UC Berkeley UC Berkeley Previously Published Works

Title

Severity of influenza-associated hospitalisations by influenza virus type and subtype in the USA, 2010-19: a repeated cross-sectional study.

Permalink

https://escholarship.org/uc/item/9nm953g7

Journal The Lancet Microbe, 4(11)

Authors

Sumner, Kelsey Masalovich, Svetlana OHalloran, Alissa <u>et al.</u>

Publication Date

2023-11-01

DOI

10.1016/S2666-5247(23)00187-8

Peer reviewed



HHS Public Access

Author manuscript *Lancet Microbe.* Author manuscript; available in PMC 2024 February 16.

Published in final edited form as: *Lancet Microbe.* 2023 November ; 4(11): e903–e912. doi:10.1016/S2666-5247(23)00187-8.

Severity of influenza-associated hospitalisations by influenza virus type and subtype in the USA, 2010–19: a repeated cross-sectional study

Kelsey M Sumner, Svetlana Masalovich, Alissa O'Halloran, Rachel Holstein, Arthur Reingold, Pam Daily Kirley, Nisha B Alden, Rachel K Herlihy, James Meek, Kimberly Yousey-Hindes, Evan J Anderson, Kyle P Openo, Maya L Monroe, Lauren Leegwater, Justin Henderson, Ruth Lynfield, Melissa McMahon, Chelsea McMullen, Kathy M Angeles, Nancy L Spina, Kerianne Engesser, Nancy M Bennett, Christina B Felsen,

This is an Open Access article under the CC BY 4.0 license.

Correspondence to: Dr Kelsey M Sumner, Influenza Division, US Centers for Disease Control and Prevention, Atlanta, GA 30329, USA rhq3@cdc.gov.

Contributors

KMS visualised and wrote the original draft of the report. KMS and SM provided formal analysis. KMS, CR, and SG contributed to conceptualisation. SM, AO'H, and MAR contributed to the methodology. SM and AO'H provided validation and verified the data. JM, RL, and HKT provided project administration. AO'H, RKH, and MM provided data curation. JM, RL, NMB, CBF, CR, and SG provided supervision. A'OH, SM, AR, PDK, NBA, RKH, JM, KY-H, EJA, KPO, MLM, LL, JH, RL, MM, CM, KMA, NLS, KE, NMB, CBF, KL, ES, AT, HKT, WS, AS, AG, RH, MAR, CR, and SG reviewed and edited the report. AR, PDK, NBA, RKH, JM, KY-H, EJA, KPO, MLM, LL, JH, RL, MM, CM, KMA, NLS, KE, NMB, CBF, KL, ES, AT, HKT, WS, AS, AG, RH, MAR, CR, and SG reviewed and edited the report. AR, PDK, NBA, RKH, JM, KY-H, EJA, KPO, MLM, LL, JH, RL, MM, CM, KMA, NLS, KE, NMB, CBF, KL, ES, AT, HKT, WS, AS, and AG were co-investigators or principal investigators for the network. All authors had full access to their site's data in the study and had final responsibility for the decision to submit for publication.

Krista Lung, Eli Shiltz, Ann Thomas, H Keipp Talbot, William Schaffner, Ashley Swain, Andrea George, Melissa A Rolfes, Carrie Reed,

Shikha Garg

Influenza Division (K M Sumner PhD, S Masalovich MS, A O'Halloran MSPH, R Holstein MPH, M A Rolfes PhD, C Reed DSc, S Garg MD), Epidemic Intelligence Service (K M Sumner), US Centers for Disease Control and Prevention, Atlanta, GA, USA; School of Public Health, University of California Berkeley, Berkeley, CA, USA (A Reingold MD); California Emerging Infections Program, Oakland, CA, USA (P D Kirley MPH); Colorado Department of Public Health and Environment, Denver, CA, USA (N B Alden MPH, R K Herlihy MD); Connecticut Emerging Infections Program, Yale School of Public Health, New Haven, CT, USA (J Meek MPH, K Yousey-Hindes MPH); Department of Medicine and Depatment of Pediatrics (E J Anderson MD), Division of Infectious Diseases (K P Openo DrPH), Emory University School of Medicine, Atlanta, GA, USA; Georgia Emerging Infections Program, Georgia Department of Public Health, Atlanta, GA, USA (E J Anderson, K P Openo); Veterans Affairs Medical Center, Atlanta, GA, USA (E J Anderson, K P Openo); Maryland Department of Health, Baltimore, MD, USA (M L Monroe MPH); Michigan Department of Health and Human Services, Lansing, MI, USA (L Leegwater MPH, J Henderson MPH); Minnesota Department of Health, Saint Paul, MN, USA (R Lynfield MD, M McMahon MPH); New Mexico Department of Health, Santa Fe, NM, USA (C McMullen MS); New Mexico Emerging Infections Program, University of New Mexico, Albuquerque, NM, USA (K M Angeles MPH); New York State Department of Health, Albany, NY, USA (N L Spina MPH, K Engesser MPH); University of Rochester School of Medicine and Dentistry, Rochester, NY, USA (N M Bennett MD, C B Felsen MPH); Ohio Department of Health, Columbus, OH, USA (K Lung MPH, E Shiltz MPH); Oregon Health Authority, Salem, OR, USA (A Thomas MD); Vanderbilt University Medical Center, Nashville, TN, USA (H K Talbot MD, W Schaffner MD); Salt Lake County Health Department, Salt Lake City, UT, USA (A Swain CHES, A George MPH)

Abstract

Background—Influenza burden varies across seasons, partly due to differences in circulating influenza virus types or subtypes. Using data from the US population-based surveillance system, Influenza Hospitalization Surveillance Network (FluSurv-NET), we aimed to assess the severity of influenza-associated outcomes in individuals hospitalised with laboratory-confirmed influenza virus infections during the 2010–11 to 2018–19 influenza seasons.

Methods—To evaluate the association between influenza virus type or subtype causing the infection (influenza A H3N2, A H1N1pdm09, and B viruses) and in-hospital severity outcomes (intensive care unit [ICU] admission, use of mechanical ventilation or extracorporeal membrane oxygenation [ECMO], and death), we used FluSurv-NET to capture data for laboratory-confirmed

influenza-associated hospitalisations from the 2010–11 to 2018–19 influenza seasons for individuals of all ages living in select counties in 13 US states. All individuals had to have an influenza virus test within 14 days before or during their hospital stay and an admission date between Oct 1 and April 30 of an influenza season. Exclusion criteria were individuals who did not have a complete chart review; cases from sites that contributed data for three or fewer seasons; hospital-onset cases; cases with unidentified influenza type; cases of multiple influenza virus type or subtype co-infection; or individuals younger than 6 months and ineligible for the influenza vaccine. Logistic regression models adjusted for influenza season, influenza vaccination status, age, and FluSurv-NET site compared odds of in-hospital severity by virus type or subtype. When missing, influenza A subtypes were imputed using chained equations of known subtypes by season.

Findings—Data for 122 941 individuals hospitalised with influenza were captured in FluSurv-NET from the 2010–11 to 2018–19 seasons; after exclusions were applied, 107 941 individuals remained and underwent influenza A virus imputation when missing A subtype (43·4%). After imputation, data for 104 969 remained and were included in the final analytic sample. Averaging across imputed datasets, $57 \cdot 7\%$ (weighted percentage) had influenza A H3N2, 24·6% had influenza A H1N1pdm09, and $17 \cdot 7\%$ had influenza B virus infections; $16 \cdot 7\%$ required ICU admission, $6 \cdot 5\%$ received mechanical ventilation or ECMO, and $3 \cdot 0\%$ died (95% CIs had a range of less than $0 \cdot 1\%$ and are not displayed). Individuals with A H1N1pdm09 had higher odds of in-hospital severe outcomes than those with A H3N2: adjusted odds ratios (ORs) for A H1N1pdm09 versus A H3N2 were $1 \cdot 42$ (95% CI $1 \cdot 32 - 1 \cdot 52$) for ICU admission; $1 \cdot 79$ ($1 \cdot 60 - 2 \cdot 00$) for mechanical ventilation or ECMO use; and $1 \cdot 25$ ($1 \cdot 07 - 1 \cdot 46$) for death. The adjusted ORs for individuals infected with influenza B versus influenza A H3N2 were $1 \cdot 06$ (95% CI $1 \cdot 01 - 1 \cdot 12$) for ICU admission, $1 \cdot 14$ ($1 \cdot 05 - 1 \cdot 24$) for mechanical ventilation or ECMO use, and $1 \cdot 18$ ($1 \cdot 07 - 1 \cdot 31$) for death.

Interpretation—Despite a higher burden of hospitalisations with influenza A H3N2, we found an increased likelihood of in-hospital severe outcomes in individuals hospitalised with influenza A H1N1pdm09 or influenza B virus. Thus, it is important for individuals to receive an annual influenza vaccine and for health-care providers to provide early antiviral treatment for patients with suspected influenza who are at increased risk of severe outcomes, not only when there is high influenza A H3N2 virus circulation but also when influenza A H1N1pdm09 and influenza B viruses are circulating.

Introduction

In the USA, in the decade before the COVID-19 pandemic (2010–20), influenza virus infection was estimated to cause on average up to 710 000 admissions to hospital and 52 000 deaths each year.¹ The burden of influenza varies greatly from season to season² because of differences in circulating influenza virus types and subtypes as well as differences in influenza vaccine coverage and effectiveness,³ with some years seeing lower vaccine effectiveness and higher hospitalisation and mortality rates when influenza A H3N2 viruses are predominant.^{4,5} Despite the effect of virus type or subtype on the yearly influenza burden, little is known about the relative severity of influenza by virus type and subtype in hospitalised individuals. Previous studies^{6–12} have assessed risk factors for severe influenza

in hospitalised individuals by virus type; however, many of these studies only focused on adults,^{6–8,12} were set in Europe or Australia,^{6–9,11,12} or assessed only one season;¹⁰ thus, they are not necessarily representative of influenza severity in children and adults throughout other regions. It is important to understand the severity of influenza infections that lead to hospitalisations across influenza types and subtypes to inform public health messaging and hospital preparation for upcoming influenza seasons.

Using hospitalisation data captured during the 2010–11 to 2018–19 influenza seasons in the US Influenza Hospitalization Surveillance Network (FluSurv-NET), we investigated the relative severity of influenza virus infection by virus type and subtype in both children and adults admitted to hospital with laboratory-confirmed influenza in the USA. Specifically, we described the demographic characteristics and clinical features and complications of individuals hospitalised with laboratory-confirmed influenza according to infecting virus type or subtype and in-hospital severity; and assessed the association between influenza virus type or subtype and in-hospital severity of disease.

Methods

Study design

FluSurv-NET is a population-based surveillance system that actively captures laboratoryconfirmed, influenza-associated hospitalisations for individuals of all ages in more than 250 acute-care hospitals in the USA. The system represents over 29 million individuals, or approximately 9% of the US population. FluSurv-NET cases are hospitalised residents of the surveillance catchment area with a positive influenza virus test within 14 days before or during their hospital stay and an admission date between Oct 1 and April 30 of an influenza season. Influenza test results are ordered at the treating health-care provider's discretion. Positive test results can be from a rapid antigen test, molecular testing, viral culture, or direct or indirect fluorescent staining assay. Surveillance staff identify cases by querying infection control and hospital discharge databases within the FluSurv-NET catchment area hospitals as well as state-wide health information exchanges, electronic laboratory reporting databases, and reportable condition databases. Medical record abstraction is performed by trained surveillance staff using a standard case report form to collect demographic information, underlying conditions, interventions, and outcomes identified while individuals are in the hospital.

For this analysis, we captured data for influenza-associated hospitalisations from the 2010– 11 to 2018–19 influenza seasons from FluSurv-NET sites in the following states: California, Colorado, Connecticut, Georgia, Maryland, Michigan, Minnesota, Ohio, Oregon, New Mexico, New York, Tennessee, and Utah. Sites that contributed data for three or fewer influenza seasons were excluded from this analysis. For the 2010–11 to 2016–17 influenza seasons, complete medical record abstraction was performed for all identified FluSurv-NET cases. For the 2017–18 and 2018–19 influenza seasons, to reduce the burden on site staff, selected sites had complete medical record abstraction only performed for all individuals younger than 50 years, those who died (at any age) during hospitalisation or within 30 days of discharge, and for a random sample of individuals aged 50 years and older.¹³ Sampling weights were created to adjust analyses by the probability of an individual being selected

for complete medical record abstraction, where each individual's sampling weight was the inverse of the probability of selection for complete abstraction. Additional details on the FluSurv-NET sampling scheme have been previously published.¹⁴

Additionally, individuals were excluded if they had hospital-onset influenza (defined as a positive influenza test >3 days after hospital admission), did not have a complete chart review because they were not sampled, influenza type could not be identified, were co-infected with multiple influenza virus types or subtypes, or were younger than 6 months at the time of hospitalisation and ineligible for influenza vaccination. Co-infections with other viruses were not assessed so this was not an exclusion criterion.

FluSurv-NET sites obtained participant and ethics approvals from their respective state health department and academic partner Institutional Review Boards (IRBs) as needed. The US Centers for Disease Control and Prevention (CDC) determined this activity met the requirement for public health surveillance; therefore, the CDC's IRB approval was not required.

Procedures

The main exposure of interest was influenza virus type or subtype, categorized as A H3N2, A H1N1pdm09, or B. The main outcomes of interest were: (1) admission to an intensive care unit (ICU); (2) receipt of invasive mechanical ventilation or extracorporeal membrane oxygenation (ECMO); and (3) death during hospitalisation.

The following covariates were assessed as potential risk factors associated with severe, influenza-associated, in-hospital outcomes: age (6 months–17 years, 18–49 years, 50–64 years, or 65 years), sex, race and ethnicity (White, non-Hispanic; Black, non-Hispanic; Native American or Alaskan Native, non-Hispanic; Asian or Pacific Islander, non-Hispanic; multiracial, non-Hispanic; Hispanic or Latino; or unknown), presence of an underlying medical condition (no comorbid conditions or 1 comorbid condition: lung disease, metabolic disease, blood disorder, cardiovascular disease, neuromuscular disorder, neurologic disease, immunocompromised condition, renal disease, liver disease, obesity, and extreme obesity [among those aged 2 years]), pneumonia during current hospitalisation, seasonal influenza vaccination status (defined as current season vaccination 14 days before a positive influenza test among individuals 6 months old), antiviral treatment, and influenza season.

Statistical analysis

Descriptive tables were created to compare risk factors for severe in-hospital outcomes stratified by influenza virus type or subtype, with counts presented unweighted and percentages weighted by FluSurv-NET's sampling design. Next, we assessed the adjusted association between influenza virus type or subtype and severe in-hospital outcomes by controlling for confounding variables; potential confounding factors were examined and chosen using directed acyclic graph analysis (appendix p 12). Confounders identified were the individual's age (categorised as mentioned previously), receipt of the season's influenza vaccine, influenza season, and FluSurv-NET site.

Because influenza A subtype, seasonal influenza vaccination status, ICU admission, mechanical ventilation or ECMO use, and death had missing observations, we conducted analyses to determine whether multiple imputation was appropriate. We first assessed the pattern of missing data and the mechanism of missingness across influenza A subtype using a χ^2 test with Rao and Scott's second-order correction. We also compared covariate distributions among individuals with missing and observed A subtype data and then regressed the indicator of missing A subtype on other variables in the dataset. After multiple imputation, we compared the parameter estimates produced by logistic regression between complete case analysis and the analysis of multiply imputed data. The proportion of unimputed and imputed A subtypes by season was also compared.

We used multiple imputation by chained equations to impute missing data for variables included in the adjusted analysis model with a focus on imputing influenza A subtype.¹⁵ Race and ethnicity data and presence of underlying medical conditions were also imputed because they were included in the imputation model and had missing observations. The imputation model included all outcome variables (ICU admission, mechanical ventilation or ECMO, and death), confounding covariates from the analysis models (individual's age, categorised into finer groups: 0-4 years, 5-17 years, 18-49 years, 50-64 years, 65-74 years, 75-84 years, and 85 years, seasonal influenza vaccination status, influenza season, and FluSurv-NET site), and other variables associated with A subtype missingness (sex, race and ethnicity, presence of an underlying medical condition, influenza antiviral use, pneumonia presence, and month of hospital admission [grouped as October to December, January, February, and March to April]), and the sampling weight for the individual. These additional variables were selected based on their correlation with other variables in the dataset (ie, tetrachoric and polychoric correlation coefficients 0.3) and their association with indicators for missingness (ie, assessed as statistically significant using logistic regression). Covariates in the imputation model with missing observations were imputed sequentially using logistic regression, with the covariate with the least amount of missing data imputed first. 30 imputed datasets were created. After imputation, we derived a three-category version of the influenza type or subtype variable encompassing influenza A H3N2, A H1N1pdm09, and B viruses. Influenza B lineage was not captured by FluSurv-NET and thus not imputed.

After imputation, to investigate the adjusted odds ratio (OR) between influenza type or subtype and severe in-hospital outcomes, we ran logistic regression models accounting for the complex survey design and adjusted for influenza season, seasonal influenza vaccination status, individual's age, and FluSurv-NET site on each imputed dataset; parameter estimates were summarised using Rubin's rule.¹⁶ Each model was weighted using sampling weights created from sampling strata on the basis of whether an individual died, the individual's age, the influenza season, and FluSurv-NET site. The adjusted association was assessed for effect measure modification on the multiplicative scale by individual's age, influenza season, and seasonal influenza vaccination status; to do so, we compared adjusted ORs and 95% CIs stratified by age, season, or seasonal influenza vaccination status with the full model adjusted OR.

Sensitivity analyses were conducted to evaluate whether having a pre-existing medical condition or receipt of influenza antiviral drugs changed results, with an additional covariate

for each added to the adjusted logistic regression models. Models that were additionally adjusted for having a pre-existing medical condition were also assessed stratified by age. To assess the performance of the imputation model, we performed complete case analysis of the adjusted association between influenza type or subtype with severe in-hospital outcomes and compared results with effect estimates produced using the imputed datasets. Data analyses were performed using SAS version 9.4 software (SAS Institute, Cary, NC, USA) and the 'mice' and 'survey' packages in R version 4.0.3 software (R Core Team, Vienna, Austria).^{15,17} For all statistical analyses, statistical significance was assessed at an α level of 0.05. STROBE and RECORD reporting guidelines were followed in the generation of this report.

Role of the funding source

The funder of the study had a role in study design, data collection, data analysis, data interpretation, and writing of the report.

Results

Data for 122 941 hospitalised individuals with laboratory-confirmed influenza virus infection were captured in FluSurv-NET from the 2010–11 to 2018–19 seasons (figure 1). After exclusions were applied, 107 941 individuals remained in the dataset (figure 1); about half of these individuals were female and most had at least one comorbid condition (table 1). 25.9% (weighted percentage) of individuals were hospitalised with influenza A H3N2, 11.9% with influenza A H1N1pdm09, 43.4% with influenza A unknown subtype, 18.2% with influenza B, 0.4% with influenza A and B, and 0.3% with influenza of unknown virus type (table 1). Specific comorbid conditions across influenza virus type or subtype are described in the appendix (p 2). ICU admission occurred in 16.2% (weighted percentage), mechanical ventilation or ECMO use in 6.2%, and death in 2.7% of 107 941 hospitalisations.

Influenza A subtype was imputed for 43·4% (weighted percentage) of hospitalisations. The pattern of influenza A subtype missingness was non-monotone and probably not missing completely at random (appendix p 4). After influenza A subtype imputation, data for 104 969 remained and were included in the final analytic sample (figure 1). After imputation, averaging across imputed datasets, 57·7% (weighted percentage) of hospitalised individuals had A H3N2, 24·6% had A H1N1pdm09, and 17·7% had influenza B infections; 16·7% required ICU admission, 6·5% received mechanical ventilation or ECMO, and 3·0% died (95% CIs had a range of less than 0·1% and are not displayed). Distributions of imputed influenza virus type or subtype varied by outcome and age group (figure 2).

Using the imputed datasets, and adjusting for influenza season, seasonal influenza vaccination status, the individual's age, and FluSurv-NET site, we found that individuals with influenza A H1N1pdm09 virus infections had higher odds of in-hospital severe outcomes of ICU admission, mechanical ventilation or ECMO, and death than those with A H3N2 virus infection (table 2, appendix p 6). Individuals infected with influenza A H3N2 virus also had higher odds of in-hospital severe outcomes than those with influenza A H3N2 virus (table 2, appendix p 6).

Comparing the association of influenza B virus versus A H3N2 virus and severe in-hospital outcomes across age groups, higher odds of mechanical ventilation or ECMO use were observed in children aged 6 months–17 years and adults aged 18–49 years (table 2). Higher odds of death comparing individuals hospitalised with influenza B versus A H3N2 virus were observed in children aged 6 months to 17 years and adults aged 65 years and older (table 2). By contrast, no association was found comparing odds of death in individuals aged 6 months to 17 years and aged 65 years and older hospitalised with A H1N1pdm09 versus A H3N2 virus; however, the adjusted OR was significant for death in individuals aged 18–49 and 50–64 years.

In analyses stratified by influenza vaccination status, individuals who did not receive the seasonal influenza vaccine had higher odds of death when infected with influenza A H1N1pdm09 or B virus compared with A H3N2 virus (table 2). Associations were less strong for those who had received the seasonal influenza vaccine (table 2). Occasionally, variation in the association of influenza type or subtype with severe in-hospital outcomes was observed across influenza seasons (table 2).

In sensitivity analyses, when adjusting for the presence of at least one underlying medical condition, model results were similar for the outcomes of ICU admission, mechanical ventilation or ECMO use, and death (appendix p 8) even after stratifying by age (appendix p 9). Adjusting for antiviral use also produced results in the same direction of effect across all outcomes, including ICU admission, mechanical ventilation or ECMO use, and death (appendix p 8).

In a complete case sensitivity analysis using the original, unimputed dataset, increased odds of severe in-hospital outcomes were observed when comparing individuals with influenza A H1N1pdm09 with those with A H3N2 virus (adjusted ORs: ICU admission 1.40, 95% CI 1.29-1.51; mechanical ventilation or ECMO use 1.77, 1.58-1.99; death 1.30, 1.10-1.54) but not when comparing influenza B with A H3N2 virus (adjusted ORs: ICU admission 0.95, 95% CI 0.89-1.01; mechanical ventilation or ECMO use 0.99, 0.90-1.09; death 1.14, 1.00-1.30; appendix pp 10–11).

Discussion

Across nearly 105 000 influenza hospitalisations and nine influenza seasons from 2010 to 2019 in the USA, we found that more than one in six individuals hospitalised with influenza had severe outcomes. Although influenza A H3N2 virus-predominant seasons are typically associated with more hospitalisations overall, we found that in-hospital severity was greater among individuals infected with influenza A H1N1pdm09 virus or influenza B virus. Higher in-hospital severity in individuals with A H1N1pdm09 occurred across all age groups when severity was measured as ICU admission, or mechanical ventilation or ECMO use, although results were variable for B and when severity was measured as in-hospital death.

Previous studies noted more severe influenza illness when adults^{7,8,12} or both children and adults^{9–11,18,19} were hospitalised with influenza A H1N1pdm09 virus compared with A H3N2 virus; however, some of these studies found that results varied depending on

the severity outcome measured^{7,18} or age of participants.⁸ We observed similar variability across age and outcome measures, where all ages had increased odds of ICU admission and mechanical ventilation or ECMO use with A H1N1pdm09 virus versus A H3N2 virus, but neither young children nor older adults had increased odds of death; this difference in the association between virus type or subtype and death across age groups could be partly explained by immune imprinting of the specific influenza viruses that individuals were first exposed to in childhood.^{20,21} An evaluation of the effect of immune imprinting on influenza severity was outside the scope of this study but could be investigated in future work. The increased odds of ICU admission and mechanical ventilation or ECMO use that we noted across all ages hospitalised with influenza A H1N1pdm09 virus infections compared with influenza A H3N2 virus infections could be related to the increased infiltration of pneumocytes and intra-alevolar macrophages with influenza A H1N1pdm09 virus, causing alveoli inflammation and more severe disease manifestation, as has been noted in animal models.^{22,23} Future studies could further investigate the biological mechanisms for increased severity of influenza A H1N1pdm09 compared with A H3N2 virus in hospitalised individuals with influenza.

We noted that children younger than 18 years and adults aged 65 years and older infected with influenza B viruses had higher odds of death compared with those infected with A H3N2. Increased odds of death in children with influenza B have been detected before; during the 2010–11 to 2017–18 US influenza seasons, a review²⁴ showed that the proportion of paediatric deaths in individuals infected with influenza B virus was disproportionately higher than the fraction of circulating viruses that were influenza B. In Greece, older adults have also accounted for a disproportionately high number of deaths from influenza B virus infections, as noted over nine seasons.¹¹ Other previous studies found no association with disease severity between infections with influenza B virus versus influenza A^{25,26} or A H3N2 viruses;¹⁰ however, these studies did not present comparisons of the influenza type–severity relationship stratified by age groups.^{10,25,26} We concluded that the relative severity between influenza B and A H3N2 virus infections might be heterogeneous by age, as our age-stratified estimates were statistically significant only for selected age groups and severity measures.

Typically, lower vaccine effectiveness and subsequently higher hospitalisation rates are associated with influenza A H3N2-predominant seasons.⁴ Although we noted a higher number of hospitalisations from infections with influenza A H3N2 virus in our study, we found that once individuals were hospitalised, those with influenza A H1N1pdm09 and B virus infections were more likely to have severe outcomes. Future work could evaluate how the likelihood of hospitalisation compared with in-hospital outcomes differs by influenza type or subtype. Although the reasons for increased odds of severe illness when infected with A H1N1pdm09 or B virus are unclear, it is important for individuals to take precautions to protect themselves against influenza virus infection not only when A H3N2 virus is predominant, but also when A H1N1pdm09 and B viruses are circulating. During the influenza season, increased subtyping to aid early identification of the influenza virus type or subtype infecting most hospitalised individuals could inform planning for hospitals as well as public health messaging aimed at health-care providers around vaccination and appropriate antiviral treatment.^{10,27} Finally, we noted higher odds of death in unvaccinated

individuals who were hospitalised with influenza A H1N1pdm09 or B compared with A H3N2 virus. No association between influenza type or subtype and odds of death was noted in vaccinated individuals, highlighting the importance of receiving the annual influenza vaccine to protect against severe illness.

This study had some limitations. FluSurv-NET, which relies on clinical testing for influenza, probably under-ascertains influenza-associated hospitalisations, as we know that not all individuals hospitalised with acute respiratory illness during the influenza season are tested for influenza.²⁸ Any case under-ascertainment that might have occurred was probably irrespective of influenza type or subtype and we do not believe it affected results. Coinfection with other viruses was not assessed and could have affected an individual's likelihood of having a severe in-hospital outcome. Influenza A subtype information was missing for more than 40% of hospitalised individuals and required imputation, which could have misclassified influenza A subtypes; however, sensitivity analyses found similar model results comparing influenza A H1N1pdm09 with A(H3N2) proportions by season, indicating that misclassification was probably low. Previous studies have successfully imputed the high missingness of influenza A subtype in FluSurv-NET.^{13,14,29} Additionally, differences in vaccine effectiveness by season could have varied by type or subtype. Although we were able to adjust for influenza vaccination and season, we were unable to adjust estimates for annual vaccine effectiveness. Vaccine effectiveness is typically lower for influenza A H3N2 than for A H1N1pdm09 virus,³⁰ which, when not accounted for, would increase the likelihood of individuals having more severe A H3N2 virus infections and bias our results towards the null. Influenza B lineage was not captured by FluSurv-NET, so differences in influenza severity by B lineage could not be assessed. Finally, we were unable to assess the relative risk of hospitalisation with A H3N2 virus compared with A H1N1pdm09 or B because we did not have population level data on prevalence of infection by influenza type or subtype. Future analyses could assess the relative risk of hospitalisation by influenza subtype.

Overall, differences in in-hospital disease severity were noted by influenza type and subtype, with higher odds of severe disease in individuals hospitalised with influenza A H1N1pdm09 and B viruses than in those with influenza A H3N2 virus. Thus, it is important for individuals to receive an annual influenza vaccine and for health-care providers to prescribe early antiviral treatment for patients with suspected influenza who are at higher risk for complications, not only when there is high influenza A H3N2 virus circulation but also when influenza A H1N1pdm09 and B viruses are circulating. Early season detection of influenza type and subtype can inform public health messaging and vaccination campaigns to reduce influenza-associated hospitalisations and severe in-hospital outcomes; this is especially important now with co-circulation of other respiratory viruses, such as SARS-CoV-2 and respiratory syncytial virus, adding to the respiratory disease burden in hospital settings.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

This work was funded by grant CK17-1701 from the CDC through an Emerging Infections Program cooperative agreement, grant 5U38HM000414 from a 2008-13 Influenza Hospitalization Surveillance Project (IHSP) cooperative agreement, grant 5U38OT000143 from a 2013-18 IHSP cooperative agreement, and grant 5NU38OT000297 from a 2018-23 IHSP cooperative agreement. The project was supported by CTSA award UL1 TR002243 from the National Center for Advancing Translational Sciences (Vanderbilt University Medical Center, Nashville, TN, USA). The findings and conclusions in this report are those of the authors and do not necessarily represent the views of the CDC. Some authors are federal employees of the United States Government, and this work was prepared as part of their official duties. Title 17 U.S.C. 105 provides that "copyright protection under this title is not available for any work of the United States Government". We thank the hospitals at participating sites for their efforts to record influenza hospitalisations. We would like to acknowledge the infection preventionists for their reporting efforts and the microbiology laboratories for submitting influenza specimens to their state public health laboratories for subtyping. We thank Sherry Quach, Gretchen Rothrock, Jeremy Roland, Joelle Nadle, Ashley Coates, and Susan Brooks at the California Emerging Infections Program (Oakland, CA, USA); Darcy Fazio, Amber Maslar, Adam Misiorski, and Tamara Rissman with the Connecticut Emerging Infections Program and Yale School of Public Health (New Haven, CT, USA); Emily Fawcett, Olivia Almendares, Megan Riley Lasure, Jeremiah Williams, Katelyn Ward, and Taylor Eisenstein with the Foundation for Atlanta Veterans Education and Research, the Georgia Emerging Infections Program, Georgia Department of Public Health, and Atlanta Veterans Affairs Medical Center (Atlanta, GA, USA); Patricia Ryan, David Blythe, Alicia Brooks, Robert Sunkel, Elisabeth Vaeth, Molly Hyde, Brian Bachaus, Emily Blake, and Cindy Bryant with the Maryland Emerging Infections Program and Maryland Department of Health (Baltimore, MD, USA); Jim Collins, Shannon Johnson, Justin Henderson, Sue Kim, Val Tellez Nunez, Alyanna Melicor, Libby Reeg, Chloe Brown, and Sanchitha Meda with the Michigan Department of Health and Human Services (Lansing, MI, USA); Craig Morin with the Minnesota Department of Health FluSurv-NET team and Minnesota Department of Health Public Health Laboratory (St Paul, MN, USA); Daniel M Sosin, Chad Smelser, Yomei Shaw, Sunshine Martinez, Jasmyn Sanchez, Mark Montoya, and Sri Chinta with the New Mexico Department of Health (Santa Fe, NM, USA); Sarah A Khanlian, Nancy Eisenberg, Molly Bleecker, Mayvilynne Poblete, and Zachary Landis with the New Mexico Emerging Infections Program (Health Sciences Center, Albuquerque, NM, USA); Marc Martinez, Caroline McCahon, Saif Shaman, Yassir Talha, Jennifer Akpo, Alesia Reed, and Murtada Khalifa with the CDC Foundation and New Mexico Department of Health (Santa Fe, NM, USA); Adam Rowe, Jemma Rowlands, and Suzanne McGuire with the New York State Department of Health; Sophrena Bushey, Virginia Cafferky, Maria Gaitan, Christine Long, and Thomas Peer with the University of Rochester School of Medicine and Dentistry (Rochester, NY, USA); Laurie Billing and Nicholas Fisher with the Ohio Department of Health (Columbus, OH, USA); Sam Hawkins with the Public Health Division of the Oregon Health Authority; Tiffanie Markus, Katie Dyer, Karen Leib, Terri McMinn, Danielle Ndi, Gail Hughett, Bentley Akoko, Kathy Billings, Anise Elie, and Manideepthi Pemmaraju of Vanderbilt University Medical Center (Nashville, TN, USA); and Amanda Carter, Andrea Price, Andrew Haraghey, Ashton Bruno, Emily Roberts, Kristen Olsen, Mary Hill, Melanie Crossland, Ryan Chatelain, and Holly Staten with the Salt Lake County Health Department (Salt Lake City, UT, USA).

Funding

The US Centers for Disease Control and Prevention.

Declaration of interests

EJA has consulted for Pfizer, Sanofi Pasteur, GlaxoSmithKline (GSK), Janssen, Moderna, and Medscape; serves on a safety monitoring board for Kentucky BioProcessing and Sanofi Pasteur; and serves on a data adjudication board for WCG and ACI Clinical. EJA's institution receives funds to conduct clinical research unrelated to this manuscript from MedImmune, Regeneron, PaxVax, Pfizer, GSK, Merck, Sanofi Pasteur, Janssen, and Micron, and has also received funding from the US National Institutes of Health to conduct clinical trials of COVID-19 vaccines. ES notes the agency grant funding supporting the data collection for this project, and agency grant funding from the US Centers for Disease Control and Prevention (CDC) to support other epidemiology projects. HKT, WS, NBA, JM, KY-H, RKH, and RL have received CDC grant funding. JH and LL have received grants from the Michigan Department of Health and Human Services. ES reports grants from the Council for State and Territorial Epidemiologists (CSTE) during the conduct of the study and grants from CDC outside the submitted work. AG reports grants from CSTE. All other authors declare no competing interests.

Data sharing

Individuals interested in receiving a limited dataset can submit a brief proposal to the corresponding author (KMS) for review and consideration by the CDC and FluSurv-NET site partners.

References

- 1. Centers for Disease Control and Prevention. Disease burden of flu. 2022. https://www.cdc.gov/flu/ about/burden/index.html (accessed Aug 22, 2022).
- Biggerstaff M, Kniss K, Jernigan DB, et al. Systematic assessment of multiple routine and near real-time indicators to classify the severity of influenza seasons and pandemics in the United States, 2003–2004 through 2015–2016. Am J Epidemiol 2018; 187: 1040–50. [PubMed: 29053783]
- 3. Xu X, Blanton L, Isa Abd Elal A, et al. Update: influenza activity in the United States during the 2018–19 season and composition of the 2019–20 influenza vaccine. Morb and Mortal Wkly Rep 2019; 68: 544–51.
- 4. Tenforde MW, Chung J, Smith ER, et al. Influenza vaccine effectiveness in inpatient and outpatient settings in the United States, 2015–2018. Clin Infect Dis 2021; 73: 386–92. [PubMed: 32270198]
- Vestergaard LS, Nielsen J, Krause TG, et al. Excess all-cause and influenza-attributable mortality in Europe, December 2016 to February 2017. Euro Surveill 2017; 22: 30506. [PubMed: 28424146]
- Chagvardieff A, Persico N, Marmillot C, Badiaga S, Charrel R, Roch A. Prospective comparative study of characteristics associated with influenza A and B in adults. Med Mal Infect 2018; 48: 180–87. [PubMed: 29258804]
- Martinez A, Soldevila N, Romero-Tamarit A, et al. Risk factors associated with severe outcomes in adult hospitalized patients according to influenza type and subtype. PLoS One 2019; 14: e0210353. [PubMed: 30633778]
- Derqui N, Nealon J, Mira-Iglesias A, Diez-Domingo J, Mahe C, Chaves SS. Predictors of influenza severity among hospitalized adults with laboratory confirmed influenza: analysis of nine influenza seasons from the Valencia region, Spain. Influenza Other Respir Viruses 2022; 16: 862–72. [PubMed: 35411561]
- Delgado-Sanz C, Mazagatos-Ateca C, Oliva J, Gherasim A, Larrauri A. Illness severity in hospitalized influenza patients by virus type and subtype, Spain, 2010–2017. Emerg Infect Dis 2020; 26: 220–28. [PubMed: 31961295]
- Chaves SS, Aragon D, Bennett N, et al. Patients hospitalized with laboratory-confirmed influenza during the 2010–2011 influenza season: exploring disease severity by virus type and subtype. J Infect Dis 2013; 208: 1305–14. [PubMed: 23863950]
- Lytras T, Andreopoulou A, Gkolfinopoulou K, Mouratidou E, Tsiodras S. Association between type-specific influenza circulation and incidence of severe laboratory-confirmed cases; which subtype is the most virulent? Clin Microbiol Infect 2020; 26: 922–27. [PubMed: 31760112]
- Minney-Smith CA, Selvey LA, Levy A, Smith DW. Post-pandemic influenza A/H1N1pdm09 is associated with more severe outcomes than A/H3N2 and other respiratory viruses in adult hospitalisations. Epidemiol Infect 2019; 147: e310. [PubMed: 31775940]
- Chow EJ, Rolfes MA, O'Halloran A, et al. Respiratory and nonrespiratory diagnoses associated with influenza in hospitalized adults. JAMA Netw Open 2020; 3: e201323. [PubMed: 32196103]
- Chow EJ, Rolfes MA, O'Halloran A, et al. Acute cardiovascular events associated with influenza in hospitalized adults: a cross-sectional study. Ann Intern Med 2020; 173: 605–13. [PubMed: 32833488]
- van Buuren S, Groothuis-Oudshoorn K. mice: multivariate imputation by chained equations in R. J Stat Softw 2011; 45: 1–67.
- Rubin DB. Multiple imputation for nonresponse in surveys. New York (NY): John Wiley & Sons, 1987.
- 17. Lumley T Analysis of complex survey samples. J Stat Softw 2004; 9: 19.
- Park JE, Ryu Y. Transmissibility and severity of influenza virus by subtype. Infect Genet Evol 2018; 65: 288–92. [PubMed: 30103034]
- Caini S, Kroneman M, Wiegers T, El Guerche-Séblain C, Paget J. Clinical characteristics and severity of influenza infections by virus type, subtype, and lineage: a systematic literature review. Influenza Other Respir Viruses 2018; 12: 780–92. [PubMed: 29858537]
- 20. Verma N, Dimitrova M, Carter DM, et al. Influenza virus H1N1pdm09 infections in the young and old: evidence of greater antibody diversity and affinity for the hemagglutinin globular head

domain (HA1 domain) in the elderly than in young adults and children. J Virol 2012; 86: 5515–22. [PubMed: 22379097]

- Gostic KM, Bridge R, Brady S, Viboud C, Worobey M, Lloyd-Smith JO. Childhood immune imprinting to influenza A shapes birth year-specific risk during seasonal H1N1 and H3N2 epidemics. PLoS Pathog 2019; 15: e1008109. [PubMed: 31856206]
- 22. Guarner J, Falcon-Escobedo R. Comparison of the pathology caused by H1N1, H5N1, and H3N2 influenza viruses. Arch Med Res 2009; 40: 655–61. [PubMed: 20304252]
- Itoh Y, Shinya K, Kiso M, et al. In vitro and in vivo characterization of new swine-origin H1N1 influenza viruses. Nature 2009; 460: 1021–25. [PubMed: 19672242]
- 24. Doyle JD, Campbell AP. Pediatric influenza and illness severity: what is known and what questions remain? Curr Opin Pediatr 2019; 31: 119–26. [PubMed: 30531402]
- 25. Oh YN, Kim S, Choi YB, Woo SI, Hahn YS, Lee JK. Clinical similarities between influenza A and B in children: a single-center study, 2017/18 season, Korea. BMC Pediatr 2019; 19: 472. [PubMed: 31796033]
- 26. Sharma Y, Horwood C, Hakendorf P, Thompson C. Clinical characteristics and outcomes of influenza A and B virus infection in adult Australian hospitalised patients. BMC Infect Dis 2020; 20: 913. [PubMed: 33261559]
- 27. US Centers for Disease Control and Prevention. People at higher risk of flu complications. 2022. https://www.cdc.gov/flu/highrisk/index.htm (accessed Sept 28, 2022).
- Tenforde MW, Campbell AP, Michaels MG, et al. Clinical influenza testing practices in hospitalized children at United States medical centers, 2015–2018. J Pediatric Infect Dis Soc 2022; 11: 5–8. [PubMed: 34643241]
- Holstein R, Dawood FS, O'Halloran A, et al. Characteristics and outcomes of hospitalized pregnant women with influenza, 2010 to 2019: a repeated cross-sectional study. Ann Intern Med 2022; 175: 149–58. [PubMed: 34958603]
- Belongia EA, McLean HQ. Influenza vaccine effectiveness: defining the H3N2 problem. Clin Infect Dis 2019; 69: 1817–23. [PubMed: 31102401]

Research in context

Evidence before this study

Influenza virus type or subtype circulation and hospitalisation burden varies by season, but information is scarce regarding how severe influenza outcomes, and thus use of intensive care unit resources, differ by influenza type or subtype once individuals are admitted to hospital. We did a PubMed search of "influenza" AND "severity" AND ("type" OR "subtype") of studies published from database inception to Dec 31, 2022, with no language restrictions. 2066 articles met the PubMed search criteria; eight of these were research articles that investigated severe influenza in hospitalised individuals by virus type. These articles were predominantly set in Europe and Australia. The previous studies mainly focused on adults, smaller populations, or few seasons.

Added value of this study

We built on the previous studies by assessing the association between influenza virus type or subtype and in-hospital severity for nearly 105 000 patients of all ages with influenza admitted to hospital across nine influenza seasons in the USA.

Implications of all the available evidence

We showed that, although influenza A H3N2 virus-predominant seasons are typically associated with more hospitalisations overall, in-hospital severity was greater in individuals infected with influenza A H1N1pdm09 or influenza B viruses compared with influenza A H3N2 virus. Thus, it is important for individuals to receive an annual influenza vaccine and for health-care providers to provide early antiviral treatment for high-risk patients with suspected influenza, not only when there is high circulation of A H3N2 virus, but also when A H1N1pdm09 and B viruses are circulating.

Author Manuscript

Author Manuscript



Figure 1: Study profile

Data are for individuals hospitalised with laboratory-confirmed influenza captured in the US Influenza Hospitalization Surveillance Network, 2010–19. Main analysis model covariates that were imputed were influenza A virus subtype, seasonal influenza vaccination status, mechanical ventilation or use of extracorporeal membrane oxygenation, admission to an intensive care unit, and death. Race and ethnicity, and presence of at least one underlying medical condition, were part of the imputation model and imputed due to missingness.

Sumner et al.





Data are for all captured hospitalisations, those admitted to an ICU, those who received mechanical ventilation or ECMO, and those who died in the hospital. Across each age group, the mean proportion of hospitalised individuals with influenza A H3N2, A H1N1pdm09, and influenza B virus infections is shown across all hospitalisations and across the presence or absence of each severity measure. The mean percentages of influenza types and imputed subtypes were calculated across the 30 imputed data sets; 95% CIs had a range of less than 0.1% and are not displayed in the figures. ICU=intensive care unit. ECMO=extracorporeal membrane oxygenation.

Table 1:

Characteristics of individuals hospitalised with laboratory-confirmed influenza in the USA, 2010–19, by infecting virus type and unimputed subtype *

	Total cases (n=107 941)	Influenza A (n=88 276)		Influenza B (n=18 909)	Influenza A and B (n=441)	Unknown type (n=315)
		H3N2 (n=28 409)	H1N1pdm09 (n=13 597)	Unknown A subtype (n=46 270)	-		
Demographic cha	racteristics						
Age, years							
0–17	13 236 (11·2%)	3080 (10·1%)	2121 (15.1%)	4749 (9.3%)	3090 (14.5%)	75 (15.7%)	121 (37.5%)
18–49	19 438 (16·5%)	3869 (12·7%)	3752 (26.7%)	8327 (16.3%)	3360 (15.7%)	90 (18.8%)	40 (12.4%)
50-64	23 422 (21·0%)	5013 (17·1%)	4295 (31.3%)	9833 (20.4%)	4139 (21.5%)	95 (21.7%)	47 (14.8%)
65	51 845 (51·2%)	16 447 (60·1%)	3429 (26.9%)	23 361 (54·0%)	8320 (48.3%)	181 (43.7%)	107 (35.3%)
Sex							
Male	49 814 (46·1%)	12 977 (45·6%)	6626 (48.7%)	21 132 (45·7%)	8720 (46.0%)	198 (43.6%)	161 (51-1%)
Female	58 127 (53·9%)	15 432 (54·4%)	6971 (51-3%)	25 138 (54·3%)	10 189 (54·0%)	243 (56.4%)	154 (48.9%)
Race or ethnicity							
White, non- Hispanic	61 951 (58·1%)	17 179 (61·4%)	7360 (54-4%)	26 563 (58·0%)	10 451 (56·2%)	243 (52.7%)	155 (49.5%)
Black, non- Hispanic	21 682 (19·7%)	4936 (16·9%)	2972 (21.7%)	9533 (20·2%)	4067 (20.8%)	103 (23.9%)	71 (22.0%)
Native American or Alaskan Native, non-Hispanic	718 (0.7%)	137 (0.5%)	118 (0.9%)	362 (0.8%)	90 (0.4%)	8 (1.7%)	3 (0.9%)
Asian or Pacific Islander, non-Hispanic	4808 (4.8%)	1155 (4.1%)	544 (4.1%)	2239 (5.3%)	833 (5.1%)	23 (5.8%)	14 (4.3%)
Multiracial, non-Hispanic	391 (0.3%)	86 (0.3%)	53 (0.4%)	174 (0.4%)	75 (0.4%)	2 (0.4%)	1 (0.3%)
Hispanic or Latino	9753 (8.9%)	2232 (7.8%)	1548 (11.2%)	4103 (8.7%)	1789 (9.3%)	41 (11.0%)	40 (13.3%)
Unknown	8638 (7.6%)	2684 (9.1%)	1002 (7.3%)	3296 (6.7%)	1604 (7.9%)	21 (4.4%)	31 (9.6%)
Influenza season							
2010-11	5442 (4.6%)	1653 (5.4%)	935 (6.7%)	1871 (3.7%)	917 (4.3%)	27 (5.7%)	39 (12-1%)
2011-12	2294 (1.9%)	950 (3.1%)	312 (2.2%)	755 (1.5%)	260 (1.2%)	8 (1.7%)	9 (2.8%)
2012-13	11 088 (9·4%)	3816 (12·5%)	165 (1.2%)	4725 (9.3%)	2311 (10.8%)	28 (5.9%)	43 (13·3%)
2013-14	9255 (7.9%)	456 (1.5%)	4460 (31.8%)	3250 (6.4%)	1011 (4.7%)	34 (7.1%)	44 (13.6%)
2014–15	17 235 (14·6%)	6805 (22·3%)	16 (0.1%)	7919 (15.5%)	2356 (11.0%)	103 (21.6%)	36 (11·2%)
2015-16	8506 (7.2%)	427 (1.4%)	2952 (21.0%)	2924 (5.7%)	2129 (10.0%)	51 (10.7%)	23 (7.1%)

	Total cases (n=107 941)	Influenza A (n=88 276)			Influenza B (n=18 909)	Influenza A and B (n=441)	Unknown type (n=315)	
		H3N2 (n=28 409)	H1N1pdm09 (n=13 597)	Unknown A subtype (n=46 270)				
2016–17	16 992 (14·4%)	5964 (19·6%)	117 (0.8%)	7199 (14.1%)	3604 (16.9%)	63 (13·2%)	45 (13.9%)	
2017-18	20 996 (24·6%)	5269 (22·9%)	1078 (9.2%)	8837 (24.4%)	5685 (37.7%)	88 (25.6%)	39 (13·2%)	
2018–19	16 133 (15·3%)	3069 (11·2%)	3562 (27.0%)	8790 (19.5%)	636 (3.4%)	39 (8.6%)	37 (12.7%)	
Underlying medica	al condition							
No comorbid conditions	13 901 (12·3%)	3039 (10·3%)	2166 (15.6%)	5644 (11.6%)	2905 (14.5%)	69 (14·5%)	78 (24·2%)	
1 comorbid conditions	93 751 (87·4%)	25 297 (89·5%)	11 417 (84.3%)	40 475 (88·0%)	15 954 (85·2%)	372 (85.5%)	236 (75.2%)	
Unknown	289 (0.3%)	73 (0.2%)	14 (0.1%)	151 (0.4%)	50 (0.3%)	0 (0.0%)	1 (0.6%)	
Seasonal influenza	vaccination*							
No	44 087 (40·6%)	10 219 (35·8%)	7273 (54·2%)	18 363 (39·5%)	7911 (41.2%)	185 (42-4%)	136 (44.8%)	
Yes	48 212 (46·3%)	14 208 (51·6%)	4561 (34.8%)	21 155 (47·2%)	7981 (44.0%)	173 (41·2%)	134 (45.7%)	
Unknown	13613 (13·1%)	3443 (12·5%)	1458 (11-1%)	5921 (13.3%)	2699 (14.8%)	64 (16·4%)	28 (9.5%)	
Influenza complic	cations and inter	ventions						
Pneumonia during	current hospitalis	sation						
No	84 122 (77·9%)	22 810 (80·3%)	9766 (72.0%)	36 151 (78·0%)	14 842 (78·5%)	317 (72.5%)	236 (74.9%)	
Yes	23 819 (22·1%)	5599 (19·7%)	3831 (28.0%)	10 119 (22·0%)	4067 (21.5%)	124 (27.5%)	79 (25.1%)	
Antiviral treatment	t							
No	15 331 (13·5%)	4811 (16·2%)	2171 (15.6%)	4908 (10.1%)	3306 (16.4%)	66 (14.1%)	69 (21.7%)	
Yes	92 610 (86·5%)	23 598 (83·8%)	11426 (84.4%)	41 362 (89·9%)	15 603 (83·6%)	375 (85.9%)	246 (78.3%)	
Influenza disease severity								
ICU admission								
No	89 381 (83·2%)	23 778 (84·1%)	10 425 (76.8%)	38 868 (84·4%)	15 702 (83·5%)	341 (77.9%)	267 (84.5%)	
Yes	17 971 (16·2%)	4466 (15·3%)	3131 (22.9%)	7133 (15.0%)	3097 (15.9%)	97 (21.4%)	47 (14.9%)	
Unknown	589 (0.6%)	165 (0.5%)	41 (0.3%)	269 (0.6%)	110 (0.6%)	3 (0.6%)	1 (0.6%)	
Mechanical ventilation or ECMO								
No	10 0228 (93·1%)	26 571 (93·7%)	12 068 (88.8%)	43 275 (93·7%)	17 620 (93·4%)	397 (90.8%)	297 (93.8%)	
Yes	6968 (6.2%)	1613 (5.6%)	1467 (10.8%)	2669 (5.5%)	1161 (5.9%)	41 (8.6%)	17 (5.6%)	
Unknown	745 (0.7%)	225 (0.8%)	62 (0.5%)	326 (0.8%)	128 (0.7%)	3 (0.6%)	1 (0.6%)	
Death, n (%)								
No	10 4360 (96·9%)	27 445 (96·8%)	13166 (96.9%)	44 783 (97·1%)	18 248 (96·9%)	412 (93.9%)	306 (96.9%)	

	Total cases (n=107 941)	Influenza A (n=88 276)			Influenza B (n=18 909)	Influenza A and B (n=441)	Unknown type (n=315)
		H3N2 (n=28 409)	H1N1pdm09 (n=13 597)	Unknown A subtype (n=46 270)			
Yes	3227 (2.7%)	859 (2.8%)	405 (2.9%)	1323 (2.6%)	606 (2.8%)	26 (5.4%)	8 (2.5%)
Unknown	354 (0.3%)	105 (0.3%)	26 (0.2%)	164 (0.3%)	55 (0.3%)	3 (0.6%)	1 (0.6%)
Days hospitalised, median (IQR)	3 (3)	3 (3)	3 (4)	3 (3)	3 (3)	3 (4)	3 (3)

Data are n (%) unless otherwise stated. Counts are presented unweighted, whereas percentages were weighted using complex survey weights. Median and IQR presented were weighted using complex survey weights. Data presented are before imputation. ICU=intensive care unit. ECMO=extracorporeal membrane oxygenation.

* Only captured for individuals aged 6 months or older; 2029 observations from infants younger than 6 months were excluded.

Table 2:

Adjusted odds ratios of severe outcomes by influenza type and imputed influenza A virus subtype for individuals hospitalised with laboratory-confirmed influenza in the USA, 2010–19

	ICU admission		Mechanical ventilatio	n or ECMO use	Death	
	A H1N1pdm09 vs A H3N2	B vs A H3N2	A H1N1pdm09 vs A H3N2	B vs A H3N2	A H1N1pdm09 vs A H3N2	B vs A H3N2
Overall	1.42 (1.32–1.52)	1·06 (1·01– 1·12)	1.79 (1.60–2.00)	1·14 (1·05– 1·24)	1.25 (1.07–1.46)	1·18 (1·07– 1·31)
Age						
6 months to 17 years	1.26 (1.08–1.47)	0·92 (0·82– 1·04)	1.67 (1.26–2.22)	1·30 (1·04– 1·61)	1.31 (0.54–3.16)	2·55 (1·42– 4·57)
18–49 years	1.43 (1.26–1.63)	1·10 (0·98– 1·23)	1.79 (1.47–2.18)	1·24 (1·04– 1·48)	1.81 (1.14–2.87)	1·39 (0·93– 2·06)
50-64 years	1.43 (1.25–1.64)	1·05 (0·95– 1·17)	1.77 (1.47–2.14)	1·08 (0·93– 1·26)	1.49 (1.13–1.98)	1·11 (0·88– 1·41)
65 years	1.49 (1.32–1.67)	1·12 (1·03– 1·21)	1.83 (1.52–2.20)	1·10 (0·97– 1·25)	1.05 (0.87–1.27)	1·15 (1·03– 1·30)
Seasonal influer	nza vaccination status					
No	1.41 (1.29–1.54)	1·06 (0·98– 1·14)	1.86 (1.61–2.14)	1·17 (1·05– 1·32)	1.49 (1.19–1.87)	1·27 (1·07– 1·51)
Yes	1.42 (1.28–1.58)	1·07 (0·99– 1·16)	1.68 (1.44–1.95)	1·12 (0·99– 1·26)	1.06 (0.87–1.31)	1·12 (0·98– 1·29)
Influenza seasor	n					
2010-11	1.73 (1.41–2.11)	0·94 (0·74– 1·20)	2.01 (1.50–2.68)	1·10 (0·78– 1·55)	1.28 (0.81–2.03)	1·20 (0·72– 2·01)
2011-12	1.19 (0.83–1.72)	1·29 (0·87– 1·90)	1.44 (0.82–2.52)	1·70 (0·94– 3·07)	1.24 (0.49–3.12)	2·08 (0·88– 4·91)
2012-13	1.35 (0.97–1.87)	1·03 (0·90– 1·18)	1.44 (0.90–2.32)	1·10 (0·90– 1·34)	0.90 (0.25–3.27)	1.21 (0.88 - 1.68)
2013-14	1.39 (1.09–1.77)	0·94 (0·70– 1·25)	2.06 (1.38–3.08)	1·36 (0·86– 2·13)	0.93 (0.55–1.60)	0·71 (0·38– 1·32)
2014–15*	1.29 (0.39–4.24)	1·09 (0·96– 1·23)	1.79 (0.44–7.19)	1·13 (0·94– 1·35)	0.001 (0.00–348.24)	1·07 (0·81– 1·41)
2015-16	1.27 (0.98–1.64)	0·97 (0·74– 1·26)	1.71 (1.17–2.48)	1·04 (0·70– 1·55)	1.25 (0.70–2.22)	1·13 (0·63– 2·04)
2016-17	1.60 (1.13–2.27)	1·00 (0·90– 1·11)	1.53 (0.87–2.70)	0·99 (0·84– 1·18)	0.86 (0.27–2.72)	1·17 (0·93– 1·47)
2017-18	1.46 (1.25–1.71)	1·14 (1·02– 1·26)	1.81 (1.39–2.35)	1·25 (1·06– 1·48)	1.34 (0.95–1.90)	1·24 (1·05– 1·47)
2018–19	1.35 (1.20–1.51)	1·19 (0·94– 1·51)	1.69 (1.40–2.04)	1·02 (0·68– 1·56)	1.25 (0.98–1.59)	1·24 (0·77– 1·99)

Data are adjusted odds ratios (95% CI), stratified by age, receipt of seasonal influenza vaccine, and influenza season. Odds ratios and corresponding 95% CIs were estimated using logistic regression models accounting for survey weights and imputation for each outcome separately. Results were pooled across 30 imputed datasets using Rubin's rules. Each model included influenza type and subtype, age (6 months to 17 years, 18–49 years, 50–64 years, and 65 years), seasonal influenza vaccination status, influenza season, and FluSurv-NET site. ICU=intensive care unit. ECMO=extracorporeal membrane oxygenation.

* Very few influenza A H1N1pdm09 viruses circulated during the 2014–15 influenza season.