Influenza and Other Respiratory Viruses Weekly Report

California Influenza Surveillance Program

)CDPH

Highlights (Week 7: February 9, 2020 – February 15, 2020)

Statewide Activity

No Activity	Sporadic	Local	Regional	Widespread
		• Deaths	381* since Sept	. 29, 2019
		• <u>Outbrea</u>	aks: 111 since S	Sept. 29, 2019
Regions		► <u>Labora</u>	tory: 29.0% flu	positive
Elevated	Activity	▶ <u>Outpati</u>	i ent ILI: Above	e expected levels
		▶ <u>Hospita</u>	alizations: Abo	ove expected levels
	-		d deaths from death c and links for more info	
		Key mes	sages:	
	A l	• Flu activity	remains elevated in	California.
	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	• Flu A (H1)p	odm09 viruses are	now the predominate
S. S. S.				oria) viruses are still
			ember 29, 2019, 12 pediatric deaths ha	
			spread; it's not too	
_			>6 months of age n	
		• Pregnant w	omen, children und	der 5, adults 65+,
		and people	with chronic condi	tions are at high risk
		for flu-relat	ted complications.	

**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.

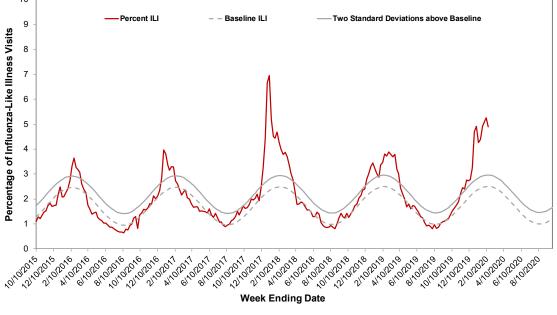
#### A. Outpatient, Inpatient, and Death Data

#### 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 83 enrolled sentinel providers have reported data for Week 7. Based on available data, the percentage of visits for ILI during Week 7 was 4.9% compared to Week 6 (5.3%) and is above expected levels for this time of year (Figure 1).





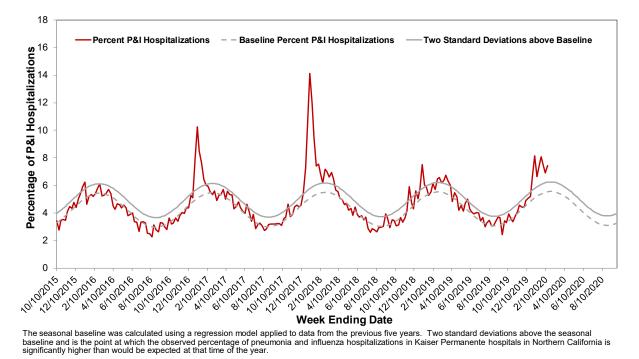
The seasonal baseline was calculated using a regression model applied to data from the previous five years. Two standard deviations above the seasonal baseline is the point at which the observed percentage of ILI is significantly higher than would be expected at that time of year.

#### 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.

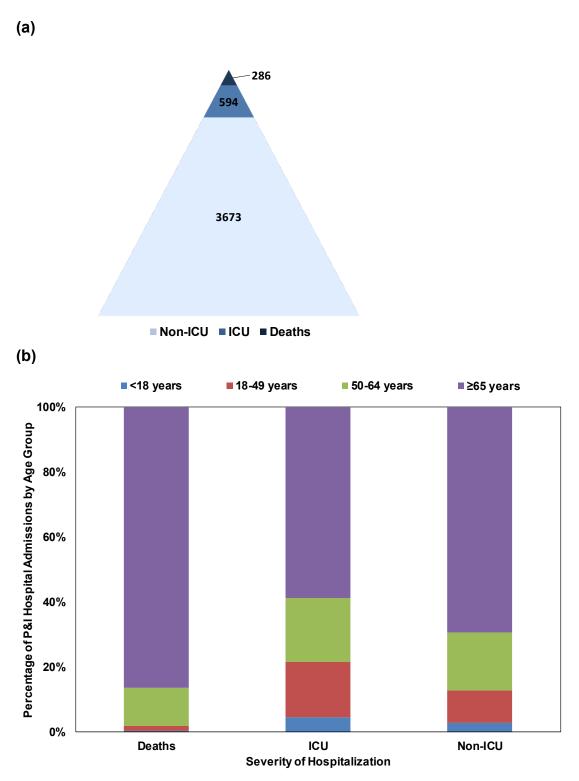
The percentage of admissions for pneumonia and influenza (P&I) in Kaiser Permanente facilities in northern California during Week 7 was 7.4% compared to Week 6 (6.9%) and is above expected levels for this time of the year (Figure 2).

### Figure 2. Percentage of P&I Admissions in Kaiser Permanente Northern California Hospitals, 2015–2020



The majority of admissions for pneumonia and influenza did not result in intensive care unit (ICU) admission or death; however, 594 ICU admissions and 286 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.

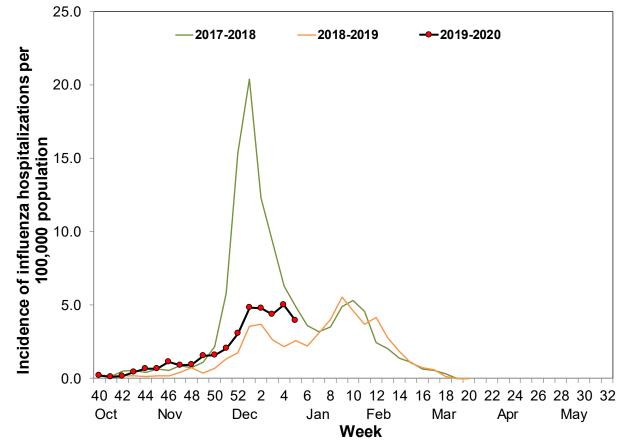
Figure 3. Number (a) and age group distribution (b) of non-ICU, ICU, and deaths associated with P&I Admissions in Kaiser Permanente Northern California hospitals, 2019–2020 season to date



#### 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 lower in week 5 (3.95) compared to Week 4 (5.01) (Figure 4). Data for the most recent two weeks are not presented because results are still being collected and are likely to change.

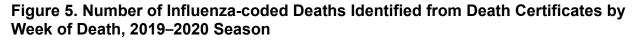


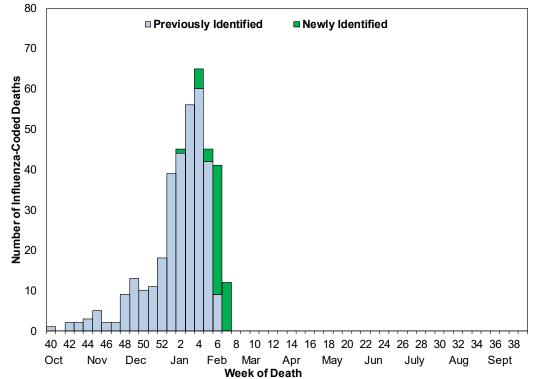


#### 4. Influenza Mortality Surveillance from Death Certificates

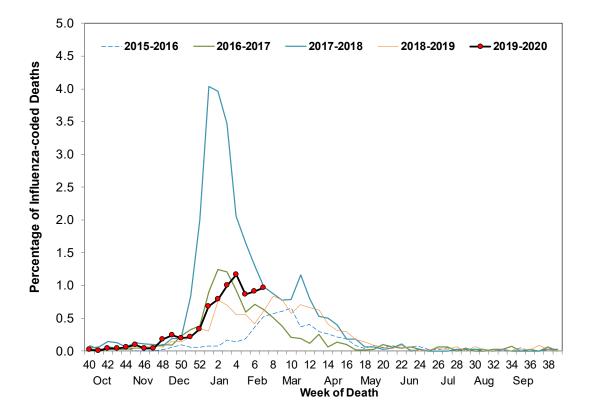
Deaths occurring in California among residents who had influenza noted in any cause of death field on the death certificate (text or coded) are defined as "influenza-coded deaths." The percentage of influenza-coded deaths is calculated by dividing the number of influenza-coded deaths by the total number 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 7, 53 new influenza-coded deaths were identified. To date during the 2019–2020 influenza season, 381 influenza-coded deaths have been identified (Figure 5). The percentage of deaths coded as influenza during Week 7 was 1.0% compared to 0.9% during Week 6 (Figure 6).





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

To date, more deaths have occurred among persons  $\geq$ 65 years of age (63.3%) 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 (36.7%) 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).

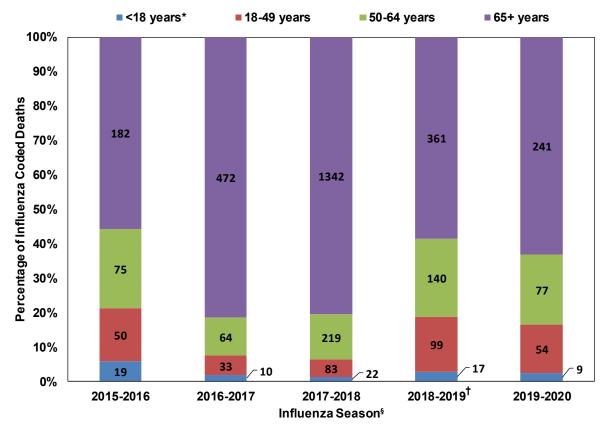


Figure 7. Age Distribution of Influenza-coded Deaths Occurring in California among California Residents, 2015–2016 Season through 2019–2020 Seasons

* Methods used to identify pediatric influenza-coded deaths on death certificates differ from those used to identify the influenzaassociated 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; mixed influenza B (Victoria) and influenza A (H1N1)pdm09 season

#### 5. Laboratory-Confirmed 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 February 15, 2020 (Week 7). Methods used to identify pediatric influenza-coded deaths on death certificates differ from those used to identify the influenza-associated pediatric deaths presented below and might not include the same individuals.

No laboratory-confirmed influenza-associated fatalities in children <18 years of age were reported to CDPH during Week 7. To date, CDPH has received 12 reports of laboratory-confirmed influenza-associated deaths among children <18 years of age during the 2019–2020 influenza season.

#### B. Laboratory Update – Influenza

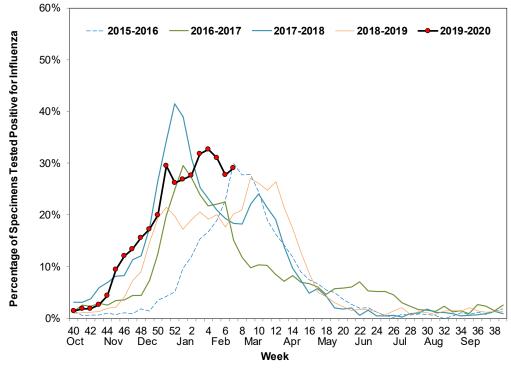
#### <u>1. Respiratory Laboratory Network (RLN) and Clinical Sentinel Laboratory Surveillance</u> <u>Results</u>

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 7 (29.0%) was higher than Week 6 (27.7%) (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.







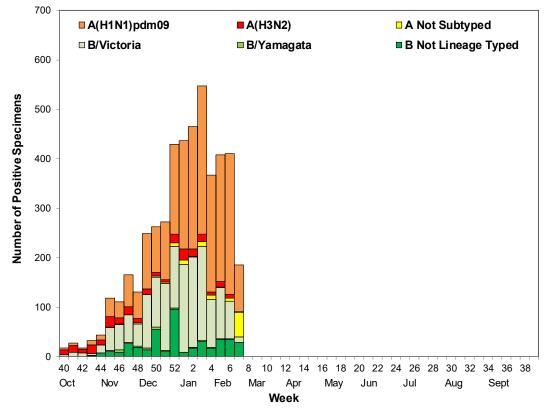


 Table 1. Respiratory Specimens Testing Positive for Influenza — Clinical Sentinel

 Laboratories, Current Week and Season to Date

	Current Week Number	Current Week Percent	Season to Date Number	Season to Date Percent
Number of Specimens Tested	5,679		85,002	
Influenza Positive	1,647	29.0	18,991	22.3
Α	1,242	75.4 [*]	10,318	54.3 [*]
В	405	24.6*	8,673	45.7 [*]

* Percent of specimens positive for influenza

Table 2. Respiratory Specimens Testing Positive for Influenza by Influenza Typeand Subtype — Respiratory Laboratory Network, Current Week and Season toDate

	Current Week Number	Current Week Percent	Season to Date Number	Season to Date Percent
Influenza Positive	185	Fercent	4,693	Fercent
Α	145	78.4*	2,697	57.5 [*]
A (H1)pdm09	94	64.8†	2,358	87.4†
A (H3)	2	1.4†	236	8.8†
A, not subtyped	49	33.8†	103	3.8†
В	40	21.6*	1,996	42.5*
B Victoria	11	27.5 [‡]	1,548	77.6‡
B Yamagata	0	0.0 [‡]	32	1.6 [‡]
B, not lineage typed	29	72.5 [‡]	416	20.8 [‡]

* Percent of specimens positive for influenza

† Percent of influenza A positives

‡ Percent of influenza B positives

#### 2. Antiviral Resistance Testing

Of the influenza specimens tested by the CDPH-VRDL to date this season, none have been found to be resistant to Oseltamivir (Table 3).

#### Table 3. Number of Specimens Tested for Oseltamivir Resistance, 2019–2020

	Oseltamivir Resistance
Influenza A (H1)pdm09	0/56
Influenza A (H3)	0/32
Influenza B	0/71

#### 3. 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 <u>vaccine</u> effectiveness estimates are needed to determine how much protection was provided to the population by vaccination.

Twenty-three influenza A (H3N2) viruses from California have been genetically characterized to date during the 2019–2020 influenza season, of which eight were

antigenically characterized. One influenza A (H3N2) virus was a member of the 3c.3a clade and 22 viruses were members of the 3c.2a1 subclade. Of the eight influenza A (H3N2) viruses antigenically characterized, four (50%) were well-inhibited by A/Kansas/14/2017-like (H3N2) antisera and four (50%) were poorly inhibited by A/Kansas/14/2017-like (H3N2) antisera (Table 4a).

Forty-four influenza A (H1N1)pdm09 viruses from California have been genetically characterized to date during the 2019–2020 influenza season, of which nine were antigenically characterized. All 44 influenza A (H1N1)pdm09 viruses were members of the 6B.1A clade. Of the nine influenza A (H1N1)pdm09 viruses antigenically characterized, nine (100%) were well-inhibited by A/Brisbane/02/2018-like (H1N1) antisera (Table 4b).

Fifty-two influenza B (Victoria) viruses from California have been genetically characterized to date during the 2019–2020 influenza season, of which eight were antigenically characterized. Five influenza B (Victoria) viruses were members of the V1A.1 subclade and 47 were members of the V1A.3 subclade. Of the eight influenza B (Victoria) viruses antigenically characterized, four (50%) were well-inhibited by B/Colorado/06/2017-like (Victoria) antisera and four (50%) were poorly inhibited by B/Colorado/06/2017-like (Victoria) antisera (Table 4c).

Six influenza B (Yamagata) viruses from California have been genetically characterized to date during the 2019–2020 influenza season, of which one was antigenically characterized. All six influenza B (Yamagata) viruses were members of the Y3 clade. The antigenically characterized virus was well inhibited by B/Phuket/3073/2013-like (Yamagata) antisera (Table 4d).

Table 4 a–d. Influenza virus antigenic characterization and genomic sequencing by influenza A subtype and influenza B lineage type — California, 2019–2020 influenza season

	Virus Clade/Subclade		
Antigenic Characterization	3C.2a1	3C.3a*	Total
Well inhibited by A/Kansas/14/2017*	3	1	4
Poorly inhibited by A/Kansas/14/2017*	4	0	4
Not antigenically characterized	15	0	15
Total A (H3N2)	22	1	23

#### a. Influenza A (H3N2) viruses

* 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

	Virus Clade		
Antigenic Characterization	6B.1	6B.1A*	Total
Well inhibited by A/Brisbane/02/2018*	0	9	9
Poorly inhibited by A/Brisbane/02/2018*	0	0	0
Not antigenically characterized	0	35	35
Total A (H1N1)pdm09	0	44	44

* 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

	Virus Subclade		
Antigenic Characterization	V1A.1*	V1A.3	Total
Well inhibited by B/Colorado/06/2017*	0	4	4
Poorly inhibited by B/Colorado/06/2017*	0	4	4
Not antigenically characterized	5	39	44
Total B (Victoria)	5	47	52

* 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

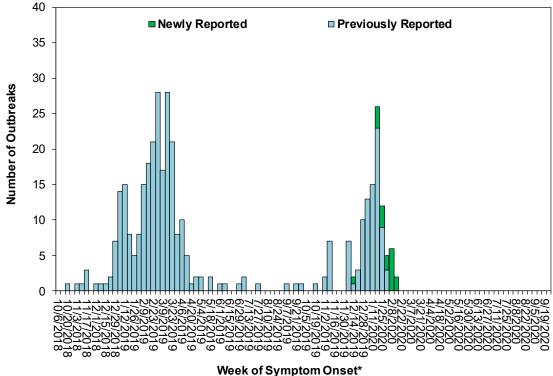
	Virus Clade	
Antigenic Characterization	Y3*	Total
Well inhibited by B/Phuket/3073/2013*	1	1
Poorly inhibited by B/Phuket/3073/2013*	0	0
Not antigenically characterized	5	5
Total B (Yamagata)	6	6

* 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

Seventeen laboratory-confirmed influenza outbreaks were reported to CDPH during Week 7. To date, 111 laboratory-confirmed influenza outbreaks have been reported to CDPH for the 2019–2020 season.





*Earliest date associated with the outbreak was used for outbreaks without reported date of first patient's symptom onset.

#### 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 14 border region sentinel providers reported data during Week 7. The total number of patients screened by all sentinel sites for ILI during Week 7 was 12,913. Outpatient ILI activity was 3.9% during Week 7. ILI activity for the California border region during Week 7 was higher when compared to activity for the same week during the 2017–2018 and 2018–2019 seasons (Figure 11). All influenza syndromic data summarized for the border region represent a subset of CDC influenza sentinel providers in California.

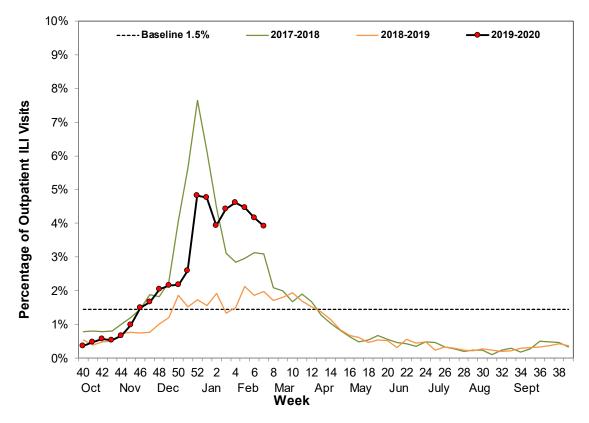


Figure 11. Percentage of Influenza-like Illness Visits among Patients Seen by Sentinel Providers — California Border Region, 2017–2020

#### 2. Virologic Surveillance Update

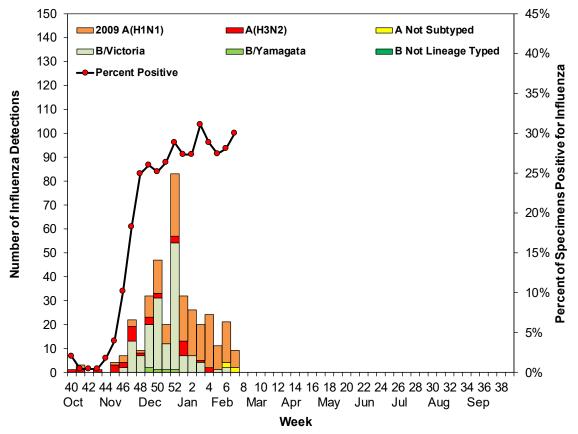
During Week 7, 856 respiratory specimens were tested from border region sentinel clinical laboratories; of these, 257 (30.0%) tested positive for influenza (200 [77.8%] influenza A and 57 [22.2%] influenza B). Cumulatively this season, a total of 12,102 respiratory specimens were tested from border region sentinel clinical laboratories; of these, 2,838 (23.5%) tested positive for influenza (1,431 [50.4%] influenza A; 1,407 [49.6%] influenza B).

During Week 7, nine influenza positive specimens were reported from border region RLN laboratories; of which nine (100.0%) were influenza A. Of the nine specimens that tested positive for influenza A at RLN laboratories, seven (77.8%) were subtyped as A (H1)pdm09 and two (22.2%) had no further subtyping performed. Cumulatively this season, a total of 372 influenza positive specimens have been detected at border region RLN laboratories; of which, 212 (57.0%) were influenza A and 160 (43.0%) were influenza B. Of the 212 specimens that tested positive for influenza A at RLN laboratories, 176 (83.0%) were subtyped as A (H1)pdm09, 32 (15.1%) were subtyped as A (H3), and four (1.9%) had no further subtyping performed. Of the 160 specimens that tested positive for influenza B, five (3.1%) were lineage typed as B (Yamagata) and 155 (96.9%) 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

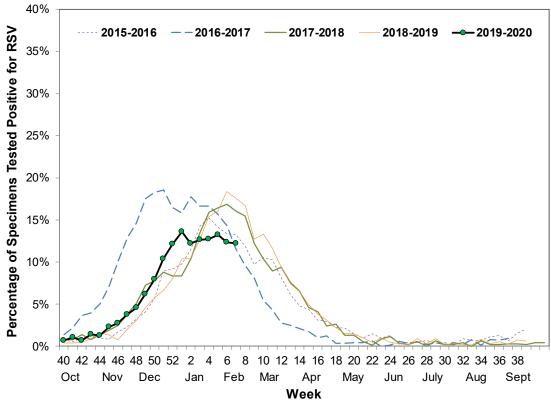
Currently, as mandated under Section 2500 of the California Code of Regulations, deaths among children aged 0–4 years with laboratory-confirmed respiratory syncytial virus (RSV) are reportable to CDPH. The weekly influenza report includes confirmed deaths formally reported to CDPH through February 15, 2020 (Week 7).

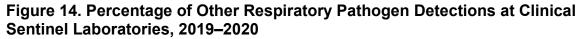
No laboratory-confirmed RSV-associated deaths among children <5 years of age were reported to CDPH during Week 7. To date, CDPH has received no reports of laboratory-confirmed RSV-associated deaths among children <5 years of age during the 2019–2020 influenza season.

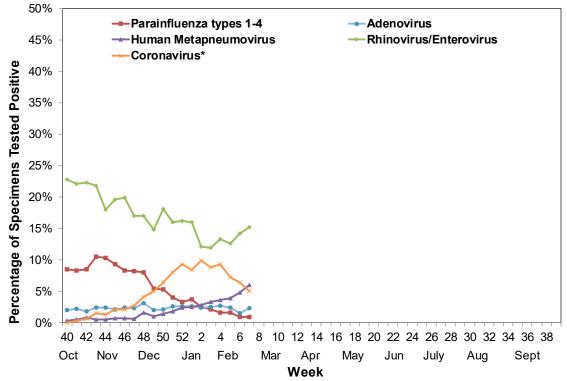
#### 2. Other Respiratory Virus Laboratory Update

During Week 7, 5,045 specimens were tested for RSV and 616 (12.2%) were positive, which is similar compared to Week 6 (12.3%) (Figure 13). During Week 7, adenovirus, human metapneumovirus, and rhinovirus/enterovirus activity increased; and, coronavirus and parainfluenza virus activity decreased (Figure 14).

Figure 13. Percentage of RSV Detections at Clinical Sentinel Laboratories, 2015–2020







^{*}Coronaviruses identified include common human coronaviruses 229E, NL63, OC43, and HKU1 and do NOT include the 2019 novel coronavirus associated with the outbreak in China

#### 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 laboratoryconfirmed 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 <u>InfluenzaSurveillance@cdph.ca.gov</u>. This account is monitored daily by several epidemiologists.

To obtain additional information regarding influenza, please visit the <u>CDPH influenza</u> <u>website</u>

(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 <u>CDPH influenza</u> <u>website</u>

(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 <u>FluVlew</u> (https://www.cdc.gov/flu/weekly/index.htm) and <u>FluView Interactive</u> (https://www.cdc.gov/flu/weekly/fluviewinteractive.htm) websites.

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