The burden of influenza A and B in Mexico from the year 2010 to 2013: An observational, retrospective, database study, on records from the Directorate General of Epidemiology database

Ricardo Cortes-Alcala, Gaël Dos Santos, Rodrigo DeAntonio, Raghavendra Devadiga, Cuitlahuac Ruiz-Matus, Maria E. Jimenez-Corona, Jose A. Diaz-Quinonez, Luis Romano-Mazzotti, Maria Yolanda Cervantes-Apolinar, and Pablo Kuri-Morales

ABSTRACT

Despite vaccination programs, influenza still represents a significant disease burden in Mexico. We conducted an observational, retrospective analysis to better understand the epidemiological situation of the influenza virus in Mexico. Analysis of the seasonal patterns of influenza A and B were based on the Directorate General of Epidemiology dataset of influenza-like illness(ILI), and severe acute respiratory infection(SARI) that were recorded between January 2010 and December 2013. Our objectives were 1) to describe influenza A and B activity, by age group, and subtype and, 2) to analyze the number of laboratory-confirmed cases presenting with ILI by influenza type, the regional distribution of influenza, and its clinical features. Three periods of influenza activity were captured: August 2010–January 2011, December 2011–March 2012, and October 2012–March 2013. Cases were reported throughout Mexico, with 50.3% (n = 10,320) of cases found in 18–49 year olds. Over the entire capture period, a total of 76,085 ILI/SARI episodes had swab samples analyzed for influenza, 27% were positive. During the same period, influenza A cases were higher in the 18–49 years old, and influenza B cases in both 5–17 and 18–49 age groups. Peak activity occurred in January 2012 (n = 4,159) and December 2012 (n = 348) for influenza A and B respectively. This analysis confirms that influenza is an important respiratory pathogen for children and adults in Mexico despite vaccination recommendations. School-age children and adolescents were more prone to influenza B infection; while younger adults were susceptible to both influenza A and B viruses. Over the seasons, influenza A and B co-circulated.

Introduction

In 2009, Mexico experienced a severe influenza A (H1N1) outbreak, which was the premise for the first Public Health Emergency of International Concern (PHEIC) and according to the international health regulations the outbreak could be confirmed as a pandemic. This pandemic was characterized by high morbidity and mortality rates especially during its first outbreak, and there was an increase in high transmission rates between younger and relatively healthier individuals than previously observed.1–4

Vaccination against influenza began during the latter part of 2004 in children of 6 to 24 months of age, and adults 65 years and older. By winter 2006–2007 the vaccination age for children was extended to 35 months. This meant an increase from 89.8% to 91.7% of the target programmed by the Health Sector and Ministry of Health. The target of 100% was attained for people aged 60 and over.5

Prior to the 2009 influenza pandemic, the vaccination recommendation in Mexico was that the vaccine should contain two subtypes of influenza A virus and one subtype of influenza B virus and administered as an annual dose a few months before the start of winter. The recommendation was also that children should receive the fractionated vaccine. In addition, people 60 years and older and those at high risk should receive first preference for vaccination. High-risk patients include those with chronic diseases of the renal, pulmonary or
cardiovascular system and metabolic diseases including anaemia, diabetes and immunosuppression.6

Important studies have been published on the specific characteristics and outcomes of this influenza pandemic and the subsequent seasonal outbreaks of influenza in Mexico.1,3,7–11 However, the patterns of influenza occurrence during the years following the pandemic infection have been poorly documented.

In 2006, the national surveillance program database at the Directorate General of Epidemiology (DGE), a sentinel platform to conduct influenza surveillance, was implemented in Mexico. Since then, all cases of influenza like-illness (ILI) and severe acute respiratory infection (SARI), have been registered via this platform in the DGE national surveillance system.12

This observational retrospective analysis was conducted to assess the burden of influenza in Mexico, the analysis involved describing the seasonal activity patterns of influenza A and B viruses, using the DGE dataset of ILI and SARI records collected between January 2010 and December 2013.

Results

Participants

A total of 133,478 episodes of ILI and SARI were reported between January 2010 and December 2013. Of these, 57% (n = 76,085) had samples tested for influenza, and among them 73% (n = 55,558) were negative (Figure 1). The classification of these episodes by influenza diagnosis and by ILI or SARI subgroup is illustrated in Figure 1.

Viewing data from laboratory tests, the percentage of influenza positivity ranged from 9.5% in 2011 to 34.4% in 2012 based on data by year and amongst the ILI and SARI patients (see Supplementary Table 1).

Descriptive data

Half of the influenza-positive cases were in the age group 18–49 years (Table 1). One third of all influenza-positive cases were admitted to hospital as a consequence of their illness (Table 1).

Table 1 displays the main demographic characteristics of the population with influenza diagnosis (positive or negative). Table 2 displays the main demographic characteristics of analysis by diagnosis, influenza type and season stratified by season.

The monthly activity during 2010–2013 ranged between 2 to 4,159 cases (0.3–53.3% of tested samples), and from 0 to 348 cases (0%–21.8% of tested samples) for influenza A and B, respectively. The highest number of cases was recorded in January 2012 for influenza A, and in December 2012 for influenza B.

In the 2010–2011 influenza season, the highest peak of influenza-positive patients was recorded in December 2010: 44.3% (n = 1,142) of the respective ILI+SARI patients based on available influenza laboratory test results. This influenza season lasted for six months, from August 2010 to January 2011 (Figure 2).

In the 2011–2012 influenza season, the highest peak of influenza-positive patients was recorded in January 2012: 53.7% (n = 4,189), of the respective ILI+SARI patients (based on available influenza laboratory test results). This influenza season lasted for four months, from December 2011 to March 2012 (Figure 2).

In the 2012–2013 influenza season, the highest peak of influenza-positive patients was recorded in January 2013: 30.3% (n = 695), of the respective ILI+SARI patients with available influenza laboratory test results. This influenza season lasted for six months, from October 2012 to March 2013 (Figure 2).

The highest activity for influenza B was reported during 2012–2013. Global Influenza Surveillance and Response System (GISRS) data show that the influenza B virus co-circulated with the A H3N2 virus until the 12th week of 2013.13 In the 13th week, influenza B co-circulated with the pandemic A (H1N1)pdm09 virus.13 Our dataset ended in December 2013, but the GISRS 2013–2017 data for Mexico showed that influenza B virus continued to co-circulate with either A subtype (A H3N2 or A (H1N1)pdm09).13

Influenza cases characterisation

The age groups with the highest influenza occurrence were 18–49 years for all influenza A subtypes; and 5–17, and 18–49 years old for influenza B (Table 3). On average, 22% (n = 4,041) of all 2010–2013 influenza A cases were not subtyped. Lineage characterisation was not reported for the influenza B viruses.

The influenza season with the highest number of cases were April 2011–March 2012 and April 2012–March 2013 (Table 4, Figure 2) for influenza A and B respectively.

Regional distribution and clinical features

Influenza cases were distributed across all Mexican regions. Most influenza cases were reported in Mexico’s Central region (Supplementary Table 2).

Deaths

Amongst all 2010–2013 influenza cases, 944 deaths were recorded, almost all (98%, n = 926) had influenza A (Supplementary Table 5). Of the respective ILI+SARI patients, 230 patients died in 2010 of influenza A while only 57 died in 2011. A total of 18 (2%) patients died with influenza B, no patients died in 2010, whilst 10 died in 2013 (Supplementary Table 5).

Discussion

This observational-retrospective database analysis used the data captured in the DGE database from 2010 to 2013, for investigating activity patterns and the demographic and clinical characteristics of influenza A and B occurrence. This is the first descriptive analysis to be completed on the influenza activity over 3 consecutive influenza seasons following the 2009 pandemic in Mexico. Since this pandemic, vaccination recommendations have been modified.

The current recommendation for vaccination is now that the annual vaccination should be administered to children ranging in age from 6 months to 4 years and adults of 60 years and older. As high-risk individuals now also include health workers, and individuals who are pregnant or morbidly obese. There is currently no data available on the effectiveness of the influenza vaccine in Mexico.14
However, studies have been completed on vaccine effectiveness in the US. In the 2017–2018 flu season, vaccine effectiveness against influenza A and influenza B was found overall to be about 36%, with most cases being from H3N2. Vaccine effectiveness against A(H1N1)pdm09 virus was 67%, and 42% against influenza B virus. The CDC continues to recommend vaccination since it can decrease hospitalization rates and prevent some illnesses.

In 2010, the influenza season had an early start in August due to the epidemiological change that influenza pandemic posed in the epidemic profile of influenza. WHO GISRS data for Mexico, indicated that this early peak was mainly caused by A H3N2 virus activity. In the USA, the influenza 2010–2011 season started in November in from the southeastern states, and peaked nationally in February (2011). Similarly to Mexico, the predominant virus was the A H3N2 (62% of all influenza A cases), which co-circulated with the A H1N1 and influenza B viruses. In the USA, influenza B was the predominant virus in the southeastern states during November and December 2011 and the Victoria lineage was identified in 94% of all tested influenza B viruses (2011). In Canada, A H3N2 was the predominant virus in 2010–2011.

Figure 1. Classification of the population by influenza diagnosis and by ILI and SARI episodes, 2010–2013. Acronyms: ILI, influenza-like illness; SARI, severe acute respiratory infection.

According to the GISRS data, the influenza peak recorded in 2012, consisted mainly of activity of the recrudescent pandemic A (H1N1)pdm09 virus (2012). Chowell and colleagues (2012) analyzed data from 760 A H1N1 positive respiratory-infection patients that were hospitalized in Mexico between December 2011 and February 2012. They found that this fourth wave of the pandemic virus caused more hospitalizations than the previous three waves in 2009. They also reported a shift towards older age distribution of hospitalizations and deaths. In our analysis, one third of all influenza cases were individuals hospitalized because of their illness. Given the reported peak of influenza activity in January-February 2012, we hypothesize that most of these hospitalizations were caused by this recrudescent pandemic wave. There was no such pandemic wave in the USA, in which the A H3N2 type was predominant (2010-2013).
The highest activity for influenza B virus was recorded in the influenza season that followed the peak of influenza A activity. We know, from the analysis of influenza prevalence in the USA over a period of 30 years, that influenza B prevalence is lower at periods with higher levels of influenza A prevalence.\textsuperscript{20} During or shortly following these periods, a shift in the dominant B lineage occurs.\textsuperscript{20} Immune selection may determine the evolutionary dynamics of influenza B virus and alter the lineage dominance.\textsuperscript{20} Increasing immunity to the dominant lineage reduces its frequency but also leaves room for other lineages to become dominant.\textsuperscript{25} Which lineage will predominantly circulate each season cannot be predicted with certainty, which makes developing an influenza vaccine challenging. For example, in half of the influenza seasons in the USA since 2000, and in one quarter of influenza seasons examined in 26 countries between 2000 and 2013, the dominant lineage did not match with the one included in the seasonal vaccine.\textsuperscript{21,22} However the vaccine can still provide protection against some of the co-circulating viruses present during a particular influenza season.\textsuperscript{15}

The latest available GISRS data (season 2016–2017) for Mexico showed that B/Yamagata lineage was more common than B/Victoria lineage. The recently isolated B/Yamagata/16/88

\begin{table}[h]
\centering
\caption{Summary of demographic characteristics by influenza status.}
\begin{tabular}{lcccc}
\hline
 & Influenza-positive & & Influenza-negative & \\
 & N = 20,527 & & N = 55,558 & \\
\hline
Sex, male, n (%) & 9,578 (46.7) & & 26,995 (48.6) & \\
Age, years & & & & \\
Median (range) & 29.0 (0–111) & & 26.0 (0–109) & \\
Age groups, n (%) & & & & \\
0–4 years & 2,618 (12.8) & & 16,990 (30.7) & \\
5–17 years & 3,728 (18.2) & & 5,779 (10.5) & \\
18–49 years & 10,320 (50.4) & & 19,076 (34.5) & \\
50–64 years & 2,540 (12.4) & & 6,005 (10.9) & \\
≥65 years & 1,236 (6.1) & & 7,442 (13.5) & \\
missing & 65 (0.3) & & 266 (0.5) & \\
Region, n (%) & & & & \\
Center & 6,377 (31.1) & & 16,931 (30.5) & \\
Center-West & 4,442 (21.6) & & 12,913 (23.2) & \\
North & 4,963 (24.2) & & 10,110 (18.2) & \\
South-Southwest & 4,745 (23.1) & & 15,604 (28.1) & \\
Patient type & & & & \\
Ambulatory, n (%) & 12,965 (63.2) & & 21,084 (37.9) & \\
Hospitalized, n (%) & 7,562 (36.8) & & 34,474 (62.1) & \\
\hline
\end{tabular}
\end{table}

\*age was available for 20,462 influenza-positive and 55,292 influenza-negative records.

\textbf{N} Number of ILI+SARI records with samples tested for influenza.

\textbf{n} Number of records within given category.

Acronyms: ILI, influenza-like illness; SARI, severe acute respiratory infection.

\begin{table}[h]
\centering
\caption{Summary of outcome by diagnosis, influenza type and season.}
\begin{tabular}{lcccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccccc
lineage viruses are antigenically and genetically related to B/Phuket/3073/2013. Therefore, a B/Phuket/3073/2013-like virus not included in the commonly used trivalent vaccines is one of the components of the quadrivalent vaccine recommended for use in the 2016–2017 Northern hemisphere influenza season.

In agreement with previously published findings, from across the globe, school-age children and adolescents have been found to have a higher susceptibility to influenza B virus. School-age children and adolescents are the first populations to be affected by influenza infection. Children need more time to acquire natural immunity to influenza B compared to influenza A. In a population-based study in the Netherlands 72% of children aged seven years had developed antibodies to influenza B and 100% to influenza A. Interestingly, several studies have shown that the Victoria lineage affects younger people, while Yamagata is more evenly distributed across age groups. In addition, the antibodies against both B lineages did not cross-react.

Some previously published studies on the epidemiology of influenza in Mexico, used data from hospitalized influenza cases or more frequently studies used data from the...
prospective surveillance system implemented in 2009 by the Mexican Institute for Social Security (IMSS for Spanish acronym). IMSS collects data from 1,099 primary clinics and 259 hospitals across Mexico, covering 40% of the Mexican population, while excluding the lower socio-economic population. Hence, studies using the IMSS database have an inherent limitation in that seasons with early influenza activity may serve as a proxy for estimating the time of year with the highest influenza incidence. Such information can improve preparedness for management of hospitalizations and comorbidities. Since 2004, Canada’s sentinel surveillance system has incorporated vaccine effectiveness data. Based on this dataset, vaccine effectiveness studies have been published providing highly informative results that may guide decisions in reformulating vaccines in terms of composition and in improving disease protection.

A key factor for successful influenza control and effective vaccination strategies is an effective and reliable national surveillance system. The documentation of virus activity that is performed via such systems offers a unique opportunity to characterize seasonal patterns, develop reliable estimates, and identify parameters affecting preparedness for future epidemics. An analysis on USA Medicare and Medicaid records, showed firstly that seasons with early influenza activity tend to produce more influenza cases, and secondly that the timing and absolute intensity of influenza activity may serve as a proxy for estimating the time of year with the highest influenza incidence. Such information can improve preparedness for management of hospitalizations and comorbidities. Since 2004, Canada’s sentinel surveillance system has incorporated vaccine effectiveness data. Based on this dataset, vaccine effectiveness studies have been published providing highly informative results that may guide decisions in reformulating vaccines in terms of composition and in improving disease protection.

A limitation that we had was that there is no currently available data on the effectiveness of the vaccination program in Mexico. Influenza surveillance in Mexico is based on sentinel sites and as DGE is a secondary database, underreporting is expected, as DGE is only a sample of all ILI/SARI cases over the National Healthcare System, hence the reason ILI+SARI were reported together. We state that this is a limiting factor of the analysis as patients may be recorded twice. However, our analysis indicates that younger adults are also vulnerable to both influenza A and B viruses and that only one out of five patients with ILI had received a vaccination. In addition, for an effective vaccination strategy, we need to reduce the probability of having a mismatch between the seasonal vaccine and the dominant influenza B lineage. Although our dataset did not contain lineage information, DGE has now started to characterize B strains, which will provide valuable information for a successful implementation of the vaccination strategy. A limitation that we had was that there is no currently available data on the effectiveness of the vaccination program in Mexico.
2014 (based on the WHO defined influenza seasons\textsuperscript{23}) to help characterize potential seasonal patterns, however this data was not available, therefore we characterized the data based on observed seasons within the data we had available. Whilst this is not optimal it is unlucky to affect our findings due to seasonal distribution peaks throughout each period.

**Conclusions**

Using a validated epidemiological surveillance system, representative of the Mexican population, this study provides unique meaningful information on the influenza activity in Mexico over three consecutive influenza seasons. This analysis confirms that influenza is an important respiratory pathogen for children and adults in Mexico despite the fact that an influenza vaccination program has been implemented since 2004. School-age children and adolescents were more prone to influenza B infection while younger adults were susceptible to both influenza A and B viruses. Over the three analyzed seasons, influenza B virus continuously co-circulated with influenza A virus. However, lineage characterization for influenza B viruses was not available, such description is needed to strengthen the knowledge of influenza B strains characteristics for immunization strategies in the country. Increased knowledge of lineage characteristics would lead to improved vaccines.

**Methods**

**Study design and setting**

The study was conducted in accordance with all applicable ethics requirements, including patient privacy and the guiding principles of the International Ethical Guidelines for Epidemiological Studies. Due to the nature of the study (i.e., retrospective by nature) and the anonymisation process prior to data extraction, as per research standards and local regulations, no ethics committee approval was required to conduct the analysis. The dataset was anonymised before data extraction, so it was not possible to make a link between the extracted data and specific individuals. Therefore, no informed consent was required from patients. Regarding the surveillance activities in the country, participants agreed with the terms of the standardised healthcare and the subsequent sample analysis and report. The data were analysed between December 2014 and November 2015.

**Population data**

All January 2010–December 2013 DGEs ILI and SARI records were eligible for inclusion in the analysis dataset.

**Description of the DGE database**

Since 2006, a sentinel surveillance program has continued to be conducted in Mexico in order to collect data on influenza. The Weekly Report System for New Cases (SUAVE for Spanish acronym) and the Influenza Monitoring Healthcare Units (USMI for Spanish acronym), were developed to support this program. SUAVE is a weekly, routine, passive, population-based surveillance system, based on the International Classification of Diseases (ICD) coding system, where 50 diagnoses are broken down into 13 disease groups.\textsuperscript{39} The USMIs consist of 583 designated primary healthcare facilities and general hospitals (including second and third-level hospitals) found throughout Mexico. Their role is to use systematic and standardized procedures to collect samples of persons with ILIs.\textsuperscript{40} The USMI conducts sentinel surveillance according to the World Health Organisation (WHO) recommendations. The USMI surveillance system is synonymous with the US and Canadian systems.\textsuperscript{39} Since October 2009, all confirmed influenza cases: ILI, and SARI are registered via SUAVE and USMI in a unique informative platform. From this platform, the information is centralised at the DGE national surveillance program database.\textsuperscript{12} Before being entered into SUAVE and UNMIs information is checked locally for consistency where the number of cases must be equal or higher in the SUAVE due to the more general symptom recording procedure of SUAVE.

Throat and nasopharyngeal swab samples were collected and tested for the presence of influenza A and B viruses in adults and infants (including very young children), respectively. Reverse transcriptase-polymerase chain reaction (rt-PCR) analysis was used to confirm cases,\textsuperscript{39} and results were recorded in the database.

**Case definitions**

ILI: Fever (≥38°C), cough and headache accompanied by one or more of the following symptoms: rhinorrhea, coryza, arthralgia, myalgia, fatigue, sore throat, chest pain, abdominal pain, nasal congestion and diarrhea. For patients under five years of age, signs of irritability were recorded instead of headache.

SARI: Shortness of breath, history of fever (≥38°C) and cough, and any of the following symptoms: malaise, chest pain, tachypnea or acute respiratory distress syndrome. Influenza-related pneumonia and chronic disease exacerbations were included.

Influenza confirmed case: Patients who tested positive (in rt-PCR) for influenza.

**Objectives**

This analysis aimed to primarily describe influenza A and B cases, by age group and A-subtype/strain B-lineage during the period analysed.

Additionally, we aimed to describe the number of ILI cases attributable to influenza A and B, the regional distribution of influenza A and B cases, the frequency of clinical features and outcomes (clinical symptoms, duration of illness, complications) experienced by patients with ILI/SARI who were positive for influenza A or B, and finally the proportion of influenza cases among ILI and SARI cases.

Age at date of symptom onset was computed as the difference between the date of onset of the first symptom and the date of birth, and was measured in years for patients older than one year, and in months for patients younger than one year.

The age groups analysed were: 0–4, 5–17, 18–49, 50–64, and ≥65 years.
Data collection and procedures
All influenza A and B, ILI and SARI cases in the DGE database were extracted to a Microsoft Excel file and converted into Statistical Analysis System datasets for analysis. Demographic data, date of symptom onset, geographic region (Center, Center-West, North, and South-SouthEast), clinical features and outcomes (clinical symptoms, duration of illness, complications) for all patients with ILI caused by influenza A and B were extracted and anonymised by DGE personnel and then transferred to the sponsor. The data were then analysed.

Statistical analyses
Descriptive statistics were used including frequency tables generated for categorical variables. The mean, median, and standard deviation were calculated for continuous data. The distribution of influenza cases by month and year was tabulated by their respective frequencies and percentages, overall and by age and group and strain subtype/lineage. The proportion of cases attributable to influenza A and B was calculated with exact 95% confidence intervals (CI) for the overall sample and by age and subtype/lineage.

The statistical analyses were performed using the Statistical Analysis Systems version 9.2, the Drug and Development web portal version 3.5, and Microsoft Excel 2007.

Abbreviations
CI Confidence Interval
DGE Directorate General of Epidemiology
GISRS Global Influenza Surveillance and Response System
ILI Influenza-Like Illness
IMSS Mexican Institute of Social Security
rt-PCR reverse transcriptase Polymerase Chain Reaction
SARI Severe Acute Respiratory Infection
SUAVE Weekly Report System for New Cases
USMI Influenza Monitoring Healthcare Units
WHO World Health Organisation

Disclosure of potential conflicts of interest
RCA, RD, GDS, YCA and RDA are employees of the GSK group of companies. GDS was employed by Business & Decision Life Sciences on behalf of GSK Vaccines at the time of the study and is now an employee of the GSK group of companies. YCA, GDS, and RDA, report holding shares in the GSK group of companies as part of their employee remuneration. CRM, MJJC, JDQ and PKB have nothing to disclose.

Financial disclosure & funding source
GlaxoSmithKline GSK Biologicals SA funded this study (GSK Study identifier: 201075) and was involved in all stages of study conduct, including analysis of the data. GlaxoSmithKline GSK Biologicals SA also covered all costs associated with the development and publication of this manuscript.

Acknowledgments
Authors would like to acknowledge and thank Gabriela Nucamendi-Cervantes, Irma Lopez-Martinez, and Gisela Barrera-Badillo, from the Directorate General of Epidemiology, for their hard work in the development of the study. Authors would also like to thank Business & Decision Life Sciences platform for editorial assistance, manuscript coordination, and writing support, on behalf of GSK. Pierre-Paul Prevot coordinated manuscript development and editorial support; and Athanasia Benekou provided medical writing support.

References
1. Echevarría-Zuno S, Grajales-Muñiz C, Robles-Pérez E, González-León M, Ortega-Alvarez MC, González-Bonilla C, Rascón-Pacheco RA, Borja-Aburto VH. Infection and death from influenza A H1N1 virus in Mexico: a retrospective analysis. Lancet. 2009;374(9694):2072-9. doi:10.1016/S0140-6736(09)61638-X. PMID:19913290
2. Domínguez-Cherit G, Lapinsky SE, Macias AE, Pinto R, Espinosa-Perez L, de la Torre A, Poblano-Morales M, Baltazar-Torres JA, Bautista E, Martínez A et al. Critically Ill patients with 2009 influenza A (H1N1) in Mexico. JAMA. 2009;302:1880–7. doi:10.1001/jama.2009.1536. PMID:19822626
3. Chowell G, Echevarría-Zuno S, Viboud C, Simonsen L, Tamerius J, Miller MA, Borja-Aburto VH. Characterizing the Epidemiology of the 2009 Influenza A/H1N1 Pandemic in Mexico. doi:10.1371/journal.pmed.1000436
4. Rothberg MB, Haessler SD, Brown RB. Complications of Viral Influenza. American Journal of Medicine. 2008;121:258-64. doi:10.1016/j.amjmed.2007.10.040. PMID:18374680
5. Secretaría de Salud. Vacunación universal y semanas nacionales de salud lineamientos generales. Centro Nacional para la Salud de Niños, Niñas y Adolescentes de México. 2008
6. Secretaría de Salud. NORMA Oficial Mexicana NOM-036-SSA2-2002, Prevención y control de enfermedades. Aplicación de vacunas, toxoides, sueros, antitoxinas e inmunoglobulinas en el humano. Available at http://www.salud.gob.mx/unidades/cdmi/nom036ss2002.html. 2002
7. Dávila J, Chowell G, Borja-Aburto VH, Viboud C, Grajales Muñiz C, Miller M. Substantial Morbidity and Mortality Associated with Pandemic A/H1N1 Influenza in Mexico, Winter 2013-2014: Gradual Age Shift and Severity. PLoS Curr. 2014;6;4
8. Chowell G, Echevarría-Zuno S, Viboud C, Simonsen L, Grajales Muñiz C, Rascón Pacheco RA, González León M, Borja-Aburto VH. Recrudescent wave of pandemic A/H1N1 influenza in Mexico, winter 2011-2012: Age shift and severity. PLoS Curr. 2012;4:RN1306. doi:10.1371/currents.RRN1306. PMID:22485199
9. Borja-Aburto VH, Chowell G, Viboud C, Simonsen L, Miller MA, Grajales-Muñiz C, González-Bonilla CR, Díaz-Quinonez JA, Echevarría-Zuno S. Epidemiological characterization of a fourth wave of pandemic A/H1N1 influenza in Mexico, winter 2011-2012: age shift and severity. Arch. Med. Res. 2012;43:563–70. doi:10.1016/j.armed.2012.09.005. PMID:23079035
10. Chowell G, Echevarría-Zuno S, Viboud C, Simonsen L, Miller MA, Fernández-Gárarte I, González-Bonilla C, Borja-Aburto VH. Epidemiological Characteristics and Underlying Risk Factors for Mortality during the Autumn 2009 Pandemic Wave in Mexico. PLoS One. 2012;7:e41069. doi:10.1371/journal.pone.0041069. PMID:22815917
11. Chowell G, Viboud C, Simonsen L, Miller MA, Echevarría-Zuno S, González-León M, Aburto VH. Impact of antiviral treatment and hospital admission delay on risk of death associated with 2009 A/H1N1 pandemic influenza in Mexico. BMC Infect. Dis. 2012;12:97. doi:10.1186/1471-2334-12-9. PMID:22520743
12. Secretaría de Salud. Manual de Procedimientos Estadizados para la Vigilancia Epidemiológica de la Influenza. Secretaría de Salud. Subsecretaría de Prevencion y promocion de la Salud Direccional General de Epidemiología, 2012
13. FluNet. WHO | FluNet - CHARTS. WHO Available at: http://www.who.int/influenza/gisrs_laboratory/flu/flu/Charts/en/. (Accessed: 18th December 2017)
14. Secretaría de Salud. NOM-036-SSA2-2012, Prevención y control de enfermedades. Aplicación de vacunas, toxoides, faboterápicos e inmunoglobulinas en el humano. 2012. Available at: https://www.gob.mx/
15. Hanney B, Chung JR, Belongia EA, McLean HQ, Gaglani M, Murthy K, Zimmerman RK, Nowalk MP, Jackson ML, Jackson LA, et al. Interim Estimates of 2017–18 Seasonal Influenza Vaccine Effectiveness — United States, February 2018. MMWR. Morb. Mortal. Wkdly. Rep. 2018;67:180–5. doi:10.15585/mmwr.mm6706a2. PMID:29447141

16. World Health Organisation. Global Influenza Surveillance and Response System GISRS. Number of specimens positive for influenza by subtype, Mexico, 2012–2013. Searchable Database 2012. Available at: http://apps.who.int/flucontrol/Default?ReportNo=7. (Accessed: 18th December 2017)

17. 2011–2012 Flu Season Draws to a Close | Spotlights (Flu) | CDC. Centers for Disease Control and Prevention 2012. Available at: https://www.cdc.gov/flu/spotlights/2011-2012-flu-season-wrapup.htm. (Accessed: 18th December 2017)

18. Skowronski DM, Janjua NZ, De Serres G, Winter AL, Dickinson JA, Gardy JL, Gubbay J, Fonseka K, Charest H, et al. A Sentinel Platform to Evaluate Influenza Vaccine Effectiveness and New Variant Circulation, Canada 2010–2011 Season. Clin. Infect. Dis. 2012;55:332–42. doi:10.1093/cid/cis341. PMID:22539661

19. Fowlkes A, Steffens A, Tette J, Lonardo SD, McHugh L, Martin K, Rubino H, Feist M, Davis C, Selzer C, et al. Incidence of medically attended influenza during pandemic and post-pandemic seasons through the Influenza Incidence Surveillance Project, 2009–13. Lancet Respir. Med. 2015;3:709–18. doi:10.1016/S2213-2600(15)00278-7. PMID:26300111

20. Chen R, Holmes EC. The Evolutionary Dynamics of Human Influenza B Virus. J. Mol. Evol. 2008;66:655–63. doi:10.1007/s00239-008-9119-2. PMID:18504518

21. Ambrose CS, Levin MJ. The rationale for quadrivalent influenza vaccines. Hum. Vaccin. Immunother. 2012;8:1–88. doi:10.4161/hv.8.1.17623. PMID:22252006

22. Caini S, Huang QS, Ciblak MA, Kusznier G, Owen R, Wangchuk S, Henriquez CM, Njouom R, Fusce RA, Yu H, et al. Epidemiological and virological characteristics of influenza B: results of the Global Influenza B Study. Influenza Other Respi. Viruses. 2015;9:3–12. doi:10.1111/irv.12139

23. World Health Organisation (WHO). Recommended composition of influenza virus vaccines for use in the 2017–2018 northern hemisphere influenza season. 2017

24. Pan Y, Zhang Y, Yang P, Qian H, Shi W, Wu S, Cui S, Zhang D, Wang Q. Epidemiological and Phylogenetic Characteristics of Influenza B Infection in Severe Acute Respiratory Infection Cases in Beijing, 2014 to 2015. Medicine (Baltimore). 2015;94:e2399. doi:10.1097/MD.0000000000002399. PMID:26717393

25. Mos AM, Muscatello DJ, Turner RM, MacIntyre CR. Epidemiology of influenza B in Australia: 2001–2014 influenza seasons. Influenza Other Respi. Viruses. 2017;11:102–9. doi:10.1111/irv.12432

26. Paiva TM, Benega MA, Silva DB, Santos KC, Cruz AS, Hortencio MF, Barbieri MT, Monteiro MM, Barbosa HA, Carvalhanas TR. Evolutionary pattern of reemerging influenza B/Victoria lineage viruses in São Paulo, Brazil, 1996–2012: Implications for vaccine composition strategy. J. Med. Virol. 2013;85:1983–9. doi:10.1002/jmv.23684. PMID:23926069

27. Sochan M, Prosenc K, Učakar V, Berginc NA comparison of the demographic and clinical characteristics of laboratory-confirmed influenza B Yamagata and Victoria lineage infection. J. Clin. Virol. 2014;61:156–60. doi:10.1016/j.jcv.2014.06.018. PMID:25034374

28. Lagacé-Wiens PRS, Rubinstein E, Gumel A. Influenza epidemiology—past, present, and future. Crit. Care Med. 2010;38:e1–e9. doi:10.1097/CCM.0b013e3181cef45a. PMID:20029350

29. Bodewes R, de Mutsert G, van der Klis FR, Ventresca M, Wilks S, Smith DJ, Koopmans M, Fouchier RA, Osterhaus AD, Rimmelzwaan GF. Prevalence of antibodies against seasonal influenza A and B viruses in children in Netherland. Clin. Vaccine Immunom. 2011;18:469–76. doi:10.1128/CVI.00396-10. PMID:21209157

30. Barr IG, Vijaykrishna D, Sullivan SG. Differential age susceptibility to influenza B/Victoria lineage viruses in the 2015 Australian influenza season. Eurosurveillance. 2016;21:30118. doi:10.2807/1560-7917.ES.2016.21.4.30118

31. Tan Y, Guan W, Lam TT, Pan S, Wu S, Zhan Y, Viboud C, Holmes EC, Yang Z. Differing Epidemic Dynamics of Influenza B Virus Lineages in Guangzhou, Southern China, 2009–2010. J. Virol. 2013;87:12447–56. doi:10.1128/JVI.01039-13. PMID:24027322

32. Vijaykrishna D, Holmes EC, Joseph U, Fourment M, Su YC, Halpin R, Lee RT, Deng YM, Gunalan V, Lin X, et al. The contrasting phylogeny of human influenza B viruses. Elite. 2015;4:e5055. doi:10.7554/eLife.05055. PMID:25594904

33. Shiari B, Drori Y, Micheli M, Friedman N, Orzitzer S, Bassal R, Glattman-Freeman A, Shohat T, Mendelson E, Hindyeh M, et al. Epidemiological and Virological Characterization of Influenza B Virus. Infections. PLoS One. 2016;11:e0161195. doi:10.1371/journal.pone.0161195. PMID:27533045

34. Cox N. Influenza seasonality: timing and formulation of vaccines. Bull. World Health Organ. 2014;92:311–311. doi:10.2471/BLT.14.139428. PMID:24839317

35. Wenger JB, Naumova EN. Seasonal Synchronization of Influenza in the United States Older Adult Population. PLoS One. 2010;5:e10187. doi:10.1371/journal.pone.0010187. PMID:20419169

36. Skowronski DM, Chambers C, Sabaidu S, De Serres G, Winter AL, Dickinson JA, Gubbay J, Fonseka K, Charest H, Krajden M, et al. Integrated Sentinel Surveillance Linking Genetic, Antigenic, and Epidemiologic Monitoring of Influenza Vaccine-Virus Relatedness and Effectiveness During the 2013–2014 Influenza Season. J. Infect. Dis. 2015;212:726–39. doi:10.1093/infdis/jiv177. PMID:25784728

37. Chambers C, Skowronski DM, Sabaidu S, Winter AL, Dickinson JA, De Serres G, Gubbay JB, Drews SJ, Martineau C, Eshaghi A, et al. Interim estimates of 2015/16 vaccine effectiveness against influenza A(H1N1)pdm09, Canada, February 2016. Eurosurveillance. 2016;21:30168. doi:10.2807/1560-7917.ES.2016.21.11.30168. PMID:27020673

38. Skowronski DM, Chambers C, Sabaidu S, Dickinson JA, Winter AL, De Serres G, Drews SJ, Jassem A, Gubbay JB, Charest H, Krajden M, et al. Integrated Sentinel Surveillance Linking Genetic, Antigenic, and Epidemiologic Monitoring of Influenza Vaccine-Virus Relatedness and Effectiveness During the 2013–2014 Influenza Season. J. Infect. Dis. 2015;212:726–39. doi:10.1093/infdis/jiv177. PMID:25784728