital COVID-19+ cohorts: Electronic patient records were reviewed. Only COVID-19 cases confirmed by RT-PCR of nasopharyngeal swab were included. First documented oral or tympanic temperature on admission was used to reflect temperature during acute infection.

(d) Community “app” COVID-19+ cohort: Self-reported temperature measurements were extracted for all adult participants self-reported to have tested positive for COVID-19, who had recorded at least one temperature measurement using the app. Maximum temperature was used for analysis, to most likely reflect temperature during acute illness.

#### Analyses:

Descriptive statistics were used to describe demographic and key clinical characteristics of the study cohorts.

Differences within the unaffected volunteer cohort (a) and the two hospitalised COVID-19+ cohorts (b&c) were compared using the Fisher exact test for categorical and Wilcoxon signed rank for continuous variables.

To analyse associations between age and temperature in the unaffected volunteer cohort (a), a linear mixed effect model was used, with age, gender and BMI set as fixed factors and

accounting for family and zygosity as random factors. In hospital COVID-19+ cohorts (b&c), multivariable linear regression was used to model the effects of age, sex and BMI on temperature recorded on admission. In the community-based COVID-19+ cohort (d), multivariable linear regression was used to analyse associations between age, sex and BMI and maximum self-recorded temperature in the context of confirmed COVID-19. For each cohort, two analyses were performed, the first including age and sex as predictor variables, the second including age, sex and BMI. Participants with missing temperature data were excluded from analysis. Participants with missing BMI data were only excluded for analyses using BMI. Residuals were checked to confirm homoscedasticity in all samples. P-values were Bonferroni-adjusted for multiple testing.

In COVID-19+ hospital and community cohorts (b,c&d), logistic regression was performed to ascertain associations between recorded fever (defined as temperature  $\geq 37.8^{\circ}\text{C}$ ) and age, adjusted for sex and BMI. A complete case approach was used, as low levels of missing data were expected. P-values were not adjusted since multiple comparisons were not applied.

Receiver Operating Characteristic (ROC) analysis was performed on a combined dataset of unaffected healthy volunteers (a) and hospitalised COVID-19+ participants (b&c) to identify a temperature threshold in older adults ( $\geq 65$  years of age) with similar sensitivity and specificity for discriminating COVID-19 to a temperature  $\geq 37.8^{\circ}\text{C}$  in adults  $<65$  years of age.

Genetic and environmental contributions of basal temperature in TwinsUK cohort was estimated using a univariate model (32). The model decomposes the observed variance of the basal temperature into additive genetic variance (A), shared environment variance (C) and non-shared environment variance (E), deducing the genetic influence of a trait and by comparison of the different nested models. The analysis was carried out using Open Mx (32) for R on residuals of linear regression adjusted for age and sex and BMI. Heritability of self-reported fever was carried out in 3099 COVID Symptom Study app participants who were also enrolled in TwinsUK [291(9.4%) of whom were reporting fever], using biometric modelling for the liability threshold model, which, for a categorical trait, assumes an underlying continuous liability that follows a normal distribution (33).

Data analysis and graphics were performed in the R statistical environment (version 4.0) using the Tidyverse (34), Open Mx (32) and lme4 (35) packages, in Stata (version 15.0) and in Python (version 3.9).

#### Ethics:

TwinsUK main ethics was reviewed and approved by the NHS London – London-Westminster Research Ethics Committee (REC reference EC/04/015), and by Guy's and St Thomas' NHS Foundation Trust Research and Development (R&D) in 2012. TwinsUK BioBank was approved by NHS North West - Liverpool East Research Ethics Committee (REC reference 19/NW/0187), IRAS ID 258513.

The Covid Symptom Study App Ethics has been approved by KCL ethics Committee REMAS ID 18210, review reference LRS-19/20-18210. All subscribers provided consent when signing up for the app.

COVIDCollab project service evaluation approved by Guy's and St Thomas' NHS Trust audit leads. Reference number 10777.

The University Hospitals Birmingham (UHB) validation was performed as part of a service evaluation with a (CARMS-16005), after approval on routinely collected anonymised dataset.

## **Results**

Baseline characteristics and temperature measurements for all cohorts are shown in Table 1. 6/520 patients in the London hospital cohort and 81/757 in the Birmingham hospital cohort had missing temperature data and were excluded from analysis. Of included participants, 28/514 in the London hospital cohort and a further 26/676 in the Birmingham hospital cohort had missing BMI data and were excluded from the analyses including BMI. 5/3972 in the app cohort had sex “other” and were excluded from sex-adjusted analysis only.

## Regression Analysis

Results of the linear mixed effect model in the TwinsUK sample and linear regression models in the community-based and hospital cohorts are displayed in Table 2. The assumption of homoscedasticity of residuals was not violated for the cohorts used in regression analysis. Increasing age was associated with both lower basal temperature, and lower temperature during illness, whilst increasing BMI was associated with higher basal and illness temperature.

Logistic regression, exploring the relationship between age and temperature  $\geq 37.8^{\circ}\text{C}$ , demonstrated that for each additional year of age, participants with confirmed COVID-19 were 1% less likely to show a fever in both the app (Odds Ratio 0.99 (95% Confidence Interval 0.98-0.99);  $P < 0.001$ ) and the Birmingham hospital cohorts (OR 0.99 (95% CI 0.98-0.99);  $P = 0.040$ ) (Table 3).

## ROC analysis

A temperature of  $37.8^{\circ}\text{C}$  had a sensitivity of 35.2% and specificity of 100% for discriminating between the unaffected volunteer cohort and hospitalised COVID-19+ patients aged  $< 65$  years of age, in a combined dataset of unaffected healthy twin volunteers and hospitalised COVID-19+ participants. ROC analysis performed on a combined dataset demonstrated that a temperature of  $37.4^{\circ}\text{C}$  in adults  $\geq 65$  years of age had the most closely matched sensitivity (36.3%) and specificity (98.9%) to that of  $37.8^{\circ}\text{C}$  in adults  $< 65$  years of age for discriminating COVID-19+

patients from unaffected volunteers. ROC curves for adults <65 and  $\geq 65$  years of age are shown in Figure 1, with sensitivity and specificity of different temperature thresholds in eTable 1. We note a slightly smaller area under the curve (AUC) in older versus younger adults which may point towards reduced discriminatory performance in older adults. AUC was similar in sub-analyses of adults  $\geq 75$  and  $\geq 85$  years of age, although these numbers should be cautiously interpreted given small numbers of adults in older age groups (eTable 2).

#### Heritability analysis:

Results for heritability analysis are summarised in Table 4.

Analysis of monozygotic twin intra-pair correlation showed a clear heritability signal in comparison to dizygotic twin pairs for basal temperature. When adjusted for age and sex, genetic factors accounted for 47% (95% CI 18-57%) of variance in mean temperature in the best model (ACE) as compared to the other models (Saturated, ACE, CE, E) (Table 4). When adjusted for age, sex and BMI, genetic factors accounted for 44% (95% CI = 15-57%) of variance in mean temperature in the best model (ACE) (Table 4).

Heritability of fever, using twins who are app participants, did not demonstrate a genetic component to temperature response to infection, which may be explained by predominantly environmental factors driving exposure to SARS-CoV-2 (Table 4).

## **Discussion**

### Sources of variability in temperature: Age, BMI and heritability.

In this study, associations between age and temperature were replicated across healthy volunteer, hospitalised COVID-19 and community-based COVID-19 cohorts. The observed associations persisted after adjustment for confounding by sex and BMI in the healthy volunteer, the larger Birmingham hospital cohort, and community-based cohorts. Temperature regulation is affected by many physiological processes, including vasomotor sweating functions, skeletal muscle response, temperature perception, and physical behaviours, many of which have been described in previous studies and reviews (1,3,5). These processes change with ageing, and our observation that increasing age reduces both basal temperature and temperature in response to SARS-CoV-2 infection supports this. Case reports and small cohort studies have previously shown that older adults, especially those with frailty or multiple chronic conditions, may not display a clinically significant fever response to infection (13,25). This may reflect both a lower baseline temperature in older adults, as well as a lower temperature change in response to infection. Age-related reductions in pro-inflammatory mediators in response to infection, such as interleukin-1 and tumour necrosis factor, may also have an effect.

The possible association between higher BMI, often considered a pro-inflammatory state, and an increased basal and fever temperature would appear to concord with this finding for age. The



observed associations between BMI and temperature in unaffected volunteer and community-based COVID-19 cohorts were apparent after adjustment for age. Some earlier studies, often with smaller, specific samples, demonstrated a conflicting association between BMI and temperature (36). However, our observation is supported by the recent, large CoLaus study of 4224 men and post-menopausal women (37). Here, other markers of obesity and insulin metabolism were also associated with higher temperature.

In this analysis, after adjustment for both age and BMI, we demonstrate a clear heritable component to basal temperature, pointing to other biological variables which may underlie temperature variation. Genetic factors may also play a crucial role in temperature variability among individuals of different ages and with different BMI. Physiological factors related to age and temperature control may also be heritable. In contrast to basal temperature, “fever”, defined here as a temperature of 37.8°C (100.0°F) or above, did not demonstrate a significant genetic component, suggesting that environmental factors (in this case exposure to the SARS-CoV-2 virus) may be the main driver of fever within this population.

#### Fever Threshold for Discriminating Infection in Older Adults:

Results of ROC analysis, combining unaffected volunteers and hospitalised COVID-19 patients, supports observations from previous studies and concerns from frontline clinicians that current fever threshold definitions may be inappropriate for older adults (8,12). A lower temperature of

37.4°C (97.3°F) in older adults aged 65 years and over had a similar sensitivity for discriminating COVID-19 patients from healthy volunteers as the widely utilised threshold of 37.8°C in older adults. Our observation is supported by a previous study that demonstrated optimised sensitivity and specificity thresholds of 37.9°C for younger adults and 37.3°C for older adults for discriminating influenza infection (38). Our finding may have important utility for screening in clinical settings, with a lower threshold in older adults to promptly detect infections and implement effective containment measures. However, it should be noted that both in our study and the previous influenza study using a ROC analysis, AUCs of between 0.66-0.73 suggest that the discriminatory power of temperature alone for identifying infection is relatively poor. Furthermore, temperature thresholds identified in both studies for younger and older adults did not have a high sensitivity for discriminating infection. Using temperature alone to identify infection risks missing cases and we would advocate incorporating temperature measurements as part of a wider clinical assessment.

#### Implications and generalisability:

Associations observed in this study have important clinical implications for case detection and surveillance of infectious diseases, including COVID-19. Internationally, to our knowledge there are currently no countries or organisations that have adopted age-related temperature thresholds for defining fever, highlighting the international relevance of this research. There is a risk of cases being missed in older adults if inappropriately high fever thresholds are used as a screening or diagnostic tool for COVID-19 infection. This may be compounded in those with lower BMI, for example as a result of sarcopenia, who have lower temperatures. Whilst this study did not

intend to analyse associations between temperature and mortality, we know that older adults experience a higher fatality rate from COVID-19 (20,21,39). Our findings support use of a lower temperature threshold for older adults to initiate appropriate testing and isolation precautions in hospital settings. This may be relevant for care homes and long-term care facilities, with older populations vulnerable to outbreaks of respiratory disease. Observations from our community-based cohort may have relevance when considering methods for community case detection and limitation of spread (such as quarantining or screening for travel). Frailty, when assessed by specialists, may be a better predictor of disease outcomes than age or comorbidity alone (39). However, an age threshold has utility for those in clinical settings lacking the specialist input required for accurate frailty assessment.

### Strengths and Limitations

Our study combines data from three different types of research cohort: a classic research cohort (TwinsUK), routinely collected clinical data from hospital cohorts, and a cohort of community ‘citizen scientists’. We are not aware of any previous studies using an epidemiological approach across multiple large cohorts to analyse temperature at baseline and in febrile illness. The TwinsUK cohort benefits from a large number of observations, collected using rigorous study procedures. The associations observed in this cohort were reassuringly recapitulated in the “real-life” hospital and community-based data. This concordance increases generalisability from the academic sphere to clinical and community-based settings.

For historical reasons, the TwinsUK cohort is predominantly female. Females generally show greater individual temperature variability than males (40). The app cohort was also predominantly female. However, both hospital cohorts were predominantly male, reflecting the fact that men are more severely affected by COVID-19 (19-21). However, our models did not show any association with sex and recorded temperature, suggesting that these differences in cohort structure did not affect results. The TwinsUK cohort had a lower average BMI, possibly also reflecting that those with higher BMI are more likely to be unwell with COVID-19. However, we adjusted for BMI in our analyses, and the relationship between age and temperature was maintained in all of our analyses. Since the ROC method does not allow for adjustment for confounding variables, some of the difference between the COVID positive and COVID negative cohorts may be attributable to BMI. We used age in our ROC analysis as this can be used by clinicians as part of initial clinical assessment. BMI is not commonly or reliably assessed at triage. Importantly, the temperature threshold identified distinguishes between the two cohorts and is therefore relevant for clinical practice.

This study used data from a large population of individuals reporting on a mobile app. Information was self-reported and therefore there may be inaccuracies in logging of temperature and BMI. Although the app cohort only contained people with self-reported COVID-19 testing status, it is possible that maximal temperature logged in the app was not synchronous with COVID-19 infection or could be the result of another infection. Only those reporting “fever or chills” were prompted to enter a temperature measurement. Some may not have logged on the day of their maximal temperature. App users did under-represent older age groups (proportion  $\geq 65$  years 11.7%, compared to 18.3% of the UK population (41). 4.8% of app users with a

confirmed positive COVID-19 test were  $\geq 65$  years of age, which may reflect shielding in this group. This is supported by the fact that UK Office for National Statistics data showed lower rates of antibody seropositivity in the older, non-patient facing population during the study period (42). Sampling using a mobile app will under-represent individuals without mobile devices, which often includes those from more deprived backgrounds and marginalised groups, known to be at greater risk for severe disease. Furthermore, community testing for COVID-19 was not universally available in the UK during the period of data collection. The method for temperature measurement was not documented, although oral thermometers are most widely available in the UK. However, these limitations would be most likely to diminish power to detect effects, but would not alter the key association seen between age and temperature, which is replicated across cohorts.

Hospitalised patients present at different stages of illness; hence, temperature on admission will not always capture the most febrile component of infection. Some patients may have received antibiotics, antipyretics such as paracetamol or corticosteroids prior to initial temperature recording. BMI data was missing for a small proportion (<5%) of patients - likely not missing at random, as there may have been difficulty ascertaining accurate heights and weights for example due to extremes of body habitus or severe frailty.

## Conclusion

The significant associations observed between age and basal temperature are important in highlighting effects of ageing on thermoregulation and the immune response. The observed heritability of basal temperature in TwinsUK unaffected participants suggests that biological factors underlie differences in temperature regulation. These require further elucidation. As well as demonstrating lower temperatures in older adults, we report lower temperature with decreasing BMI. These two factors may compound each other in older, frail patients with sarcopaenia.

Our findings support a lower temperature threshold of 37.4°C (97.3°F) for identifying possible COVID-19 infection in older adults aged 65 and over. This has important implications for case detection, surveillance and isolation and could be incorporated into vital signs assessments for older people.

## **Data sharing**

TwinsUK data used in this study is available upon reasonable request to TwinsUK.

COVIDCollab data are available upon application to the Geriatric Medicine Research Collaborative <https://www.gemresearchuk.com/>.

App data used in this study are available to bona fide researchers through UK Health Data Research using the following link <https://healthdatagateway.org/detail/9b604483-9cdc-41b2-b82c-14ee3dd705f6>

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### **Conflicts of Interest**

None declared

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## Tables

**Table 1.**

	<b>(a) Unaffected volunteers (n=1089)</b>	<b>(b) London hospitalised patients (n= 514)</b>	<b>(c) Birmingham hospitalised patients (n=676)</b>	<b>(d) Community “app” participants (n=3967)</b>
	<i>Unaffected Volunteers</i>	<i>COVID-19 +</i>	<i>COVID-19 +</i>	<i>COVID-19 +</i>
<b>Age (years)</b>	56.6/16.8	61.9/17.1	67.8/17.0	42.9/12.8
<b>Sex (females)</b>	914 (84%)	202 (39%)	321 (43%)	2770 (70%)
<b>BMI (kg/m<sup>2</sup>)</b>	25.8/5.2	28.5/6.8 (n=486)	29.2/7.2 (n=646)	28.2/6.2
<b>Temperature (°C)</b>	36.6/0.4*	37.5/1.1**	36.9/0.9**	37.6/1.0***





	p<0.001	p<0.001	p=1.000	p= 1.000	p=1.000	p=1.000	p=0.043	p=0.031
<b>BMI</b>	<b>0.100</b>		<b>0.0157</b>		<b>0.0087</b>		<b>0.0136</b>	
	---	(0.032)	---	(0.007)	---	(0.005)	---	(0.003)
		p=0.001		p= 0.101		p=0.49		p<0.001

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**Table 3.**

	<b>(b) London Hospital (n= 514)</b>	<b>(c) Birmingham Hospital (n=676)</b>	<b>(d) Community “app” (n=3967)</b>
	<i>COVID-19 PCR+</i>	<i>COVID-19 PCR+</i>	<i>Self-report COVID-19 PCR+</i>
	Temperature on Admission $\geq 37.8^{\circ}\text{C}$	Temperature on Admission $\geq 37.8^{\circ}\text{C}$	Self-reported temperature $\geq 37.8^{\circ}\text{C}$
<b>Age</b>	0.99 (0.98-1.00)	0.99 (0.98-0.99)	0.99 (0.98-0.99)
<b>OR (95% CI)</b>	P= 0.45	P=0.040	P<0.001

**Table 4.**

Trait	Model	Variance components			Model fit			Model comparison			
		A	C	E	es	-2LL	AIC	$\chi^2$	df	p	base
Temperature	Saturated	-	-	-	10	921.88	-	-	-	-	-
							1236.111				
(Adjusted for Age and Sex)	<b>ACE</b>	<b>0.47[0.18- 0.57]</b>	<b>0.03[0-0.28]</b>	<b>0.49[0.42- 0.57]</b>	<b>4</b>	<b>923.73</b>	-	<b>1.85</b>	<b>6</b>	<b>0.933</b>	<b>Saturated</b>
	AE	0.50[0.43- 0.57]	-	0.49[0.42- 0.57]	3	923.79	-	0.06	1	0.810	ACE
	CE	-	0.42[0.34- 0.48]	0.57[0.51- 0.65]	3	934.40	-	10.61	1	1.092e- 03	ACE
	E	-	-	1.00 [1.00- 1.00]	2	1033.16	-	109.37	1	1.346e-	AE
							1140.835			25	
Temperature	Saturated	-	-	-	10	913.39	-	-	-	-	-
							1244.602				

(Adjusted for Age, Sex and BMI)	<b>ACE</b>	<b>0.44[0.15-</b>	<b>0.06[0-0.31]</b>	<b>0.49[0.42-</b>	<b>4</b>	<b>915.40</b>	<b>-</b>	<b>2.01</b>	<b>6</b>	<b>0.918</b>	<b>Saturated</b>
		<b>0.57]</b>		<b>0.57]</b>			<b>1254.594</b>				
	AE	0.50[0.43-	-	0.49[0.42-	3	915.61	-	0.21	1	0.646	ACE
		0.57		0.57]			1256.384				
	CE	-	0.24[0.35-	0.57[0.50-	3	924.83	-	9.22	1	2.139e-	ACE
			0.49]	0.64]			1247.169			03	
	E	-	-	1.00 [1.00-	2	1025.86	-	110.25	1	8.646e-	AE
				1.00]			1148.136			26	
Fever	<b>Saturated</b>	-	-	-	13	1853.05	-4318.95	-	-	-	-
(Adjusted for Age, Sex and BMI)	<b>ACE</b>	<b>0 [0-0]</b>	<b>0.21 [0-0.38]</b>	<b>0.79 [0.62-</b>	<b>8</b>	<b>1856.93</b>	<b>-4329.07</b>	<b>3.88</b>	<b>7</b>	<b>0.793</b>	<b>Saturated</b>
				<b>0.98]</b>							
	AE	0.2 [0-0.21]	-	0.8 [0.66-	7	1858.20	-4329.80	1.27	1	0.260	ACE
				0.99]							
	<b>CE</b>	<b>-</b>	<b>0.21 [0.02-</b>	<b>0.79 [0.62-</b>	<b>7</b>	<b>1856.93</b>	<b>-4331.07</b>	<b>0.00</b>	<b>1</b>	<b>1.000</b>	<b>ACE</b>



**Table 1.** Baseline characteristics and temperature measurements of (a) unaffected volunteers; (b) London and (c) Birmingham hospitalised cohorts with RT-PCR-confirmed COVID-19 and (d) Community-based “app” cohort with self-reported confirmed COVID-19. Categorical variables presented as count (%) and continuous variables as mean (standard deviation). All presented as mean/standard deviation. \*mean value of 3 recordings of basal tympanic temperature; \*\* first recorded temperature on hospital admission; \*\*\* maximum temperature self-recorded on app.

**Table 2.** Results from linear mixed effect model for (a) TwinsUK unaffected volunteers; multivariable linear regression for hospitalised cohorts from (b) London and (c) Birmingham with confirmed COVID-19 infection and (d) Community-based “app” cohort with self-reported confirmed COVID-19 infection. Beta coefficients are reported in bold and standard errors in brackets; P-values are Bonferroni adjusted at 5%.

**Table 3.** Results of logistic regression for (b) London Hospital cohort and (c) Birmingham Hospital cohort, adjusted for sex and BMI and (d) Community-based “app” cohort. Odds ratios (ORs) are reported in bold, Confidence Intervals (CI) in brackets. Fever is defined as (b) recorded temperature  $\geq 37.8^{\circ}\text{C}$  and (c) self-reported temperature  $\geq 37.8^{\circ}\text{C}$ ; P-values are not adjusted for multiple testing.

**Table 4.** Results of heritability analysis for (i) basal temperature adjusted for age and sex; (ii) basal temperature adjusted for age, sex and BMI; and (iii) fever, adjusted for age, sex and BMI.

**Figure 1.** ROC curves for adults <65 years of age (orange) and  $\geq 65$  years of age (green); the red star indicates the point characterised by a True Positive Rate of 0.36 and False Positive Rate of 0.01.

Figure 1

