











Burden of Influenza in Hospitalized Mexican Patients During the 2018-2019 Season

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Site presentation

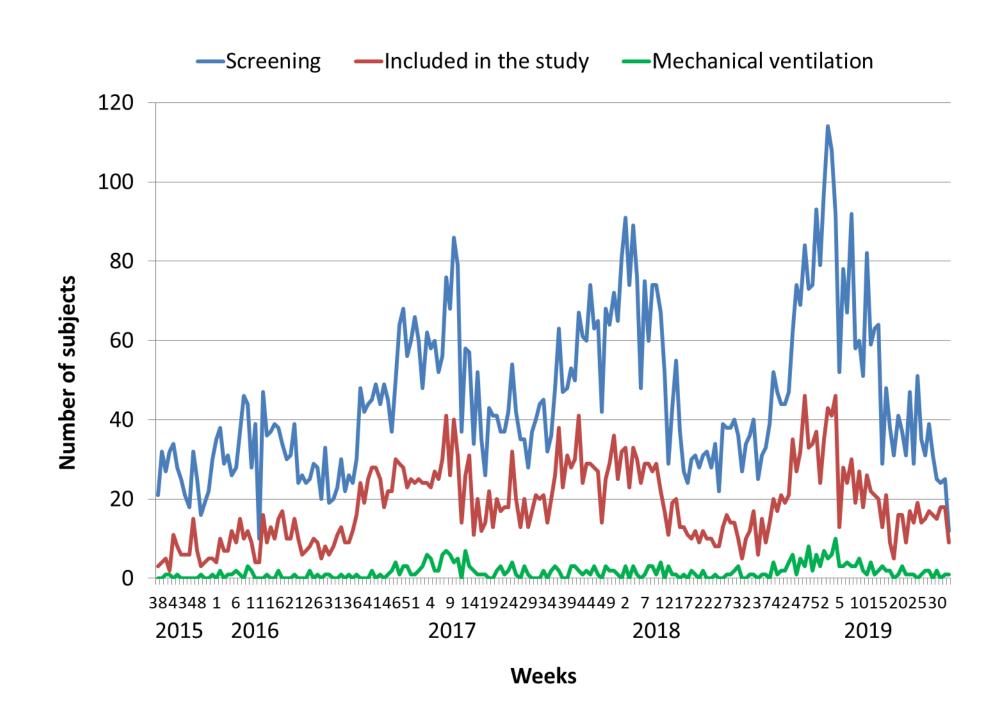
To establish the Mexico branch of the GIHSN, 11 hospitals have been included since September 2015, 5 in the Mexico City area and 6 in the following Mexican States: Durango, Oaxaca and Chiapas. Potential study patients are assessed throughout the whole calendar year. Influenza season is defined as the first two weeks in which influenza is being identified and there is a continuous increase thereafter. The end of the season occurs with the decline of influenza positive cases followed by twoinfluenza-free weeks. Hospitalized patients ≥I month of age (no upper age limit) are eligible for the study.

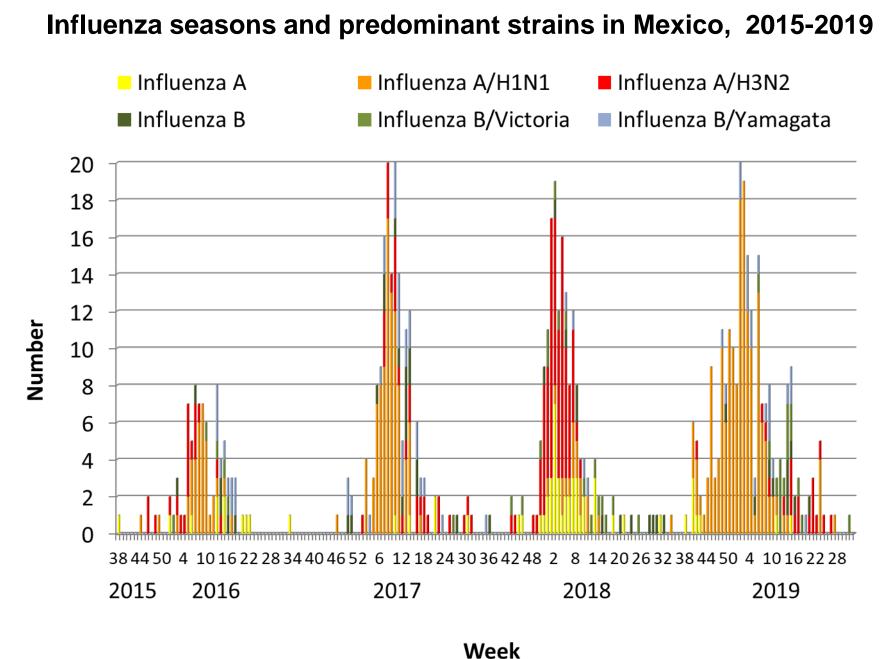
Methods

Hospitalized, non-institutionalized patients residing in a pre-defined hospital catchment area, reporting symptoms of a predefined set of conditions described as possibly associated with a recent influenza infection within 7 days prior to admission and not having been discharged from a hospital in the previous 30 days are included in the study. Trained doctors and nurses collect relevant information by a combination of face-to-face interview of patients and attending physicians, and by reviewing clinical records. Nasopharyngeal swabs are collected from all patients meeting the inclusion criteria and tested by reverse transcription-polymerase chain reaction (RT-qPCR) for influenza and a multiplex PCR (Respifinder®) for other viruses and bacteria. Influenza-positive samples are sub-typed. Patients are considered vaccinated if they had received the current season's influenza vaccine at least 14 days before symptom onset. The study was approved by a Central Research Ethics Committee. Informed written consent/assent when appropriate is required for enrollment.

Consecutive influenza seasons: 2015-2019

Patients screened, enrolled in the study and admitted to the ICU/required invasive respiratory support in the GIHSN-Mexico, 2015-2019





Confirmed Influenza cases and deaths in sentinel hospitals in Mexico, 2010-2019 Data reported to the Mexico National Epidemiology Influenza Reporting System

Season	2010- 2011	2011- 2012	2012-2013	2013-2014	2014- 2015	2015- 2016	2016- 2017	2017- 2018	2018- 2019
Predominant subtype	H3N2	H1N1	H3N2	H1N1	H3N2	H1N1	H1N1	H3N2	H1N1
Influenza cases	3133	7267	2840	8974	2752	9270	5934	3510	6518
Influenza Deaths	43 (1.3%)	343 (4.7%)	49 (1.7%)	1104 (12.3%)	82 (2.9%)	674 (7.2%)	508 (8.5%)	152 (4.3%)	745 (11.4%)
Data reported by sentinel sites participating in the GIHSN-Mexico									
Influenza cases						82	156	159	226
Influenza Deaths						6 (7.3%)	13 (8.3%)	10 (6.3%)	39 (17.2%)

Results form the 2018-2019 influenza season

A total of 2227 hospitalized patients with community acquired respiratory illness were assessed for eligibility during the 2018-2019 season. Of them, 866 (39%) met selection criteria and consented participation: 441 (51%) patients were ≥5 years old and 425 (49%) patients were <5 years old. In the ≥5 years old group 244 (55%) were male; in the <5 years old group 220 (52%) were male.

Of 866 enrolled patients, 226 (26%) tested positive for influenza. Influenza subtype A(H1N1)pdm09 predominated during the season [163 (70%) cases], followed by influenza A(H3N2) [16 (7%) cases] and non-typeable influenza A [7 (3%) cases]. Influenza B tested positive in 45 (19.5%) patients; of them 20 (9%) were subtype B-Yamagata, 15 (7%) B-Victoria and 10 (4%) non-typeable influenza B.

Among influenza positive patients there were 49 (22%) admissions to the ICU, 70 (31%) required invasive mechanical ventilation and 39 (17%) died; 85% of the patients were not vaccinated.

The table and the graphs below show the cumulative frequency and seasonal distributions of respiratory viruses in hospitalized patients participating in the study.

Pathogen	Number (%)				
Adenovirus	47 (1.3)				
Bocavirus	47 (1.3)				
B. pertussis	13 (0.3)				
Chlamydophila pneumoniae	0				
Coronavirus	80 (2.1)				
Influenza	463 (12.4)				
Legionella	1 (0.1)				
Metapneumovirus	274 (7.3)				
Mycoplasma pneumoniae	5 (0.1)				
Parainfluenza	185 (4.9)				
VSR	453 (12.1)				
Rhinovirus	643 (17.2)				
Co-infections (≥2 pathogens)	697 (18.7)				
Negative	824 (22.1)				

The phylogenetic tree (shown below) is based on the hemagglutinin (HA)

genes of influenza A(H1N1) collected during consecutive seasons; genetic

Clade 6B.1

Clade 6B

JINCMNSZ/BMN570/2019(H1N1)

VMexico/NCMNSZ/RFB864 Feb 2012 —— A/Mexico/INCMNSZ/RFB682 Jan 2012

A/Mexico/INCMNSZ/RF007 Apr 2010

A/Mexico/NCMNSZ/VR7907 Nov 2009

A/California/VRDL64/Aug 2009 A/Mexico/INCMNSZ/VR9443 Jan 2010

A/Finland/576/Jun 2009

A/New York/34/Apr 2009 A/New York/19/2009

A/Mexico City/WRAIR1691N/Sep 200

A/Mexico/4486/2009 Mexico/4269/2009

—— A/Mexico/4635/2009

—— A/Mexico/nDRE13494/200 — A/California/05/2009 A/Mexico/4627/2009

> A/Mexico/4108/2009 A/Mexico/nDRE4487/2009

A/Mexico/4603/2009

A/Mexico/3955/2009

A/California/07/Apr 2009 A/California/04/2009

Antigenic Regions Aligment

Influenza A H1N1 pmd. Segment 4 (Haemmaglutinin)

A/Mexico/INCMNSZ/RFB570 Jan 2012

A/Mexico/INCMNSZ/RF103 Aug 2010

A/Texas/05/2009

A/Mexico/InDRE4114/2009

A/Mexico/NCMNSZ/RFB584 Jan 2017

A/CHRISTCHURCH/16/2010 NIB-74XP

the phylogenetic three.

2018-2019 strain

—— A/Mexico/INCMNSZ/RFH460 Dec 201: — A/Mexico/INCMNSZ/RFH697 Jan 2014

fexico/INCMNSZ/RFH184 Dec 201

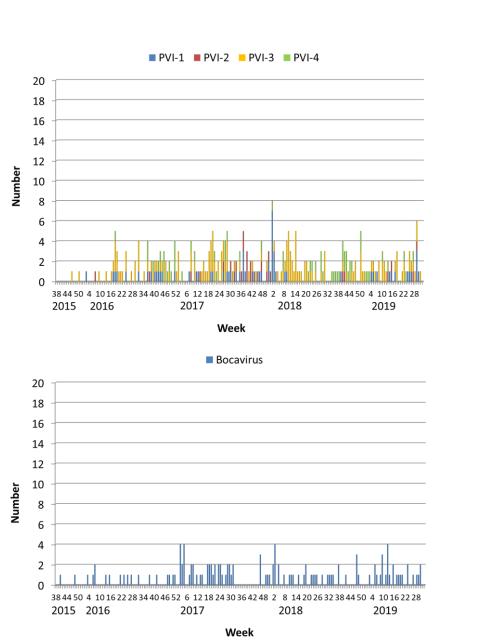
VMexico/INCMNSZ/RFH130 Dec 2013

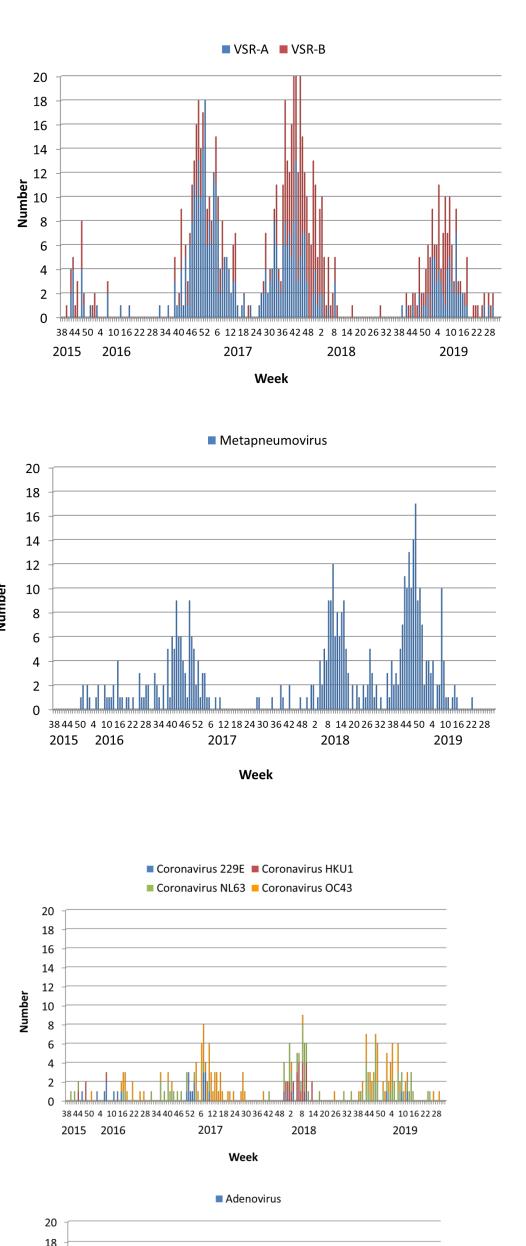
xico/INCMNSZ/RFH776 Jan 2014 A/Mexico/INCMNSZ/RFH619 Jan 2014

—— A/Mexico/INCMNSZ/RFI269 Mar 2014 —— A/Mexico/INCMNSZ/RFH415 Dec 2013 A/Mexico/INCMNSZ/RFH477 Dec 2013 Mexico/INCMNSZ/RFH926 Feb 2014 xico/INCMNSZ/RFH874 Jan 2014 exico/INCMNSZ/RFH542 Jan 201

Influenza H1N1

differences are compared to the A/California/04/2009 consensus sequence. Amino acid substitutions delineating major branches are shown. Date of collection follows names of viruses. Clade 6B.1 predominated during the 2018-2019 season. Amino acid substitutions are shown in the table below ■ PVI-1 ■ PVI-2 ■ PVI-3 ■ PVI-4





The GIHSN-Mexico has successfully implemented a hospital surveillance system for severe influenza and other respiratory viruses. Data obtained provides an indication of changes in the epidemiology of severe influenza, and potentially contributes to the refinement of clinical care guidelines and policy.

Challenges include: the sustainability of the surveillance system and timely dissemination of the results to the scientific community and public health agencies.

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Key aspects & challenges