DATA NOTE

The Avon Longitudinal Study of Parents and Children - A resource for COVID-19 research: Questionnaire data capture

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Abstract
The Avon Longitudinal Study of Parents and Children (ALSPAC) is a prospective population-based cohort study which recruited pregnant women in 1990-1992. The resource provides an informative and efficient setting for collecting data on the current coronavirus 2019 (COVID-19) pandemic. In early March 2020, a questionnaire was developed in collaboration with other longitudinal population studies to ensure cross-cohort comparability. It targeted retrospective and current COVID-19 infection information (exposure assessment, symptom tracking and reported clinical outcomes) and the impact of both disease and mitigating measures implemented to manage the COVID-19 crisis more broadly. Data were collected on symptoms of COVID-19 and seasonal flu, travel prior to the pandemic, mental health and social, behavioural and lifestyle factors. The online questionnaire was deployed across the parent and offspring generations between the 9th April and 15th May 2020. 6807 participants completed the questionnaire (2706 original mothers, 1014 original fathers/partners, 2973 offspring (mean age ~28 years) and 114 partners of offspring). Eight (0.01%) participants (4 G0 and 4 G1) reported a positive test for COVID-19, 77 (1.13%; 28 G0 and 49 G1) reported that they had been told by a doctor they likely had COVID-19 and 865 (12.7%; 426 G0 and 439 G1) suspected that they have had COVID-19. Using algorithmically defined cases, we estimate that the predicted proportion of COVID-19 cases fell between 1.03% - 4.19% depending on timing of measurement during the period of reporting.

Data from this first questionnaire will be complemented with at least two more follow-up questionnaires, linkage to health records and results of biological testing as they become available. Data has been released as: 1) a standard dataset containing all participant responses with key sociodemographic factors and 2) as a composite release coordinating data from the existing resource, thus enabling bespoke research across all areas supported by the study.
Keywords
ALSPAC, Children of the 90s, birth cohort study, COVID-19, coronavirus, online questionnaire, mental health

This article is included in the Avon Longitudinal Study of Parents and Children (ALSPAC) gateway.

This article is included in the Coronavirus (COVID-19) collection.

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Introduction

The coronavirus disease 2019 (COVID-19) pandemic is a rapidly developing global health challenge. There is marked heterogeneity in disease prevalence, severity and outcome both within and across populations. In part, this may be driven by the interplay between environmental, social and host factors such as age and pre-existing comorbidities which predispose or protect against infection or modify disease outcomes. Understanding this interplay requires studies with detailed environmental, health, lifestyle, and biological data – ideally measured within the context of longitudinal data and with prospective collection opportunities.

Alongside the health implications of the virus itself, the response to the pandemic is likely to affect health and wellbeing. Mitigation measures have resulted in far-reaching changes to daily activity (exemplified by the nationwide ‘lockdown’ strategy implemented in the UK from 23rd March 2020) which are likely to have an impact on nearly all aspects of work, family life, recreation and, potentially, health. Any adverse effects of these mitigation strategies may themselves be heterogeneous, with certain groups at higher risk of adverse effects. Understanding the effect of mitigation strategies on health and identifying the social, environmental, or other factors which help reduce their impact will be an important part of planning both ongoing and future COVID-19 mitigation strategies as the pandemic develops.

The Avon Longitudinal Study of Parents and Children (ALSPAC) is a unique three-generational study, comprising ‘G0’: the cohort of original pregnant women, the biological father and other carers/partners; ‘G1’: the cohort of index children and ‘G2’: the cohort of offspring of the index children. The study has a wealth of biological, genetic and phenotypic data across these generations[1,2]. ALSPAC has an opportunity to capture information across key parts of the population in light of the COVID-19 pandemic – in particular the contrast between those in higher risk (G0 mean age: ~58years) and lower risk (G1 mean age: ~28yrs) groups. We were well placed to collect data quickly and could be updated and repeated. For these reasons, we chose to use an online only data collection approach for this, restricting our invites to those participants with a valid email address (and coordinated with a systematic communications/outreach campaign). The questionnaire was developed and deployed using REDCap (Research Electronic Data CAPture tools); a secure web application for building and managing online data collection exercises, hosted at the University of Bristol.

Content design

Content was initially developed internally, we then consulted with a network of 16 UK and international longitudinal population studies and partners through a process facilitated by Wellcome (see acknowledgements). This resulted in a core set and a recommended set of questions about health, behaviour, social, economic and environmental impact of COVID-19. Whilst we were able to align many features of the ALSPAC questionnaire and the Wellcome core questionnaire, there are some areas of divergence. This is because the ALSPAC questionnaire was deployed while the Wellcome core questionnaire was still being finalized, and because we chose to use mental health measures that we have used previously to facilitate longitudinal analyses rather than the Wellcome recommended measures. It is worth noting that ALSPAC is the contact point for using the Wellcome questionnaire and we can provide data dictionaries in either REDCap or Qualtrics (via Generation Scotland) on request. This work is therefore part of a coordinated effort to generate and promote a Wellcome Trust supported core questionnaire. This is now complete and available and access to this can be organised through ALSPAC. The questionnaire has not been formally validated, however, extensive testing by our ethics committee, participant advisory group and ALSPAC staff led to clarity of wording and ensuring REDCap functionality worked as expected.

The questionnaire included 4 sections, and captured information on the following:

A. General health, recent travel and seasonal symptoms
- Conditions making people high risk
- Frailty assessed using PRISMA -7
- Regular medications (prescription and over the counter)
- Home country and travel outside that country since October 2019

Wellcome Open Research 2020, 5:127 Last updated: 11 JUN 2020
• Symptoms of COVID-19 and negative control symptoms since October 2019
• Tested/diagnosed with COVID-19

B. Behaviour as a result of COVID-19
• Self-isolation and reasons why
• Behaviour changes prior to lockdown
• Lifestyle changes since lockdown
• Social contacts and methods of communication

C. Impact of the pandemic
• Worries during the pandemic
• Depression assessed using the Short Moods and Feelings questionnaire (SMFQ; 7)
• Anxiety assessed using the General Anxiety Disorder-7 questionnaire (GAD7; 8)
• Well-being assessed using the Warwick-Edinburgh Mental Wellbeing Scales (WEMWBS; 9)

D. About you during the pandemic
• Understanding of official guidance on COVID-19
• Time spent talking/reading about or listening to information about COVID-19
• Living arrangements
• Healthcare worker and keyworker status
• Effect of pandemic on plans to have children (G1 only)
• Free text inviting participants to provide details of other ways they have been affected by the pandemic

The final questionnaire (REDCap PDF) used is available with the associated data dictionary (which includes frequencies of all variables that are available) and both are available as extended data.

Invitation and reminder strategy
Between the 9th and 15th April 2020, all participants (G0, G1 and G1 partners enrolled as part of G2 (children of the Children of the 90s)) for whom we had an active email address were sent an invitation to complete the questionnaire. Participants were not contacted if our administrative database record indicated that they were deceased, had withdrawn from the study, had declined further contact or had declined questionnaires. The questionnaire survey was live on the online platform for just over one month. After 2 weeks, any non-responders were sent a reminder email to complete the questionnaire. In addition, traditional (print, radio, tv) & social media (Facebook, Instagram and Twitter) were used to inform participants that the questionnaire was live, asking them to contact us if they had not received it and to encourage completion. These communication channels were also used to encourage re-engagement of friends and family back into the study. Unlike our standard questionnaires (usually completed annually) we did not provide any incentive for completion; however, we did offer a prize draw (three prizes of £100) for those who completed their questionnaire by 11th May.

Response rate
A total of 12,520 invitations were sent out and responses were received from 6811 participants (overall response rate of 54%). Over 4,000 participants completed the questionnaire within the first week of the invitation (see Figure 1), this represents a snapshot of participant experience around 3 weeks after the start of ‘lockdown’. The questionnaire was closed on 15th May 2020.

Overall, female participants were more likely to respond, and this was particularly true of the younger generation. A potential explanation for the difference in male response rates between younger and older participants we note that G0 fathers included in this data collection exercise are already relatively engaged with the study as they had to enrol in their own right in 2011 as part of our ‘Focus on Fathers’ data collection (previously they were only invited to participate via the mother). The response from G1 males was disappointing and we aim to re-engage this group using targeted social media and other activities in the future. Table 1 summarises the response rate within each group organised by cohort structure.

Characteristics of responders according to key variables that will be released with the complete dataset can be seen in Table 2.

Key results
Participants were asked whether they thought they have had COVID-19. Options were: ‘Yes, confirmed by a positive test’, ‘Yes, suspected by a doctor but not tested’, ‘Yes, my own suspicions’ or ‘No’. Overall 8 respondents reported that they had tested positive to COVID-19 (when combined with participant group this information is potentially disclosive it will therefore be combined with ‘yes, suspected by a doctor’ to create a new category in the released datasets ‘Yes, tested positive or suspected by a doctor’). Table 3 summarises the responses to this question by cohort structure. Due to small numbers we have had to combine the first two categories together for the released dataset.

Menzi and Colleagues recently published a model combining symptoms to predict ‘probable infection’ using data collected from an app-based symptom tracker. This algorithm uses four symptoms: loss of smell and taste, severe or significant persistent cough, severe fatigue and skipped meals (coded as 1 if present and 0 otherwise), together with age and sex (1 male; 0 female). We had slight difference in wording and thus the algorithm (using the same weightings) applied was as follows:

\[-1.32 - (0.01 \times \text{age}) + (0.44 \times \text{sex}) + (1.75 \times \text{loss of smell or taste}) + (0.31 \times \text{new persistent cough}) + (0.49 \times \text{severe fatigue}) + (0.39 \times \text{decreased appetite}).\]
Probable COVID-19 cases were obtained by applying an \( \exp(x)/[1+(\exp(x))] \) transformation and coding values >0.5 as probable cases. We applied this algorithm to our monthly symptom data and the number of predicted ‘cases’ each month are shown in Figure 2. We used Stata v.15.0 and our Stata do file is available as extended data. Peak predicted cases were seen in March (4.19%) with similar proportions seen in all other months (2.10% - 2.38%), except November which was substantially lower (1.03%). Our figures were consistently lower than the 5.36% of responders to the app who were reported as likely being infected by the virus. Table 4 summarises these predicted cases of COVID-19 each month according to self-reported infection status. There is a clear temporal trend such that the proportion of predicted cases increases over time in those who self-reported that they had COVID-19.

Whilst we have presented these results as a mark of the type of analysis one can undertake using these data, we note that these predictions are subject to important assumptions. First, the baseline risk of having COVID-19 (intercept term in the model) is assumed to be the same in the ALSPAC study population as the Menni study population. This assumption may be invalid as there are fewer reported COVID-19 cases in South West England compared to other regions of the UK, potentially over-estimating prevalence in ALSPAC. Secondly, the symptoms we termed ‘new persistent cough’, ‘severe fatigue’ and

<table>
<thead>
<tr>
<th>Cohort Group</th>
<th>Eligible</th>
<th>Responded</th>
</tr>
</thead>
<tbody>
<tr>
<td>G0 Mothers</td>
<td>4590</td>
<td>2706 (59%)</td>
</tr>
<tr>
<td>G0 Fathers/partners</td>
<td>1803</td>
<td>1014 (56%)</td>
</tr>
<tr>
<td>G1 Offspring daughters</td>
<td>3617</td>
<td>2126 (59%)</td>
</tr>
<tr>
<td>G1 Offspring sons</td>
<td>2225</td>
<td>847 (38%)</td>
</tr>
<tr>
<td>G1 Offspring partners (female)</td>
<td>103</td>
<td>63 (61%)</td>
</tr>
<tr>
<td>G1 Offspring partners (male)</td>
<td>182</td>
<td>51 (28%)</td>
</tr>
<tr>
<td>TOTAL</td>
<td>12520</td>
<td>6807 (54%)</td>
</tr>
</tbody>
</table>

1| valid email address, marked as contactable for questionnaires
2| Proportions of those invited (i.e. eligible)
Table 2. Summary of key characteristics for those who responded; n (%) for categorical variables or mean (sd) for continuous variables.

<table>
<thead>
<tr>
<th></th>
<th>Mothers</th>
<th>Fathers/partners</th>
<th>Offspring</th>
<th>Offspring partners</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age (years)</td>
<td>57.9 (4.44)</td>
<td>60.8 (5.17)</td>
<td>27.6 (0.54)</td>
<td>29.9 (4.37)</td>
</tr>
<tr>
<td>Latest BMI</td>
<td>26.3 (5.01)</td>
<td>27.4 (3.98)</td>
<td>24.6 (5.25)</td>
<td>27.1 (4.89)</td>
</tr>
<tr>
<td>Latest Systolic BP</td>
<td>119.4 (14.10)</td>
<td>132.9 (13.77)</td>
<td>115.2 (10.83)</td>
<td>116.3 (12.44)</td>
</tr>
<tr>
<td>Latest Diastolic BP</td>
<td>70.6 (9.33)</td>
<td>77.2 (9.03)</td>
<td>66.7 (7.77)</td>
<td>65.9 (9.97)</td>
</tr>
<tr>
<td>Education level</td>
<td>1385 (53.7%)</td>
<td>674 (70.4%)</td>
<td>1793 (77.8%)</td>
<td>26 (60.5%)</td>
</tr>
<tr>
<td>≥A level</td>
<td>197 (7.4%)</td>
<td>38 (4.1%)</td>
<td>197 (7.4%)</td>
<td>38 (4.1%)</td>
</tr>
<tr>
<td>Ethnicity</td>
<td>2533 (98.4%)</td>
<td>948 (99.2%)</td>
<td>2559 (96.6%)</td>
<td>Not available</td>
</tr>
</tbody>
</table>

Data taken from the most recent clinic that individual attended where available
Data taken from pregnancy questionnaires for G0 and from most recent questionnaire for G1 where available
Data taken from pregnancy questionnaires for all

Table 3. Participant response to whether they have had COVID-19.

<table>
<thead>
<tr>
<th></th>
<th>G0 - parents</th>
<th>G1 – offspring (+partners)</th>
<th>Total</th>
</tr>
</thead>
<tbody>
<tr>
<td>Yes, positive test</td>
<td>4 (0.01%)</td>
<td>4 (0.01%)</td>
<td>8 (0.01%)</td>
</tr>
<tr>
<td>Yes, doctor suspected, no test</td>
<td>28 (0.76%)</td>
<td>48 (1.56%)</td>
<td>76 (1.12%)</td>
</tr>
<tr>
<td>Yes, own suspicions</td>
<td>426 (11.5%)</td>
<td>439 (14.2%)</td>
<td>865 (12.8%)</td>
</tr>
<tr>
<td>No</td>
<td>3229 (87.4%)</td>
<td>2584 (83.8%)</td>
<td>5734 (84.6%)</td>
</tr>
</tbody>
</table>

Figure 2. Predicted cases (% of population) of COVID-19 per month according to symptoms reported for those months using Menni et al algorithm. 
‘decreased appetite’ are assumed to capture the same information as the symptoms used in the Menni study but may in truth have subtly different meaning, leading to either over or under-estimation of prevalence in the ALSPAC study. Thirdly, the association of these symptoms with COVID-19 (fixed effects in the model) is assumed to be the same in the ALSPAC study population as the Menni study population. This assumption might be violated if the pattern of symptom presentation varies in different groups of people, leading to over or under-estimation of prevalence in the ALSPAC study.

**Strengths and limitations of the data**

The primary strengths of this data are the timelines within which the collection occurred, the retrospectively available and prospectively continuing longitudinal collection forming the context for these new data and the potential for cross-cohort comparisons with a set of measures aligned to other UK studies. We believe the timeframe is important, as the data collected here reflects the feelings of the cohort early on in mitigation during a period of stringent lockdown measures. In addition to this, given that the questionnaire was active for over a month it will be important to take date of completion into account for certain analyses, as responses may differ between early- versus late-completers (e.g. late completers experienced an additional month of lockdown measures and have adapted more to the new circumstances). Our second COVID-19 data collection, funded and planned to start at the end of May will provide invaluable comparisons as (at the time of writing) some lockdown features are starting to be relaxed (certain work sectors being encouraged to return to work etc). Through the Wellcome coordinated group developing a core set of questions we will be able to make valuable comparisons with other cohort groups of different ages, backgrounds and from different countries and cultures.

We had an excellent response rate; however, it should be noted that invitations were only sent to those participants for whom we had a valid email address and we have limitations to cover-age. Our usual questionnaire strategy is to first send emails to encourage online completion. We then follow up as part of the reminder process and send paper questionnaires through the post. This online only strategy will have particularly affected G0 mothers who historically have tended to use paper questionnaires more than other sub-groups and for whom we are least likely to hold a current email address. However, the pandemic has led to a number of participants reaching out and getting in touch to provide these details, and indeed to re-engage with the study having dropped out previously.

In some cases, the data recorded is potentially identifiable. We went through each variable one by one and made decisions about whether to combine categories. We have combined categories where we feel the data is at high risk of potential disclosure.

A further key limitation of this data is the reduced response rate in our male G1 participants (28% of all G1 responders were male), even compared to previous data collection exercises in this group where around a third of G1 responders have been male.

Finally, the collection was of course subject to the frequency of COVID-19 infection in the population and sample. Very few participants reported a positive test to COVID-19 and the frequency of algorithmically assigned case status suggests a

<p>| Table 4. Predicted cases of COVID-19 each month using Menni et al. algorithm according to self-reported infection status (note that valid self-report infection status data available for n=6664 which is why the total column is higher than the sum of the other columns). |
|-------------------------------|-------------------------------|-------------------------------|-------------------------------|-------------------------------|-------------------------------|</p>
<table>
<thead>
<tr>
<th>Tested positive (0.01%)</th>
<th>Doctor suspected (1.13%)</th>
<th>Participant suspected (12.7%)</th>
<th>Not had (84.6%)</th>
<th>TOTAL</th>
</tr>
</thead>
<tbody>
<tr>
<td>April 2020 n=153</td>
<td>0</td>
<td>29 (18.9%)</td>
<td>70 (45.6%)</td>
<td>54 (35.3%)</td>
</tr>
<tr>
<td>March 2020 n=271</td>
<td>0</td>
<td>37 (13.7%)</td>
<td>139 (51.3%)</td>
<td>95 (35.1%)</td>
</tr>
<tr>
<td>Feb 2020 n=137</td>
<td>0</td>
<td>3 (2.2%)</td>
<td>56 (40.9%)</td>
<td>78 (56.9%)</td>
</tr>
<tr>
<td>Jan 2020 n=145</td>
<td>0</td>
<td>0</td>
<td>51 (35.2%)</td>
<td>94 (64.8%)</td>
</tr>
<tr>
<td>Dec 2019 n=136</td>
<td>0</td>
<td>1 (0.7%)</td>
<td>40 (29.4%)</td>
<td>95 (69.9%)</td>
</tr>
<tr>
<td>Nov 2019 n=64</td>
<td>0</td>
<td>1 (1.6%)</td>
<td>15 (23.4%)</td>
<td>48 (75.0%)</td>
</tr>
<tr>
<td>Oct 2019 n=166</td>
<td>0</td>
<td>1 (0.6%)</td>
<td>23 (13.9%)</td>
<td>142 (85.5%)</td>
</tr>
</tbody>
</table>
frequency of symptomatic presentation which is consistent with regional estimates\(^1\). This is partly due to UK policy not to test widely at the time that data was being collected but also the fact that the majority of our participants remain in the South-West of England which has had the lowest COVID-19 death rate in the country (and potentially the lowest infection rate\(^2\)). All symptoms were self-reported but linkage to health records and our potential plans to test participants will assist in contributing to the evidence around true symptoms of COVID-19.

**Data availability**

**Underlying data**

ALSPAC data access is through a system of managed open access. The steps below highlight how to apply for access to the data included in this data note and all other ALSPAC data:

1. Please read the ALSPAC access policy\(^1^5\) which describes the process of accessing the data and samples in detail, and outlines the costs associated with doing so.

2. You may also find it useful to browse our fully searchable research proposals database\(^6\), which lists all research projects that have been approved since April 2011.

3. Please submit your research proposal\(^1^7\) for consideration by the ALSPAC Executive Committee. You will receive a response within 10 working days to advise you whether your proposal has been approved.

Please note that a standard COVID-19 dataset will be made available at no charge (see description below); however, costs for required paperwork and any bespoke datasets required additional variables will apply.

**COVID-19 Questionnaire 1 Data File**

Data from the first ALSPAC COVID-19 questionnaire is available in two ways.

1. A freely available standard set of data containing all participants together with key sociodemographic variables (where available) is available on request (see data availability section). Subject to the relevant paperwork being completed (costs may apply to cover administration) this dataset will be made freely available to any bona fide researcher requesting it. Variable names will follow the format covid1\(_{xxxx}\) where \(_{xxxx}\) is a four-digit number. A full list of variables released is available as extended data\(^8\). Frequencies of variable names will follow the format covid1\(_{xxxx}\) where \(_{xxxx}\) is a four-digit number. A full list of variables released is available as extended data\(^8\). Frequencies of variable data and details of any coding/editing decisions and derived variables are also available in the data dictionary (see extended data\(^8\)).

2. Formal release files have been created for G0 mothers, G0 fathers and G1 participants in the usual way and now form part of the ALSPAC resource (Due to the small number of G1 partners contributing we will not be formally releasing this data, however, it may be available on request for specific G2 projects). These datasets (or sections therein) can be requested in the usual way. Variable names will replicate those in 1) above but as each variable in ALSPAC is uniquely defined we have added digits to denote the source of the variable. For example, in dataset 1, the age of the participant at completion (in years) is denoted by covid1\(_{9650}\). In the mother’s dataset this will be denoted by covid1m\(_{9650}\), for fathers/partner this will be covid1p\(_{9650}\) and for the G1 generation it will be covid1p\(_{9650}\). Frequencies for all variables for each participant group are available in the data dictionary in the usual way\(^9\).

Text data and other potentially disclosive information will not be released until they have been coded appropriately. Table 5 describes the data that is withheld at the time of first release. Text from Section C, question 1 and Section D, other are being thematically coded by qualitative researchers, the remainder will be coded as it is required/requested. Data will be incorporated back into both file sets as they become available.

**Extended data**

Open Science Framework: ALSPAC COVID-19 Q1. https://doi.org/10.17605/OSF.IO/ZU8WY\(^1^0\)

This project contains the following extended data:

- ALSPAC COVID Q1 FINAL.pdf (The final questionnaire; REDCap PDF)
- ALSPAC\_COVID1\_varlist.pdf (List of variable names and labels)
- ALSPAC\_CovidQ1\_data dictionary.pdf (Associated data dictionary including frequencies of all variables that are available)
- ALSPAC\_symptom\_algorithm\_14052020.do (Stata script for obtaining probable COVID-19 cases using the Menni algorithm)

Data are available under the terms of the Creative Commons Attribution 4.0 International license (CC-BY 4.0).

**Consent**

Completion of the questionnaire was optional and choosing to complete the questionnaire is considered informed consent for the questionnaire.

Ethical approval for the study was obtained from the ALSPAC Ethics and Law Committee and the Local Research Ethics Committees. Informed consent for the use of data collected via questionnaires and clinics was obtained from participants following the recommendations of the ALSPAC Ethics and Law Committee at the time. Study participants have the right to withdraw their consent for elements of the study or from the study entirely at any time. Full details of the ALSPAC consent procedures are available on the study website\(^1^9\).
Table 5. Data from questions that will not be released until coded.

<table>
<thead>
<tr>
<th>Question number</th>
<th>Question text</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Section A</strong></td>
<td></td>
</tr>
</tbody>
</table>
| 1               | Please tell us the type of:  
|                 | • Organ transplant  
|                 | • Diabetes  
|                 | • Heart disease or heart problems  
|                 | • Other lung condition  
|                 | • Cancer  
|                 | • Condition affecting the brain and nerves  
|                 | • Psychiatric disorder  
|                 | Please can you tell us why your immune system is weakened? |
| 3               | For each medication:  
|                 | • Name of medication  
|                 | • Amount  
|                 | • How often  
|                 | • Reason for taking |
| 5               | Which country do you live in?  
|                 | If travelled outside home country:  
|                 | • Country and region/city/resort  
|                 | • Date arrived/arrived  
|                 | • Purpose of trip |
| 6               | What kind of other medical attention did you access?  
|                 | What other medication did you take? |
| 9               | Date first told had COVID-19 |
| **Section B**   |               |
| 1               | When did you start self-isolating?  
|                 | Other reason for self-isolating |
| 2               | Other reason for changing normal day to day behaviour |
| **Section C**   |               |
| 1               | What other reason causing worry |
| **Section D**   |               |
| 7               | Other type of accommodation lived in |
| Other           | Is there anything else you would like to tell us about how the pandemic has affected you? |

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Professor George Davey Smith (University of Bristol)
Professor Peter Vickerman (University of Bristol)
Dr Robert Aldridge (University College London)
Professor David Porteous (University of Edinburgh, Gen Scot)
Honorary Professor Rosie McEachan (University of Bradford, Born in Bradford)
Professor Goncalo Abecasis (Regeneron)
Professor Naomi Allen (University of Oxford, UKBB)
Dr Mary de Silva (Wellcome)
Dr Bruna Galobardes (Wellcome)
Dr Claire Steves (Kings College London, TwinsUK)
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