Influenza and Other Respiratory Viruses
Weekly Report
California Influenza Surveillance Program

Highlights (Week 50: December 8, 2019 – December 14, 2019)

Statewide Activity

- **Deaths:** 39* since Sept. 29, 2019
- **Outbreaks:** 10 since Sept. 29, 2019
- **Laboratory:** 21.0% flu positive
- **Outpatient ILI:** Above expected levels
- **Hospitalizations:** Within expected levels

*Influenza-coded deaths from death certificates

Click on images and links for more information

Key messages:

- Flu activity continues to increase in California.
- Flu B viruses are predominating in California, but A (H1)pdm09 are also circulating.
- Haven’t gotten a flu shot yet? Get immunized now to protect you and your loved ones this holiday season.
- Take action to stop the spread of flu: wash hands often, cover coughs and sneezes, and stay home when sick.

Note: This report includes data from many sources of influenza surveillance and it should be viewed as a preliminary “snapshot” of influenza activity for each surveillance week. Because data are preliminary, the information may be updated in later reports as additional data are received. These data should not be considered population-based or representative of all California public health jurisdictions.
A. Outpatient, Inpatient, and Death Data

1. Influenza Sentinel Providers

Sentinel providers (physicians, nurse practitioners, and physician assistants) situated throughout California report on a weekly basis the number of patients seen with influenza-like illness (ILI) and the total number of patients seen for any reason. ILI is defined as any illness with fever (≥100°F or 37.8°C) AND cough and/or sore throat (in the absence of a known cause other than influenza).

A total of 85 enrolled sentinel providers have reported data for Week 50. Based on available data, the percentage of visits for ILI during Week 50 was 3.0% compared to Week 49 (2.9%) and is above expected levels for this time of year (Figure 1).

Figure 1. Percentage of Influenza-like Illness Visits Among Patients Seen by California Sentinel Providers, 2015–2020

The seasonal baseline was calculated using a regression model applied to data from the previous five years. Two standard deviations above the seasonal baseline is the point at which the observed percentage of ILI is significantly higher than would be expected at that time of year.

2. Kaiser Permanente Hospitalization Data

Inpatients at Kaiser Permanente facilities with an admission diagnosis including the keywords “flu,” “influenza,” “pneumonia,” or variants of the keywords are defined as pneumonia and influenza (P&I)-related admissions. The number of P&I admissions is divided by the total number of hospital admissions occurring in the same time period to estimate the percentage of P&I admissions. Admissions for pregnancy, labor and delivery, birth, and outpatient procedures are excluded from the denominator.

The percentage of admissions for pneumonia and influenza (P&I) in Kaiser Permanente facilities in northern California during Week 50 was 5.1% compared to Week 49 (4.9%) and is within expected levels for this time of the year (Figure 2).
The seasonal baseline was calculated using a regression model applied to data from the previous five years. Two standard deviations above the seasonal baseline and is the point at which the observed percentage of pneumonia and influenza hospitalizations in Kaiser Permanente hospitals in Northern California is significantly higher than would be expected at that time of the year.

The majority of admissions for pneumonia and influenza did not result in intensive care unit (ICU) admission or death; however, 222 ICU admissions and 126 deaths have occurred among persons with P&I admission diagnoses (Figure 3a). The majority of P&I admissions occurred among persons ≥65 years of age across all severity categories, especially among deaths (Figure 3b). Please note that pneumonia and influenza admissions serve as a proxy for influenza activity, but do not necessarily represent laboratory-confirmed influenza infections.

Figure 3. Number (a) and age group distribution (b) of non-ICU, ICU, and deaths associated with P&I Admissions in Kaiser Permanente Northern California hospitals, 2019–2020 season to date
3. Influenza-Associated Hospitalizations, California Emerging Infections Program

The California Emerging Infections Program (CEIP), Influenza Surveillance Network (FluSurv-NET) conducts population-based surveillance for laboratory-confirmed influenza-associated hospitalizations among patients of all ages in Alameda, Contra Costa, and San Francisco counties.

The incidence of influenza-associated hospitalizations per 100,000 population was the same in week 48 (0.89) compared to Week 47 (0.89) (Figure 4). Data for the most recent two weeks are not presented because results are still being collected and are likely to change.
4. Influenza Mortality Surveillance from Death Certificates

Deaths occurring in California among residents who had influenza noted in any cause of death field on the death certificate (text or coded) are defined as influenza-coded deaths. The percentage of influenza-coded deaths is calculated by dividing the number of influenza-coded deaths by the total number of all cause deaths during the same period. Influenza-coded deaths are not necessarily laboratory-confirmed and are an underestimate of all influenza-associated deaths.

During Week 50, 15 new influenza-coded death were identified. To date during the 2019–2020 influenza season, 39 influenza-coded deaths have been identified (Figure 5). The percentage of deaths coded as influenza during Week 50 was 0.2% compared to 0.3% during Week 49 (Figure 6).
Figure 5. Number of Influenza-coded Deaths Identified from Death Certificates by Week of Death, 2019–2020 Season

Note: Coding of deaths can be delayed by several weeks. Influenza-coded deaths will be included once enough information is available to identify them.

Figure 6. Percentage of Influenza-coded Deaths Occurring in California among California Residents, 2015–2020

Note: Coding of deaths can be delayed by several weeks. Influenza-coded deaths will be included once enough information is available to identify them.
To date, more deaths have occurred among persons ≥65 years of age (69.2%) than among persons <65 years of age during the 2019–2020 influenza season; however, the percentage of deaths occurring among persons <65 years of age (30.8%) is consistent with other seasons during which influenza viruses other than influenza A (H3N2) have circulated in greater numbers, such as the 2015–2016 and 2018–2019 seasons (Figure 7).

Figure 7. Age Distribution of Influenza-coded Deaths Occurring in California among California Residents, 2015–2016 Season through 2019–2020 Seasons

![Bar chart showing age distribution of influenza-coded deaths](chart.png)

* Methods used to identify pediatric influenza-coded deaths on death certificates differ from those used to identify the influenza-associated pediatric deaths presented below.
† One death during the 2018–2019 influenza season has unknown age and is not included in the figure.
‡ 2015–2016 influenza season: October 4, 2015–October 1, 2016; mixed influenza A and influenza B season
2016–2017 influenza season: October 2, 2016–September 30, 2017; influenza A (H3N2) predominant season
2017–2018 influenza season: October 1, 2017–September 29, 2018; influenza A (H3N2) predominant season
2018–2019 influenza season: September 30, 2018–September 28, 2019; mixed influenza A (H1N1)pdm09 and influenza A (H3N2) season
2019–2020 influenza season: September 29, 2019–September 26, 2020; influenza B (Victoria) viruses predominating, influenza A (H1N1)pdm09 viruses are also circulating

5. Laboratory-Confirmed Severe Influenza-associated Pediatric Deaths

Influenza-associated deaths in children <18 years of age are nationally notifiable. The weekly influenza report includes confirmed deaths formally reported to CDPH through December 14, 2019 (Week 50).
One laboratory-confirmed influenza-associated fatality in a child <18 years of age was reported to CDPH during Week 50. To date, CDPH has received four reports of laboratory-confirmed influenza-associated deaths among children <18 years of age during the 2019–2020 influenza season.

B. Laboratory Update – Influenza

1. Respiratory Laboratory Network (RLN) and Clinical Sentinel Laboratory Surveillance Results

Laboratory surveillance for influenza and other respiratory viruses involves the use of data from clinical sentinel laboratories (hospital, academic, and private laboratories) and public health laboratories in the Respiratory Laboratory Network located throughout California. These laboratories report the number of laboratory-confirmed influenza and other respiratory virus detections and isolations on a weekly basis.

The overall percentage of influenza detections in clinical sentinel laboratories in Week 50 (21.0%) was higher than Week 49 (17.1%) (Figure 8). Additional details, including influenza typing and subtyping information from public health laboratories can be found in Figures 8 and 9 and Tables 1 and 2.

Neither the RLN nor CDPH-VRDL has identified any influenza viruses by polymerase chain reaction (PCR) that are suggestive of a novel influenza virus.

Figure 8. Percentage of Influenza Detections at Clinical Sentinel Laboratories, 2015–2020
Figure 9. Number of Influenza Detections by Type and Subtype Detected in the Respiratory Laboratory Network, 2019–2020

Table 1. Respiratory Specimens Testing Positive for Influenza — Clinical Sentinel Laboratories, Current Week and Season to Date

<table>
<thead>
<tr>
<th></th>
<th>Current Week Number</th>
<th>Current Week Percent</th>
<th>Season to Date Number</th>
<th>Season to Date Percent</th>
</tr>
</thead>
<tbody>
<tr>
<td>Number of Specimens Tested</td>
<td>3,628</td>
<td></td>
<td>31,494</td>
<td></td>
</tr>
<tr>
<td>Influenza Positive</td>
<td>762</td>
<td>21.0</td>
<td>3,316</td>
<td>10.5</td>
</tr>
<tr>
<td>A</td>
<td>238</td>
<td>31.2*</td>
<td>1,156</td>
<td>34.9*</td>
</tr>
<tr>
<td>B</td>
<td>524</td>
<td>68.8*</td>
<td>2,160</td>
<td>65.1*</td>
</tr>
</tbody>
</table>

* Percent of specimens positive for influenza
Table 2. Respiratory Specimens Testing Positive for Influenza by Influenza Type and Subtype — Respiratory Laboratory Network, Current Week and Season to Date

<table>
<thead>
<tr>
<th>Influenza Positive</th>
<th>Current Week Number</th>
<th>Current Week Percent</th>
<th>Season to Date Number</th>
<th>Season to Date Percent</th>
</tr>
</thead>
<tbody>
<tr>
<td>A</td>
<td>81</td>
<td>41.1*</td>
<td>506</td>
<td>49.1*</td>
</tr>
<tr>
<td>A (H1)pdm09</td>
<td>69</td>
<td>85.2†</td>
<td>359</td>
<td>70.9†</td>
</tr>
<tr>
<td>A (H3)</td>
<td>5</td>
<td>6.2†</td>
<td>129</td>
<td>25.5†</td>
</tr>
<tr>
<td>A, not subtyped</td>
<td>7</td>
<td>8.6†</td>
<td>18</td>
<td>3.6†</td>
</tr>
<tr>
<td>B</td>
<td>116</td>
<td>58.9*</td>
<td>524</td>
<td>50.9*</td>
</tr>
<tr>
<td>B Victoria</td>
<td>59</td>
<td>50.9‡</td>
<td>324</td>
<td>61.8‡</td>
</tr>
<tr>
<td>B Yamagata</td>
<td>1</td>
<td>0.9‡</td>
<td>18</td>
<td>3.4‡</td>
</tr>
<tr>
<td>B, not lineage typed</td>
<td>56</td>
<td>48.3‡</td>
<td>182</td>
<td>34.7‡</td>
</tr>
</tbody>
</table>

* Percent of specimens positive for influenza  
† Percent of influenza A positives  
‡ Percent of influenza B positives

2. Influenza Virus Characterization

Close monitoring of influenza viruses is required to better assess the potential impact on public health. CDC characterizes influenza viruses through one or more tests including genetic characterization by analyzing viral genome sequences, and antigenic characterization by hemagglutination inhibition (HI) assays and/or neutralization based Focus Reduction assays (FRA). These data are used to compare how similar, or well-inhibited, currently circulating influenza viruses are to the reference viruses. Antigenic and genetic characterization of circulating influenza viruses gives an indication of the influenza vaccines ability to induce an immune response against the wide array of influenza viruses that are co-circulating every season. However, annual vaccine effectiveness estimates are needed to determine how much protection was provided to the population by vaccination.

Twelve influenza A (H3N2) viruses from California have been genetically characterized to date during the 2019–2020 influenza season, of which one was antigenically characterized. All 12 influenza A (H3N2) viruses were members of the 3c.2a1 subclade. The antigenically characterized virus was well-inhibited by A/Kansas/14/2017-like (H3N2) antisera (Table 2a).

Eight influenza A (H1N1)pdm09 viruses from California have been genetically characterized to date during the 2019–2020 influenza season, of which one was antigenically characterized. All eight influenza A (H1N1)pdm09 viruses were members of the 6B.1A clade. The antigenically characterized virus was well-inhibited by A/Brisbane/02/2018-like (H1N1) antisera (Table 2b).

Eighteen influenza B (Victoria) viruses from California have been genetically characterized to date during the 2019–2020 influenza season, of which three were
antigenically characterized. All 18 influenza B (Victoria) viruses were members of V1A.3 subclade. Of the three influenza B (Victoria) viruses antigenically characterized, one (33.3%) was well-inhibited by B/Colorado/06/2017-like (Victoria) antisera and two (66.7%) were poorly inhibited by B/Colorado/06/2017-like (Victoria) antisera (Table 2c).

Two influenza B (Yamagata) viruses from California have been genetically characterized to date during the 2019–2020 influenza season. Both influenza B (Yamagata) viruses were members of the Y3 clade (Table 2d). Antigenic characterization information will be presented once available.

Table 2 a–d. Influenza virus antigenic characterization and genomic sequencing by influenza A subtype and influenza B lineage type — California, 2019–2020 influenza season

a. Influenza A (H3N2) viruses

<table>
<thead>
<tr>
<th>Antigenic Characterization</th>
<th>3C.2a1</th>
<th>3C.3a*</th>
<th>Total</th>
</tr>
</thead>
<tbody>
<tr>
<td>Well inhibited by A/Kansas/14/2017*</td>
<td>1</td>
<td>0</td>
<td>1</td>
</tr>
<tr>
<td>Poorly inhibited by A/Kansas/14/2017*</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Not antigenically characterized</td>
<td>11</td>
<td>0</td>
<td>11</td>
</tr>
<tr>
<td><strong>Total A (H3N2)</strong></td>
<td>12</td>
<td>0</td>
<td>12</td>
</tr>
</tbody>
</table>

* A/Kansas/14/2017 is the influenza A (H3N2) component included in the 2019–2020 quadrivalent and trivalent influenza vaccines and is a member of the 3c.3a clade

b. Influenza A (H1N1)pdm09 viruses

<table>
<thead>
<tr>
<th>Antigenic Characterization</th>
<th>6B.1</th>
<th>6B.1A*</th>
<th>Total</th>
</tr>
</thead>
<tbody>
<tr>
<td>Well inhibited by A/Brisbane/02/2018*</td>
<td>0</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>Poorly inhibited by A/Brisbane/02/2018*</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Not antigenically characterized</td>
<td>0</td>
<td>7</td>
<td>7</td>
</tr>
<tr>
<td><strong>Total A (H1N1)pdm09</strong></td>
<td>0</td>
<td>8</td>
<td>8</td>
</tr>
</tbody>
</table>

* A/Brisbane/02/2018 is the influenza A (H1N1)pdm09 component included in the 2019–2020 quadrivalent and trivalent influenza vaccines and is a member of the 6B.1A clade

c. Influenza B (Victoria) viruses

<table>
<thead>
<tr>
<th>Antigenic Characterization</th>
<th>V1A.1*</th>
<th>V1A.3</th>
<th>Total</th>
</tr>
</thead>
<tbody>
<tr>
<td>Well inhibited by B/Colorado/06/2017*</td>
<td>0</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>Poorly inhibited by B/Colorado/06/2017*</td>
<td>0</td>
<td>2</td>
<td>2</td>
</tr>
<tr>
<td>Not antigenically characterized</td>
<td>0</td>
<td>15</td>
<td>15</td>
</tr>
<tr>
<td><strong>Total B (Victoria)</strong></td>
<td>0</td>
<td>18</td>
<td>18</td>
</tr>
</tbody>
</table>

* B/Colorado/06/2017-like is the influenza B (Victoria) lineage component included in the 2019–2020 quadrivalent and trivalent influenza vaccines and is a member of the V1A.1 subclade
d. Influenza B (Yamagata) viruses

<table>
<thead>
<tr>
<th>Antigenic Characterization</th>
<th>Y3*</th>
<th>Total</th>
</tr>
</thead>
<tbody>
<tr>
<td>Well inhibited by B/Phuket/3073/2013*</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Poorly inhibited by B/Phuket/3073/2013*</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Not antigenically characterized</td>
<td>2</td>
<td>2</td>
</tr>
<tr>
<td><strong>Total B (Yamagata)</strong></td>
<td>2</td>
<td>2</td>
</tr>
</tbody>
</table>

* B/Phuket/3073/2013 is the influenza B (Yamagata) component included in the 2019–2020 quadrivalent influenza vaccine and is a member of the Y3 clade

C. Influenza-Associated Outbreaks

Four laboratory-confirmed influenza outbreaks were reported to CDPH during Week 50. To date, 10 laboratory-confirmed influenza outbreaks have been reported to CDPH for the 2019–2020 season.

**Figure 10. Number of Laboratory-Confirmed Influenza-Associated Outbreaks by Week of First Onset, 2018–2020**

D. California Border Region Influenza Surveillance Network Data

The border influenza surveillance network is comprised of outpatient provider sentinel sites whose geographical coverage extends approximately 100 kilometers (60 miles)
north of the California-Baja California border and includes Imperial and San Diego Counties, as well as some parts of Riverside County.

1. Syndromic Surveillance Update
A total of 8 border region sentinel providers reported data during Week 50. The total number of patients screened by all sentinel sites for ILI during Week 50 was 7,519. Outpatient ILI activity was 0.9% in Week 50. ILI activity for the California border region during Week 50 was lower when compared to activity for the same week during the 2017–2018 season and 2018–2019 season (Figure 11). All influenza syndromic data summarized for the border region represent a subset of CDC influenza sentinel providers in California.

![Figure 11. Percentage of Influenza-like Illness Visits among Patients Seen by Sentinel Providers — California Border Region, 2017–2020](image)

2. Virologic Surveillance Update
During Week 50, 600 respiratory specimens were tested from border region sentinel clinical laboratories; of these, 151 (25.2%) tested positive for influenza (18 [11.9%] influenza A and 133 [88.1%] influenza B). Cumulatively this season, a total of 4,199 respiratory specimens were tested from border region sentinel clinical laboratories; of these, 589 (14.0%) tested positive for influenza (79 [13.4%] influenza A; 510 [86.6%] influenza B).

During Week 50, 47 influenza positive specimens were detected at border region RLN laboratories; of which, 16 (34.0%) were influenza A and 31 (66.0%) were influenza B.
the specimens that tested positive for influenza A at RLN laboratories, 14 (87.5%) were subtyped as A (H1) pdm09 and two (12.5%) were subtyped as A (H3). Of the specimens that tested positive for influenza B at RLN laboratories, one (3.2%) was lineage typed as B (Yamagata) and 30 (96.8%) were lineage typed as B (Victoria). Cumulatively this season, a total of 126 influenza positive specimens have been detected at border region RLN laboratories; of which, 53 (42.1%) were influenza A and 73 (57.9%) were influenza B. Of the 53 specimens that tested positive for influenza A at RLN laboratories, 33 (62.3%) were subtyped as A (H1) pdm09 and 20 (37.7%) were subtyped as A (H3). Of the 73 specimens that tested positive for influenza B, three (4.1%) were lineage typed as B (Yamagata) and 70 (95.9%) were lineage typed as B (Victoria).

Laboratory data summarized in Figure 12 include data from border region influenza clinical sentinel laboratories (percentage of specimens testing positive for influenza) as well as data from border region RLN laboratories (influenza type and subtype/lineage type).

Figure 12. Number of Influenza Detections by Type and Subtype Detected in Respiratory Laboratory Network Laboratories and the Percentage of Specimens Testing Positive at Clinical Sentinel Laboratories — California Border Region, 2019–2020
E. Other Respiratory Viruses

1. Laboratory-Confirmed Severe Respiratory Syncytial Virus Case Reports

Currently, as mandated under Section 2500 of the California Code of Regulations, deaths among children aged 0–4 years with laboratory-confirmed respiratory syncytial virus (RSV) are reportable to CDPH. The weekly influenza report includes confirmed deaths formally reported to CDPH through December 14, 2019 (Week 50).

No laboratory-confirmed RSV-associated deaths among children <5 years of age were reported to CDPH during Week 50. To date, CDPH has received no reports of laboratory-confirmed RSV-associated deaths among children <5 years of age during the 2019–2020 influenza season.

2. Other Respiratory Virus Laboratory Update

During Week 50, 3,210 specimens were tested for RSV and 283 (8.8%) were positive, which is higher compared to Week 49 (6.5%) (Figure 13). During Week 50, adenovirus, coronavirus, human metapneumovirus, and rhinovirus/enterovirus activity increased; and parainfluenza virus activity remained the same (Figure 14).

Figure 13. Percentage of RSV Detections at Clinical Sentinel Laboratories, 2015–2020
Figure 14. Percentage of Other Respiratory Pathogen Detections at Clinical Sentinel Laboratories, 2019–2020

- Parainfluenza types 1-4
- Adenovirus
- Human Metapneumovirus
- Rhinovirus/Enterovirus
- Coronavirus

Percentage of Specimens Tested Positive

Week

Oct Nov Dec Jan Feb Mar Apr May Jun July Aug Sept
**Activity Levels:**

**No Activity:** No laboratory-confirmed cases of influenza and no reported increase in the number of cases of ILI.

**Sporadic:** Small numbers of laboratory-confirmed influenza cases or a single laboratory-confirmed influenza outbreak has been reported, but there is no increase in cases of ILI.

**Local:** Outbreaks of influenza or increases in ILI cases and recent laboratory-confirmed influenza in a single region of the state.

**Regional:** Outbreaks of influenza or increases in ILI and recent laboratory confirmed influenza in at least two but less than half the regions of the state with recent laboratory evidence of influenza in those regions.

**Widespread:** Outbreaks of influenza or increases in ILI cases and recent laboratory-confirmed influenza in at least half the regions of the state with recent laboratory evidence of influenza in the state.

**California Regions:**

**Northern:** Alpine, Amador, Butte, Colusa, Del Norte, El Dorado, Glenn, Humboldt, Lake, Lassen, Mendocino, Modoc, Nevada, Placer, Plumas, Sacramento, Shasta, Sierra, Siskiyou, Sutter, Tehama, Trinity, Yolo, and Yuba counties

**Bay Area:** Alameda, Contra Costa, Marin, Napa, Solano, San Francisco, San Mateo, Santa Clara, Santa Cruz, and Sonoma counties

**Central Valley:** Calaveras, Fresno, Inyo, Kings, Mono, Madera, Mariposa, Merced, Monterey, San Benito, San Joaquin, Stanislaus, Tulare, and Tuolumne counties

**Upper Southern:** Kern, Los Angeles, San Luis Obispo, Santa Barbara, and Ventura counties

**Lower Southern:** Imperial, Orange, Riverside, San Bernardino, and San Diego counties

For questions regarding influenza surveillance and reporting in California, please email InfluenzaSurveillance@cdph.ca.gov. This account is monitored daily by several epidemiologists.

To obtain additional information regarding influenza, please visit the CDPH influenza website (https://www.cdph.ca.gov/Programs/CID/DCDC/Pages/Immunization/Influenza.aspx).

A copy of the case report form for reporting any laboratory-confirmed influenza case that was either admitted to the ICU or died can be downloaded from the CDPH influenza website (https://www.cdph.ca.gov/Programs/CID/DCDC/Pages/Immunization/Influenza.aspx).

For information about national influenza activity, please visit the Centers for Disease Control and Prevention’s FluView (https://www.cdc.gov/flu/weekly/index.htm) and FluView Interactive (https://www.cdc.gov/flu/weekly/fluviewinteractive.htm) websites.