

1 **Omicron-associated changes in SARS-CoV-2 symptoms in the United Kingdom**

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10 **Running title:** Omicron-associated changes in symptoms

11

1 **ABSTRACT**

2 **Background:** The SARS-CoV-2 Delta variant has been replaced by the highly transmissible
3 Omicron BA.1 variant, and subsequently by Omicron BA.2. It is important to understand
4 how these changes in dominant variants affect reported symptoms, while also accounting
5 for symptoms arising from other co-circulating respiratory viruses.

6 **Methods:** In a nationally representative UK community study, the COVID-19 Infection
7 Survey, we investigated symptoms in PCR-positive infection episodes vs. PCR-negative study
8 visits over calendar time, by age and vaccination status, comparing periods when the Delta,
9 Omicron BA.1 and BA.2 variants were dominant.

10 **Results:** Between October-2020 and April-2022, 120,995 SARS-CoV-2 PCR-positive episodes
11 occurred in 115,886 participants, with 70,683 (58%) reporting symptoms. The comparator
12 comprised 4,766,366 PCR-negative study visits (483,894 participants); 203,422 (4%)
13 reporting symptoms. Symptom reporting in PCR-positives varied over time, with a marked
14 reduction in loss of taste/smell as Omicron BA.1 dominated, maintained with BA.2
15 (44%/45% 17 October 2021, 16%/13% 2 January 2022, 15%/12% 27 March 2022). Cough,
16 fever, shortness of breath, myalgia, fatigue/weakness and headache also decreased after
17 Omicron BA.1 dominated, but sore throat increased, the latter to a greater degree than
18 concurrent increases in PCR-negatives. Fatigue/weakness increased again after BA.2
19 dominated, although to a similar degree to concurrent increases in PCR-negatives.
20 Symptoms were consistently more common in adults aged 18-65 years than in children or
21 older adults.

22 **Conclusions:** Increases in sore throat (also common in the general community), and a
23 marked reduction in loss of taste/smell, make Omicron harder to detect with symptom-
24 based testing algorithms, with implications for institutional and national testing policies.

25

26 **Key words:** SARS-CoV-2, Omicron, symptoms

27

1 Introduction

2 Highly-transmissible SARS-CoV-2 Omicron variants, BA.1 and BA.2, emerged and become
3 dominant at the end and start of 2021 and 2022, coincident with other winter respiratory
4 viruses circulating in the Northern hemisphere, changes in symptomatology may influence
5 clinical and testing policy. Experimental and clinical data suggest Omicron has less impact on
6 the lower respiratory tract, leading to less severe disease[1–7], with the variant-defining
7 mutations potentially also affecting other symptoms.

8

9 We used the UK Covid-19 Infection Survey, a nationally representative longitudinal
10 household study[8], to investigate if SARS-CoV-2 infection symptoms have changed with the
11 Omicron variants. We compared the probability of reporting any symptoms, as well as the
12 probability of reporting specific symptoms in both SARS-CoV-2 PCR-positive infection
13 episodes and comparator PCR-negative study visits focusing on time periods when the Delta
14 variant (described previously only to August-2021[9]), Omicron BA.1 and Omicron BA.2
15 were dominant in the UK[10].

16

17 Methods

18 This analysis was based on SARS-CoV-2 PCR tests of nose and throat swabs taken regularly
19 between 1-October-2020 and 23-April-2022 from participants in the Office for National
20 Statistics (ONS) Covid Infection Survey (CIS) (ISRCTN21086382,
21 [https://www.ndm.ox.ac.uk/covid-19/covid-19-infection-survey/protocol-and-information-](https://www.ndm.ox.ac.uk/covid-19/covid-19-infection-survey/protocol-and-information-sheets)
22 [sheets](https://www.ndm.ox.ac.uk/covid-19/covid-19-infection-survey/protocol-and-information-sheets)). The survey has invited private households to enrol on a continuous basis, selected
23 at random from address lists and previous surveys to provide a representative UK sample,

1 described in detail in the Appendix of [8]. Participant characteristics and representativeness
2 are presented in detail in the Appendix of [9], illustrating the sample broadly represents the
3 wider population. Following verbal agreement to participate, a study worker visited each
4 household to take written informed consent, which was obtained from parents/carers for
5 those 2-15 years; those aged 10-15 years provided written assent. Those <2 years were not
6 eligible, to avoid asking parents to swab babies and very young children. Ethical approval
7 was provided by the South Central Berkshire B Research Ethics Committee (20/SC/0195).
8
9 Individuals were asked about demographics, symptoms, contacts and relevant behaviours
10 (<https://www.ndm.ox.ac.uk/covid-19/covid-19-infection-survey/case-record-forms>).
11 Participants ≥ 12 years self-collected nose and throat swabs following study worker
12 instructions, to reduce transmission risks. Parents/carers took swabs from children 2-11
13 years. At the first visit, participants were asked for consent for optional follow-up visits
14 every week for the next month, then monthly from enrolment. While participants were
15 offered the option of a single visit, 99% of participants participated in longitudinal sampling;
16 study samples were obtained regularly, irrespective of the presence or absence of
17 symptoms. **Table S1** provides a detailed description of the number of visits per participant,
18 median 18 (IQR 12-21) visits between 1-October-2020 and 23-April-2022.
19
20 Swabs were analysed at national Lighthouse Laboratories at Milton Keynes and Glasgow
21 using identical methodology. PCR for three SARS-CoV-2 genes (N protein, S protein and
22 ORF1ab) was performed using the Thermo Fisher TaqPath RT-PCR COVID-19 kit, and
23 analysed using UgenTec FastFinder 3.300.5, with an assay-specific algorithm and decision
24 mechanism that allows conversion of amplification assay raw data into test results with

1 minimal manual intervention. Samples are called positive if at least the N gene and/or
2 ORF1ab are detected. Although S gene cycle threshold (Ct) values are determined, S gene
3 detection alone is not considered sufficient to call a sample positive by the assay
4 manufacturer[8].

5

6 The presence of 12 specific symptoms in the previous seven days was elicited at each visit
7 from the start of the survey (cough, fever, myalgia, fatigue/weakness, sore throat, shortness
8 of breath, headache, nausea, abdominal pain, diarrhoea, loss of taste, loss of smell), as was
9 whether participants thought they had (unspecified) symptoms compatible with COVID-19.
10 Positive response to any of these questions defined “symptomatic” cases. Four additional
11 symptoms (runny nose, trouble sleeping, loss of appetite, wheezing) were added from 29
12 September-2021; as these were not elicited throughout the survey, they were considered
13 separately and not used to define symptomatic cases.

14

15 We grouped repeated PCR-positive tests into infection “episodes”[11], and included the first
16 positive study test in each episode in analysis (details in Supplementary Methods). Each
17 positive episode was characterised as wild-type/Delta/Omicron BA.2-compatible if the S-
18 gene was ever detected (by definition, with N/ORF1ab/both), or as Alpha- or Omicron BA.1-
19 compatible if positive at least once for ORF1ab+N (and never for the S-gene), otherwise
20 “other” (N-only/ORF1ab-only) depending on calendar period (**Fig.1A**). Symptom presence
21 was defined as reported symptoms at any visit within [0,+35] days of the first PCR-positive
22 test in each infection episode (i.e. spanning [-7,+35] days given the question timeframe), to
23 allow for the random sampling leading to pre-symptomatic identification of some
24 individuals, who only reported symptoms subsequently.

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As a comparator, we initially considered all visits with negative PCR tests, and then, following a previous analysis to August-2021[9], excluded visits where symptoms could plausibly be related to ongoing effects of COVID-19 or long COVID, where there was a high pre-test probability of a new COVID-19 infection that had not been detected in the study, or where symptoms were likely driven by recent vaccination (details in Supplementary Methods).

Generalised additive models (binomial distribution with complementary log-log link) were fitted to estimate the percentage of PCR-positive infection episodes and PCR-negative visits that were symptomatic, and the percentage of symptomatic PCR-positive infection episodes and symptomatic PCR-negative visits reporting each symptom separately. Models adjusted simultaneously for calendar time (smoothing spline), age (smoothing spline), sex and ethnicity (white vs non-white). From 29-September-2021 onwards fitted models with an additional interaction between age and time are used to present differences in symptoms by age.

To explore differences between Delta, Omicron BA.1, and Omicron BA.2 infections by vaccination status and infection/re-infection, we restricted PCR-positives to those occurring after 29-September-2021 and classified S-gene negatives occurring after 1-December-2021 as Omicron BA.1-compatible (34,576 infections, 20,345 [59%] symptomatics), and S-gene positives 29-September-2021 to 2-January-2022 as Delta-compatible (14,318 infections, 9,030 [63%] symptomatics) and 30-January-2022 to 23-April-2022 as Omicron BA.2-compatible (34,796 infections, 22,591 [65%] symptomatics) (excluding S-gene positives 3-29

1 January-2022 as both Delta and Omicron BA.2 infections occurred during this period and
2 genetic sequences were not available for all PCR-positives). Descriptive analyses are
3 presented of differences in symptom presence/absence and specific symptoms by variant,
4 vaccination status and infection episode. Comparisons by vaccine status are restricted to
5 participants ≥ 18 years to reduce confounding arising from lower vaccination rates in those
6 < 18 years.

7
8 All analyses were run using R 3.6.1. Generalised additive models were fitted using mgcv 1.8-
9 31; example code is provided in the Supplementary Methods. Figures were produced using
10 ggplot2 3.1.0 and cowplot 1.1.0.

12 Results

13 Between October-2020 and April-2022, 120,995 PCR-positive episodes occurred in 115,886
14 participants (median 44 years, IQR 24-61), 70,683 (58%) with reported symptoms.
15 8898/120,995 (7%) were re-infections (**Fig.S1**), 4244 (48%) with reported symptoms. The
16 comparator comprised 4,766,366 PCR-negative study visits (483,894 participants, median 55
17 years, IQR 36-68); 203,422 (4%) with reported symptoms.

18
19 While Omicron BA.1 infections dominated (19-December-2021 to 26-February-2022, when
20 $> 50\%$ of PCR-positive results were S-gene negative), the percentage of PCR-positive
21 infection episodes with reported symptoms was lower compared to much of the previous
22 time period when the Delta variant dominated (6-June-2021 to 18-December-2021,
23 **Fig.1B/C**). Reporting any symptoms increased again after Omicron BA.2 became the

1 dominant variant (27-February-2022 onwards, when >50% of PCR-positive results were S-
2 gene positive). For both Omicron BA.1 and BA.2 the mean number of symptoms reported in
3 PCR-positive infection episodes was lower than with Delta, but was higher with BA.2 than
4 BA.1. Changes in the percentage reporting any symptoms at PCR-negative visits, and the
5 mean number of symptoms reported at PCR-negative visits, were much smaller over these
6 time periods, with very slight increases from October-2021 onwards likely due in part to
7 other seasonal infections.

8
9 For specific symptoms, amongst symptomatic PCR-positive infection episodes, there was a
10 marked decline in reported loss of taste/smell for both Omicron variants, BA.1 and BA.2,
11 from high levels during the period when Delta dominated, e.g. from 44%/45% on 17
12 October-2021 (approximately peak Delta, **Fig.1A**), to 16%/13% on 2-January-2022
13 (approximately peak BA.1) with only very small changes thereafter, e.g. to 15%/12% on 27-
14 March-2022 (approximately peak BA.2). Although loss of taste/smell was also more
15 uncommon with Alpha than Delta, it was even more uncommon with Omicron BA.1/BA.2
16 than Alpha (**Fig.1D**). Loss of taste/smell remained extremely uncommon in symptomatic
17 PCR-negative visits throughout (**Fig.1D**).

18
19 There were concurrent smaller, but significant, declines in symptomatic PCR-positive
20 infection episodes with reported cough, fever, fatigue/weakness, myalgia, shortness of
21 breath and headache during December-2021, as Omicron BA.1 dominated (**Fig.1E/F/G**). As
22 Omicron BA.2 became dominant, cough and to a lesser extent fever and fatigue/weakness
23 increased again, while shortness of breath, myalgia, and headache remained at similar levels
24 to those observed with BA.1 (**Fig.1E/F/G**). The main changes in the percentages of

1 symptomatic PCR-negative visits where these specific symptoms were reported was a
2 substantial increase in cough in October-2021, which then decreased in January-2022 from
3 52% to 36%, before increasing again to 48% by 23-April-2022 (**Fig.1G**), and increases in
4 headache over December-2021 (from 30% to 35%) and in fatigue/weakness over March-
5 2022 (from 20% to 26%) (**Fig.1E**).

6
7 In contrast to these declines in other symptoms as Omicron BA.1 dominated, sore throat
8 became more commonly reported with BA.1 and increased further with BA.2, from 46% to
9 56% in symptomatic PCR-positive infection episodes during December-2021, increasing
10 further to 64% by April-2022. Similarly to cough, sore throat became more commonly
11 reported at PCR-negative visits during October-2021, if anything dropping slightly in
12 January-2022 from 43% to 33% before increasing again to 42% by 23-April-2022 (**Fig.1G**).
13 These changes were smaller in symptomatic PCR-negatives than symptomatic PCR-positives,
14 i.e., were insufficient to explain Omicron-associated increases in sore throat.

15
16 Gastrointestinal symptoms were reported infrequently in symptomatic PCR-positive
17 infection episodes regardless of variant, and were reported at similar frequencies at PCR-
18 negative visits (**Fig.S2**). Reporting of runny nose generally followed reporting of sore throat,
19 whereas other symptoms generally declined with Omicron BA.1/BA.2 (**Fig.S2**).

20
21 In those aged 18 or older, differences in symptoms comparing Delta to Omicron infections,
22 including fewer cases with loss of taste/smell and more with sore throat, were broadly
23 similar across all vaccination statuses (**Fig.2, Fig.S3**) (1,304 (2%), 606 (1%), 14,706 (22%) and
24 49,981 (75%) of PCR-positive infection episodes occurred in those unvaccinated or

1 vaccinated once, twice or three times respectively; full split by variant and evidence of
2 symptoms in **Table S2**). Similarly, changes in symptoms by variant were also relatively
3 unaffected by whether the PCR-positive infection episode was the first infection (91%) vs.
4 reinfection (9%) (**Fig.3, Fig.S4**). However, overall, symptoms were less commonly reported
5 in subsequent infections occurring from 29-September-2021 onwards (50%), compared to
6 first infections during this time period (63%), but specific symptoms were reported at
7 broadly similar frequencies in participants who were symptomatic in PCR-positive first and
8 subsequent infections with Delta and Omicron BA.1 and BA.2 variants.

9

10 There were differences in reported symptoms with these different variants by age when
11 comparing reported symptoms at the peaks of the Delta, BA.1 and BA.2 waves (**Fig.4,**
12 **Fig.S5**). Adults aged 18-65 years were more likely to report the presence of any symptoms
13 than children or adults >65 years. There was generally no evidence of difference in reporting
14 the presence of any symptoms between Delta and BA.2, but there was a lower probability
15 of reporting any symptoms with BA.1 across most ages. However, the mean number of
16 symptoms reported with both BA.1 and BA.2 were generally lower across the ages
17 compared to Delta, with the exception of the youngest and oldest for which there was no
18 evidence of difference in the mean number of symptoms between BA.1 and Delta, but a
19 higher mean number of symptoms for BA.2 vs Delta. Symptoms were less likely to be
20 reported in PCR-positive infection episodes in children than younger adults, even more so
21 with Omicron BA.1 than Delta infections and BA.2 (**Fig.S6**), whereas symptoms were most
22 likely to be reported at PCR-negative visits in children, in particular cough and fever.

23

1 Loss of taste or smell was most commonly reported with Delta infections in adults aged 18-
2 70 years, but at lower levels in older adults, and rarely in younger children; it was only seen
3 at low levels regardless of age with Omicron BA.1/BA.2 infections. Variations in the
4 percentage of symptomatic participants reporting most other specific symptoms across ages
5 were broadly similar before vs after Omicron BA.1 dominated, but slightly higher
6 percentages of symptomatic PCR-positive infection episodes in participants over 70 years
7 reported fever, headache, fatigue/weakness and muscle ache/myalgia after Omicron
8 BA.1/BA.2 dominated (**Fig.4**). Most specific symptoms were reported less frequently at
9 infections in young children than adolescents/young adults regardless of the dominating
10 variant, excepting fever which was reported significantly more with Omicron BA.1 and BA.2
11 infections in young children than adolescents/young adults, particularly for BA.2 (**Fig.S6**).
12
13 The net result of changes in the symptom profile, overall and by age, was that fever and
14 cough became most strongly associated with PCR-positivity in those reporting symptoms
15 after Omicron BA.2 became dominant, adjusting for age, sex and ethnicity (see
16 Supplementary Methods) (**Fig.S7**). Although far less strongly associated than during the
17 period when Delta was the main variant, loss of taste was still the fourth most strongly
18 associated symptom after Omicron BA.2 dominated, with fatigue/weakness also strongly
19 associated. These same four symptoms were also most strongly associated with PCR-
20 positivity when Omicron BA.1 dominated. Sore throat was positively associated with PCR-
21 positivity during the BA.2 dominant period, and to a slightly lesser degree with PCR-
22 positivity during the BA.1 dominant period, while in contrast, sore throat was less likely to
23 occur in symptomatic PCR-positives compared to symptomatic PCR-negatives in the Delta
24 period.

1

2 Discussion

3 In this study of predominantly mild community-based infection, overall Omicron BA.1 and
4 BA.2 were associated with less loss of taste, loss of smell, shortness of breath, myalgia,
5 fatigue/weakness and headache, but more sore throat, compared with Delta. The overall
6 probability of reporting any symptoms was similar for Delta and BA.2, but lower for BA.1
7 regardless of age, while the mean number of symptoms reported was generally lower for
8 both BA.1 and BA.2 compared to Delta across ages, although higher overall for BA.2 than
9 BA.1. However, this was driven by symptomatology in adults; in the youngest and oldest
10 participants, there was no evidence of difference in the percentage reporting any symptoms
11 between BA.2 and Delta, and a higher mean number of symptoms were reported with BA.2
12 in the very youngest and oldest compared to both BA.1 and Delta.

13

14 In PCR/lateral flow antigen-positive cases, the ZOE study, which relies on volunteers
15 reporting symptoms daily using an app, found a lower median number of symptoms
16 reported in infections from 28-November-2021 to 17-January-2022 (predominantly Omicron
17 BA.1) than 1-June to 27-November-2021 (predominantly Delta) matched by age, sex and
18 ethnicity in those who had had a second or third vaccine[12], with less loss of smell and
19 more sore throat being reported with Omicron BA.1, as in our study. The major strength of
20 our study is that regular PCR testing was undertaken in all participants at all visits
21 irrespective of symptoms. This provides a representative sample of PCR-negative visits
22 without SARS-CoV-2 infection for comparison with symptom rates in PCR-positives. This is
23 important because some symptoms reported in PCR-positive infections could be due to co-

1 infections with other circulating respiratory viruses. Therefore, although our study does not
2 specifically test for other viruses, we can estimate whether changes seen with Omicron BA.1
3 and BA.2 differ from underlying trends in the general population (**Fig.1D-G**), supporting
4 much of the increase in sore throat being attributable to Omicron rather than other
5 infections. We are also able to demonstrate large shifts in symptoms reported at PCR-
6 negative visits over time, with concurrent increases in cough and sore throat in October-
7 2021 likely reflecting other respiratory viruses. We also note that the probability of
8 reporting any symptoms as well as specific symptoms, varied considerably during the
9 periods when specific variants dominated, potentially reflecting how the survey captures
10 more infections earlier on when positivity is rising, and more later on as positivity is
11 decreasing[13]. We compared rates at the peak of each dominating variant to capture
12 similar phases of the epidemic, as well as considering how these changed over time.

13
14 Intriguingly, we found that the differences between variants in the probability of reporting
15 specific symptoms in symptomatic PCR-positives persisted regardless of vaccination status
16 or whether the infection was the first or subsequent, while the probability of reporting
17 symptoms was smaller for reinfections compared to first infections. A limitation is that this
18 analysis is of unadjusted percentages, and therefore the lack of observed differences by
19 vaccination status within a variant could be at least partly due to confounding with age, as
20 well as other factors such as previous infection which could lead to choosing not to get
21 vaccinated or to only get one vaccine (only 3% of the infections included in this analysis).
22 However, most symptoms were reported similarly in adults aged 18 to around 60-70 years
23 (**Fig.4**).

24

1 Other limitations of our study include the fact that we cannot have certainty in determining
2 reinfections given the data available; however, estimated reinfections were infrequent (7%),
3 even once Omicron dominated (11%) and symptom profiles were broadly similar in first and
4 subsequent infections from 29-September-2021. Another limitation is that the study does
5 not collect data on healthcare provider visits, hospitalizations, or death, to allow analysis of
6 the severity of Omicron infections beyond reported symptoms. The ZOE study found lower
7 self-reported hospitalisation rates with infections occurring during the Omicron BA.1-
8 dominant vs Delta-dominant period, and shorter duration of symptoms[12], and several
9 other studies have documented lower hospitalisation rates with Omicron BA.1[14–17].
10
11 Increases in sore throat (also commonly reported at symptomatic PCR-negative visits), and
12 the marked reduction in the previously highest specificity symptoms, namely loss of
13 taste/smell, present challenges for testing algorithms. Previously during periods when wild-
14 type, Alpha and Delta variants dominated, fever, cough or loss of taste/smell have been
15 shown to offer a good balance between sensitivity and specificity for detecting SARS-CoV-2
16 infections[9]. In the UK, for much of the pandemic to date, any of these four symptoms
17 formed a basis for the general public accessing PCR testing. However, changes in symptoms
18 with Omicron mean that symptom-based screening for testing is now much more difficult,
19 and have resulted in much broader criteria for symptoms suggestive of COVID being
20 proposed[18], albeit with likely decreased specificity. In conclusion, changes in SARS-CoV-2
21 infection symptoms mean that Omicron is harder to detect with symptom-based testing
22 algorithms with implications for institutional and national testing policies.

23

24

1 **NOTES**

2 **Contributors:** This specific analysis was designed by ASW, K-DV, KBP, PCM, NS, DWE, TH, DC,
3 TEAP. K-DV conducted the statistical analysis of the survey data. K-DV, NS, PCM, ASW
4 drafted the manuscript. All authors contributed to interpretation of the study results, and
5 revised and approved the manuscript for intellectual content. K-DV is the guarantor and
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21 Research.Support@ons.gov.uk or visit the SRS website.

22 **Disclaimer:** The lead authors affirm that the manuscript is an honest, accurate, and
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1 References

- 2 1. Kozlov M. Omicron's feeble attack on the lungs could make it less dangerous. 2022.
3 Available at: <https://www.nature.com/articles/d41586-022-00007-8>. Accessed 10
4 January 2022.
- 5 2. Diamond M, Halfmann P, Maemura T, et al. The SARS-CoV-2 B.1.1.529 Omicron virus
6 causes attenuated infection and disease in mice and hamsters. *Rev* **2021**; :1–27.
- 7 3. Willett BJ, Grove J, Maclean OA, et al. The hyper-transmissible SARS-CoV-2 Omicron
8 variant exhibits significant antigenic change, vaccine escape and a switch in cell entry
9 mechanism Brian. *medRxiv* **2022**; :1–59. Available at:
10 <https://www.medrxiv.org/content/10.1101/2022.01.03.21268111v1.full.pdf>.
- 11 4. Meng B, A.T.M FI, Abdullahi A, Goonawardane N. SARS-CoV-2 Omicron spike
12 mediated immune escape and tropism shift. *bioRxiv* **2022**; Available at:
13 <https://www.medrxiv.org/content/10.1101/2022.01.03.21268111v1.full.pdf>.
- 14 5. Bentley EG, Kirby A, Sharma P, et al. SARS-CoV-2 Omicron-B.1.1.529 Variant leads to
15 less severe disease than Pango B and Delta variants strains in a mouse model of
16 severe COVID-19. *bioRxiv* **2021**; :1–16.
- 17 6. McMahan K, Giffin V, Tostanoski LH, Chung B. Reduced Pathogenicity of the SARS-
18 CoV-2 Omicron Variant in Hamsters. *bioRxiv* **2022**;
- 19 7. Peacock TP, Brown JC, Zhou J, et al. The SARS-CoV-2 variant , Omicron , shows rapid
20 replication in human primary nasal epithelial cultures and efficiently uses the
21 endosomal route of entry . *bioRxiv* **2022**;
- 22 8. Pouwels KB, House T, Pritchard E, et al. Community prevalence of SARS-CoV-2 in
23 England from April to November, 2020: results from the ONS Coronavirus Infection
24 Survey. *Lancet Public Heal* **2021**; 6:e30–e38.

- 1 9. Vihta K-D, Pouwels KB, Peto T, et al. Symptoms and SARS-CoV-2 positivity in the
2 general population in the UK. *Clin Infect Dis* **2021**; :ciab945. Available at:
3 <https://doi.org/10.1093/cid/ciab945>.
- 4 10. UK Office for National Statistics. Coronavirus (COVID-19) Infection Survey, UK: 7
5 January 2022. Available at: [https://www.gov.uk/government/publications/tfc-](https://www.gov.uk/government/publications/tfc-children-and-transmission-update-paper-17-december-2020)
6 [children-and-transmission-update-paper-17-december-2020](https://www.gov.uk/government/publications/tfc-children-and-transmission-update-paper-17-december-2020). Accessed 9 January
7 2022.
- 8 11. Office for National Statistics. Coronavirus (COVID-19) Infection Survey: characteristics
9 of people testing positive for COVID-19, UK: 30 March 2022. *Ons* **2022**; :1–13.
- 10 12. Menni C, Valdes AM, Polidori L, et al. Symptom prevalence, duration, and risk of
11 hospital admission in individuals infected with SARS-CoV-2 during periods of omicron
12 and delta variant dominance: a prospective observational study from the ZOE COVID
13 Study. *Lancet* **2022**; 399:1618–1624.
- 14 13. ONS. Coronavirus (COVID-19) latest insights: Infections. 2022. Available at:
15 [https://www.ons.gov.uk/peoplepopulationandcommunity/healthandsocialcare/condi-](https://www.ons.gov.uk/peoplepopulationandcommunity/healthandsocialcare/conditionsanddiseases/articles/coronaviruscovid19latestinsights/infections#:~:text=Download%20the%20data-,The%20percentage%20of%20people%20testing%20positive%20for%20COVID-19%20continued,%25%20in%20the%20late)
16 [tionsanddiseases/articles/coronaviruscovid19latestinsights/infections#:~:text=Downl](https://www.ons.gov.uk/peoplepopulationandcommunity/healthandsocialcare/conditionsanddiseases/articles/coronaviruscovid19latestinsights/infections#:~:text=Download%20the%20data-,The%20percentage%20of%20people%20testing%20positive%20for%20COVID-19%20continued,%25%20in%20the%20late)
17 [oad%20the%20data-,The%20percentage%20of%20people%20testing%20positive%20for%20COVID-19%20continued,%25%20in%20the%20late](https://www.ons.gov.uk/peoplepopulationandcommunity/healthandsocialcare/conditionsanddiseases/articles/coronaviruscovid19latestinsights/infections#:~:text=Download%20the%20data-,The%20percentage%20of%20people%20testing%20positive%20for%20COVID-19%20continued,%25%20in%20the%20late). Accessed 13 May 2022.
- 19 14. Wolter N, Jassat W, Walaza S, et al. Early assessment of the clinical severity of the
20 SARS-CoV-2 omicron variant in South Africa: a data linkage study. *Lancet* **2022**;
21 399:437–446. Available at: [http://dx.doi.org/10.1016/S0140-6736\(22\)00017-4](http://dx.doi.org/10.1016/S0140-6736(22)00017-4).
- 22 15. Nyberg T, Ferguson NM, Nash SG, et al. Comparative analysis of the risks of
23 hospitalisation and death associated with SARS-CoV-2 omicron (B.1.1.529) and delta
24 (B.1.617.2) variants in England: a cohort study. *Lancet* **2022**; 399:1303–1312.

1 Available at: [http://dx.doi.org/10.1016/S0140-6736\(22\)00462-7](http://dx.doi.org/10.1016/S0140-6736(22)00462-7).

2 16. Ulloa AC, Buchan SA, Daneman N, Brown KA. Early estimates of SARS-CoV-2 Omicron
3 variant severity based on a matched cohort study, Ontario, Canada. medRxiv **2022**;

4 Available at:

5 [https://www.medrxiv.org/content/10.1101/2021.12.24.21268382v2%0Ahttps://www](https://www.medrxiv.org/content/10.1101/2021.12.24.21268382v2%0Ahttps://www.medrxiv.org/content/10.1101/2021.12.24.21268382v2.abstract)
6 [w.medrxiv.org/content/10.1101/2021.12.24.21268382v2.abstract](https://www.medrxiv.org/content/10.1101/2021.12.24.21268382v2.abstract).

7 17. Lewnard JA, Hong VX, Patel MM, Kahn R, Lipsitch M, Tartof SY. Clinical outcomes
8 associated with Omicron (B.1.1.529) variant and BA.1/BA.1.1 or BA.2 subvariant
9 infection in southern California. medRxiv **2022**;

10 18. UK Health Security Agency. What to do if you have symptoms of a respiratory
11 infection including COVID-19, or a positive COVID-19 test. 2022. Available at:

12 [https://ukhsa.blog.gov.uk/2022/04/01/what-to-do-if-you-have-symptoms-of-a-](https://ukhsa.blog.gov.uk/2022/04/01/what-to-do-if-you-have-symptoms-of-a-respiratory-infection-including-covid-19-or-a-positive-covid-19-test/)
13 [respiratory-infection-including-covid-19-or-a-positive-covid-19-test/](https://ukhsa.blog.gov.uk/2022/04/01/what-to-do-if-you-have-symptoms-of-a-respiratory-infection-including-covid-19-or-a-positive-covid-19-test/).

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1 **Figure legends**

2 **Figure 1. (A) Variants and (B)-(G) symptoms in those testing positive and negative for**

3 **SARS-CoV-2 over time in the UK.** Panel A shows the number of PCR-positive infection

4 episodes that were S-gene negative (Alpha-compatible 20 December 2020 to 5 June 2021;

5 Omicron BA.1-compatible 19 December 2021 to 26 February 2022) and S-gene positive

6 (Delta-compatible 6 June 2021 to 18 December 2021; Omicron BA.2-compatible from 27

7 February 2022 onwards). Vertical lines indicate periods when new variants came to

8 dominate based on gene positivity patterns (>50% of PCR-positives): wild type before 20

9 December 2020, then Alpha before 5 June 2021, then Delta before 19 December 2021 then

10 Omicron BA.1 before 27 February 2022; Omicron BA.2 became the dominant variant

11 afterwards. Panels B and C show the probability of reporting symptoms and the number of

12 symptoms (out of the 12 elicited throughout the study period) of all PCR-positive infection

13 episodes and all PCR-negative comparator visits. Panels D-G show the probability of specific

14 symptoms in symptomatic PCR-positive infection episodes and in symptomatic PCR-negative

15 comparator study visits, after adjustment for age, sex, ethnicity (presented at the reference

16 category age 45, male, white).

17 **Figure 2. Percentage of PCR-positives reporting symptoms by variant and by vaccination**

18 **status.** Note: Restricting to aged 18 or older. Reporting of any evidence of symptoms as well

19 as specific symptoms in symptomatic PCR-positives from 29 September 2021 onwards. Not

20 adjusted for other factors, see **Fig.4** for adjusted effect of age. Unvaccinated=before first

21 vaccination at index positive test or never vaccinated, first vaccine= 21 days after first

22 vaccination to 13 days after second, second vaccine=14 days after second vaccination to 13

23 days after third; third vaccine=14 days after third vaccination to 13 days after fourth. Fourth

24 vaccination data not shown as less than 100 infections with evidence of symptoms (**Table**

1 **S2).** The unvaccinated and first vaccine group represent only 3% of infections; these
2 participants are potentially more likely to have been previously infected (as infection may
3 have impacted subsequent vaccine uptake), and previous infection is associated with fewer
4 reported symptoms (**Fig.3**).

5 **Figure 3. Percentage of PCR-positives reporting symptoms by variant and infection/re-**
6 **infection** Note: reporting of any evidence of symptoms as well as specific symptoms in
7 symptomatic PCR-positives from 29 September 2021 onwards. Not adjusted for other
8 factors, see **Fig.4** for adjusted effect of age.

9 **Figure 4. By age, estimated percentage of PCR-positive infection episodes and comparator**
10 **PCR-negative study visits reporting symptoms and mean number of symptoms at the**
11 **peaks of Delta, Omicron BA.1 and Omicron BA.2 waves** Note: model estimates are shown
12 for reporting of any evidence of symptoms as well as specific symptoms in symptomatic
13 PCR-positive infection episodes and comparator PCR-negative study visits on 17 October
14 2021 (Delta), 2 January 2022 (when Omicron BA.1-compatible infections represented the
15 highest proportion of PCR-positives), and 27 March 2022 (when Omicron BA.2 was the
16 dominant variant). The panels in the first row show the probability of reporting symptoms
17 and the number of symptoms (out of the 12 elicited throughout the study period) in all PCR-
18 positive infection episodes and all PCR-negative comparator visits from 29 September 2021
19 onwards, estimated at three reference categories, 17 October 2021, 2 January 2022, and 27
20 March 2022. The remaining panels show the probability of reporting specific symptoms in
21 symptomatic PCR-positive infection episodes and in symptomatic PCR-negative comparator
22 study visits at these reference categories. All are adjusted for calendar date, age (allowing
23 for effect modification by calendar date by including an interaction between calendar date
24 and age), sex (reference category male), ethnicity (reference category white). See **Fig.S3** for
25 other symptoms.

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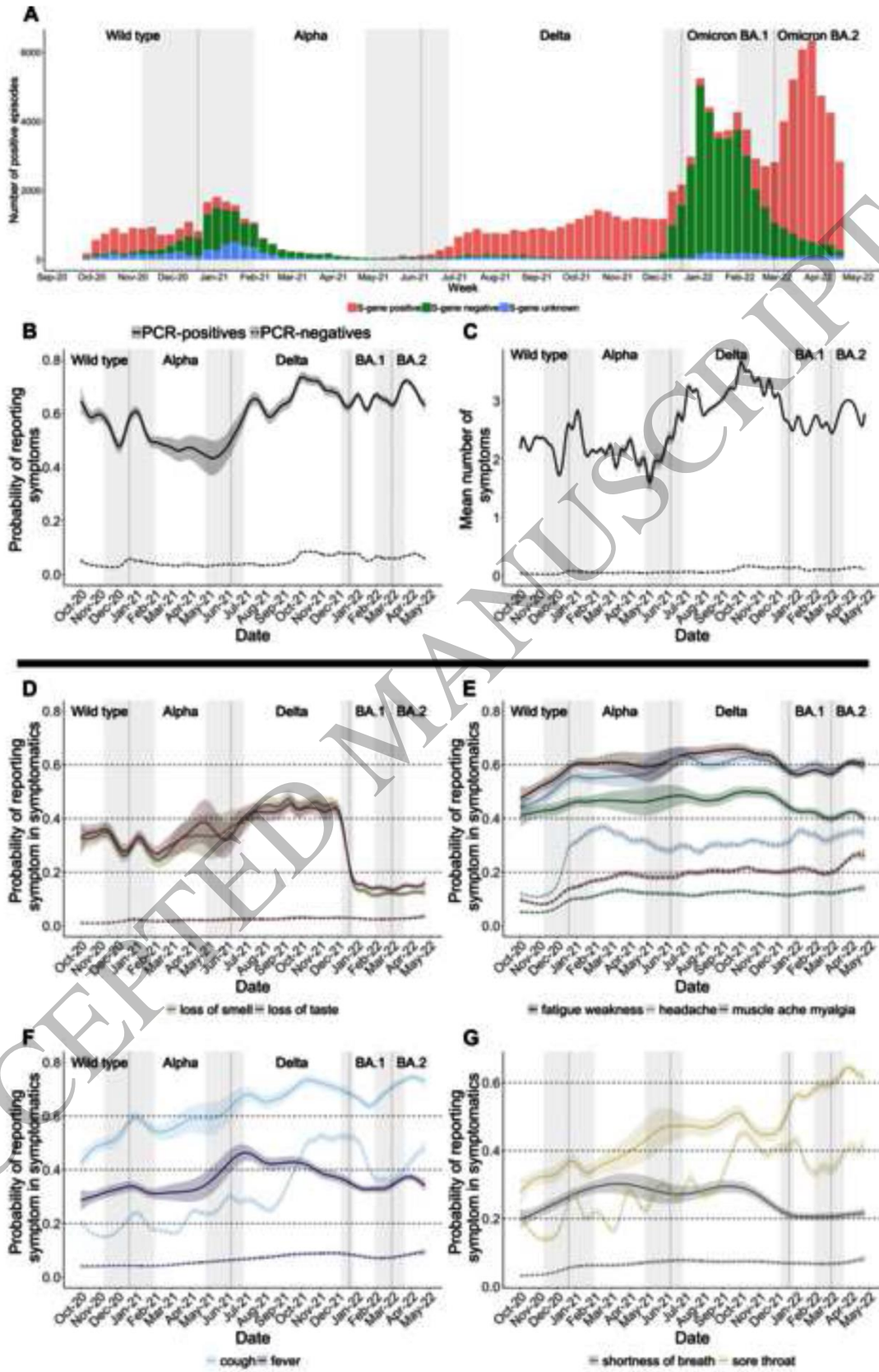


Figure 1
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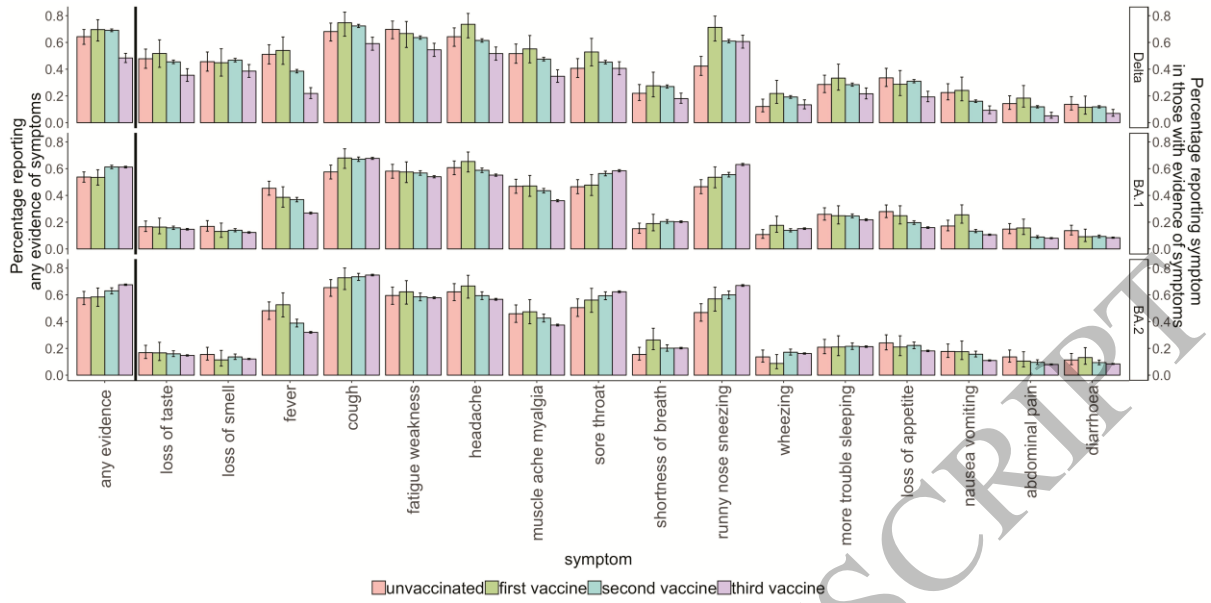


Figure 2
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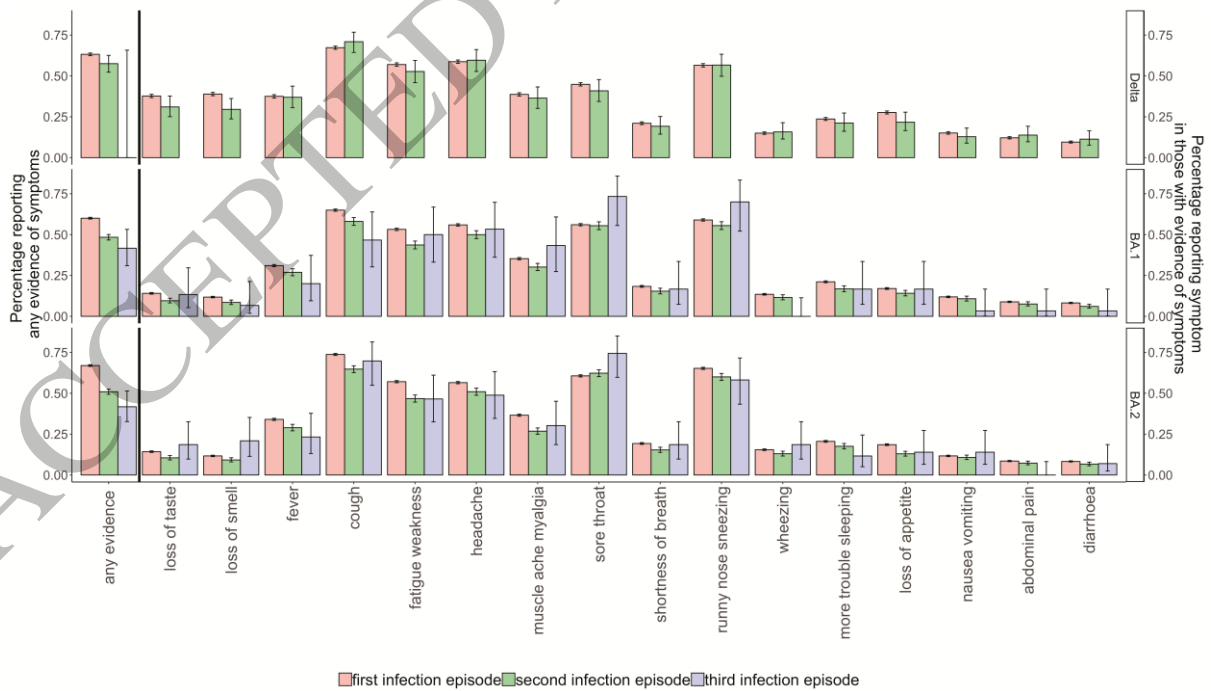
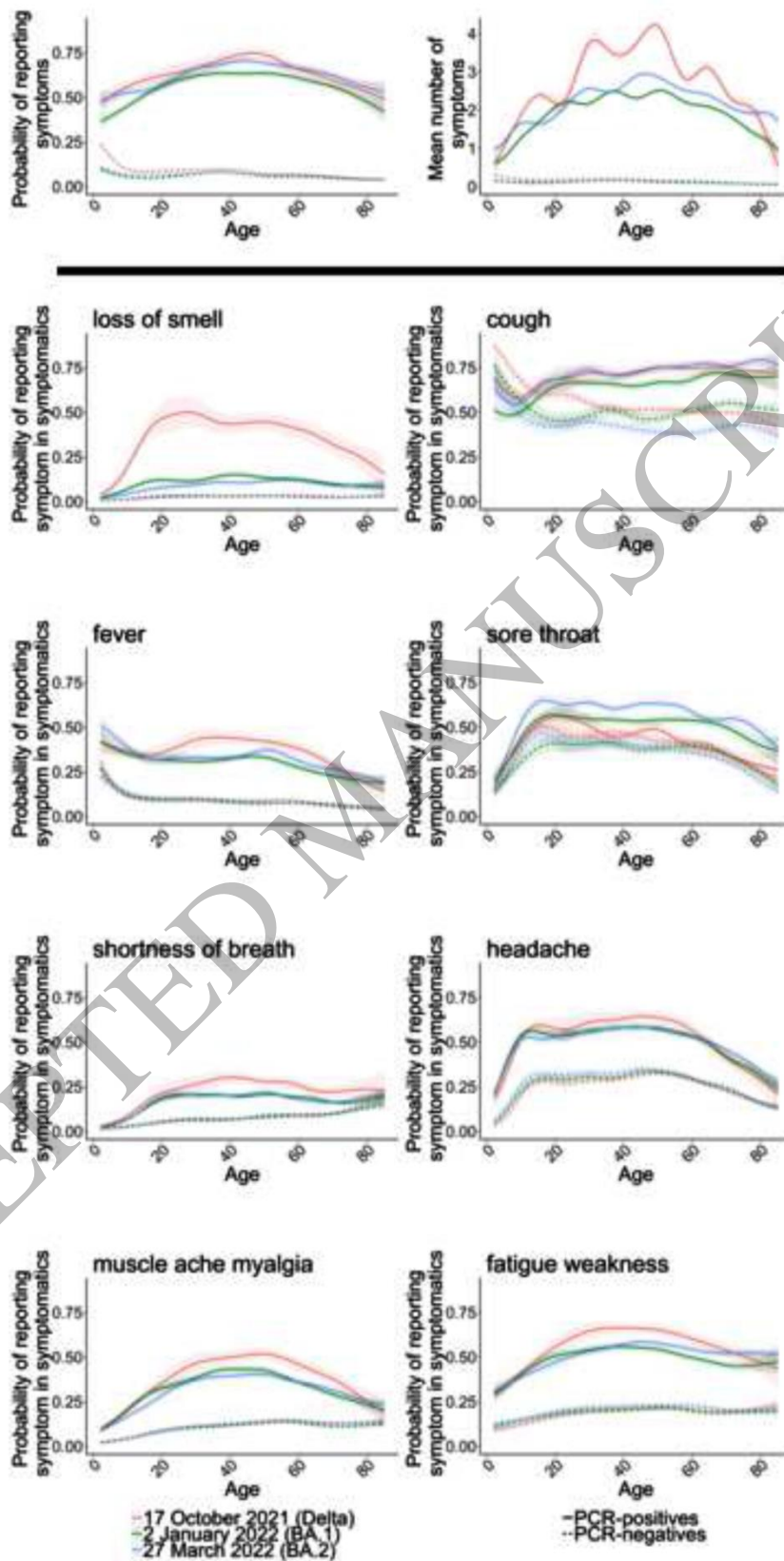


Figure 3
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Figure 4
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