



Weekly Influenza and COVID-19 Surveillance graphs

UKHSA publishes a weekly national influenza and COVID-19 surveillance report which summarises the information from the surveillance systems which are used to monitor influenza, COVID-19 and other seasonal respiratory viruses in England.

Additional figures based on these surveillance systems are included in this slide set.

The figures presented in this slide set are based on data from week 34 (between 22 August and 28 August 2022).



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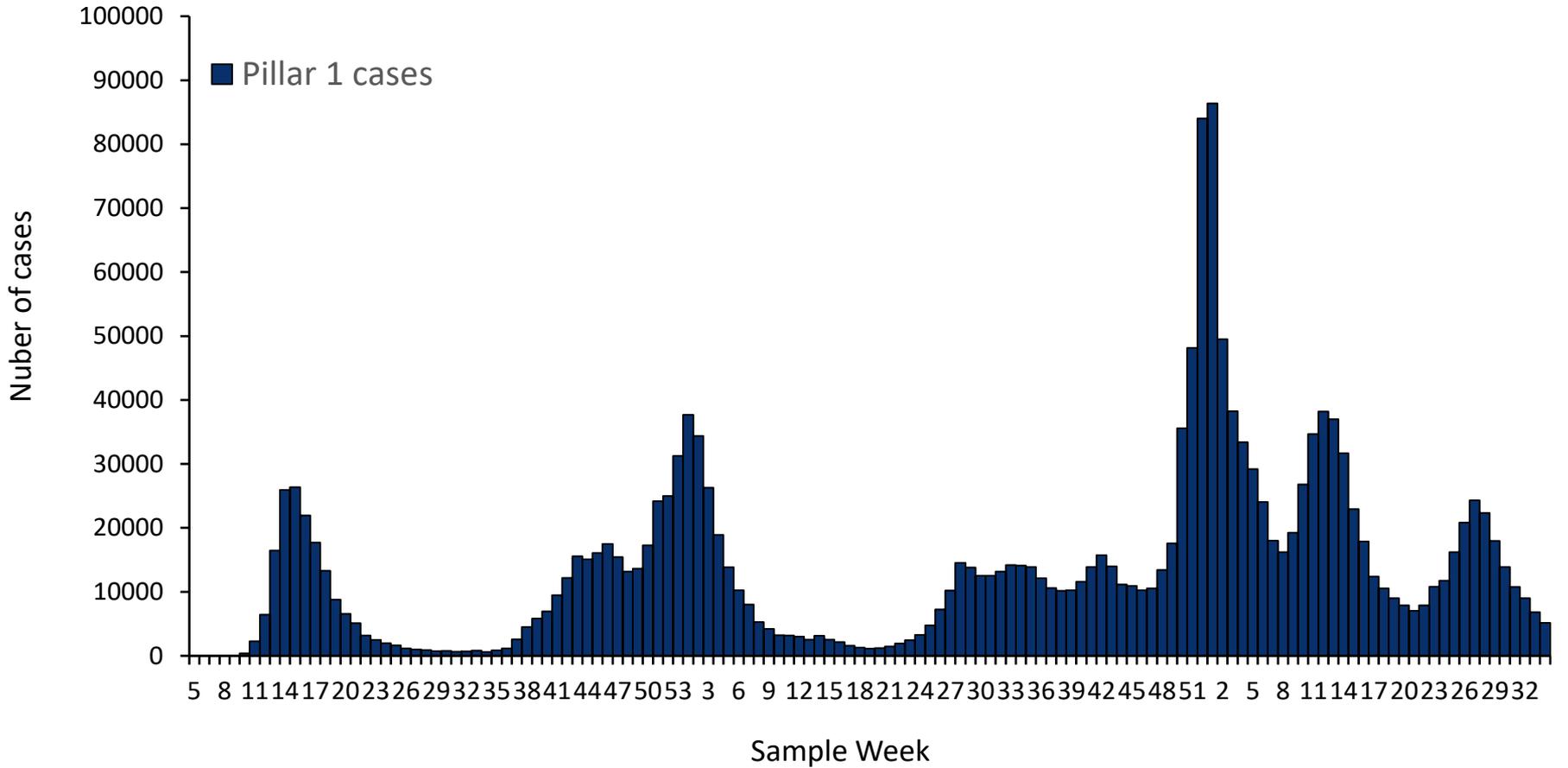
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COVID-19 Pandemic Overview



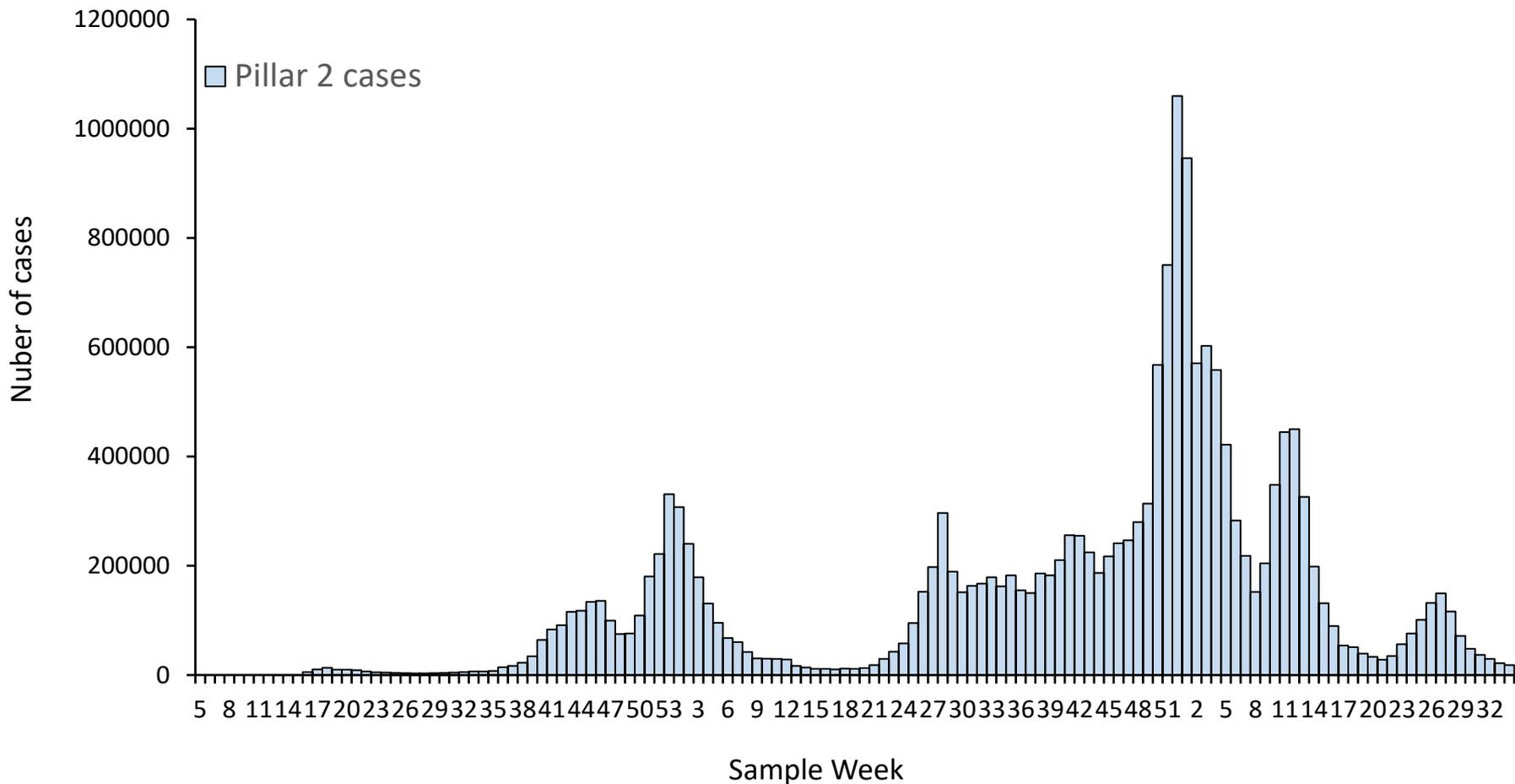
Confirmed COVID-19 episodes tested under Pillar 1, by sample week, since week

5 2020



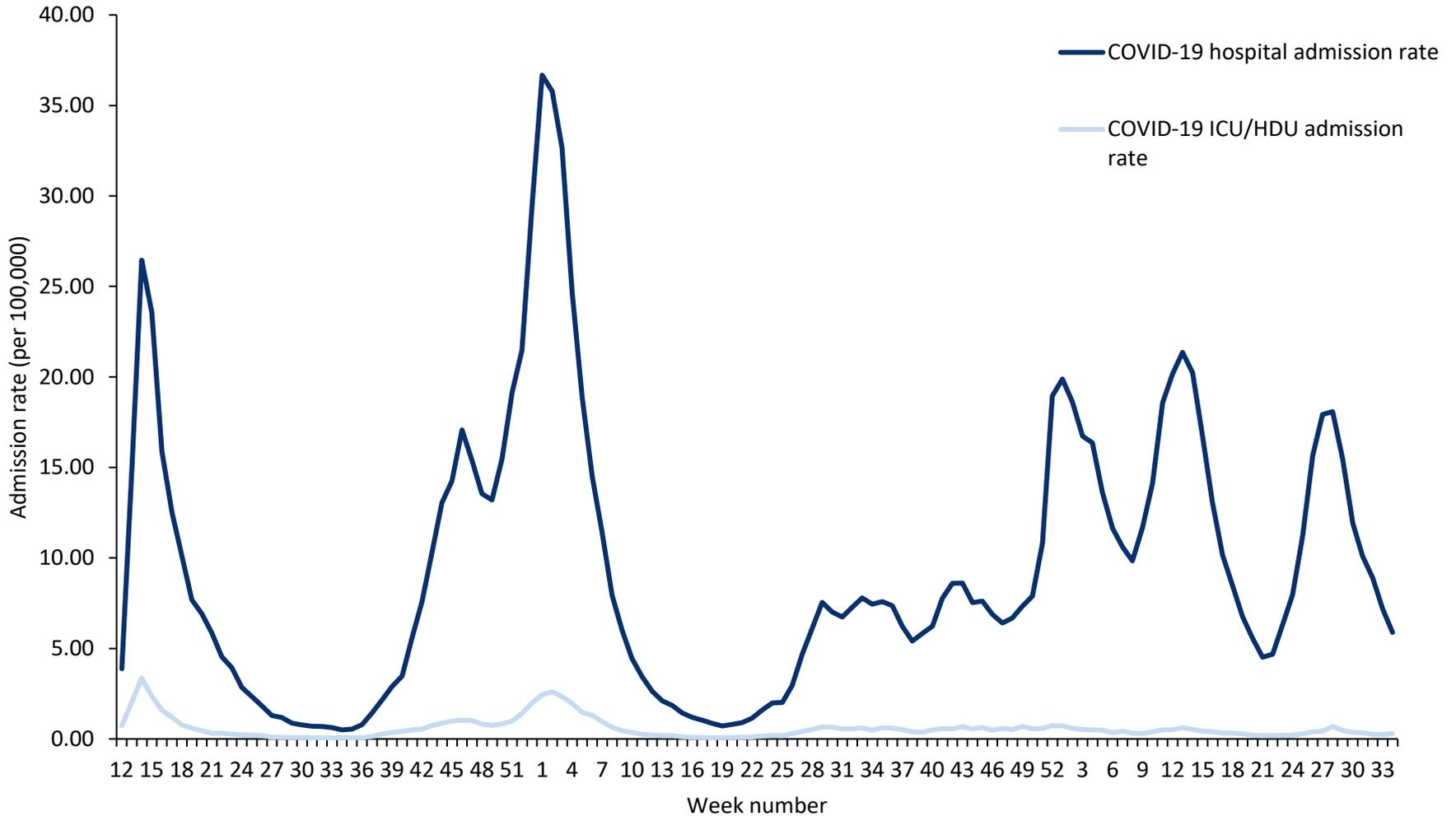


Confirmed COVID-19 episodes tested under Pillar 2, by sample week, since week 5 2020



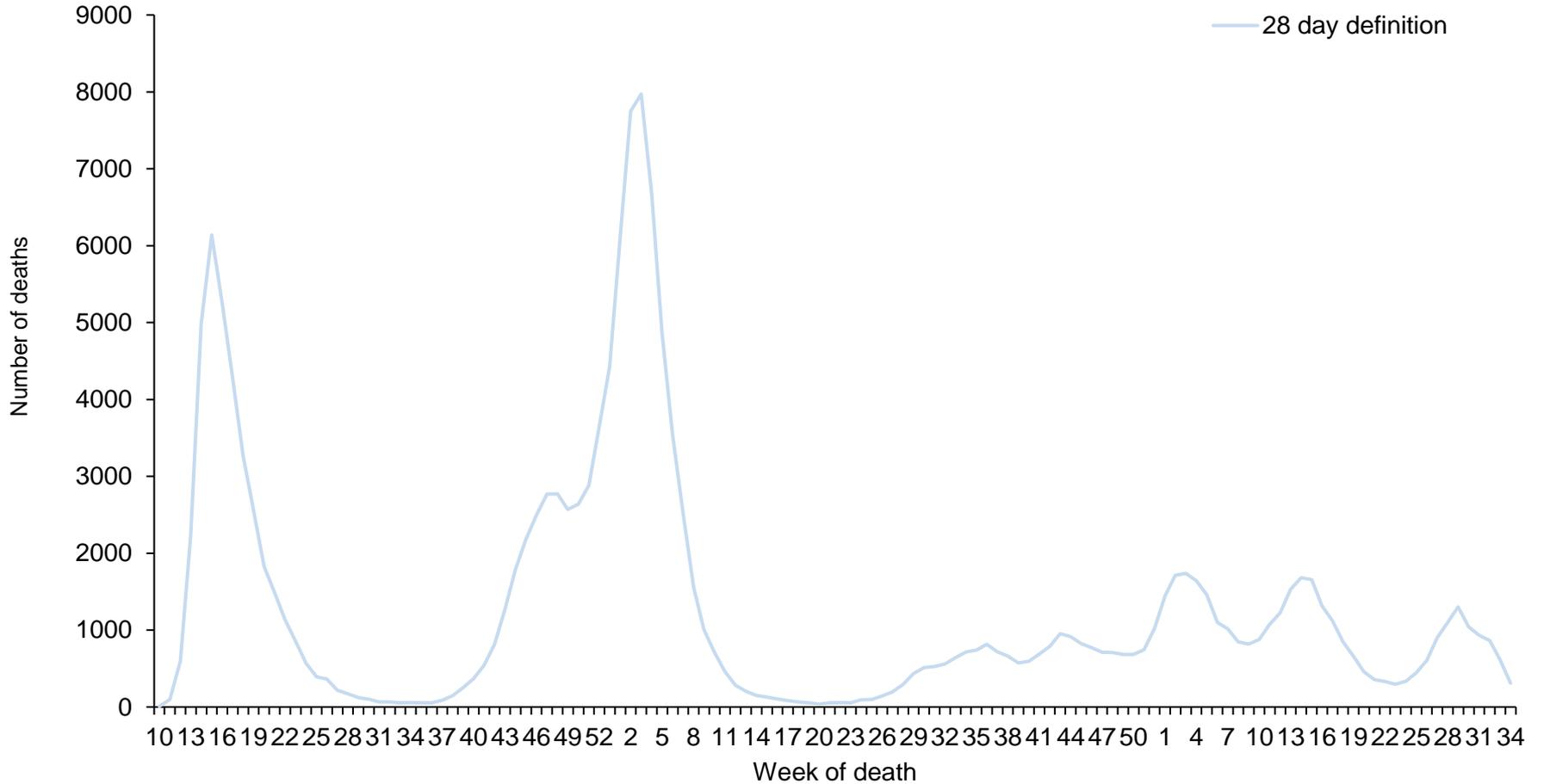


Weekly overall hospital and ICU/HDU admission rates per 100,000 of new COVID-19 cases reported through SARI Watch, England since week 12 2020





Number of deaths since week 10 2020 by week of death and time since laboratory confirmation of COVID-19, England



Since 1 April 2022, free universal symptomatic and asymptomatic testing for the general public in England is no longer available, as outlined in the plan for [living with COVID-19](#). Data should be interpreted in the context of this change to testing.</



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Confirmed COVID-19 episodes in England

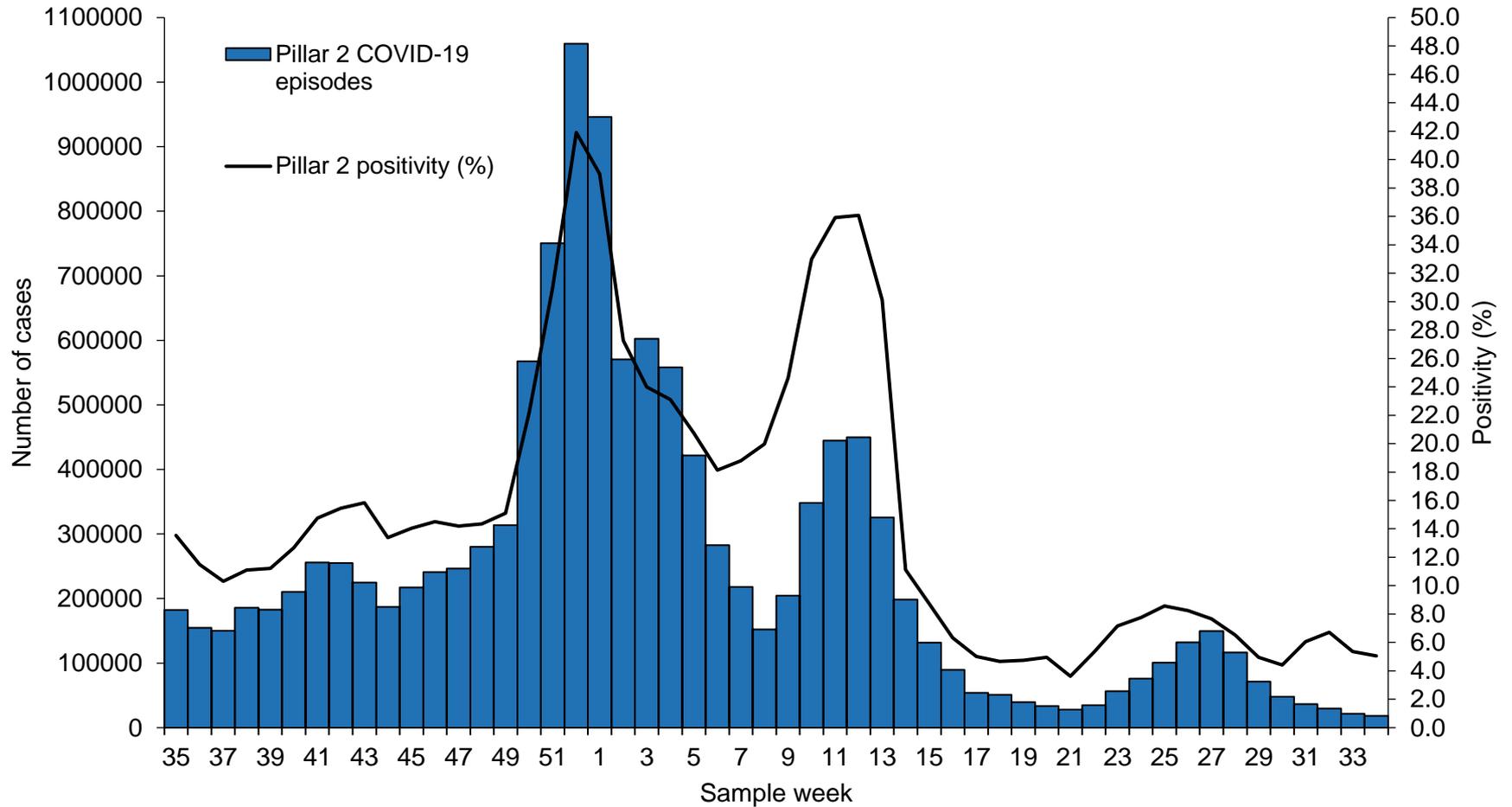


Data Information

- From the week 32 report onwards, case rates have been updated to use the latest ONS population estimates for mid-2020. Previously case rates were calculated using the mid-2019 population estimates
- From 11 January 2022 the requirement for [confirmatory PCR testing in individuals who test positive using a lateral flow device was temporarily removed](#).
- Rates by ethnicity and IMD quantile will continue to be presented using the mid-2019 estimates, until the mid-2020 estimates become available.
- From 31 January 2022, UKHSA moved all COVID-19 case reporting in England to use a new episode-based definition which includes possible reinfections. Each infection episode is counted separately if there are at least 91 days between positive test results (PCR or LFD). Each infection episode begins with the earliest positive specimen date. Further information can be found on the [UK COVID-19 dashboard](#).
- Since 1 April 2022, free universal symptomatic and asymptomatic testing for the general public in England is no longer available, as outlined in the plan for [living with COVID-19](#). As such, there will be a reduction in the reporting of data obtained through Pillar 2 from April 2022 onwards. Data in this report should be interpreted in the context of this change to testing. [Public health guidance](#) remains in place for cases and their close contacts

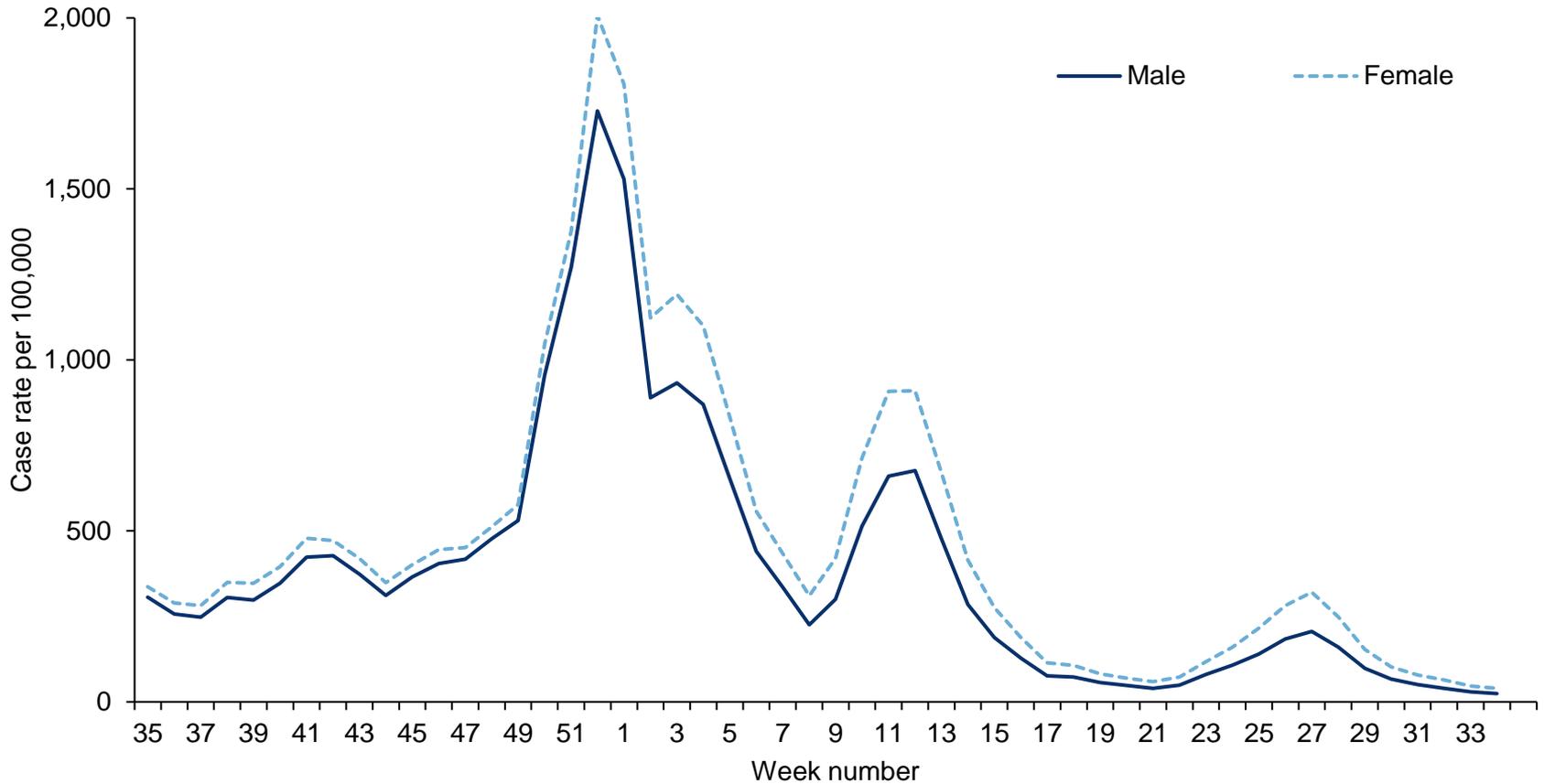


Confirmed COVID-19 episodes tested under Pillar 2, based on sample week with overall weekly PCR positivity for Pillar 2 (%)



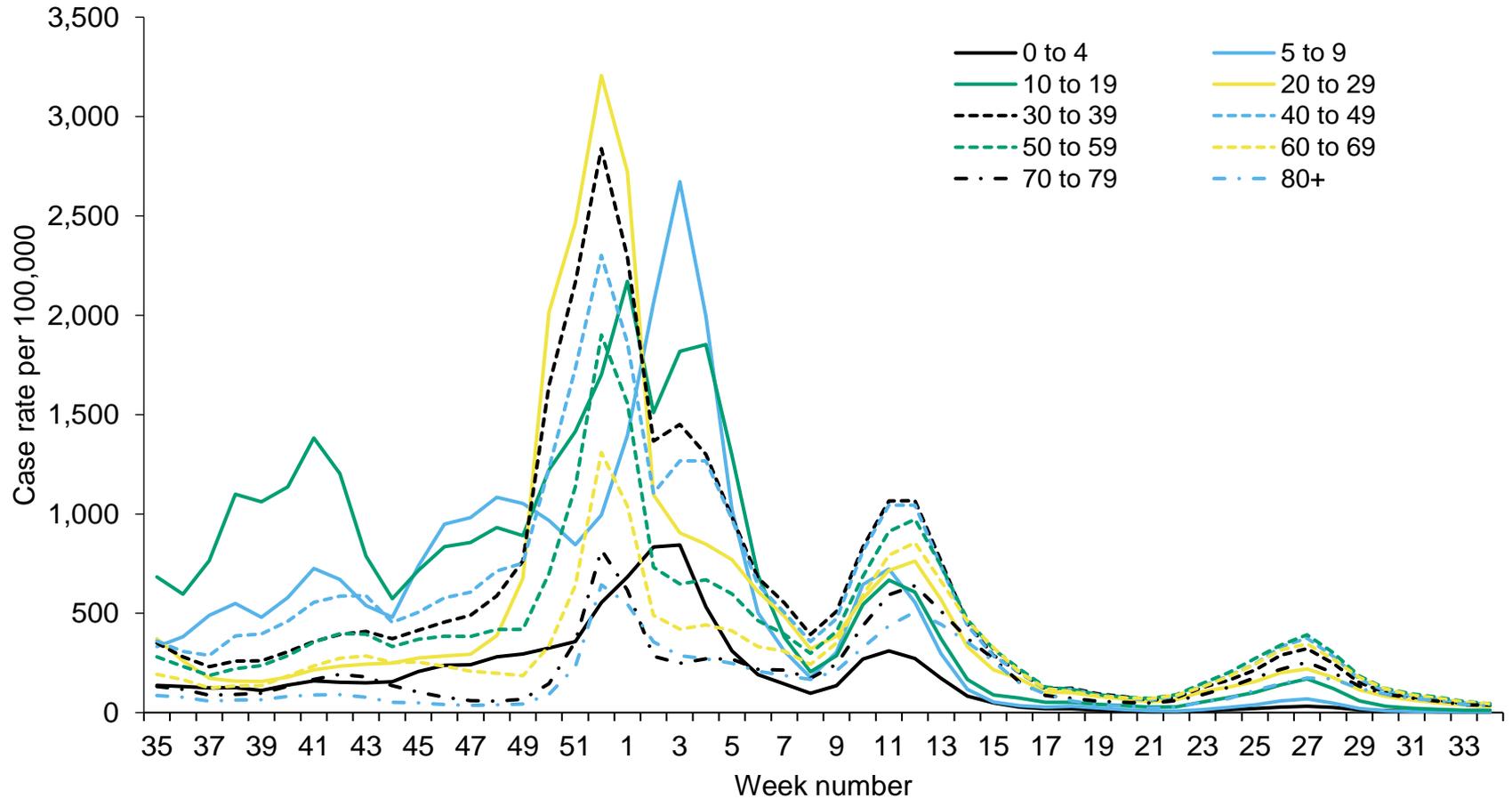


Weekly confirmed COVID-19 case rates per 100,000, by episode, tested under Pillar 2, by sex



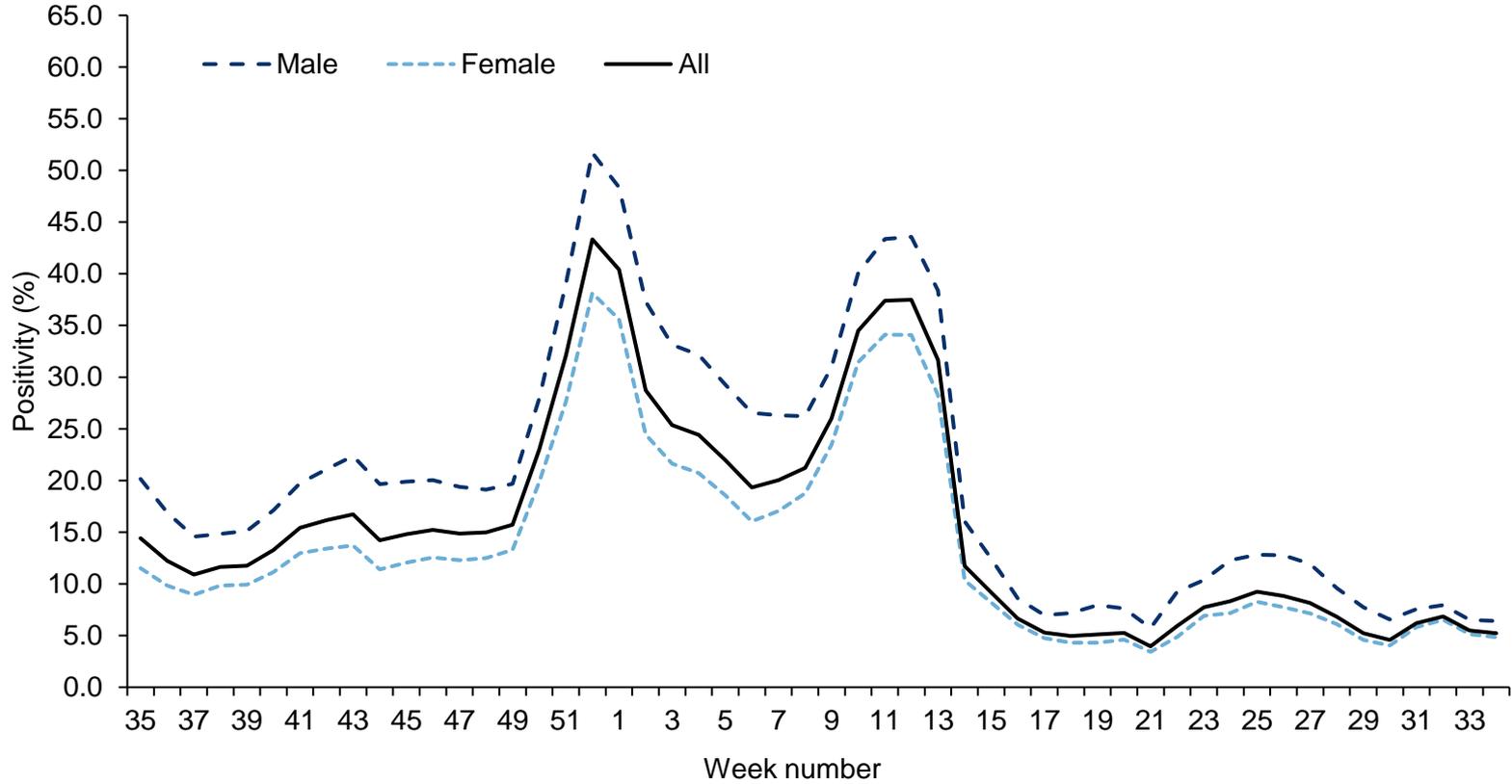


Weekly confirmed COVID-19 case rates per 100,000, by episode, tested under Pillar 2, by age group



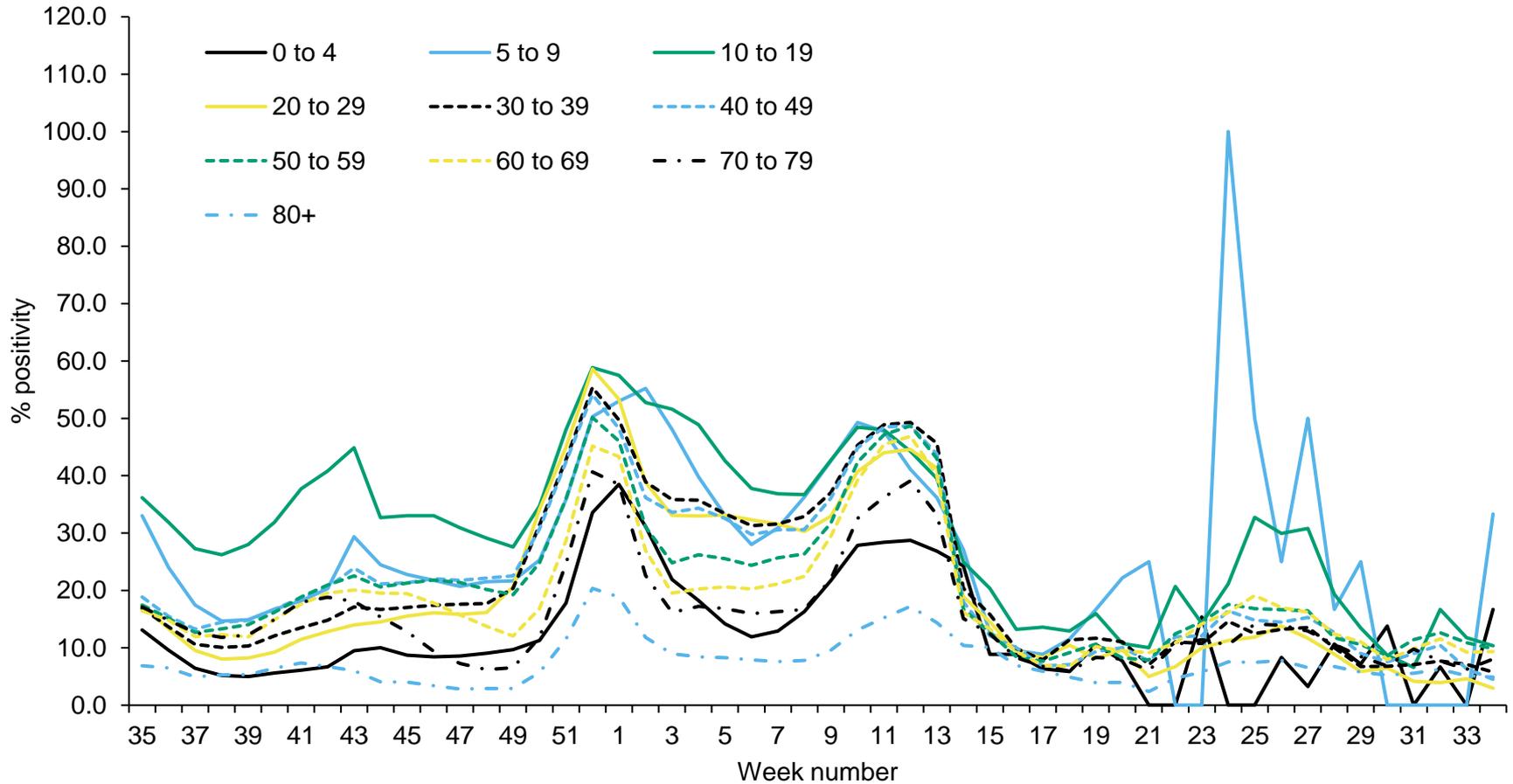


Weekly PCR positivity (%) of confirmed COVID-19 cases tested overall and by sex under Pillar 2



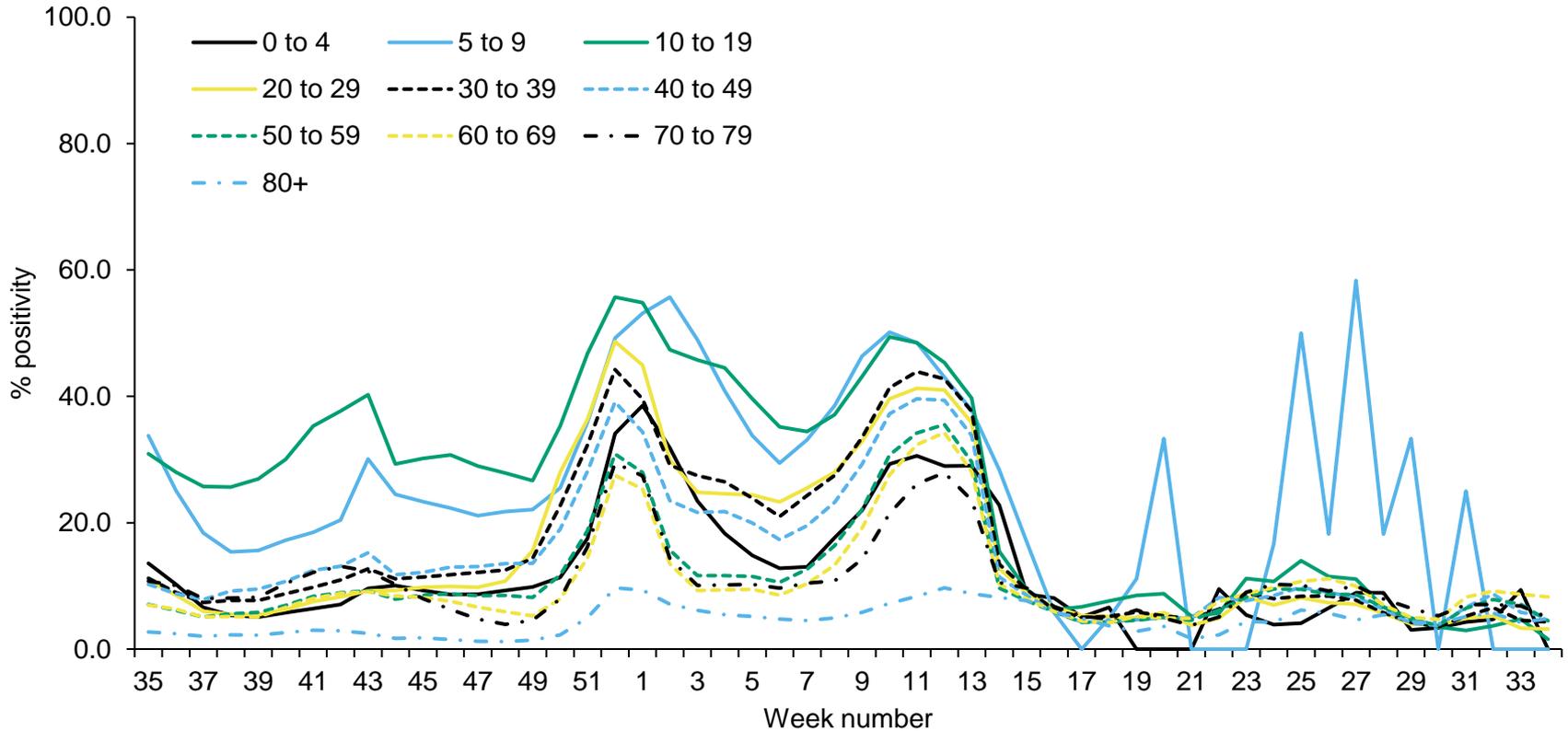


Weekly PCR positivity (%) of confirmed COVID-19 cases tested under Pillar 2, by male and age group



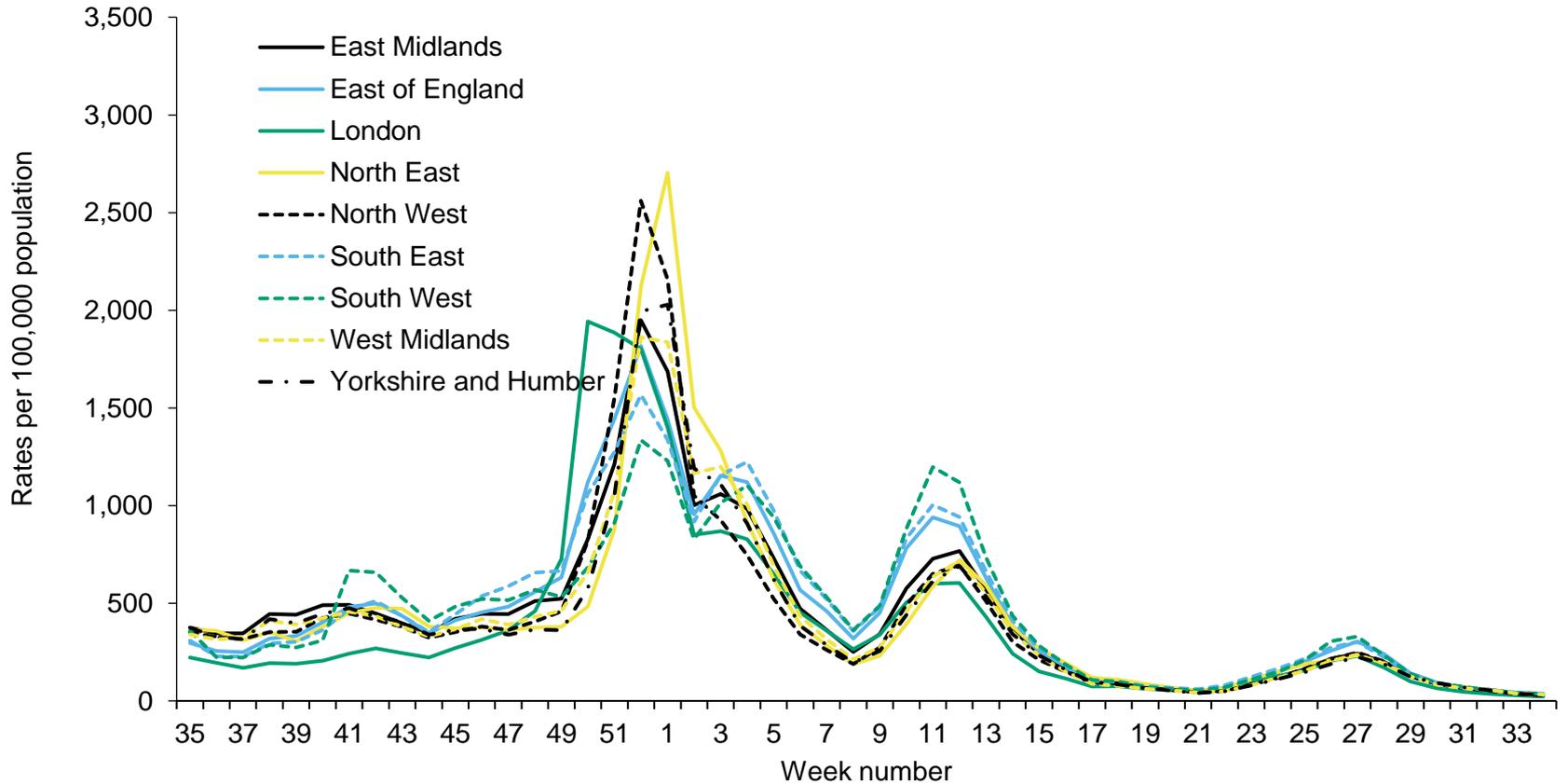


Weekly PCR positivity (%) of confirmed COVID-19 cases tested under Pillar 2, by female and age group



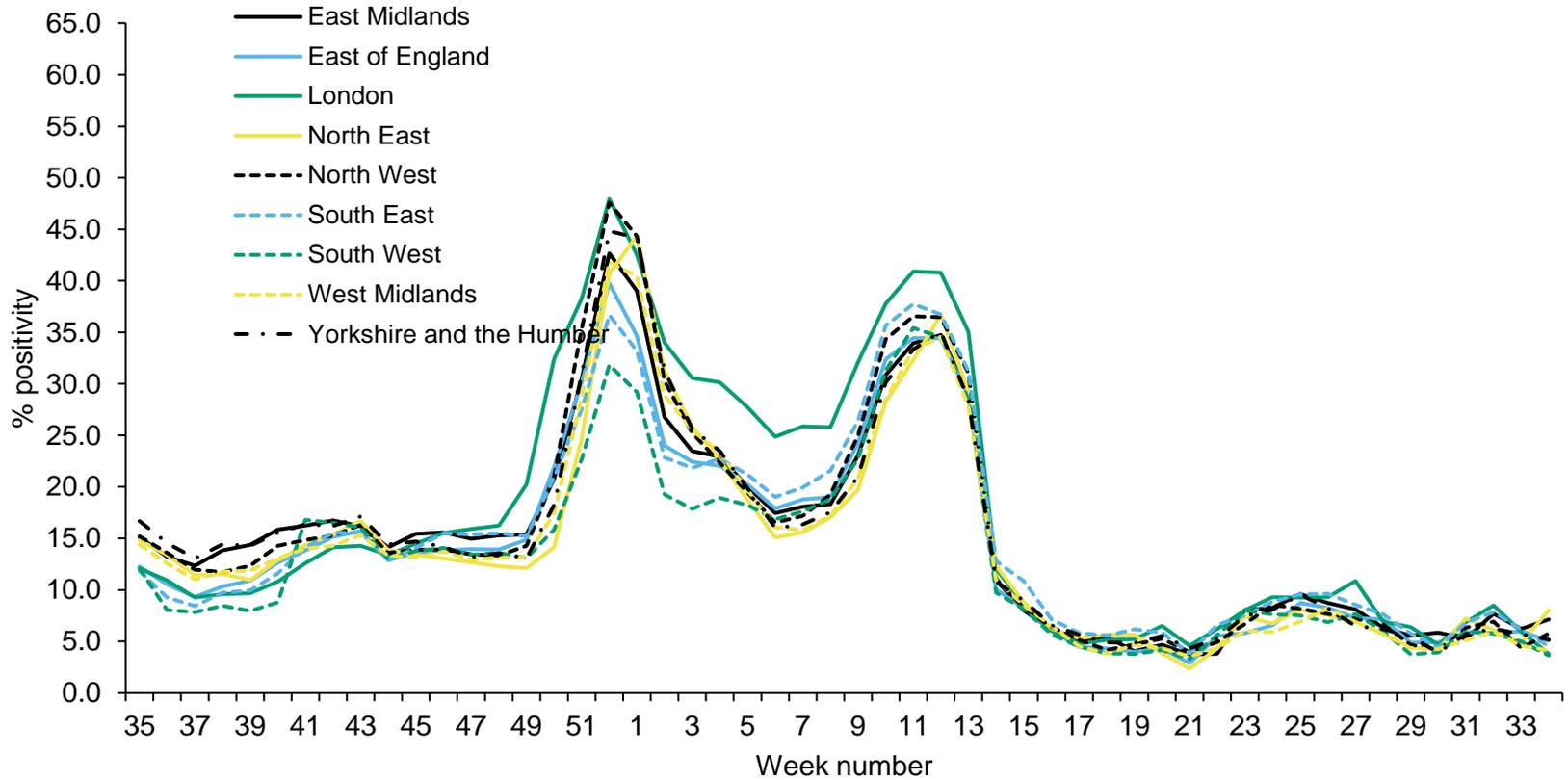


Weekly confirmed COVID-19 case rates by episode, per 100,000 population (Pillar 2), by UKHSA centres and sample week



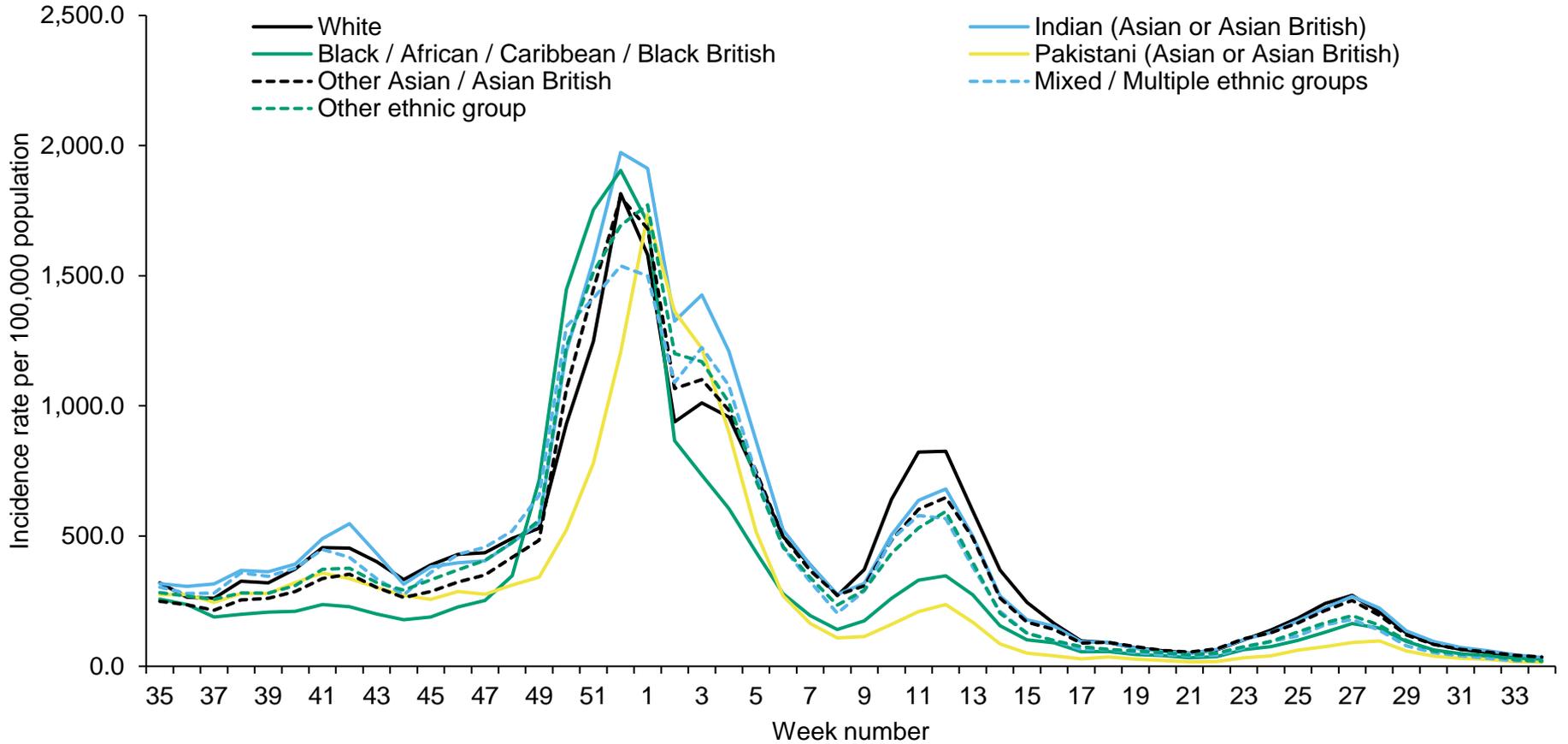


Weekly PCR positivity of confirmed COVID-19 cases tested under Pillar 2 (%) by UKHSA centres and sample week



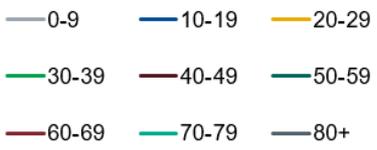
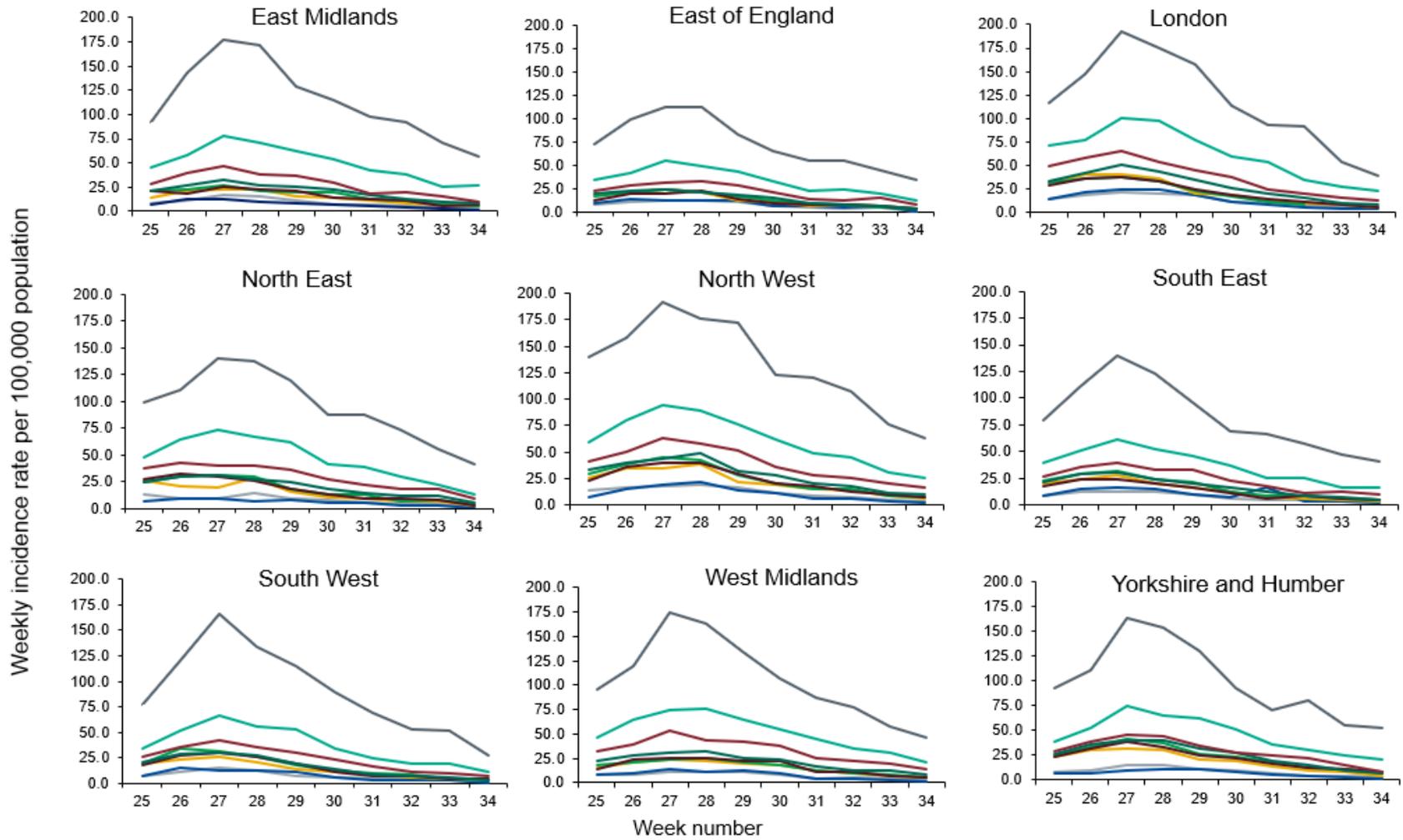


Weekly incidence per 100,000 population by ethnicity (Pillar 2), England



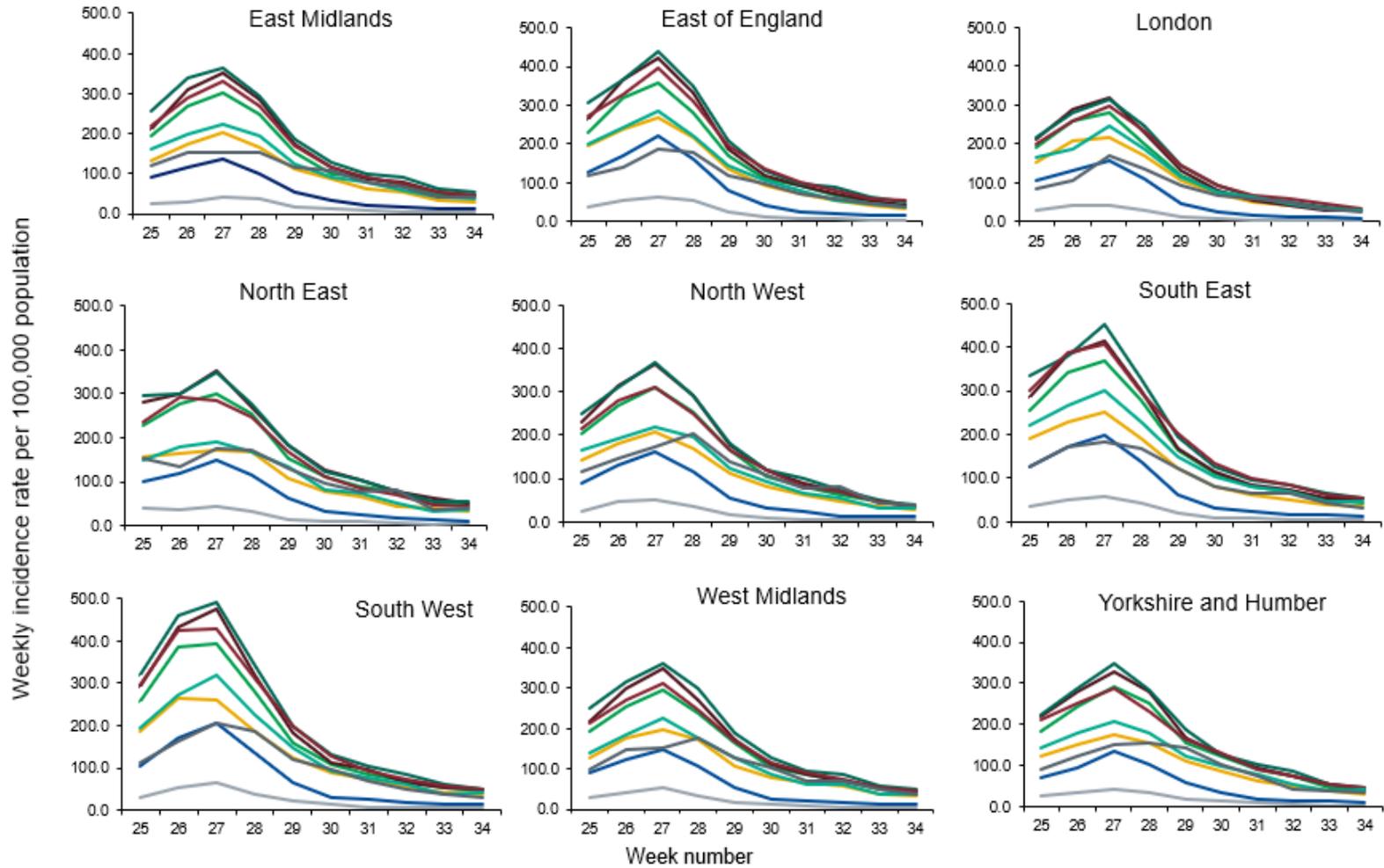


Weekly COVID-19 episodes tested under Pillar 1, per 100,000 population by age group and region, weeks 25 to 34



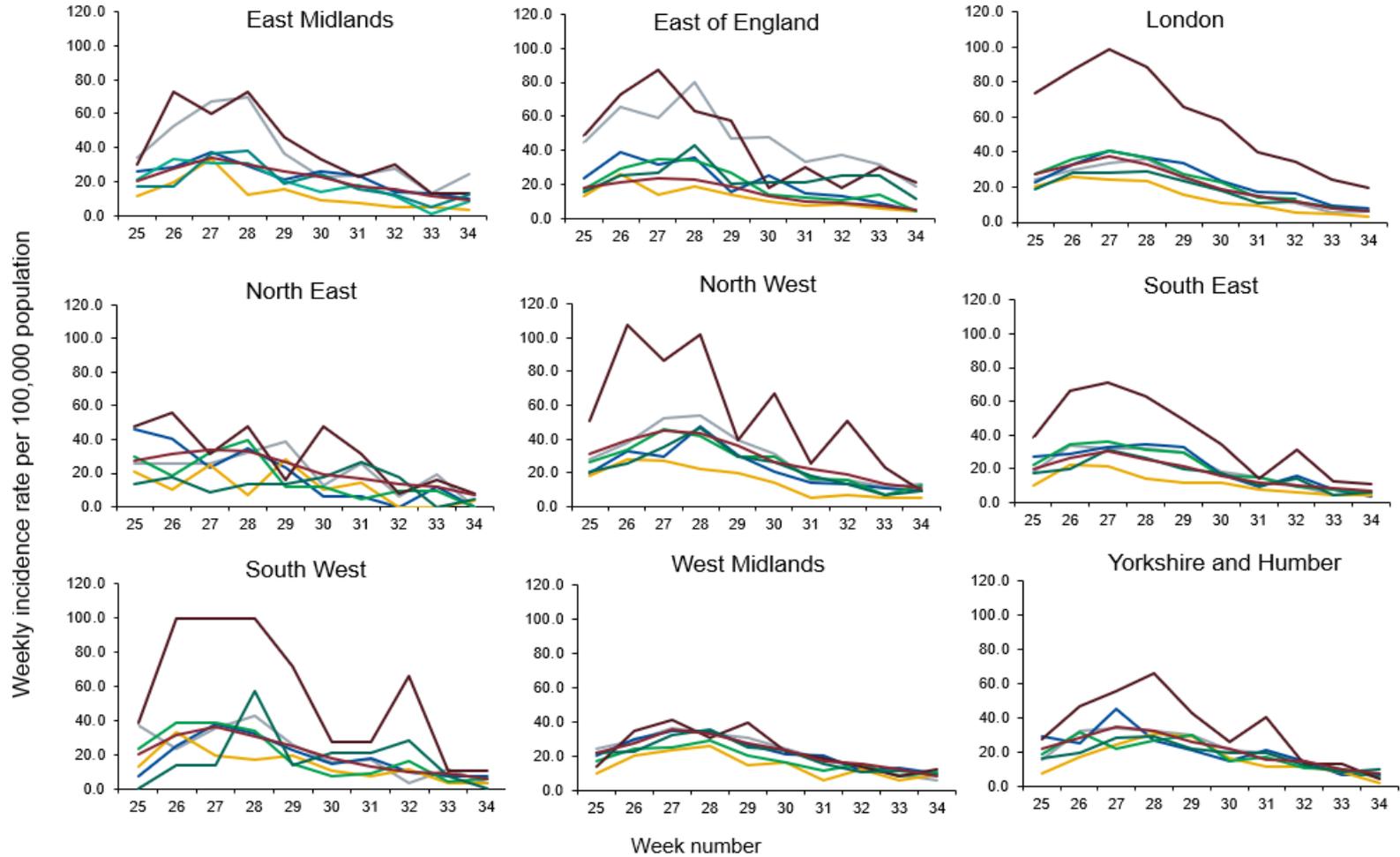


Weekly COVID-19 episodes tested under Pillar 2, per 100,000 population by age group and region, weeks 25 to 34





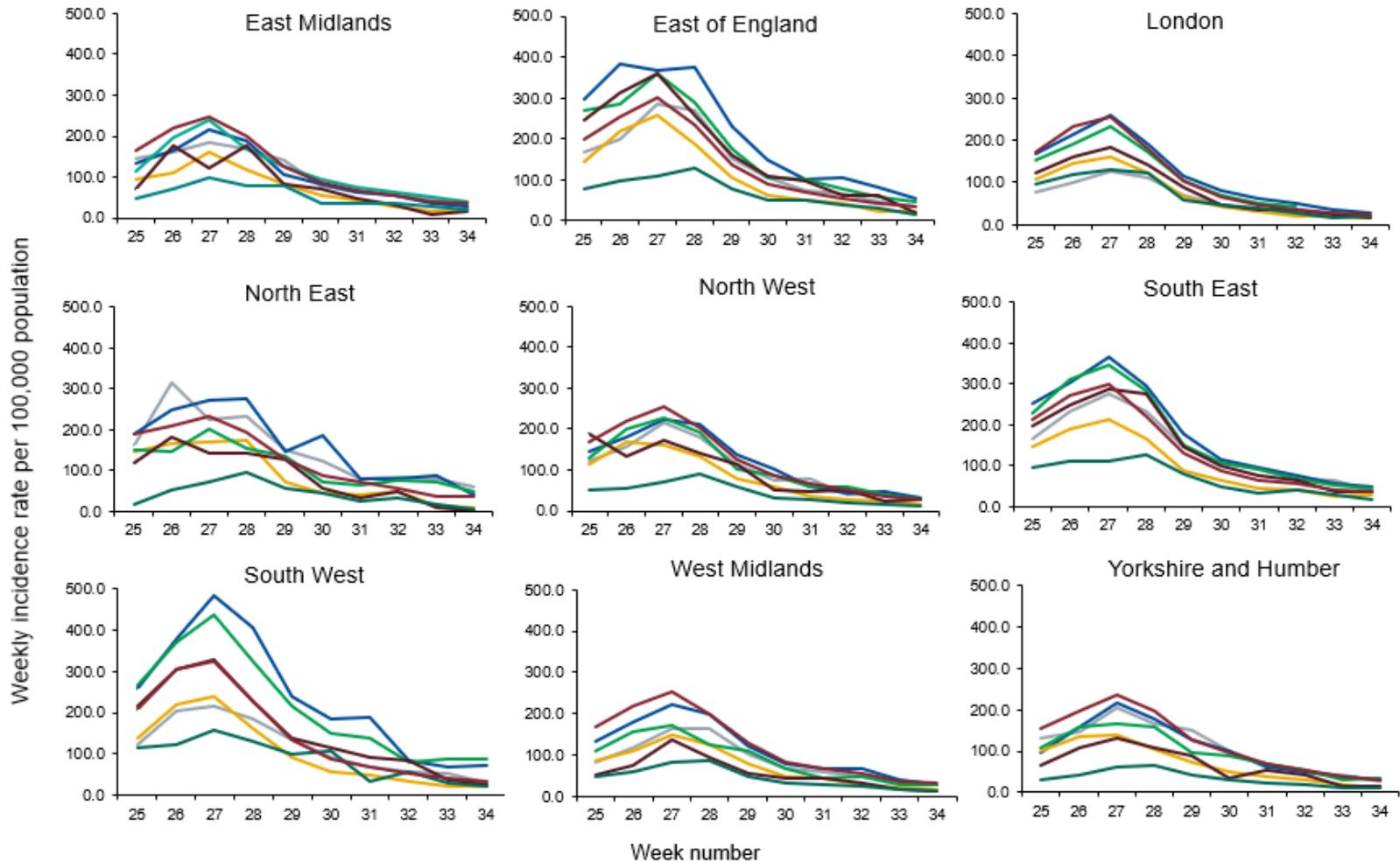
Weekly COVID-19 episodes tested under Pillar 1, per 100,000 population by ethnicity and region, weeks 25 to 34



- Black/African/Caribbean/Black British
- Indian (Asian or British)
- Mixed/Multiple Ethnic Groups
- Other Asian/Asian British
- Other ethnic group
- Pakistani (Asian or British)
- White



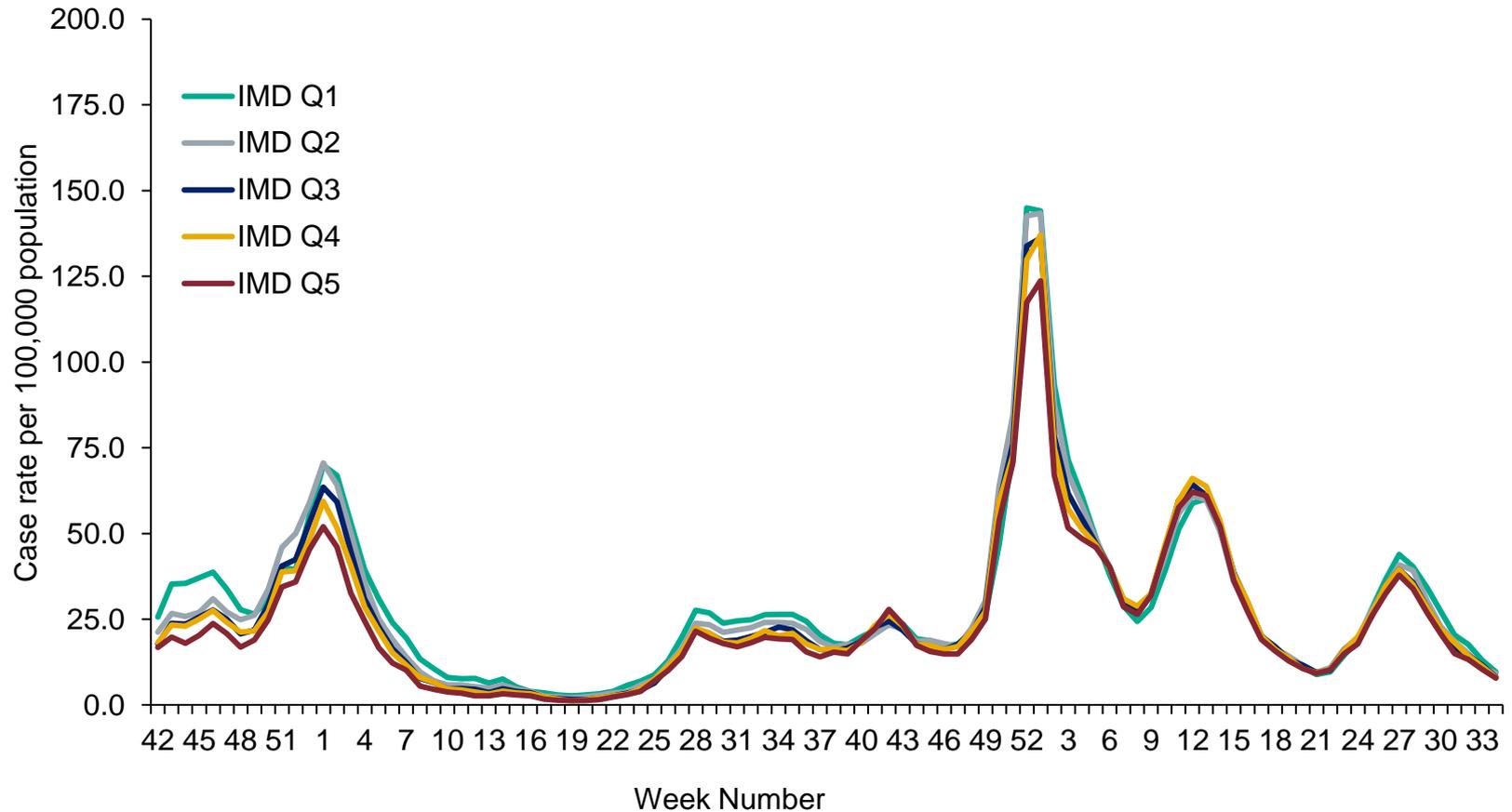
Weekly COVID-19 episodes tested under Pillar 2 per 100,000 population by ethnicity and region, weeks 25 to 34



- Black/African/Caribbean/Black British
- Indian (Asian or British)
- Mixed/Multiple Ethnic Groups
- Other Asian/Asian British
- Other ethnic group
- Pakistani (Asian or British)
- White



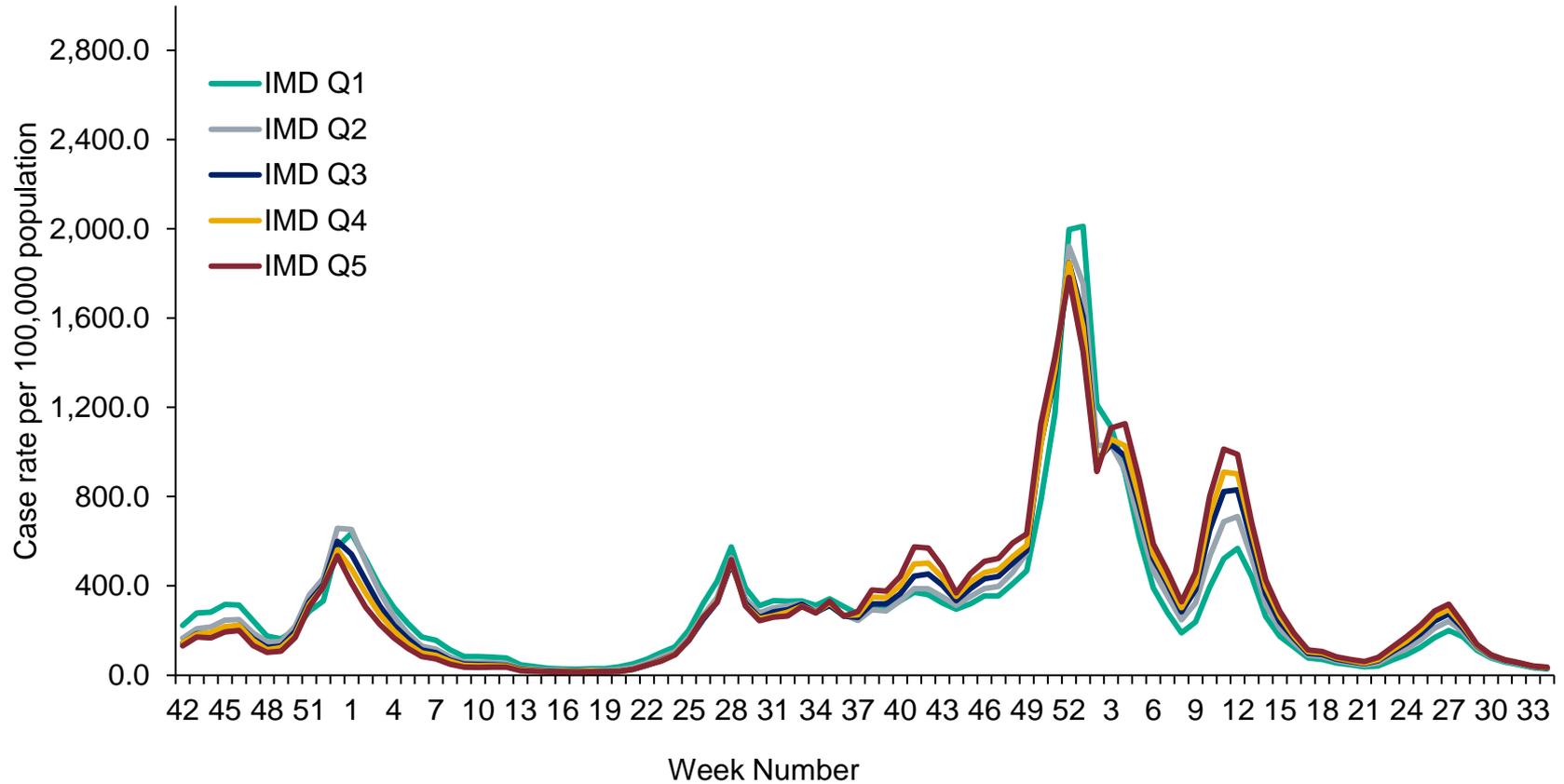
Weekly COVID-19 rate tested under Pillar 1, per 100,000 population by IMD quintile (1 being the most deprived and 5 being the least deprived)



*incidence rates have been calculated using the mid-2019 ONS population estimates

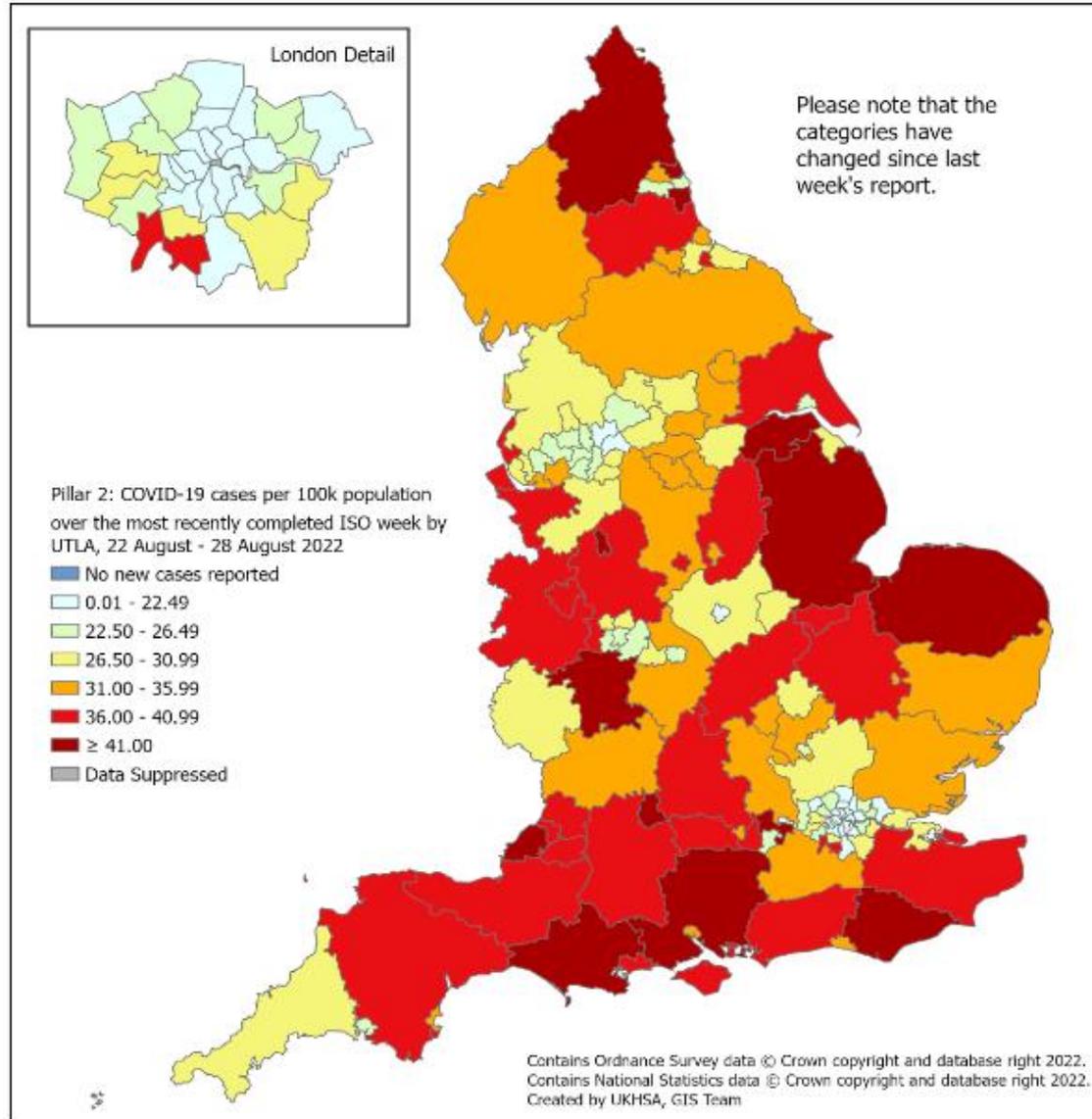


Weekly COVID-19 rate tested under Pillar 2, per 100,000 population by IMD quintile (1 being the most deprived and 5 being the least deprived)



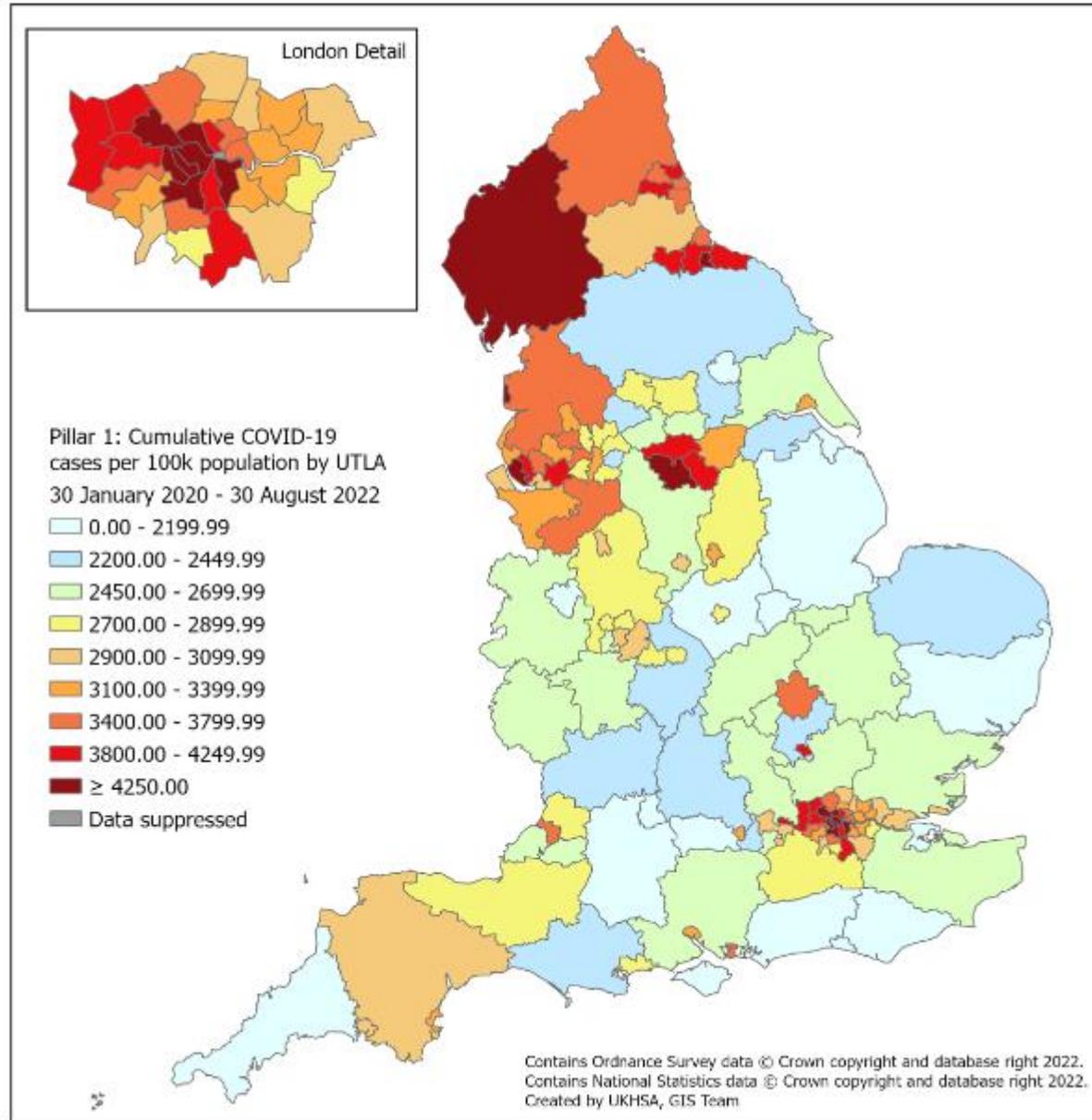


Weekly rate of COVID-19 episodes per 100,000 population (Pillar 2), by upper-tier local authority, England (box shows enlarged map of London area)



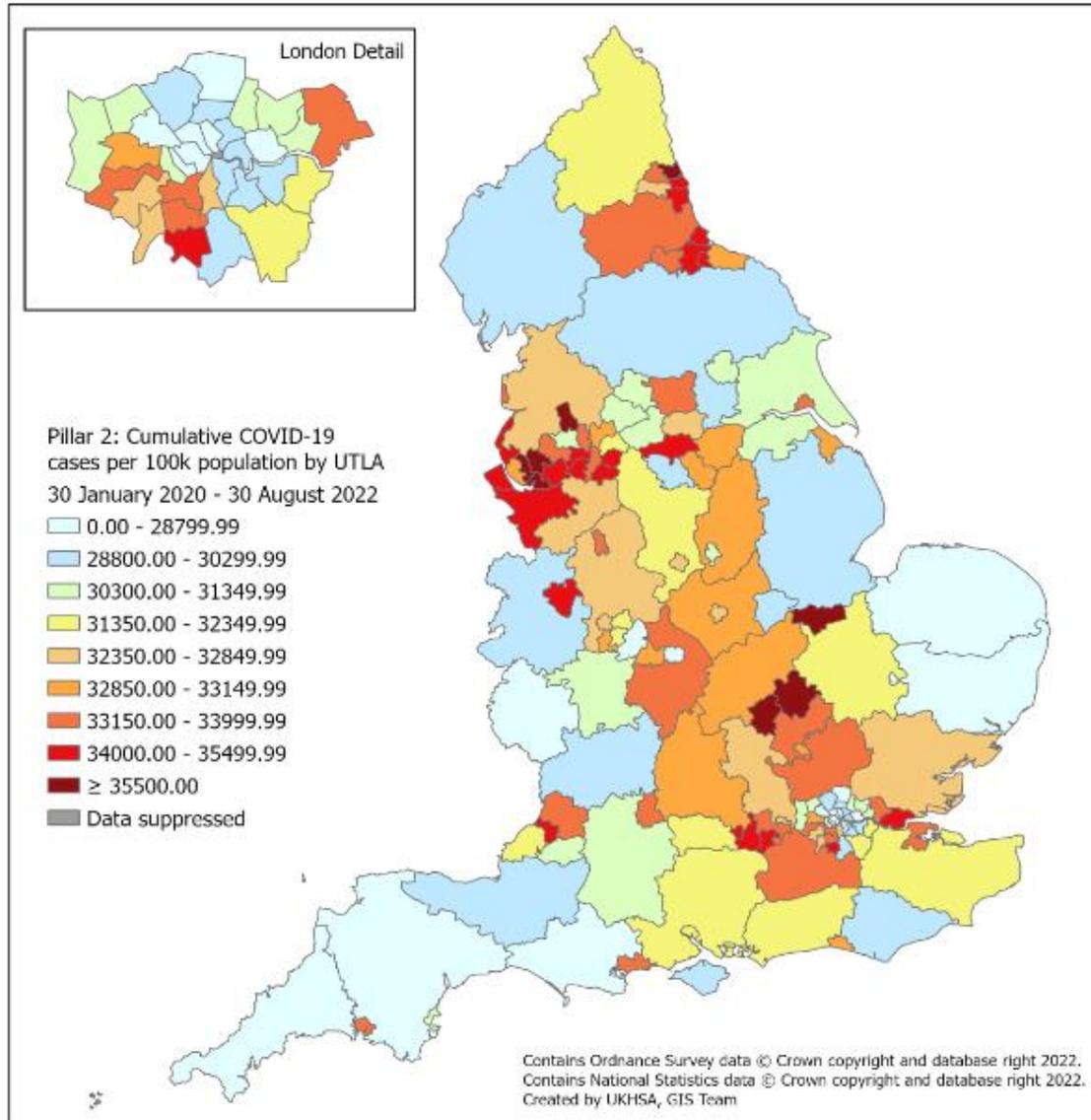


Cumulative rate of COVID-19 episodes per 100,000 population tested under Pillar 1, by upper-tier local authority, England (box shows enlarged map of London area)





Cumulative rate of COVID-19 episodes per 100,000 population tested under Pillar 2, by upper-tier local authority, England (box shows enlarged map of London area)



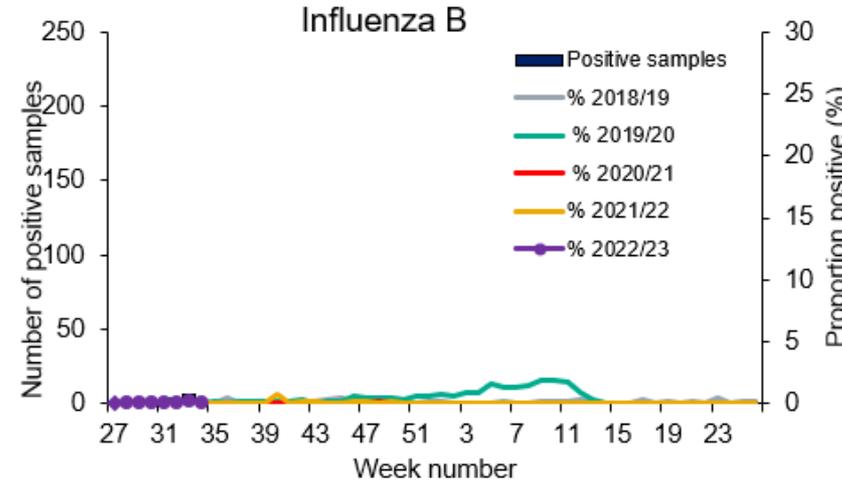
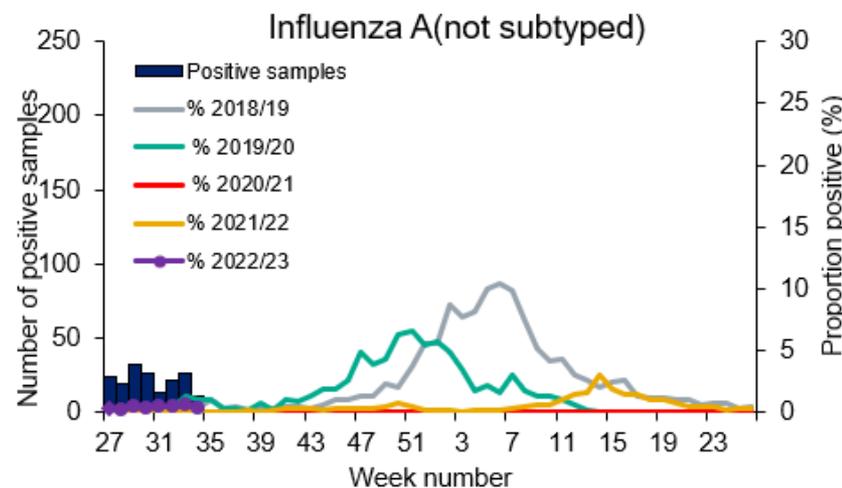
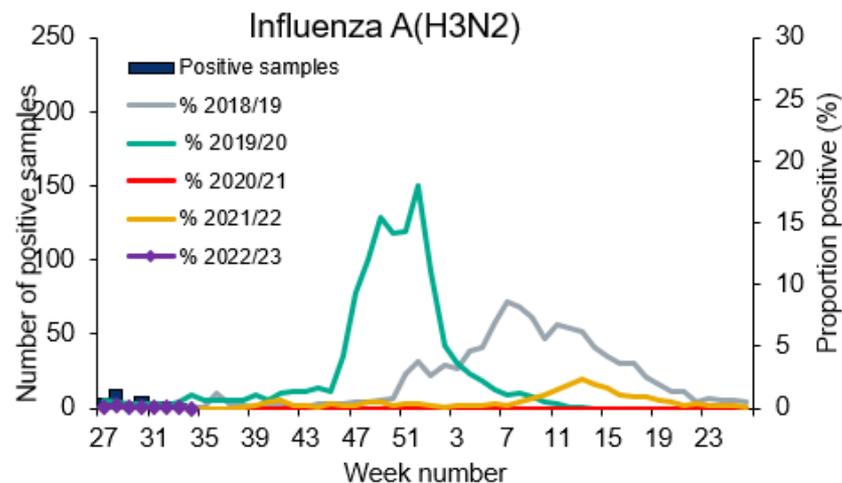
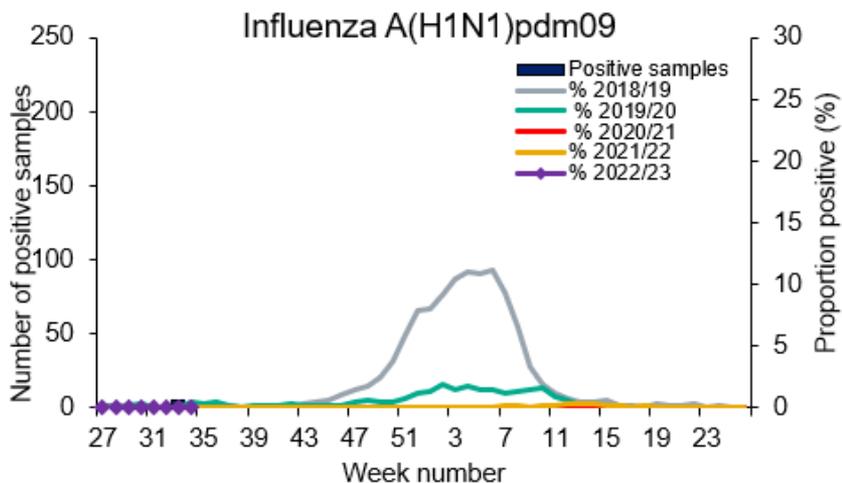


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Respiratory Datamart system (England)

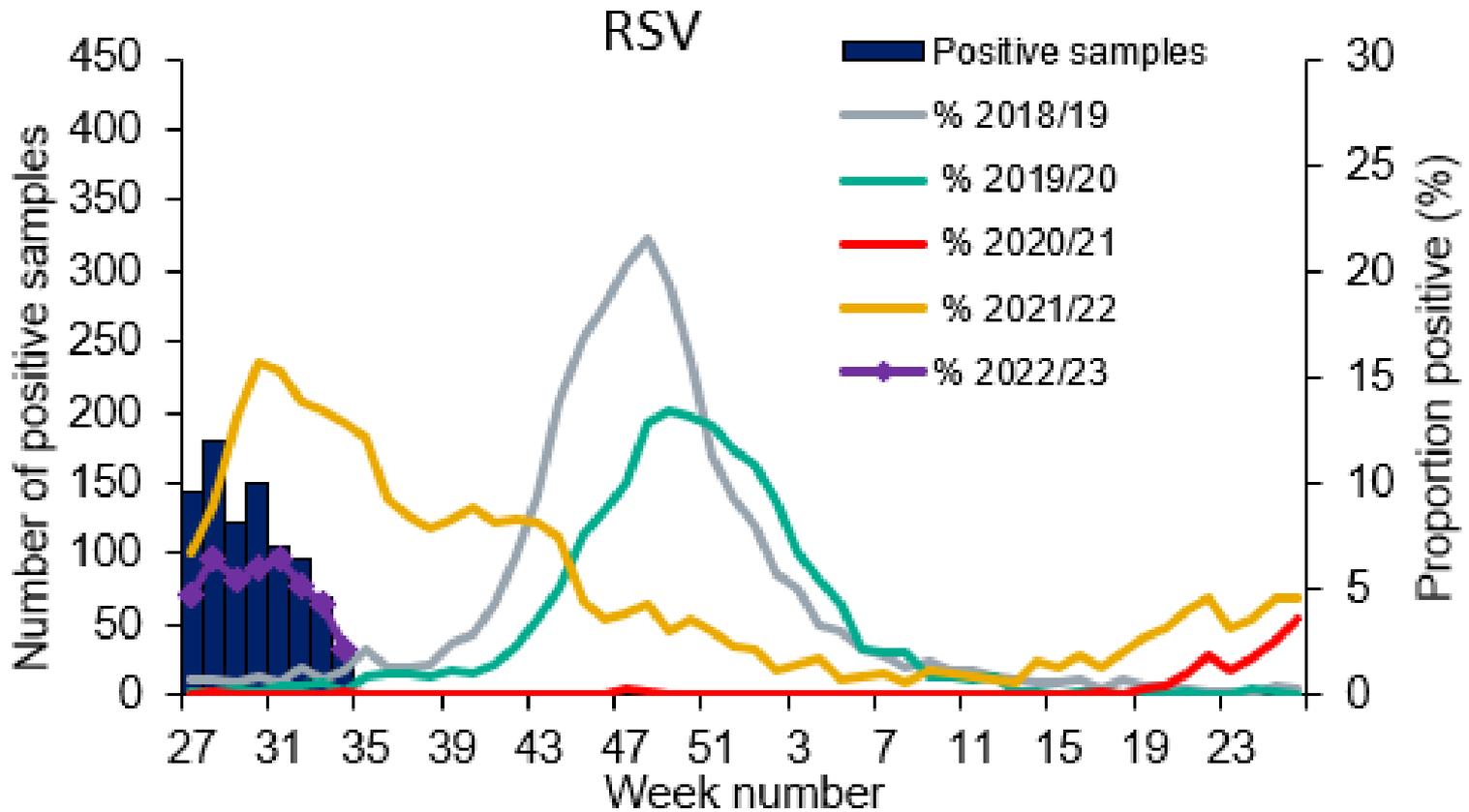


Respiratory DataMart – Influenza subtypes



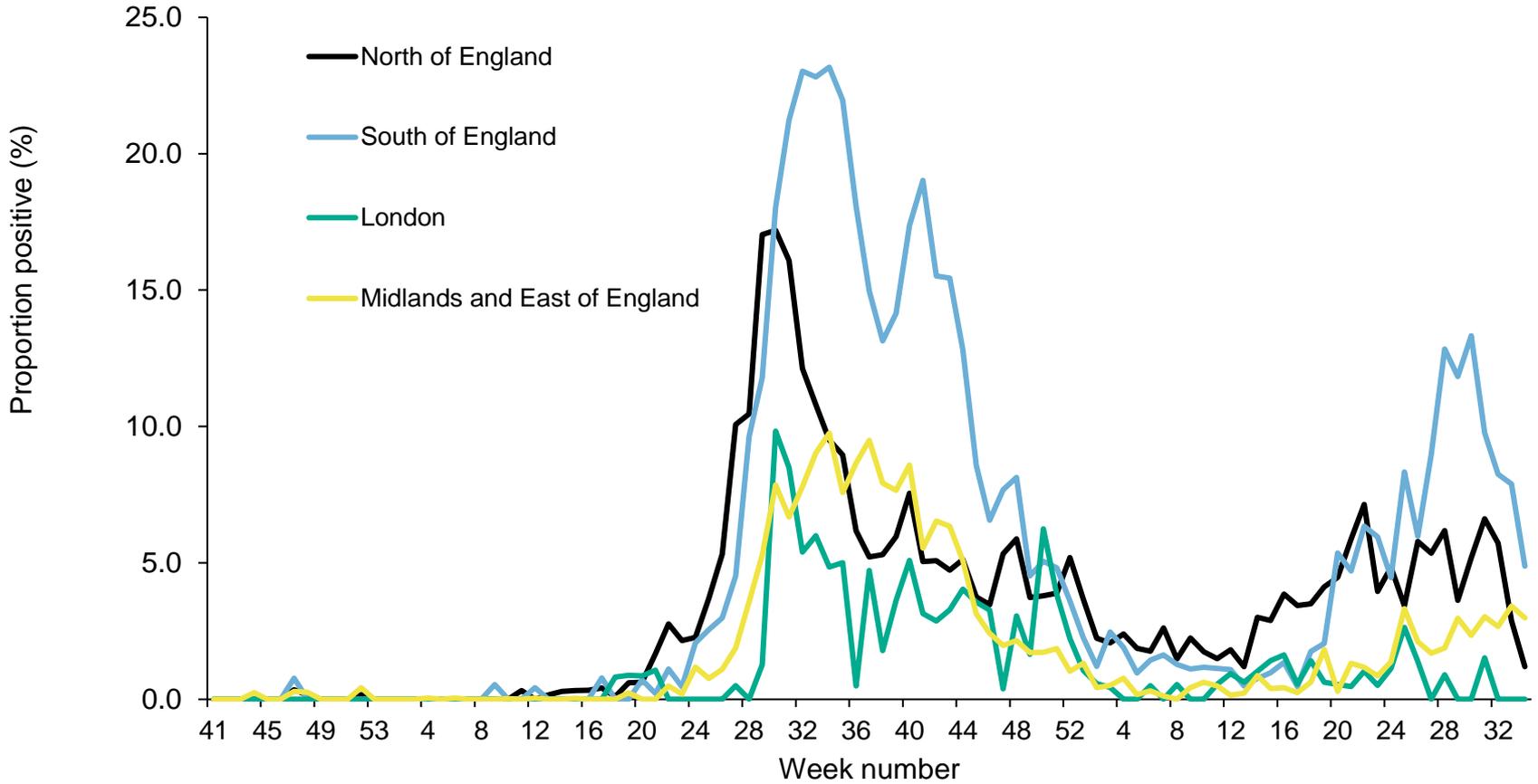


Respiratory DataMart – Respiratory syncytial virus (RSV)



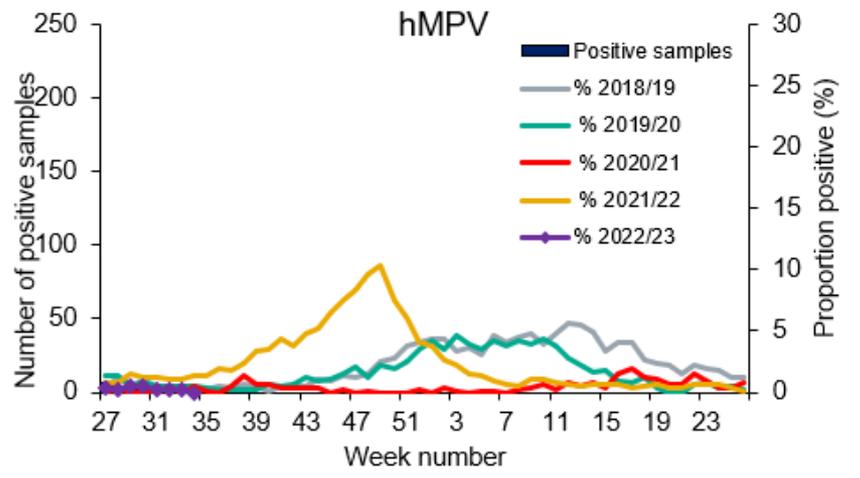
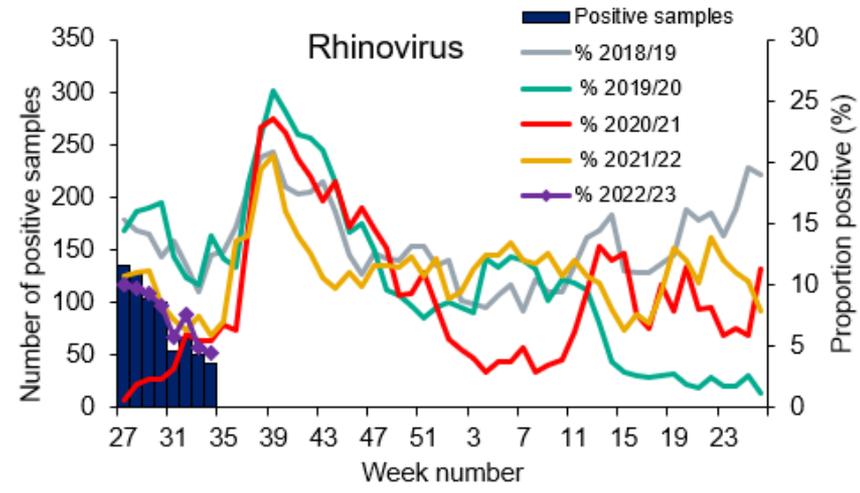
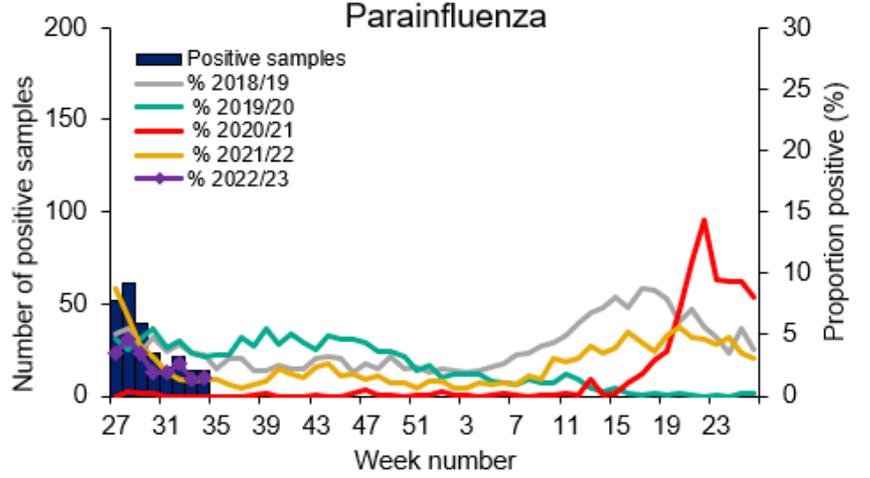
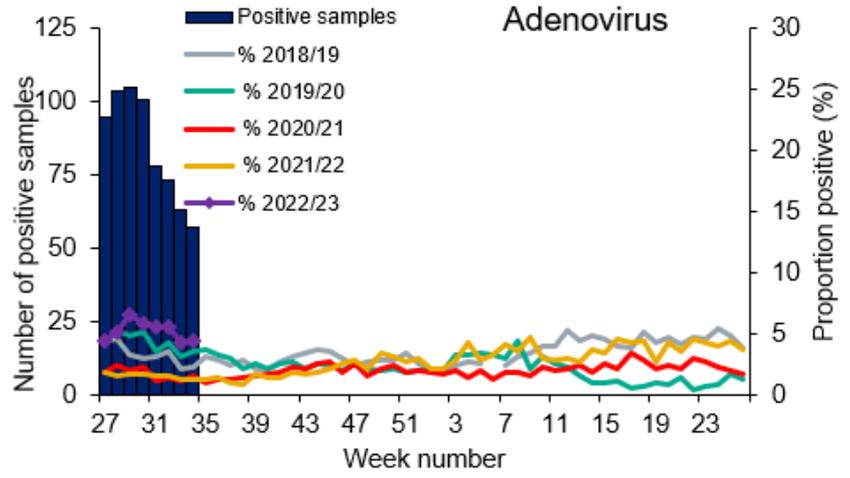


Respiratory DataMart – Respiratory syncytial virus (RSV) weekly positivity by UKHSA region





Respiratory DataMart – other respiratory viruses

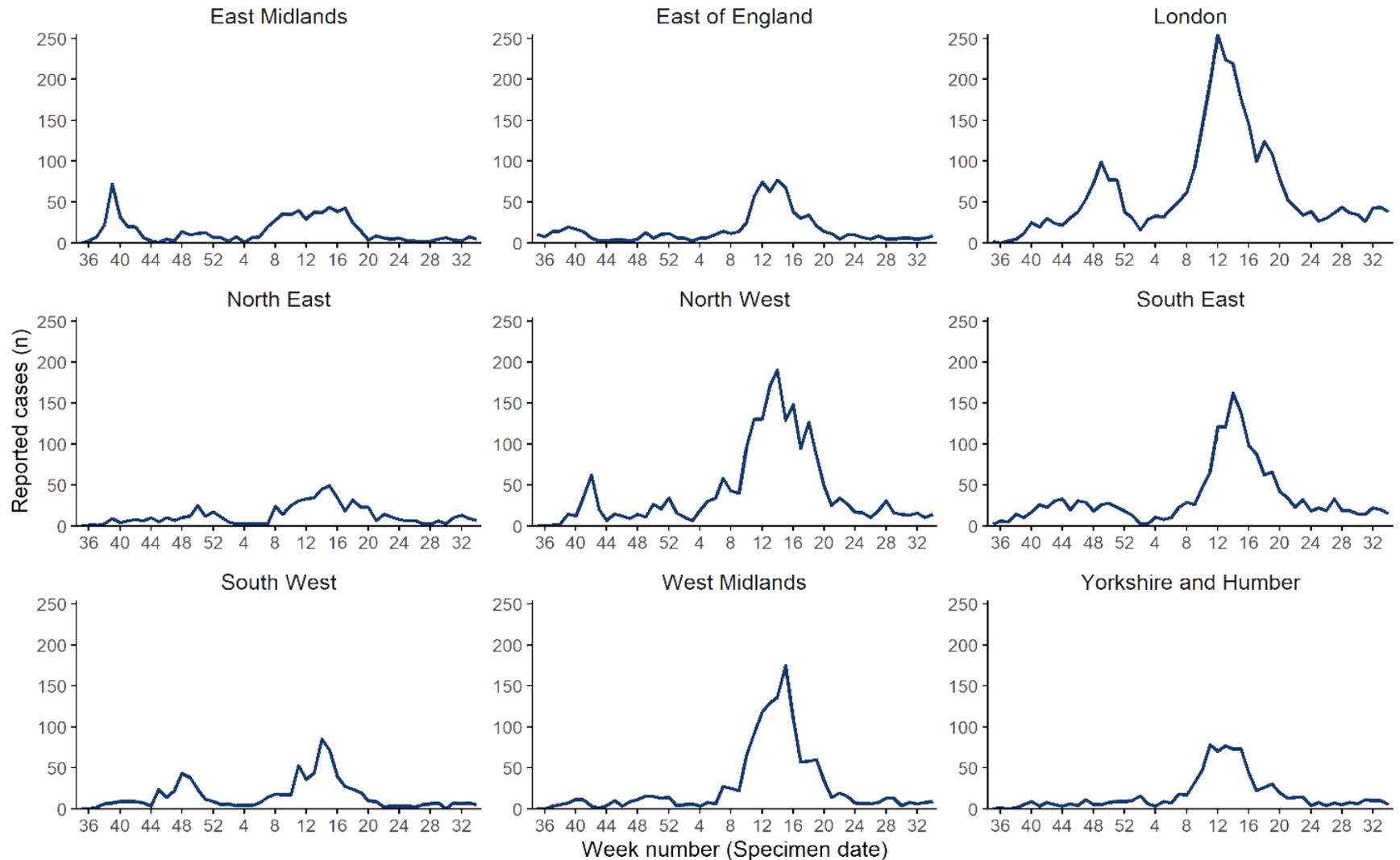




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Second generation surveillance system (SGSS)

SGSS reported Influenza A cases by region (all ages)

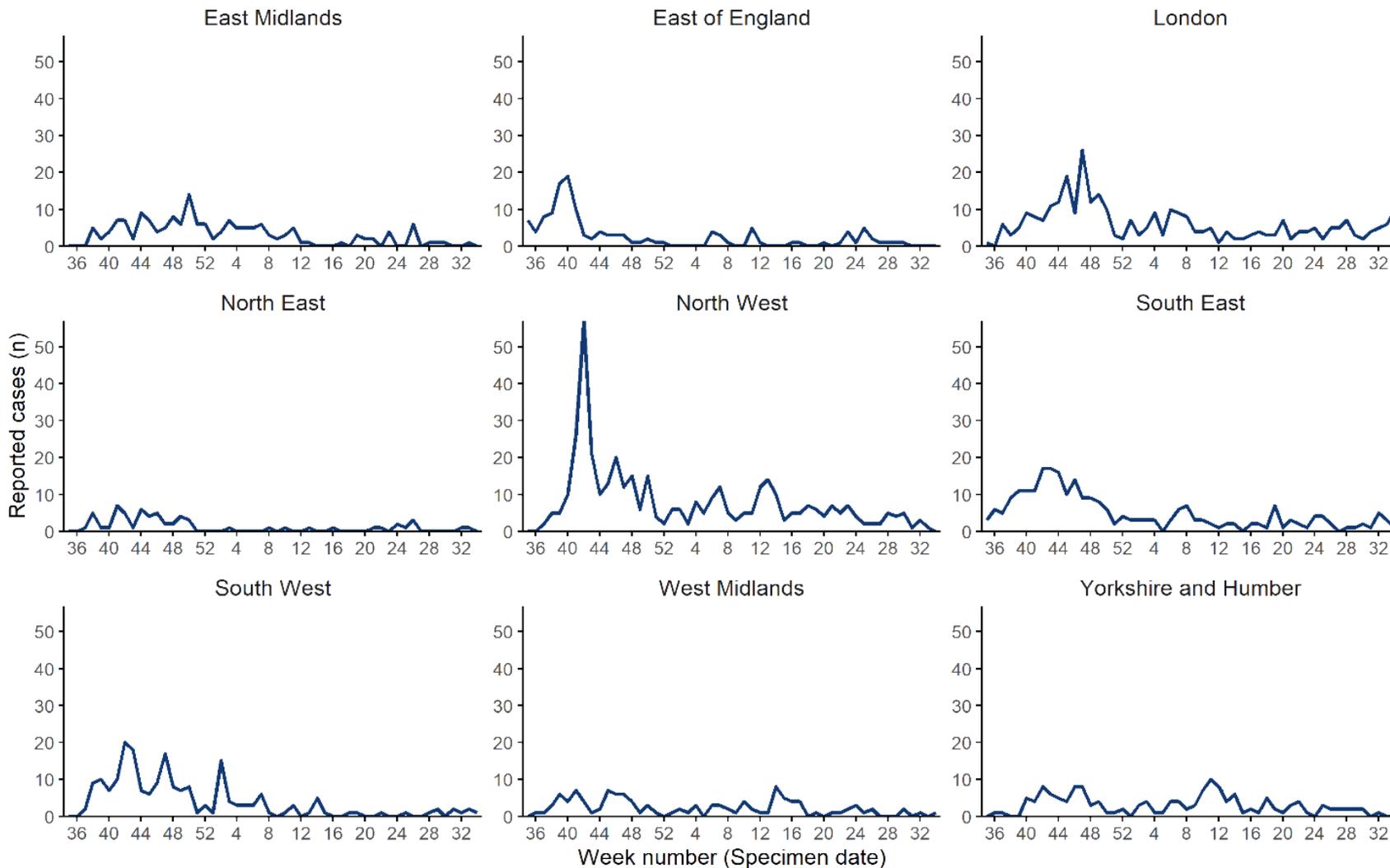


The presented figures are based on laboratory reports through SGSS. Testing and reporting procedures vary by virus, UKHSA centre and over time, including short-term trends in testing. Therefore

1 September 2020 comparisons should be done with caution.

Previously, this data was presented by report date however is now presented by specimen date.

SGSS reported Influenza B cases by region (all ages)

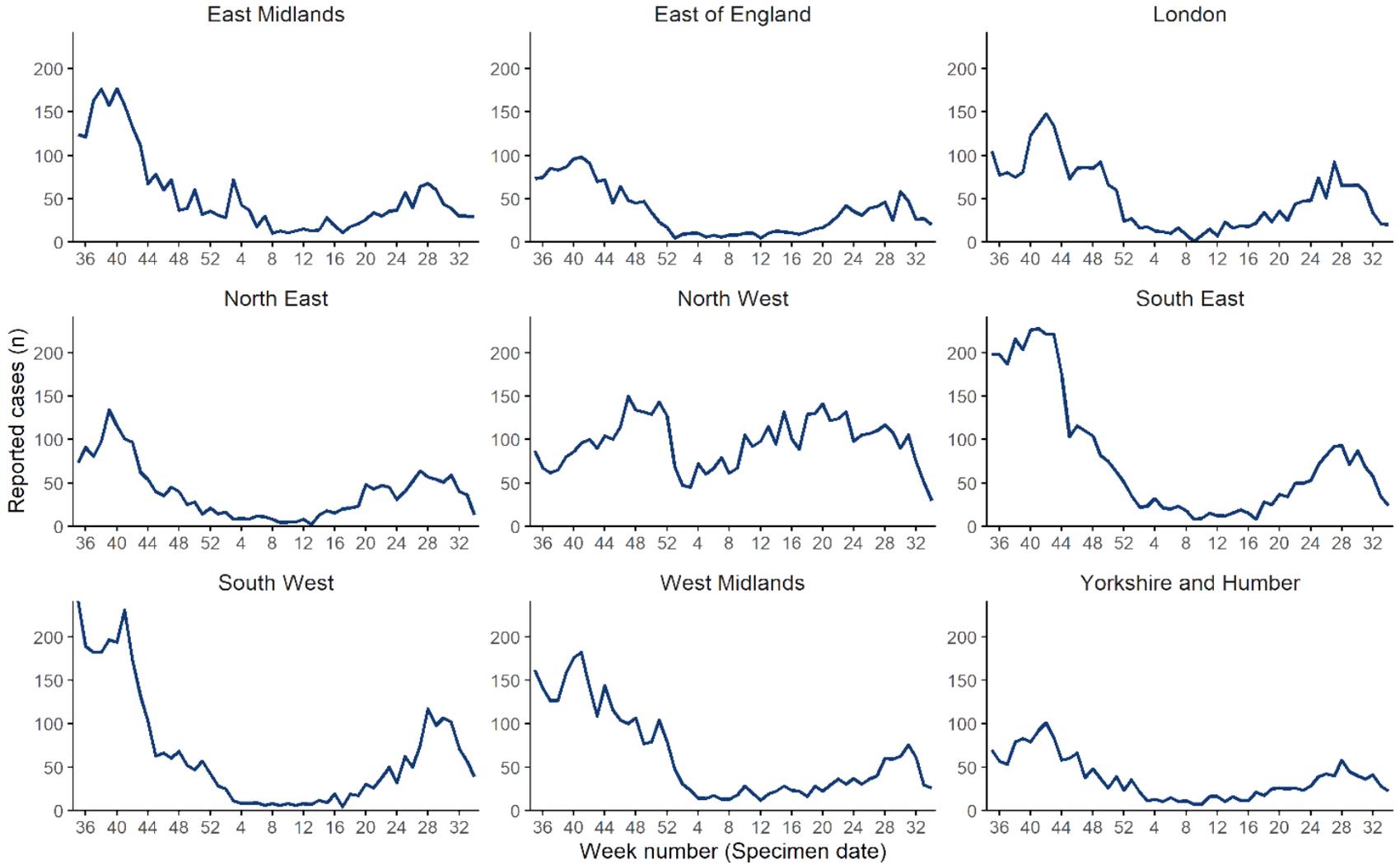


The presented figures are based on laboratory reports through SGSS. Testing and reporting procedures vary by virus, UKHSA centre and over time, including short-term trends in testing. Therefore comparisons should be done with caution.

Previously, this data was presented by report date however is now presented by specimen date.



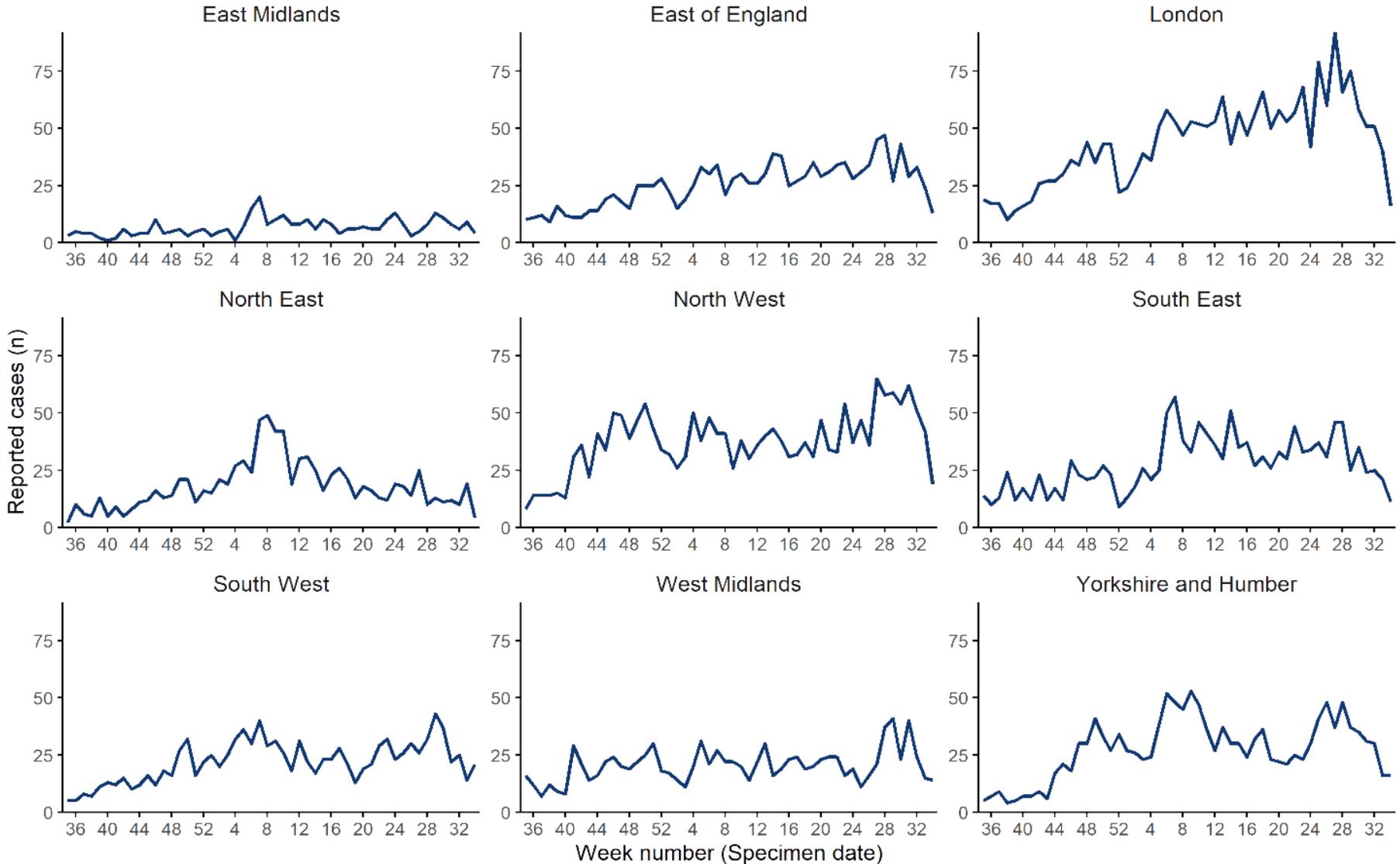
SGSS reported RSV cases by region (all ages)



The presented figures are based on laboratory reports through SGSS. Testing and reporting procedures vary by virus, UKHSA centre and over time, including short-term trends in testing. Therefore comparisons should be done with caution. Previously, this data was presented by report date however is now presented by specimen date.

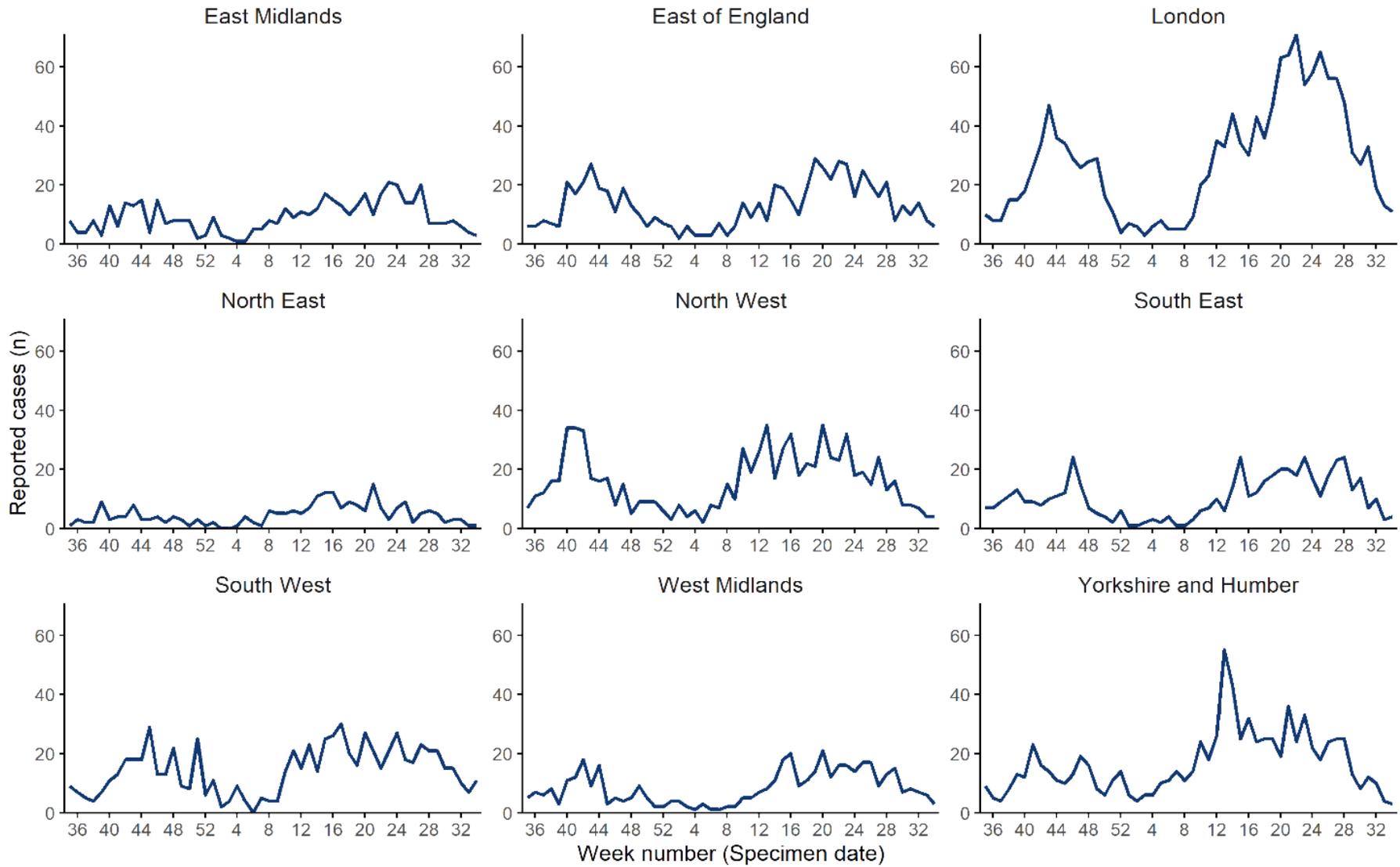


SGSS reported Adenovirus cases by region (all ages)





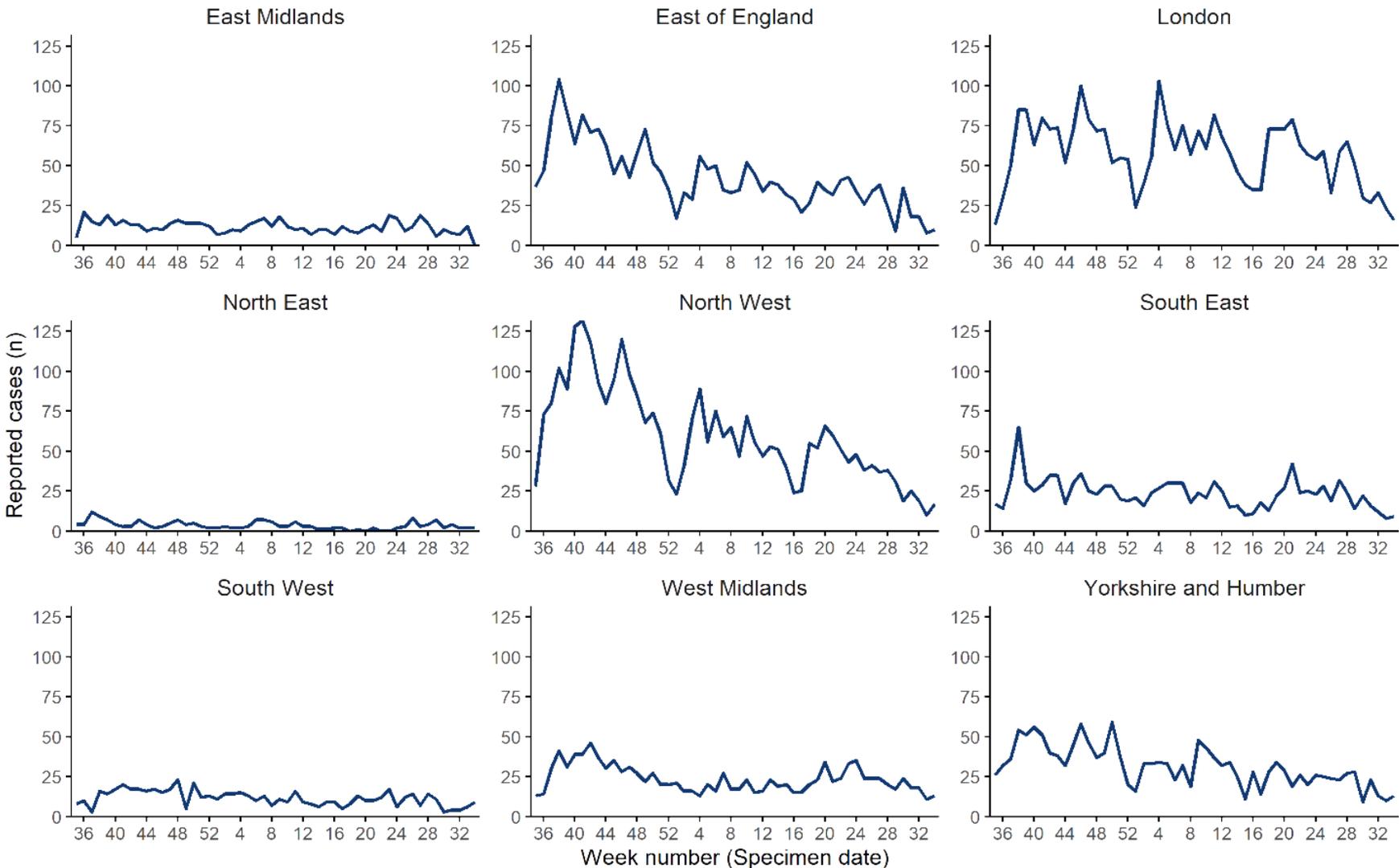
SGSS reported Parainfluenza cases by region (all ages)



The presented figures are based on laboratory reports through SGSS. Testing and reporting procedures vary by virus, UKHSA centre and over time, including short-term trends in testing. Therefore comparisons should be done with caution.



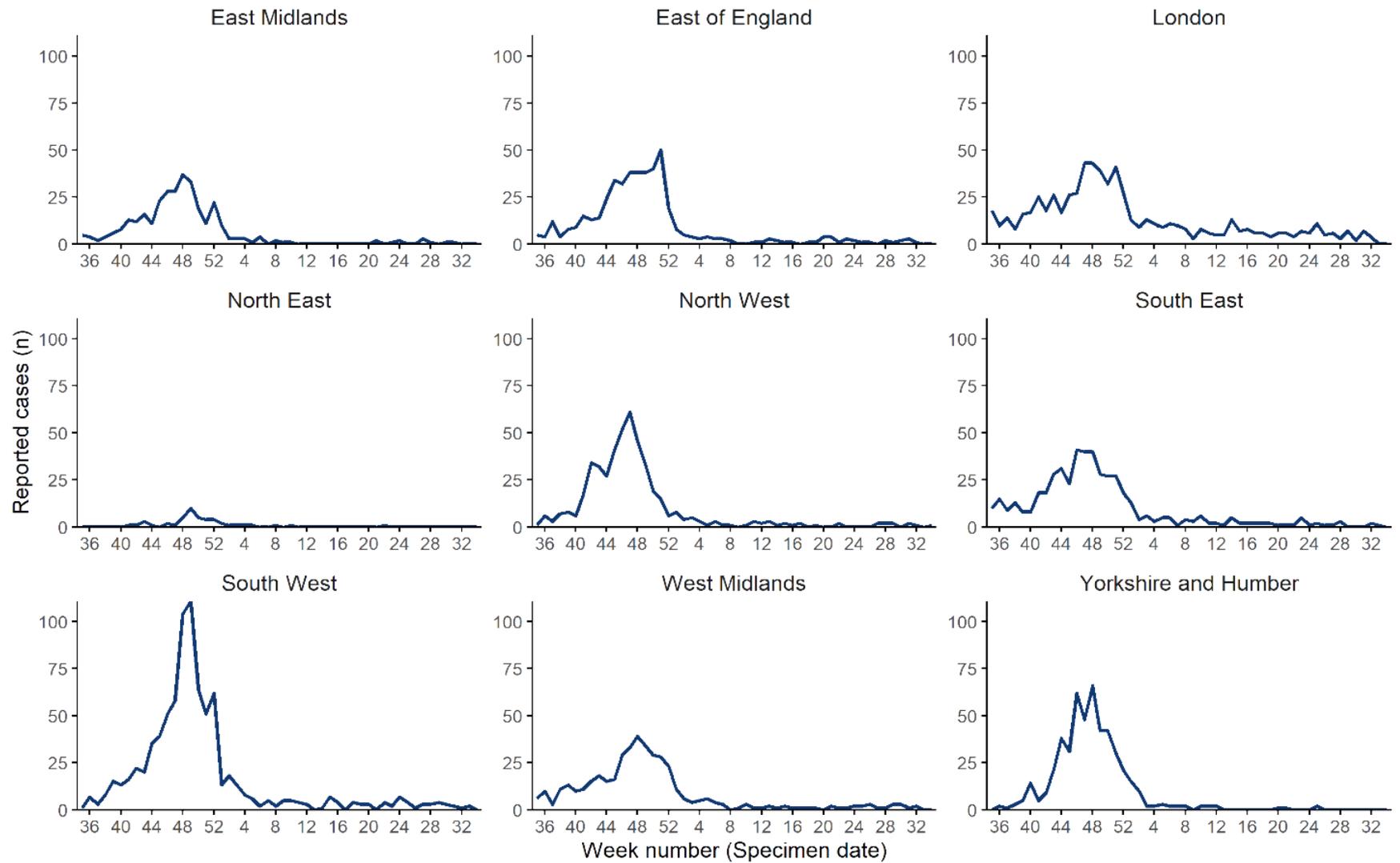
SGSS reported Rhinovirus cases by region (all ages)



The presented figures are based on laboratory reports through SGSS. Testing and reporting procedures vary by virus, UKHSA centre and over time, including short-term trends in testing. Therefore comparisons should be done with caution.



SGSS reported hMPV cases by region (all ages)



The presented figures are based on laboratory reports through SGSS. Testing and reporting procedures vary by virus, UKHSA centre and over time, including short-term trends in testing. Therefore comparisons should be done with caution.



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Community surveillance



Acute respiratory infection (ARI) outbreaks linked to educational settings

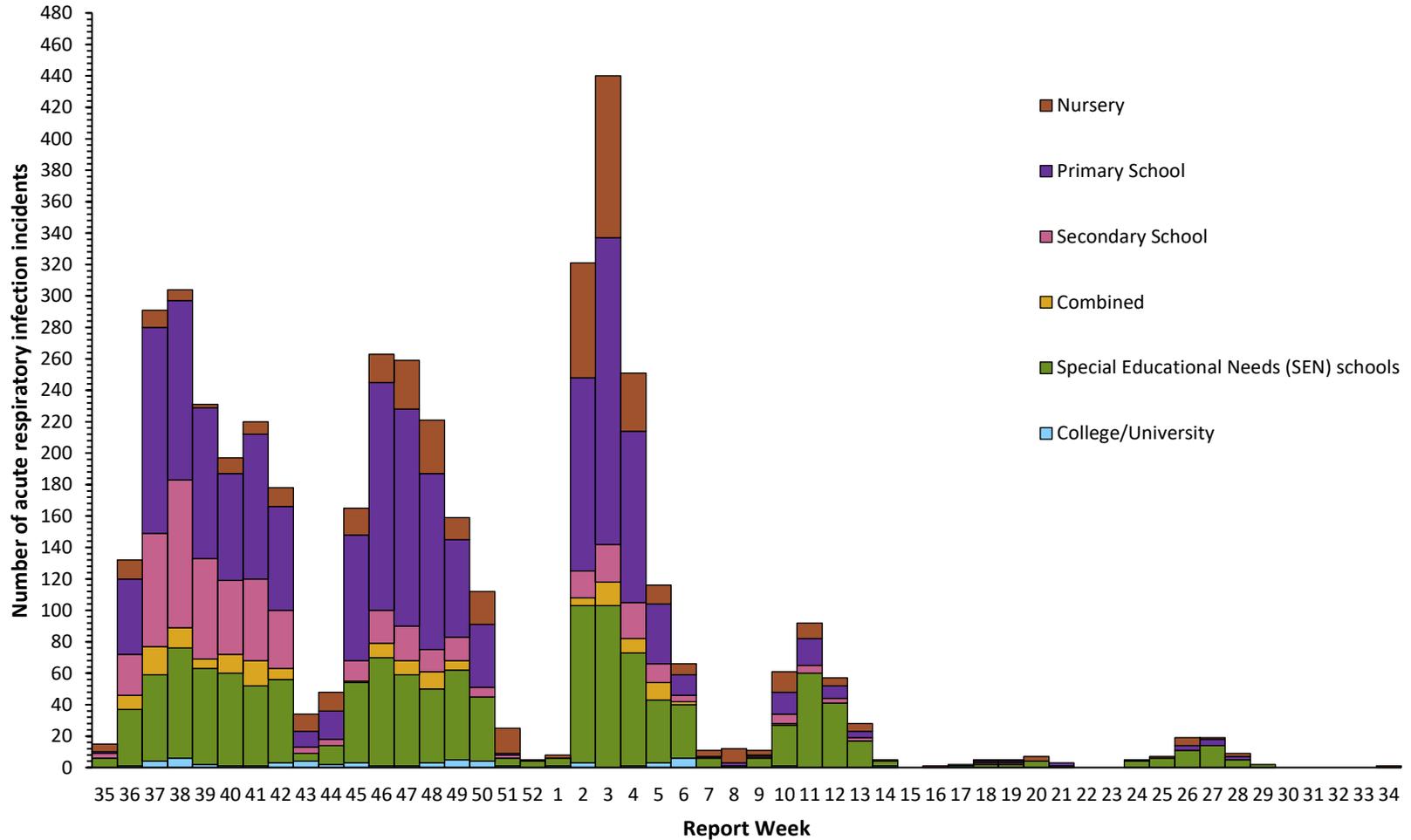
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Data Information

We report on new acute respiratory infection (ARI) incidents reported to Health Protection Teams (HPTs) and entered on HPZone in the previous reporting week in educational settings by locality. The incidents captured on HPZone represent a subset of all ongoing clusters and outbreaks in England. A variety of arrangements are in place with local authorities and other stakeholders supporting HPTs, however data may not routinely be documented on HPZone. As a result, the number of outbreaks reported for some of the regions are underestimates



Number of acute respiratory infection outbreaks reported to UKHSA by type of educational setting, England





Number of acute respiratory infection outbreaks by type of educational setting, England

End of academic year total

Week 36 2020- 34 2021

| PHE Centres | Cumulative number of suspected acute respiratory infection outbreaks by type of educational setting for the 2020/21 academic year Week 36 2020- 34 2021 | | | | | | |
|--------------|---|----------------|------------------|-----------|---|--------------------|-------------|
| | Nursery | Primary School | Secondary School | Combined | Special Educational Needs (SEN) schools | College University | Total |
| Total | 846 | 2125 | 2122 | 40 | 666 | 268 | 6067 |

Week 34 2022

Main table

| PHE Centres | Cumulative number of suspected acute respiratory infection outbreaks by type of educational setting for the 2021/22 academic year from Week 35 2021 | | | | | | |
|------------------------|---|-----------------|------------------|----------------|---|--------------------|-----------------|
| | Nursery | Primary School | Secondary School | Combined | Special Educational Needs (SEN) schools | College University | Total |
| East Midlands Centre | 72 (0) | 59 (0) | 30 (0) | 14 (0) | 170 (0) | 6 (0) | 351 (0) |
| East of England Centre | 1 (1) | 12 (0) | 8 (0) | 3 (0) | 11 (0) | 2 (0) | 37 (1) |
| London Centre | 370 (0) | 1093 (0) | 260 (0) | 59 (0) | 227 (1) | 30 (0) | 2038 (0) |
| North East Centre | 0 (0) | 2 (0) | 0 (0) | 0 (0) | 3 (0) | 0 (0) | 5 (0) |
| North West Center | 13 (0) | 31 (0) | 13 (0) | 4 (0) | 127 (0) | 7 (0) | 195 (0) |
| South East Centre | 43 (0) | 389 (0) | 127 (0) | 34 (0) | 291 (0) | 7 (0) | 891 (0) |
| South West Centre | 5 (0) | 65 (0) | 79 (0) | 37 (0) | 256 (0) | 1 (0) | 441 (0) |
| West Midlands Centre | 19 (0) | 74 (0) | 52 (0) | 7 (0) | 142 (0) | 6 (0) | 300 (0) |
| Yorkshire & the Humber | 17 (0) | 36 (0) | 27 (0) | 5 (0) | 84 (0) | 0 (0) | 169 (0) |
| Total | 540 (1) | 1761 (0) | 596 (0) | 161 (0) | 1306 (0) | 59 (0) | 4423 (1) |

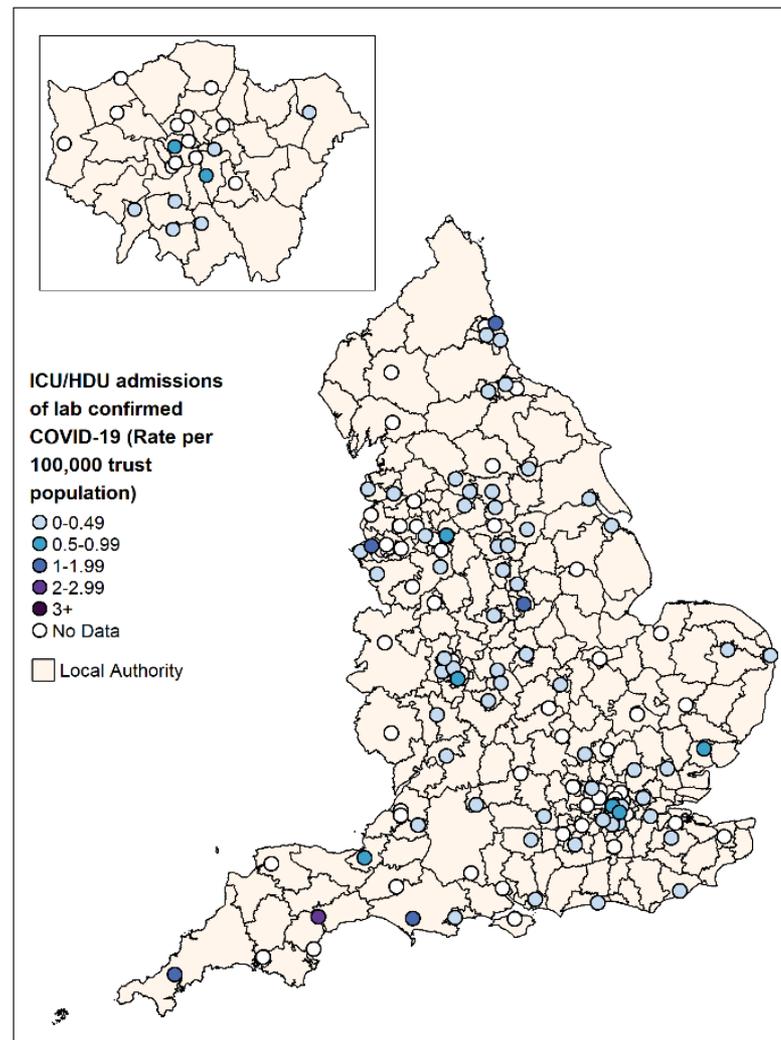
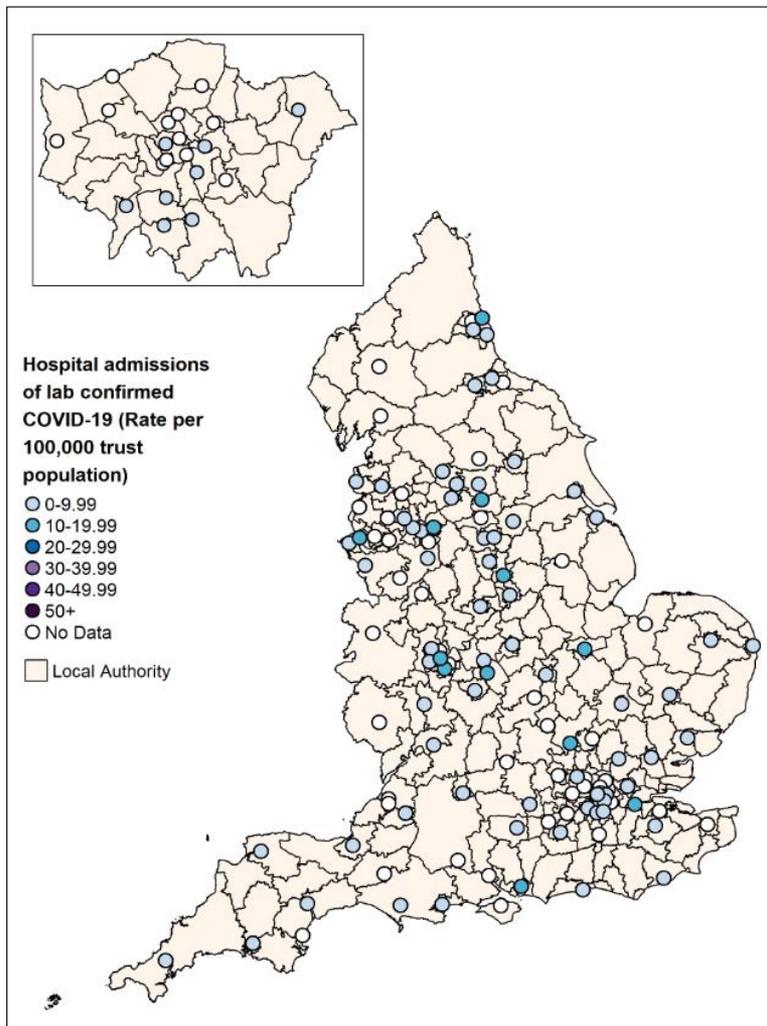
* Number of acute respiratory infection for the most recent week in brackets



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Secondary Care surveillance

Weekly admission rates for hospital and ICU/HDU laboratory confirmed COVID-19 cases reported through SARI Watch, week 34



Source: UKHSA SARI-Watch (Severe Acute Respiratory Infection-Watch, formerly CHES).

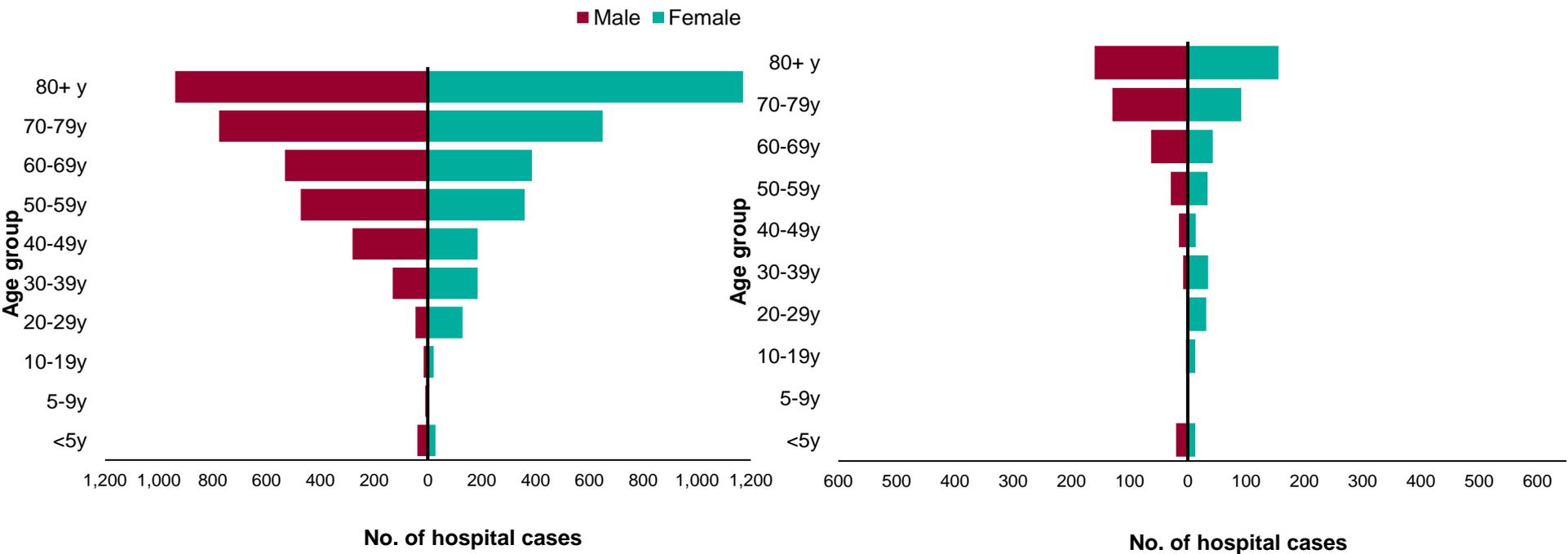
*Only NHS Acute trusts that have reported ≥ 1 day in the past week; excludes Specialist trusts. Acute NHS trusts (including Specialist trusts) reporting into SARI-Watch COVID-19 hospitalisation surveillance are typically around 100 per week. This was 86 for the hospitalisation (all levels of care) indicator in week 22 August 2022 to 28 August inclusive and 77 trusts for the ICU/HDU indicator. For the maps, as Specialist trusts are excluded, the number of trusts providing data on COVID-19 hospitalisations in week ending 28 August 2022 was 76 and 69 for ICU/HDU admissions for COVID-19



Age/sex pyramid of hospitalisations (all levels of care) for COVID-19, data from sentinel acute NHS trusts, England

(a) Peak of 2nd wave (week 53 2020 to week 3 2021) n= 6,359

(b) Most recent 4 weeks (week 31 2022 to 34 2022) n= 871



Reporting trusts=22

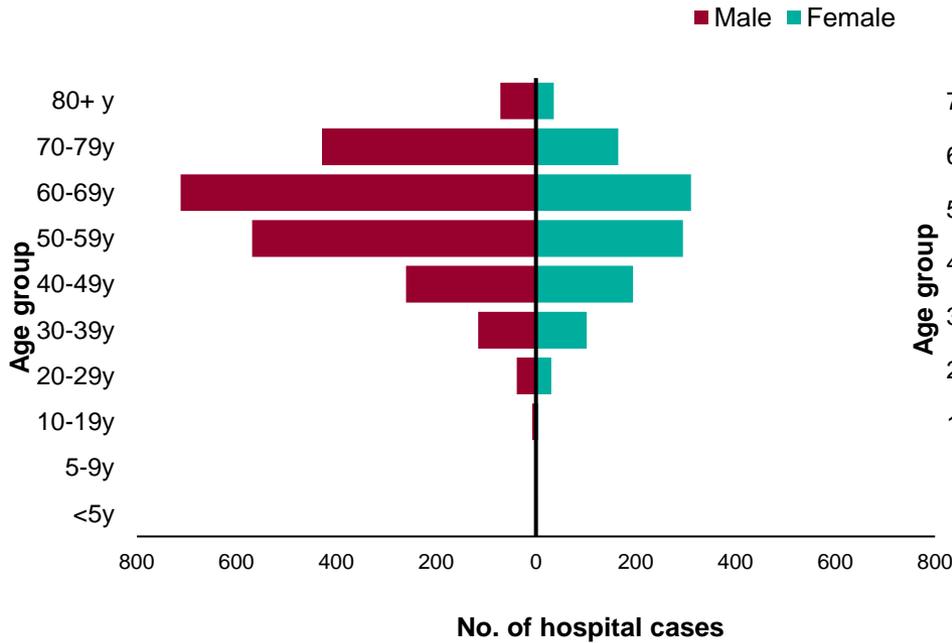
Reporting trusts=12



Age/sex pyramid for admissions to ICU/HDU for COVID-19, mandatory

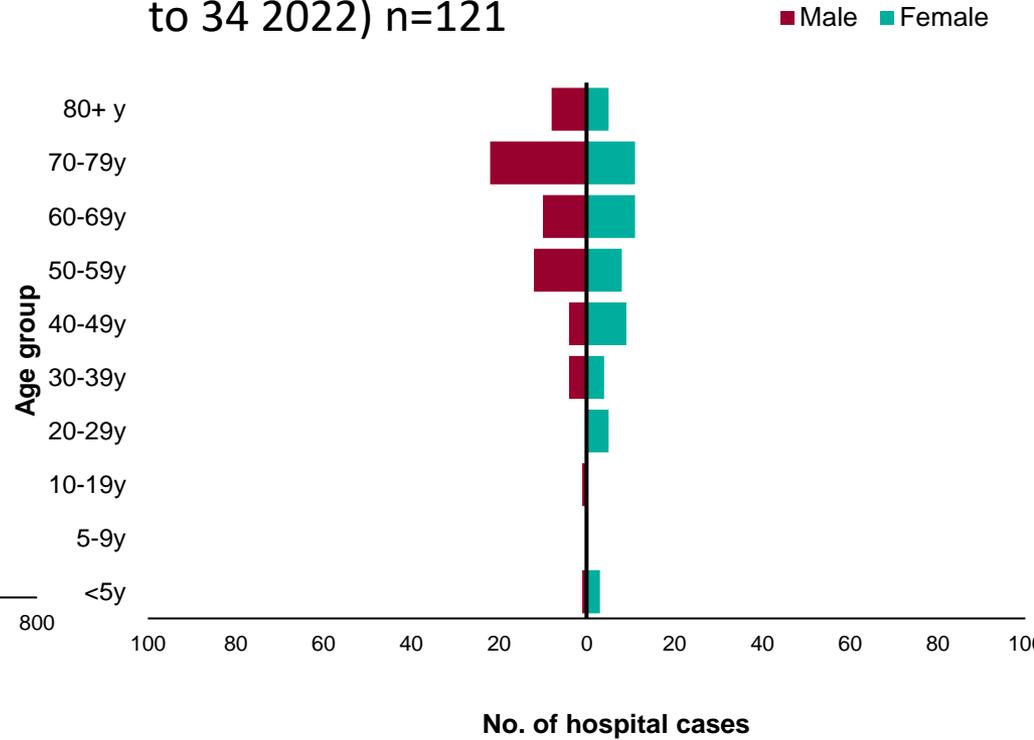
UK Health Security Agency
case level data, acute NHS trusts, England

(a) Peak of 2nd wave (week 53 2020 to week 3 2021) n= 3,349



Reporting trusts=70

(b) Most recent 4 weeks (week 31 2022 to week 34 2022) n=121

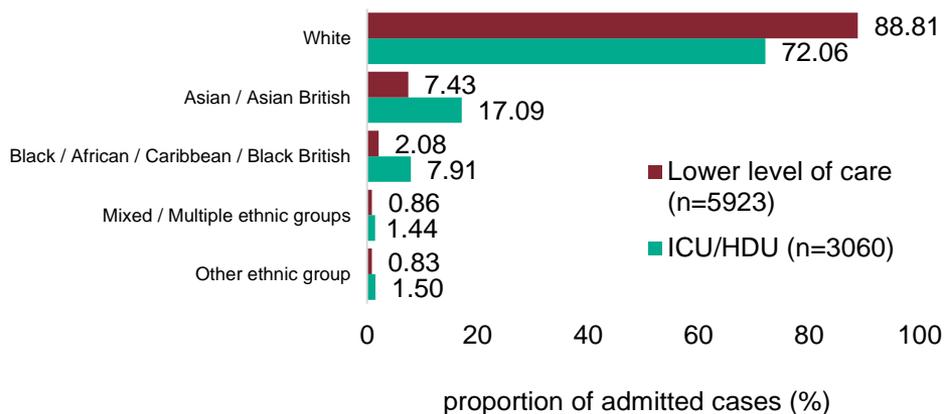


Reporting trusts=29



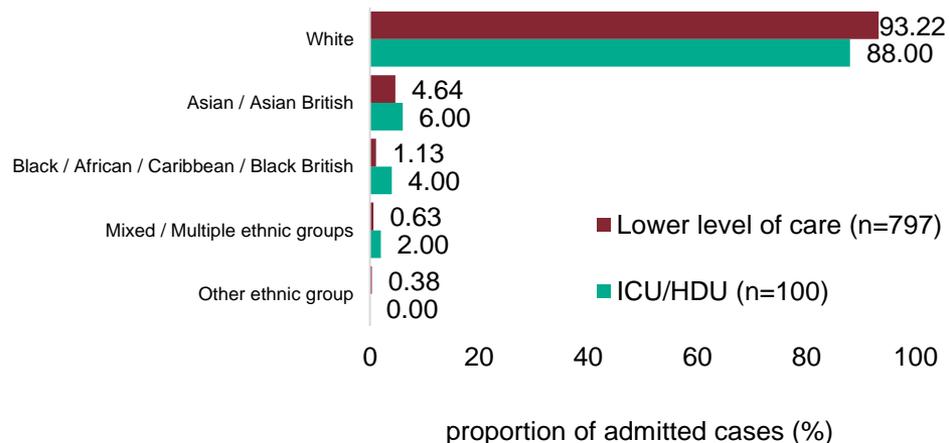
Laboratory confirmed admissions for COVID-19, to acute NHS trusts, by level of care and ethnicity

(a) Peak of 2nd wave (week 53 2020 to week 3 2021)



Reporting trusts
Lower level of care=21
ICU/HDU=68

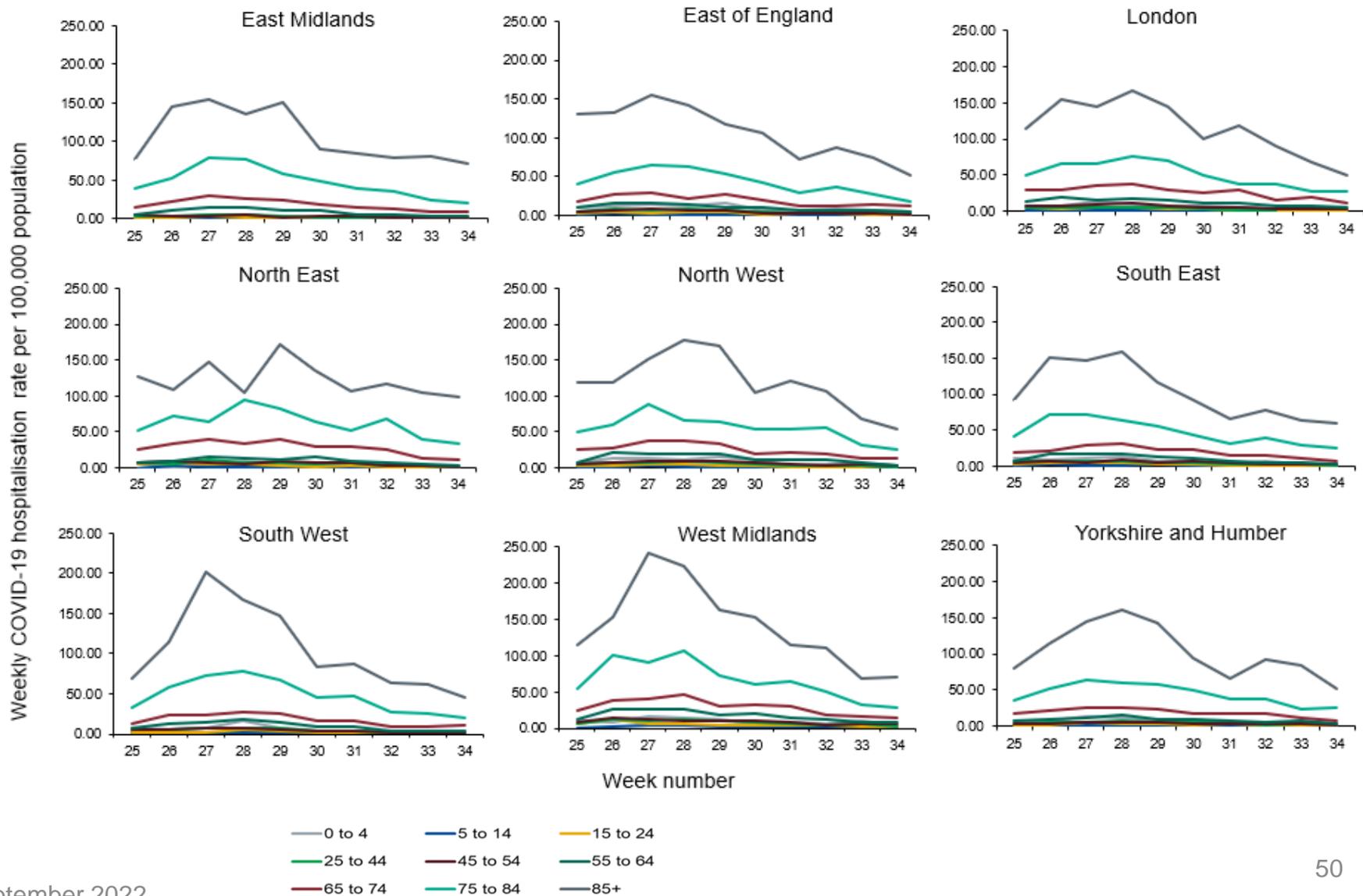
(b) Most recent 4 weeks (week 31 2022 to 34 2022)



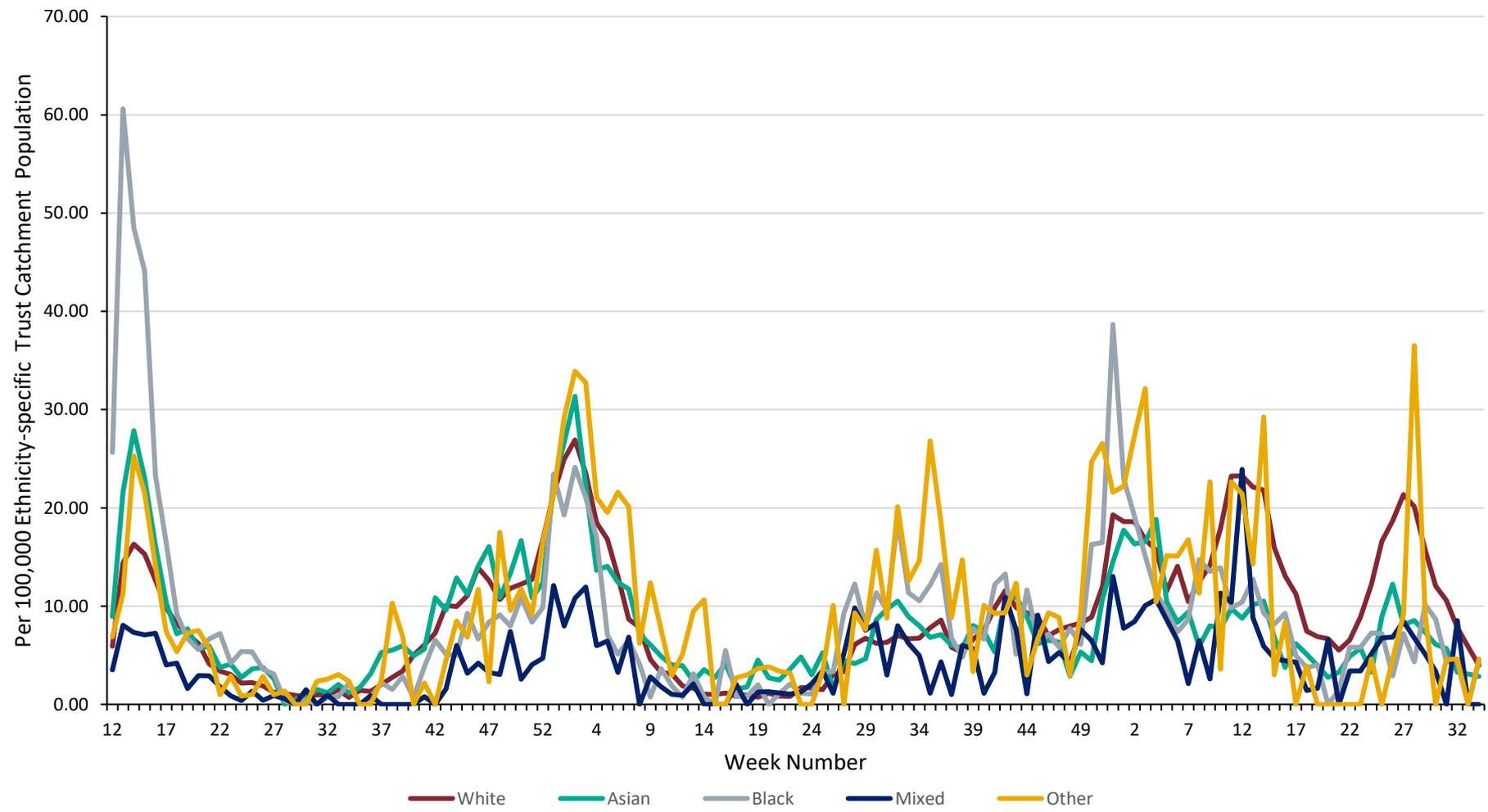
Reporting trusts
Lower level of care=12
ICU/HDU=26



Weekly COVID-19 hospitalisation rate per 100,000 trust catchment population by age group and region, weeks 25 to 34



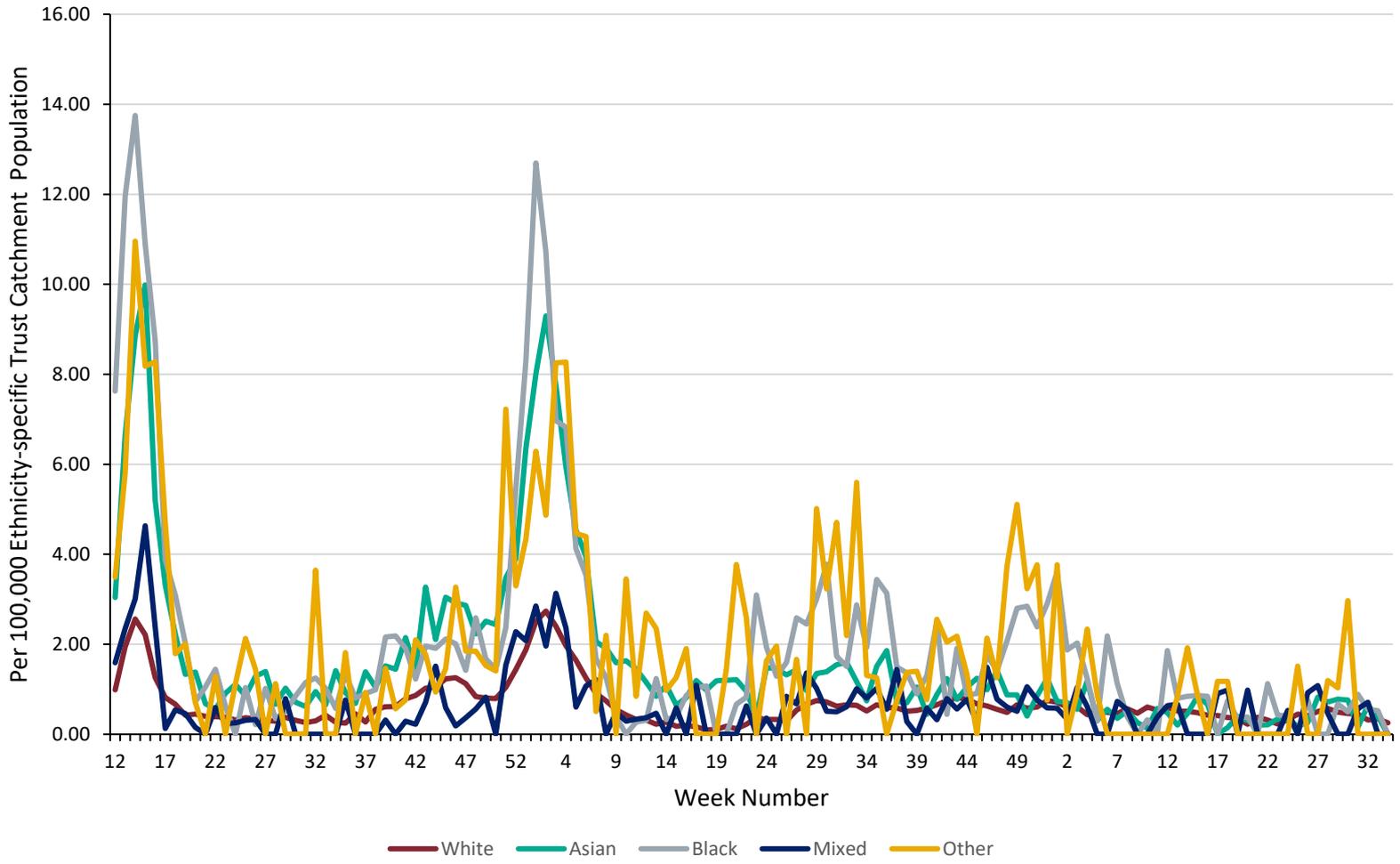
Hospital admission rate (excluding ICU/HDU) by ethnicity per 100,000 trust catchment population



Caveat: From week 24 (2021) the ethnicity analysis is based on a new method for assigning ethnicity, developed by UKHSA. The previous method used the most recent ethnicity recorded through linkage to Hospital Episode Statistics. However, this method led to unfeasibly high rates in the 'Other' ethnic group when applied to COVID-19 cases, hospitalisation or mortality. The new method uses the most frequent ethnicity recorded through linkage to Hospital Episode Statistics, unless the most frequent was 'Other' when the second most frequent was chosen.



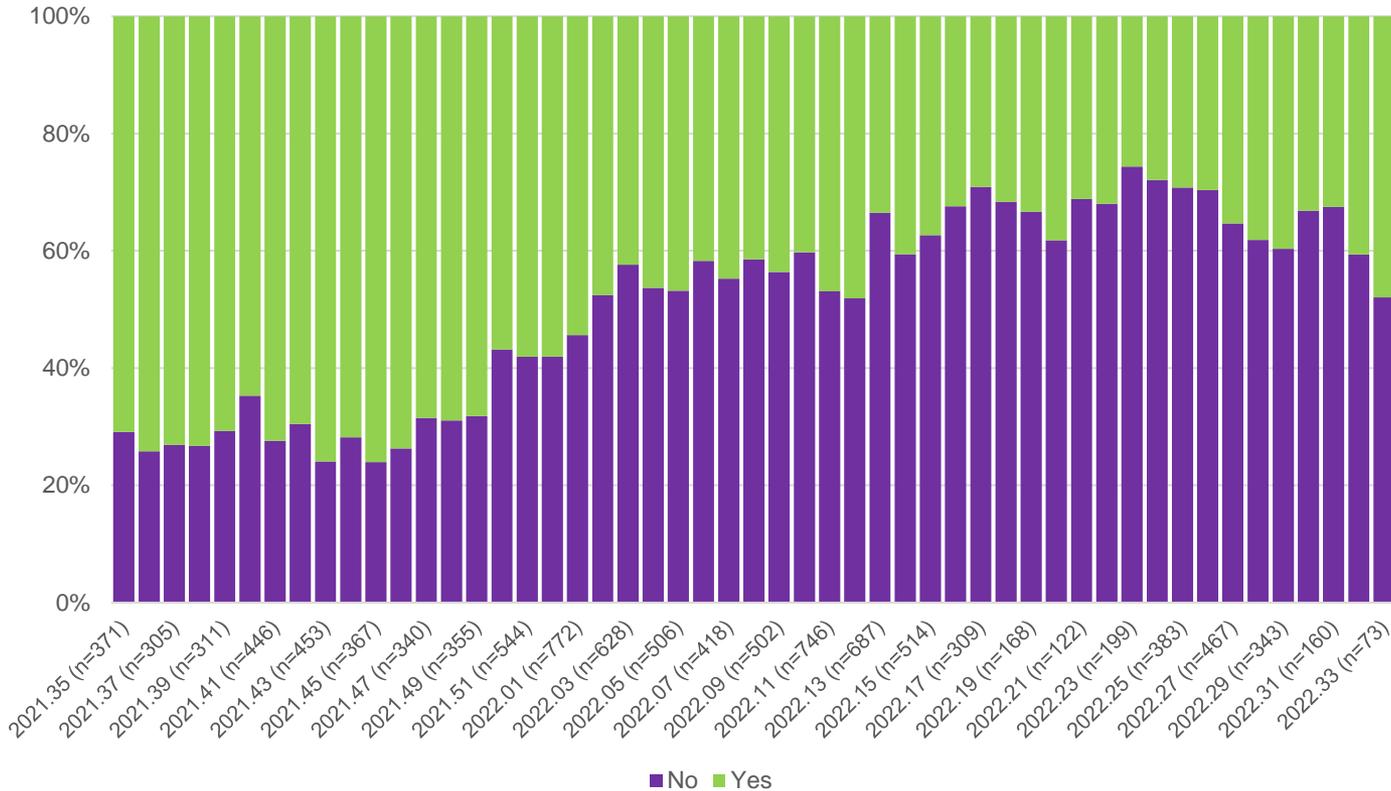
Rate of admission to ICU/HDU by ethnicity, per 100,000 trust catchment population



Caveat: From week (24 2021) the ethnicity analysis is based on a new method for assigning ethnicity, developed by UKHSA. The previous method used the most recent ethnicity recorded through linkage to Hospital Episode Statistics. However, this method led to unfeasibly high rates in the 'Other' ethnic group when applied to COVID-19 cases, hospitalisation or mortality. The new method uses the most frequent ethnicity recorded through linkage to Hospital Episode Statistics, unless the most frequent was 'Other' when the second most frequent was chosen.



COVID-19 as primary reason for admission among SARS-CoV-2 positive patient by week of admission



Notes

- 1) Case-level sentinel data from SARI-Watch, from week 35 2021 (commencing 30 August 2021) to week 33 2022 (ending 21 August 2022) inclusive
- 2) Total 27104 records in period of analysis, of which 35% (n=9498) had COVID-19 as primary reason for admission ('Yes').
- 3) SARS-CoV-2 patients with evidence of COVID-19 treatment but have 'No' or 'Unknown' for COVID-19 as primary reason for admission (n=924) are reassigned to COVID-19 as primary reason of admission ('Yes').
- 4) Reassignment increases COVID-19 as primary reason for admission ('Yes') from 9498 to 10422
- 5) 23% (6327/27104) of total records in this period have missing data on the 'Admission due to COVID-19' indicator – these are excluded from analysis
- 6) Caveats: London trusts under-represented and most recent weeks are subject to retrospective updates

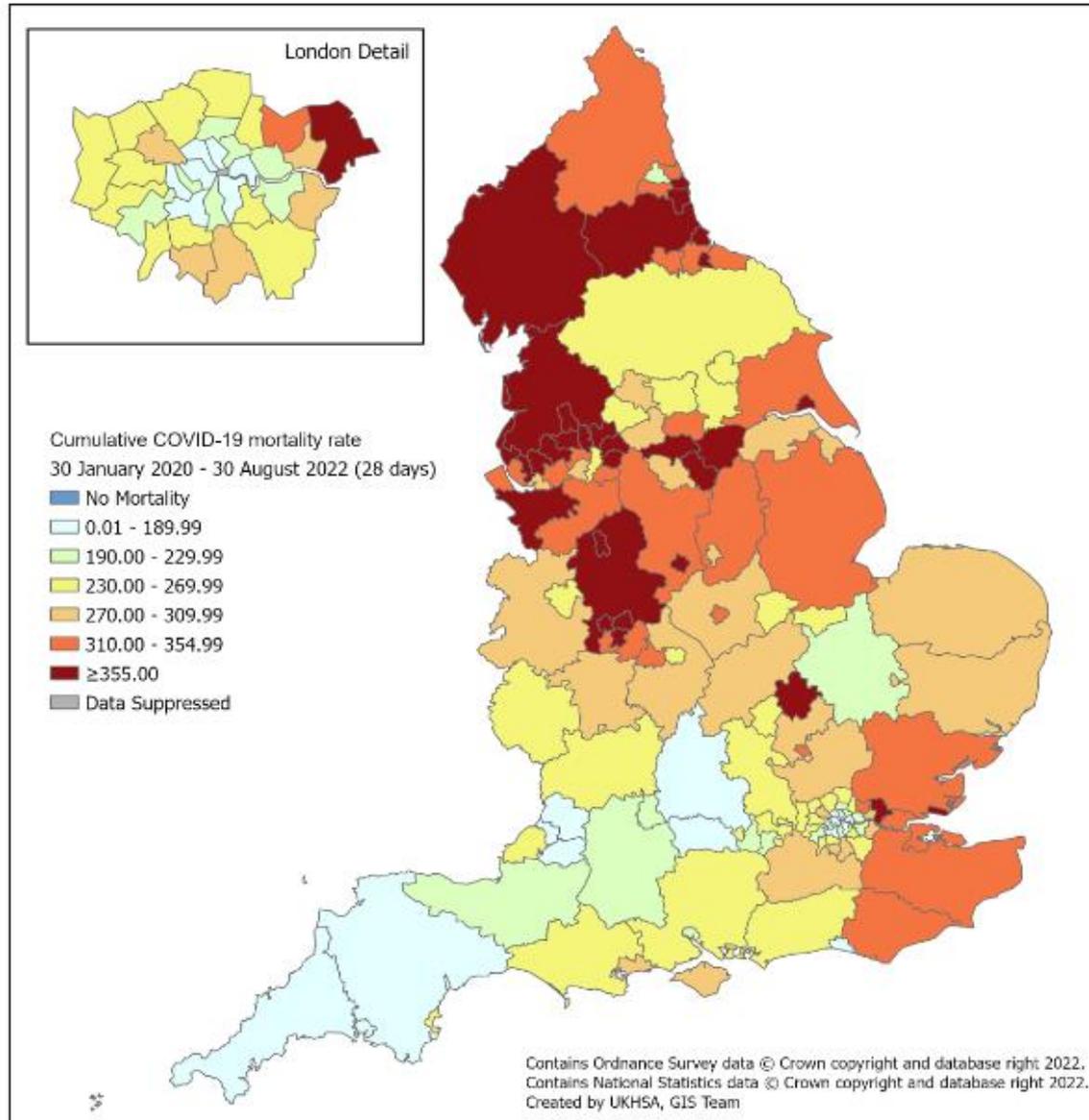


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Mortality surveillance



Cumulative mortality rate of COVID-19 cases per 100,000 population tested under Pillar 1 and 2 since the beginning of the pandemic by 28 day definition





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Co/secondary infections with COVID-19



Caveat - undertesting for other pathogens may result in an underestimate of co/secondary infection cases.

Preceding/co-/secondary infections refers to when a patient has a COVID-19 infection with one or more other pathogen (Please see Appendix 1 – Pre-/co-/secondary infection with COVID-19 definitions.)

- Preceding infection: COVID-19 acquired after another pathogen
- Co-infection: COVID-19 and other pathogen acquired at the same time
- Secondary infection: COVID-19 acquired before another pathogen

Numbers of pre-/co-/secondary infection remain low across UKHSA surveillance systems.

For patients with severe respiratory failure requiring Extra Corporeal Membrane Oxygenation (ECMO), analysis of data from five adult ECMO centres in England indicates that among patients with severe respiratory failure due to COVID-19, almost a third of these have co/secondary infections.

Published data analysis from pandemic wave 1 (W-1) indicates increased mortality associated with COVID-19 and [influenza](#), [key bacterial and fungal infections](#) and [invasive pneumococcal disease \(IPD\)](#) in comparison to patients without co/secondary infection.

[Data analysis](#) from W-1 indicates that *Aspergillus* and *candidemia* cases have increased risk of mortality in comparison to patients without co/secondary infection.



UK Health Security Agency Co/secondary infections among patients with severe respiratory failure requiring Extra Corporeal Membrane Oxygenation (ECMO)

Analysis is based on cumulative data from five adult ECMO centres in England. Data for the current and previous seasons are presented. Each season commences around October (ISO week 40) ending in September in the following year (ISO week 39).

Data for the current season (2021-22) is from 4 October 2021 to 15 May 2022 inclusive (week 40 2021 to week 19 2022). This period includes effects from the Delta and Omicron waves of the pandemic. The 2020-21 season is from 28 September 2020 to 3 October 2021 inclusive and includes effects from the Alpha and Delta waves. The 2019-20 season is from 30 September 2019 to 27 September 2020 inclusive and includes effects from the original Wuhan strain.

In the 2021-22 season, 41% (31/76) of ECMO patients admitted for severe respiratory failure due to laboratory confirmed COVID-19 had clinically significant co/secondary infections. In the previous season (2020-21) this proportion was 30% (134/402). In the 2019-20 season this proportion was 33% (79/236).

In all three seasons the majority of clinically significant co/secondary infections among respiratory failure COVID-19 cases comprised Gram-negative bacilli from the order Enterobacterales:

- 45% (14/31) in the current season 2021-22
- 32% (43/134) in 2020-21 and 46% (36/79) in 2019-20. The decrease in 2020-21 compared to 2019-20 reached borderline significance ($p=0.057$). (In the last data assessment, the decrease over this period was statistically significant but since then there were further updates to the data from ECMO centres). No change was detected in other key pathogens between these two time periods.



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Surveillance of bacterial, fungal and viral infections, in COVID-19 patients in England, Jan 2020 – June 2022

HCAI, Fungal, AMR, AMU & Sepsis Division



Updates

From 31 January 2022, UKHSA has changed the COVID-19 case definition to include multiple infection episodes. Reported co-/secondary/preceding infections in England now use the new definition, revising all cases back to the beginning of the pandemic.

The Unified Infection Dataset (UID) project has been extended to incorporate the Co- and Secondary infections with COVID-19 datasets

The following outputs included in this section have been produced via the UID, combining previously separate data pipelines

- Key HCAI bacterial and fungal specimens reported to SGSS and HCAI data capture system
- Respiratory viral specimens reported to SGSS and Respiratory Datamart
- Fungal specimens reported to mycology reference lab (MRL)

The Co- and secondary infections team have undertaken an extensive data validation exercise which has identified additional respiratory viral specimens from Respiratory Datamart and allowed us to make improvements to the methodologies. Preceding infections for all pipelines (other pathogen infections occurring before COVID-19 specimen) are now included. Please note, all cases since January 2020 have been revised in line with this validation.

Data are provisional and subject to change due to possible delayed reporting of microbiological samples



Number of COVID-19 patient-episodes with bacterial, fungal or viral infections in COVID-19 patients diagnosed in England during wave 3*, by infection type and timing of diagnosis

| Bacterial/ fungal/ viral infection by specimen type | COVID-19 patient-episodes with bacterial/ fungal/ viral infection | | Timing of bacterial/fungal/viral diagnosis in relation to COVID-19 diagnosis | | | | | | | | |
|--|---|------------------|--|----------------------|------------------|-------------|----------------------|------------------|---------------------|----------------------|------------------|
| | n | % of COVID cases | Preceding infection | | | Coinfection | | | Secondary infection | | |
| | | | n | % infections by site | % of COVID cases | n | % infections by site | % of COVID cases | n | % infections by site | % of COVID cases |
| Bacterial/fungal bloodstream & lower respiratory infection | 255 | <0.01 | 23 | 9.02 | <0.01 | 9 | 3.53 | <0.01 | 223 | 87.45 | <0.01 |
| Bacterial/fungal bloodstream infection | 8,189 | 0.06 | 3,477 | 42.46 | 0.02 | 1,631 | 19.92 | 0.01 | 3,081 | 37.62 | 0.02 |
| Bacterial/fungal lower respiratory infection | 2,646 | 0.02 | 622 | 23.51 | <0.01 | 250 | 9.45 | <0.01 | 1,774 | 67.04 | 0.01 |
| <i>Clostridioides difficile</i> infection | 1,040 | 0.01 | 419 | 40.29 | <0.01 | 102 | 9.81 | <0.01 | 519 | 49.90 | <0.01 |
| Fungal respiratory/bloodstream infection (MRL) ‡ | 198 | <0.01 | 14 | 7.07 | <0.01 | 3 | 1.52 | <0.01 | 181 | 91.41 | <0.01 |
| Other respiratory virus infection | 2,980 | 0.02 | 827 | 27.75 | 0.01 | 1,390 | 46.64 | 0.01 | 763 | 25.60 | 0.01 |
| Any site† | 15,345 | 0.10 | 5,394 | 35.15 | 0.04 | 3,386 | 22.07 | 0.02 | 6,565 | 42.78 | 0.04 |

Key findings:

- 0.1% of COVID-19 patient-episodes had a bacterial, fungal or other respiratory viral infection detected in either the 28 days prior or following (60 days following for MRL) their COVID-19 diagnosis
- Prevalence in W3 lower than W2 and W1; however, patient-episodes of COVID-19 and another key infection are still higher in W3 than W1 (15,345 vs 4,636, respectively)
- Most infections with key organisms were categorised as secondary infections (42.78%).

Please see appendix 1 for pre-/co-/secondary infection definitions with SARS-CoV-2
Please note patients can have multiple COVID-19 infection-episodes, numbers here do not reflect the number of patients.

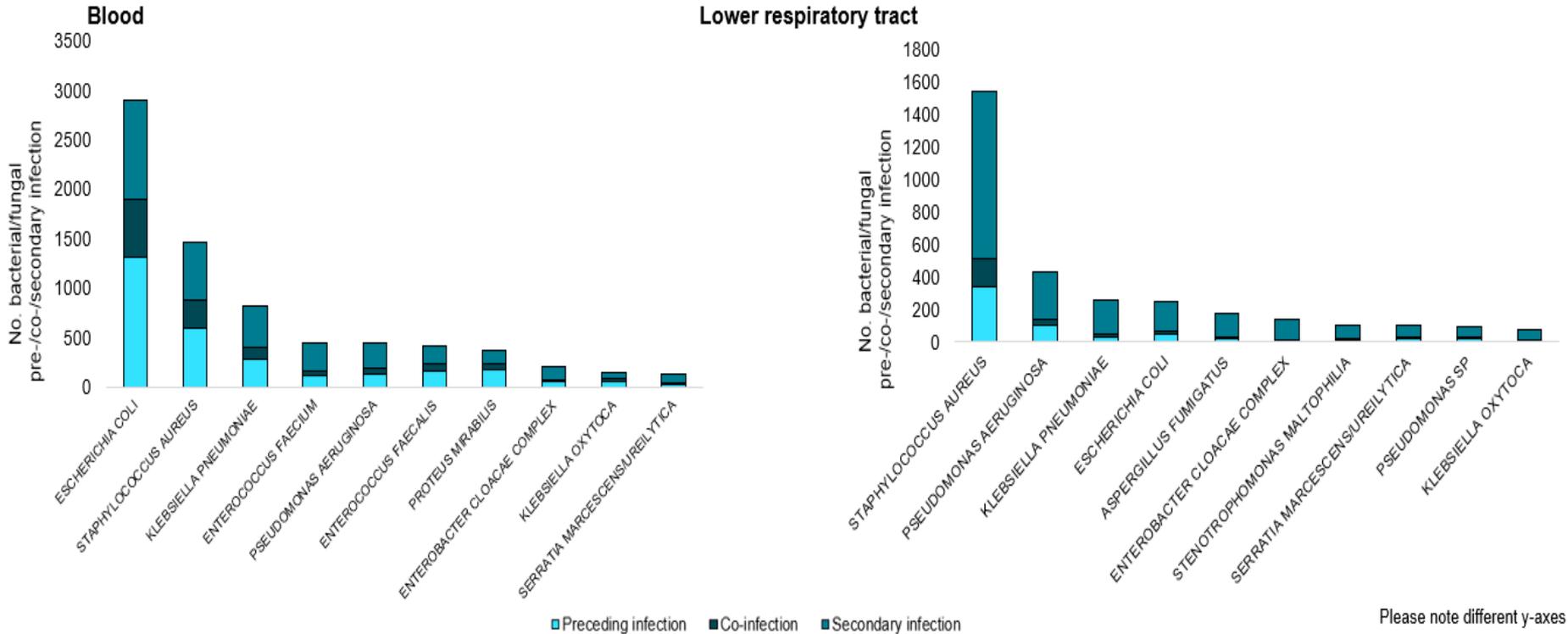
*SARS-CoV2 specimen dates from 27 Apr 2021 to 22 May 2022 (N=14,778,196). Last updated 22 Jun 2022.

‡ Definition for secondary infection differs for MRL specimens - detection within 60 days

† includes the combination Bacterial/fungal bloodstream & *Clostridioides difficile* infection (12 preceding, 1 coinfection & 20 secondary), Bacterial/fungal bloodstream, lower respiratory & *Clostridioides difficile* infection (1 secondary), & Bacterial/fungal lower respiratory & *Clostridioides difficile* infection (3 secondary)



Most frequent bacterial/fungal species in blood or lower respiratory tract specimens, by timing of diagnosis, in COVID-19 patients diagnosed in England during wave 3

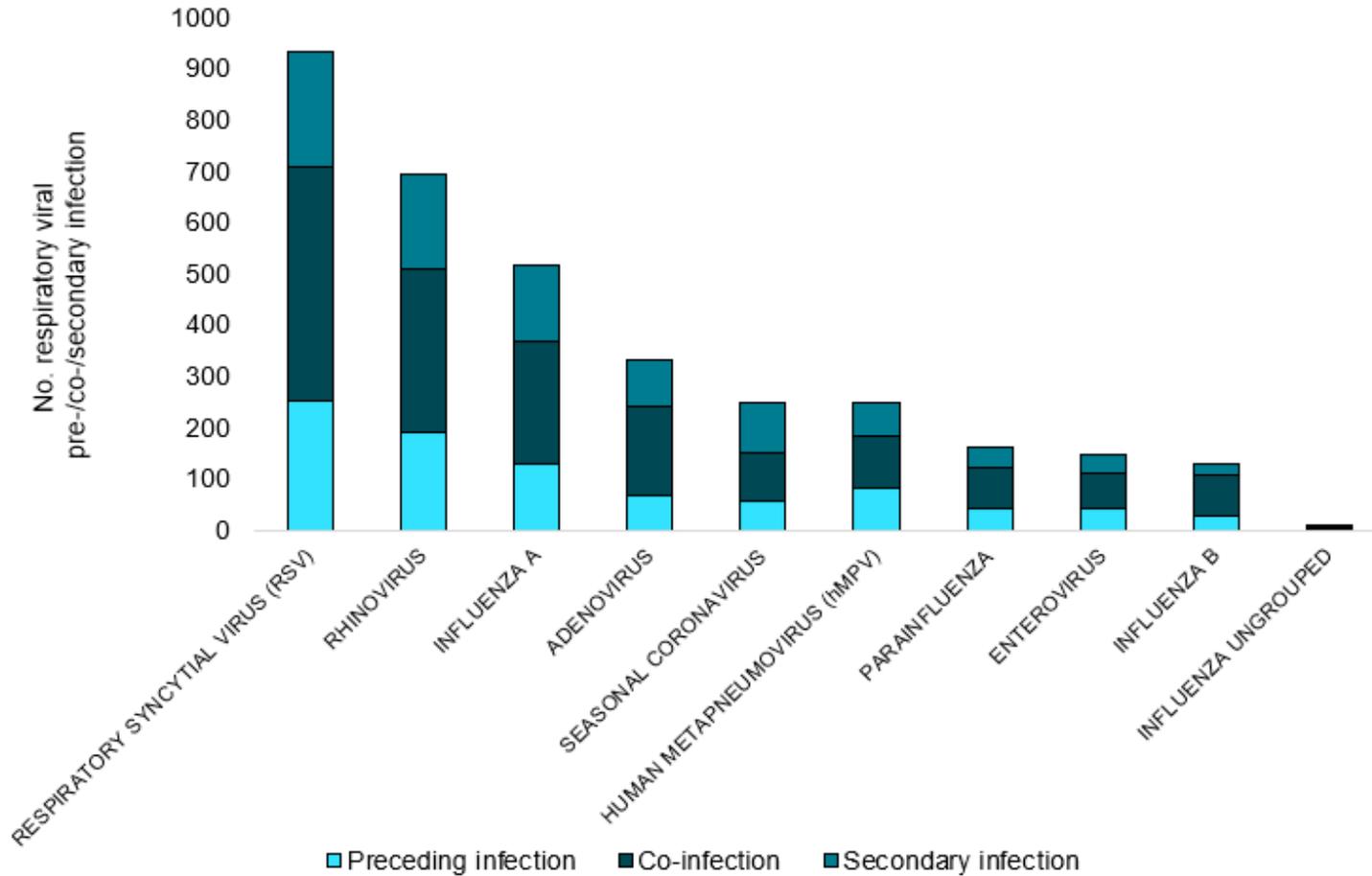


Key findings:

In wave 3, the most frequent bacterial/fungal organisms identified from blood specimens were *Escherichia coli*, *Staphylococcus aureus* and *Klebsiella pneumoniae* and from respiratory specimens were *Staphylococcus aureus*, *Pseudomonas aeruginosa* and *Klebsiella pneumoniae*.



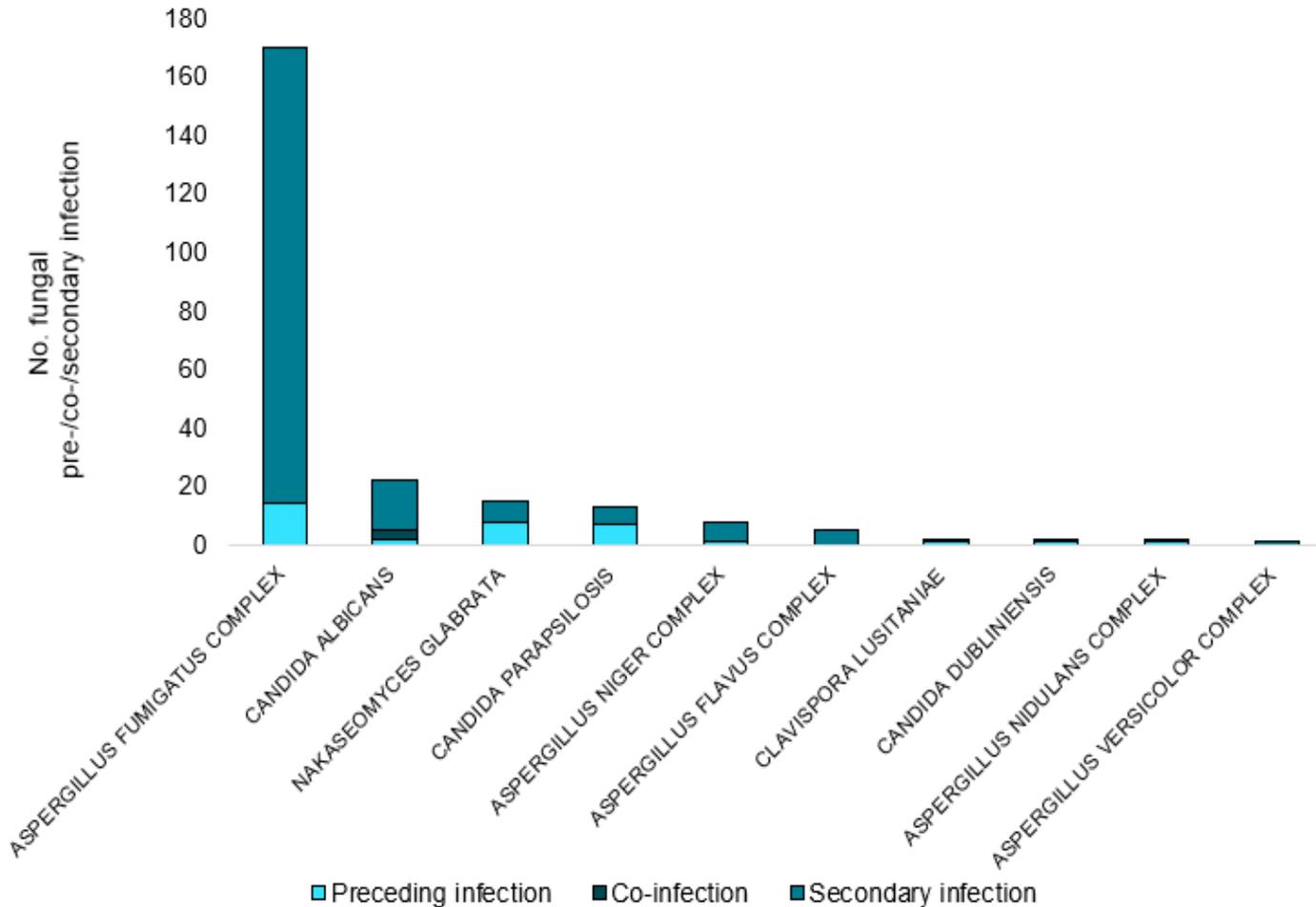
Most frequent viral specimens, by timing of diagnosis, in COVID-19 patients diagnosed in England during wave 3



Key findings:
 In wave 3, the most frequent viral organisms identified from respiratory specimens were RSV, rhinovirus and influenza A.



Most frequent fungal species (MRL), by timing of diagnosis, in COVID-19 patients diagnosed in England during wave 3



Key findings:
 In wave 3, the most frequent fungal organisms identified were *Aspergillus fumigatus* complex and *Candida albicans*.



| Bacteria/Fungi | First Wave | Second Wave | Third Wave | Total Cases |
|---|------------------------------|--------------------------------|---------------------------------|-------------|
| | (30 Jan 2020 - 28 June 2020) | (29 June 2020 – 30 April 2021) | (1 May 2021 – 24 February 2022) | |
| <i>Aspergillus fumigatus</i> isolates (azole resistant) | 46 (4) | 120 (2) | 137 (12) | 303 (18) |
| Probable/Proven cases of CAPA* | 15 | 38 | 44 | 97 |
| <i>Candida</i> spp.: Candidemia | 63 | 133 | 17 | 213 |
| <i>Bordetella pertussis</i> | 0 | 0 | 0 | 0 |
| <i>Haemophilus influenzae</i> | 3 | 2 | 0 | 5 |
| <i>Neisseria meningitidis</i> | 2 | 0 | 0 | 2 |
| <i>Streptococcus pneumoniae</i> | 40 | 45 | 14 | 99 |

*COVID-19-associated pulmonary aspergillosis

Please note fungal data refers to secondary infections only. Mycology data contains results from Mycology reference laboratory data, Candidaemia is representative of deep infection. One case of osteomyelitis, one case of ventriculitis and one case of endocarditis was documented in wave two. Fungal data are also included in the overall numbers in slides 6-8 but have been stratified here with additional details. *Bordetella pertussis* co-infection is defined as +/- 28 d Culture/PCR (based on pertussis sample date), +/- 28 Serology/Oral fluid (anti-pertussis toxin Ig) (based on pertussis symptom onset date, excluding cases without onset date). *Haemophilus influenzae*, *Neisseria meningitidis* and *Streptococcus pneumoniae* co-infection is defined as +/- 2d. *Legionella*, *Mycoplasma* and gastrointestinal infection data not included. Please note, testing in W1 was not open to the community and therefore W1 cases are predominantly hospitalised patients vs. W2 and W3.

Appendix 1: Pre-/co-/secondary infection definitions with COVID-19

The day pertains to the date of the sample collection that yielded a positive result. These definitions do not apply to persistent COVID-19 patients. Patients with persistent COVID-19 require independent clinical assessment.

| Organism | Definition co-infection with SARS-CoV-2 † | Definition of infection pre-SARS-CoV-2 infection (other pathogen is primary infection) or Definition of post SARS-CoV-2 secondary infection (SARS-CoV-2 is primary infection) |
|--|---|--|
| Influenza A | +/- 1d | 2-28d [^] |
| Influenza B | +/- 1d | 2-28d [^] |
| RSV | +/- 1d | 2-28d |
| Adenovirus | +/- 1d | 2-28d |
| Enterovirus | +/- 1d | 2-28d |
| Human metapneumovirus | +/- 1d | 2-28d |
| Parainfluenza (any subtype) | +/- 1d | 2-28d |
| Seasonal coronavirus | +/- 1d * | 2-28d |
| Rhinovirus | +/- 1d | 2-28d |
| Co-infections in ECMO patient (patients with most severe clinical respiratory signs) | | |
| ECMO patients | Individual case review | Individual case review |
| Blood stream and respiratory infections (bacterial and fungal) | | |
| <i>Achromobacter xylosoxidans</i> | +/- 1d | 2-28d |
| <i>Acinetobacter</i> spp., | +/- 1d | 2-28d |
| <i>Aspergillus</i> | +/- 1d | 2-28d (pre) 2-60d (post, continually hospitalised patients only) |
| <i>Bordetella pertussis</i> | +/- 28 d Culture/PCR (based on pertussis sample date) +/- 28 Serology/Oral fluid (anti-pertussis toxin Ig) (based on pertussis symptom onset date, excluding cases without onset date) | N/A (Pertussis presentation is often delayed) |
| <i>Burkholderia cepacia</i> | +/- 1d | 2-28d |
| <i>Candida</i> spp | +/- 1d | 2-28d (pre) 2-60d (post, continually hospitalised patients only) |
| <i>Chlamydia pneumoniae</i> | 0-7d PCR | PCR within 14-28 d (8-13d PCR*) |
| <i>Enterobacter</i> spp., | +/- 1d | 2-28d |
| <i>Enterococcus</i> spp. | +/- 1d | 2-28d |
| <i>E. coli</i> | +/- 1d | 2-28d |
| <i>Haemophilus influenzae</i> | +/- 2d | 3-28d |

Continued overleaf

Appendix 1 continued: Pre-/co-/secondary infection definitions with COVID-19

The day pertains to the date of the sample collection that yielded a positive result. These definitions do not apply to persistent COVID-19 patients. Patients with persistent COVID-19 require independent clinical assessment.

| Organism | Definition co-infection with SARS-CoV-2 † | Definition of infection pre-SARS-CoV-2 infection (other pathogen is primary infection) or Definition of post SARS-CoV-2 secondary infection (SARS-CoV-2 is primary infection) |
|--|---|--|
| Blood stream and respiratory infections (bacterial and fungal) | | |
| <i>Klebsiella</i> spp. | +/- 1d | 2-28d |
| <i>Legionella pneumophila/species</i> | Individual case review | Individual case review |
| <i>Mycoplasma pneumoniae</i> | 0-7d PCR, IgM serology 0-21d <16y | PCR within 14-28 d (8-13d PCR*) |
| <i>Neisseria meningitidis</i> | +/- 2d | 3-28d |
| <i>Pseudomonas</i> spp., | +/- 1d | 2-28d |
| <i>Serratia</i> spp., | +/- 1d | 2-28d |
| <i>Staphylococcus aureus</i> | +/- 1d | 2-28d |
| Coag-neg <i>Staphylococcus</i> (<i>S. haemolyticus</i>) | +/- 1d | 2-28d |
| <i>Stenotrophomonas</i> spp., (<i>S. maltophilia</i>) | +/- 1d | 2-28d |
| <i>Streptococcus</i> spp. ‡ | +/- 1d | 2-28d |
| <i>Streptococcus pneumoniae</i> | +/- 2d | 3-28d |
| Tuberculosis | | |
| <i>Mycobacterium tuberculosis</i> | Individual case review | Individual case review |
| Pathogens of the immunocompromised (eg HIV) | | |
| HIV | Individual case review | Individual case review |
| Gastrointestinal infections | | |
| Listeria | 0-5d * | Individual case review |
| Campylobacter | 0-5d * | Individual case review |
| Shiga toxin-producing <i>E. coli</i> (STEC) | 0-5d * | Individual case review |
| Norovirus | 0-5d * | Individual case review |
| Salmonella | 0-5d * | Individual case review |
| Shigella | 0-5d * | Individual case review |
| Anaerobes | | |
| <i>C. difficile</i> | +/- 1d | 2-28d |
| Bacteroides sp. (<i>B. fragilis</i> and non- <i>fragilis</i> Bacteroides) | +/- 1d | 2-28d |

See next slides for notes

Appendix 1 continued: Pre-/co-/secondary infection definitions with COVID-19

Notes

† From SARS-CoV-2 first detection date. Not including multiple episodes of SARS-CoV-2 per patient.

* Additional data check required. (Resistance is not detailed, data for MERS is not currently available).

^ Definition post- SARS-CoV-2 secondary infection (SARS-CoV-2 is primary infection). This has been extended from prior 14d secondary infection definition for influenza used by PHE to account for disparities in testing throughout the 28d period after SARS-CoV-2 detection.

‡ Streptococcus species includes the following groups and species:

| Group | Species/other names |
|---|---|
| Anginosus Group | <i>Streptococcus anginosus</i> ; <i>Streptococcus constellatus</i> (<i>Streptococcus constellatus</i> subspecies <i>constellatus</i> <i>Streptococcus constellatus</i> subspecies <i>pharynges</i>); <i>Streptococcus</i> Group F; <i>Streptococcus intermedius</i> ; <i>Streptococcus milleri</i> group; <i>Streptococcus sinensis</i> |
| Bovis Group | <i>Streptococcus alactolyticus</i> ; <i>Streptococcus bovis</i> untyped; <i>Streptococcus equinus</i> ; <i>Streptococcus gallolyticus</i> subspecies <i>gallolyticus</i> (<i>Streptococcus bovis</i> biotype I); <i>Streptococcus infantarius</i> (<i>Streptococcus infantarius</i> sp <i>infantarius</i> ; <i>Streptococcus bovis</i> biotype II); <i>Streptococcus lutetiensis</i> ; <i>Streptococcus infantarius</i> subspecies <i>coli</i> (<i>Streptococcus bovis</i> biotype II); <i>Streptococcus pasteurianus</i> (<i>Streptococcus bovis</i> biotype II) |
| Closely Related Genera | <i>Abiotrophia</i> spp.; <i>Aerococcus</i> spp.; <i>Faklamia</i> spp.; <i>Gemella</i> spp.; <i>Globicatella sanguinis</i> ; <i>Granulicatella</i> spp.; <i>Leuconostoc</i> spp.; <i>Pedicoccus</i> spp.; <i>Peptostreptococcus</i> spp. |
| Mitis Group | <i>Streptococcus cristatus</i> ; <i>Streptococcus mitior</i> ; <i>Streptococcus mitis</i> ; <i>Streptococcus oralis</i> ; <i>Streptococcus pseudopneumoniae</i> ; <i>Streptococcus infantis</i> ; <i>Streptococcus peroris</i> |
| Mutans Group | <i>Streptococcus mutans</i> ; <i>Streptococcus sobrinus</i> |
| Other streptococci (including but not limited to) | Anaerobic streptococcus; <i>Streptococcus acidominimus</i> ; <i>Streptococcus</i> spp., other named/not fully identified; <i>Streptococcus suis</i> ; <i>Streptococcus uberis</i> |
| Salivarius Group | <i>Streptococcus vestibularis</i> ; <i>Streptococcus thermophilus</i> |
| Sanguinis Group | <i>Streptococcus gordonii</i> ; <i>Streptococcus massiliensis</i> ; <i>Streptococcus parasanguinis</i> ; <i>Streptococcus sanguinis</i> |
| <i>Streptococcus</i> Group A | Group A; <i>Streptococcus pyogenes</i> ; <i>Streptococcus dysgalactiae</i> subspecies <i>equisimilis</i> |
| <i>Streptococcus</i> Group B | Group B; <i>Streptococcus agalactiae</i> |
| <i>Streptococcus</i> Group C | Group C; <i>Streptococcus dysgalactiae</i> subspecies <i>equisimilis</i> ; <i>Streptococcus equi</i> subspecies <i>zooepidemicus</i> |
| <i>Streptococcus</i> Group G | Group G; <i>Streptococcus canis</i> ; <i>Streptococcus dysgalactiae</i> subspecies <i>equisimilis</i> |