

Surveillance of severe influenza and COVID-19 in Sweden

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Surveillance sources for severe illness

Hospital/intensive care bed occupancy

Hospital admissions (with significant delay)

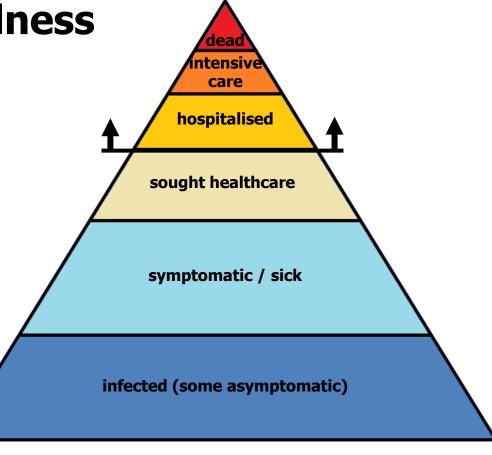
Intensive care admissions

Deceased laboratory-confirmed cases

Cause of death data (with significant delay)

• Excess mortality modelling

 Virus characterisation and sequencing, including antiviral resistance monitoring, monitoring of genetic groups





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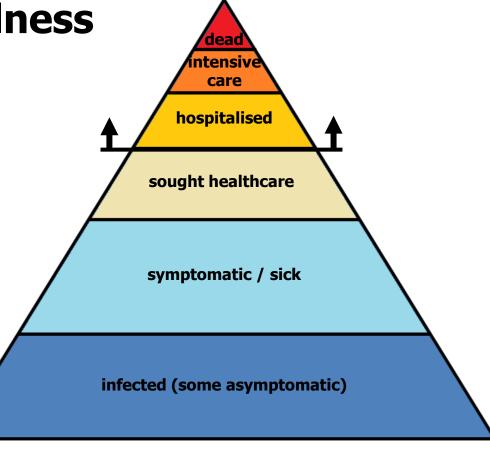
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Surveillance of intensive care admissions

- All patients admitted to intensive care with laboratory-confirmed influenza or COVID-19
 - Combined with mandatory notifications from laboratories
- Developed during the 2009 influenza pandemic
- Nearly universal coverage, ca 10 million population
- Data from the Swedish Intensive Care Registry added once care ends
 - -https://www.icuregswe.org/en/
- Year-round surveillance



Variables collected upon intake

- Virus and subtype (if available)
- Travel abroad
- Risk group status
- Vaccination status
- Antiviral treatment
- Secondary bacterial infections
- Unusual symptoms (e.g. CNS or heart symptoms)

	Riskgruppstillhörighet* (flerval) Ingen Ingen Barn under 3 år, som fötts för tidigt (före graviditetsvecka 35) Hypertoni Kronisk hjärtsjukdom, oavsett ålder Barn med medfödda eller akuta hjärtfel Kronisk lungsjukdom (inklusive astma >3 år), oavsett 1 Kronisk leversvikt	
SATEMBER VA	SIRI-protokoll för influensa och virus	7
P	eg 1 - Grunduppgifter //A*: Din egen avdelning kommer upp automatiskt när du loggar in med SITHS- kort //a*sonidentitet* (enval) // Korrekt (giltigt svenskt personnummer eller samordningsnummer):	agnation (till ex
1 1	Reservnummer: mäiligt anges:	ående uppgift
1	Om reservnummer angivits skall födelsedatum om mojnigt angivers. Datum för insjuknande*. Avser datum för första debut av symtom:	
1 1	Vårdtillfällets start på IVA*: 🏥	uknande med
	Steg 2 - Diagnostik Laboratoriefynd* (enval) Influensa A. Om Ja Subtyp efter identitet på H- och N-proteinerna. (enval) H1N1 H3N2 H5N1 H7N9 Annan Subtyp (fritext): Subtyp ej känd Influensa B Coronavirus Covid-19 (SARS-CoV-2) RS-virus (Respiratory syncytial virus - RSV) Annat agens. Vilket agens? (fritext): Ej laboratorieverifierad (vid stark misstanke om influensa eller annan svår luftburen luftvägssjukdom kan rapportering ske här, men skall (vid stark misstanke om influensa eller annan svår luftburen luftvägssjukdom kan rapportering ske här, men skall en ofullständig registrering och korrigeras alternativt raderas når laboratoriesvar anlänt) Steg 3 - Bakgrund Utlandsvistelse före insjuknandet. Avser utlandsvistelse <10 dagar för insjuknande*	I ses som
	(enval, e) aktuen om 15 m. (enval, e) aktuen om	rval)
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Variables added from the Swedish Intensive Care Registry

- Length of stay
- Interventions in intensive care (ECMO*, mechanical ventilation, etc.)
- Primary diagnosis, secondary diagnoses
- Status at end of intensive care (died, moved to other ward, discharged)

*ECMO = extracorporeal membrane oxygenation



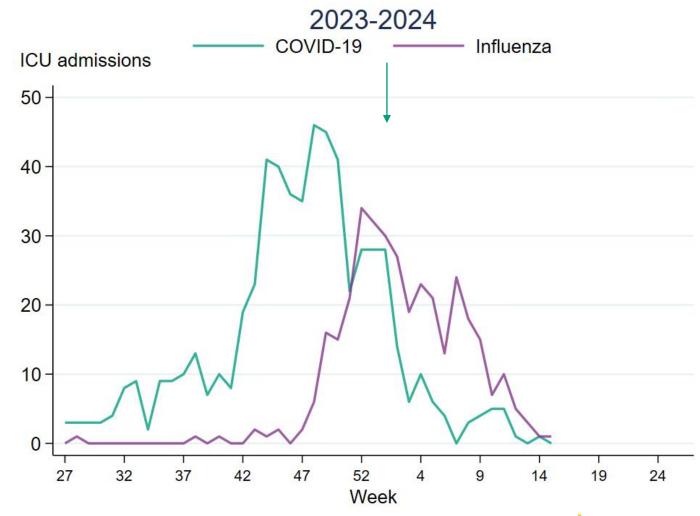
The 2023-2024 season in Sweden

Influenza (94 % flu A)

- ➤ 80 % of subtyped samples A(H1)pdm09
- ➤ Median age 61 years
 - >41 percent ≥65 years
- \triangleright Peak week 52 (n=36)

COVID-19

- ➤ Median age 71 years
 - >68 percent ≥65 years
- ➤ Peak week 49 (n=53)





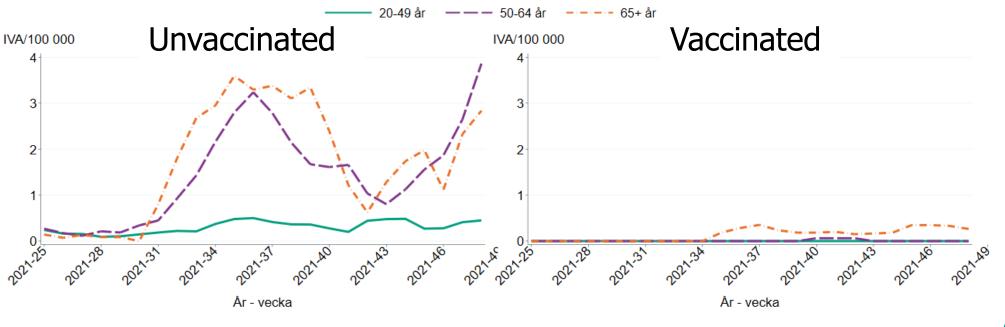
COVID-19 intensive care incidence by week and vaccination status, 2021

 Three-week moving average of ICU admissions with COVID-19 per week (weeks 25-47, 2021) Legend

green solid: 20-49 yrs

purple dashed: 50-64 yrs

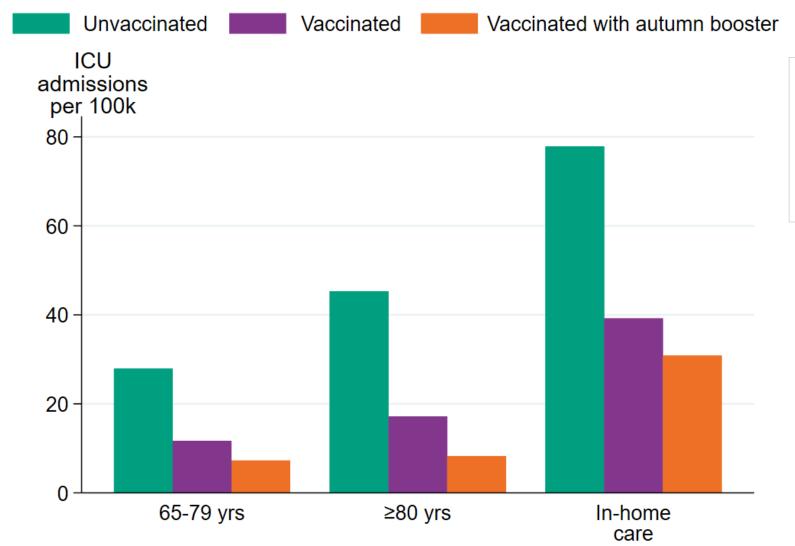
orange dot-dashed: ≥65 yrs





COVID-19 intensive care rates

week 47 2022 - week 2 2023



Legend (left to right)

green: unvaccinated

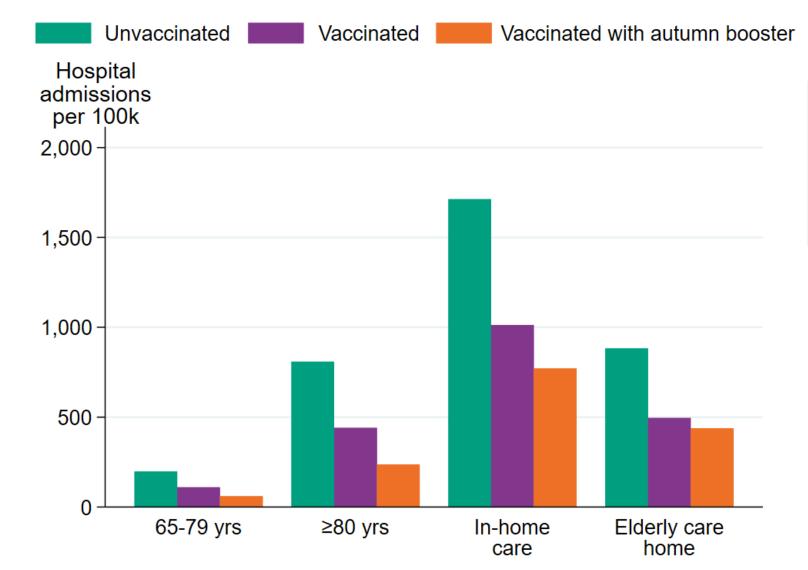
purple: vaccinated

orange: autumn booster



COVID-19 hospitalisation rates

week 47 2022 - week 2 2023



Legend (left to right)

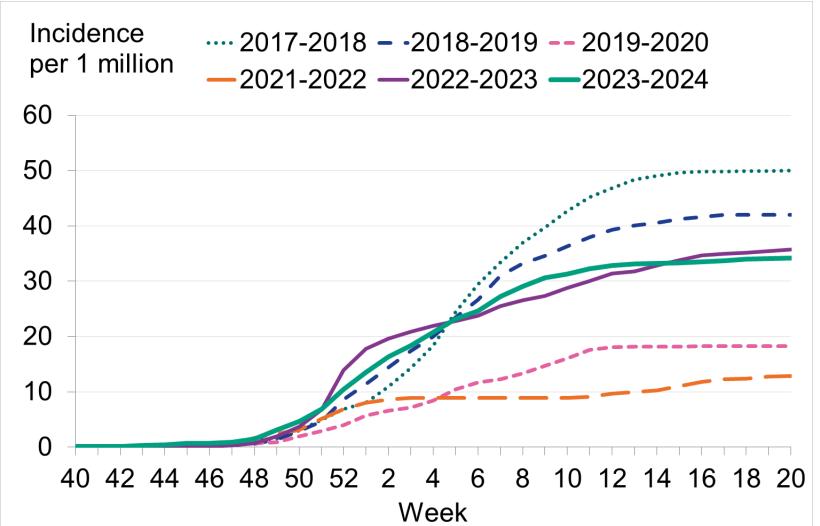
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Cumulative incidence of ICU admissions for influenza, per week and season



Dominant types

2017-18: B/Yam

2018-19: A/H1

2022-23: mixed season 2023-24: A/H1 and A/H3

2019-20: mixed season

2021-22: A/H3



Challenges and successes

Successes

- Continuous data availability, most reported within one week
- Only highest level of care
 - -Manual entry feasible
- National coverage (-ish)
- Flexible system, easy to add new pathogens
- In routine use

Challenges

- Manual data entry
 - Delays longer in lower priority periods
- Only highest level of care
 - Does not give earliest warning
- Not all patient samples are subtyped or sent for further virological analysis
- New pathogens require prioritisation
- Some data comes from patient recall (flu vaccination)

Folkhälsomvndigheten

Lessons learned and future plans

- Systems in routine use before a pandemic, work during a pandemic
- National vaccination registry important for evaluation of interventions
- Systems requiring manual data entry should focus on most prioritised pathogens, at a feasible level of care
- Integration of surveillance of COVID-19, influenza and RSV continues throughout all levels of disease severity
- Evaluation and further development, identification of gaps and overlaps
- Preparedness for the next pandemic





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