

Digital Participatory Surveillance: the ten-years experience of Influzenanet and what's next

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Workshop on Infectious Disease Surveillance
Institute of Social and Preventive Medicine (University of Bern)
November 25th, 2019

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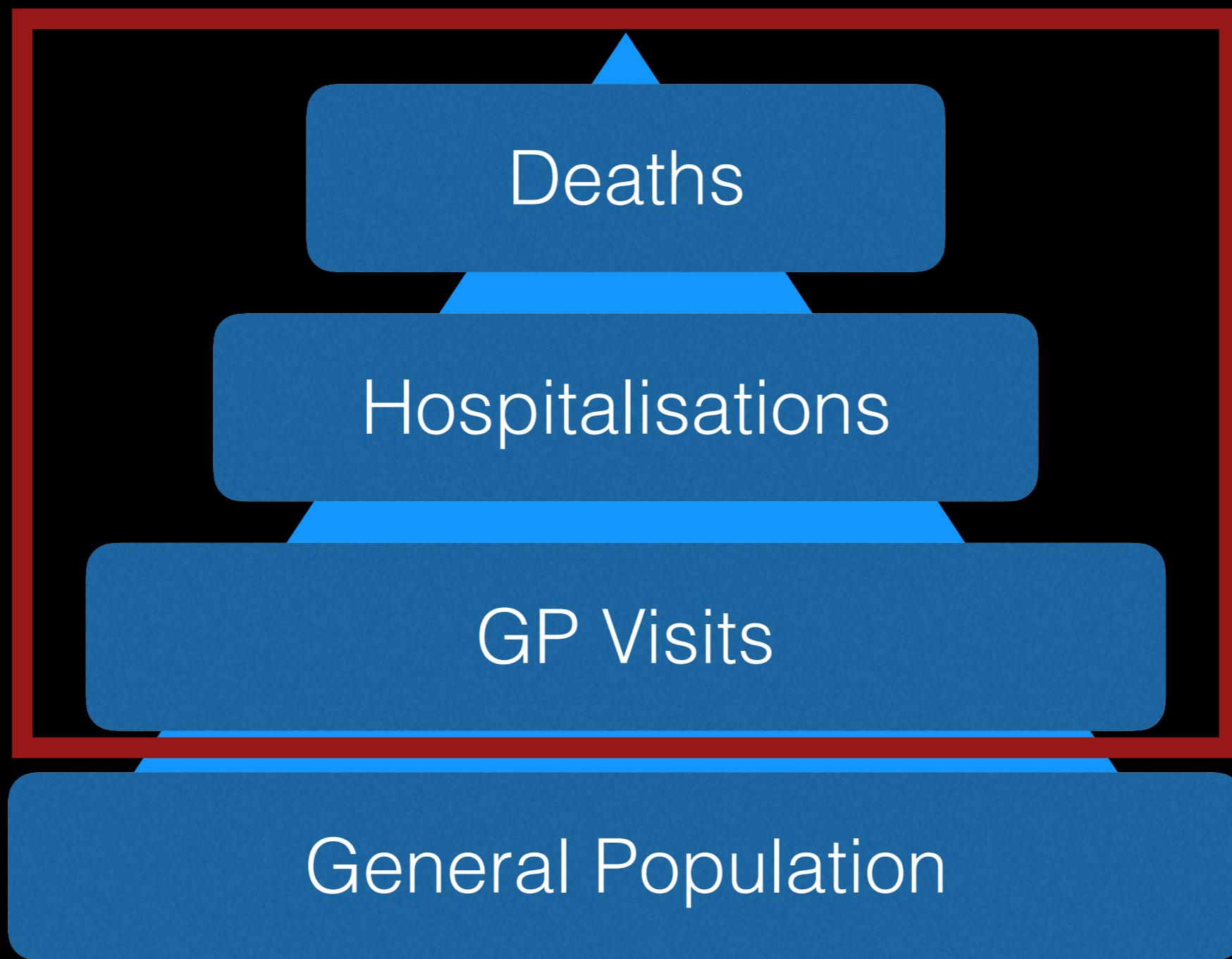
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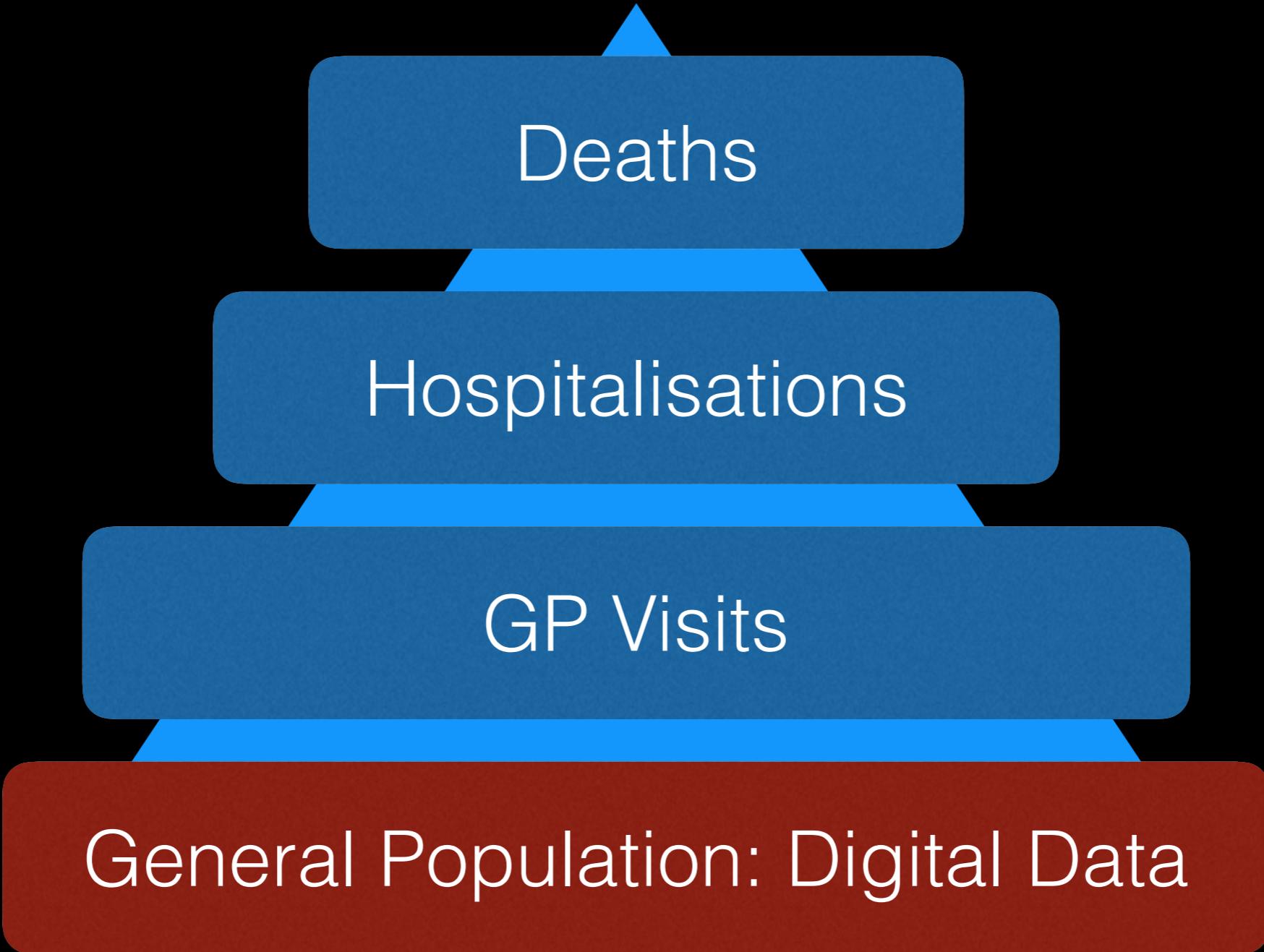
A photograph of a massive iceberg. Only a small portion of the iceberg is visible above the dark blue water, while the vast majority of it rests hidden beneath the surface in the bright blue sky.

Surveillance

Disease Burden

Surveillance





Deaths

Hospitalisations

GP Visits

General Population: Digital Data

[advanced search](#)

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RESEARCH ARTICLE

The Use of Twitter to Track Levels of Disease Activity and Public Concern in the U.S. during the Influenza A H1N1 Pandemic

Alessio Signorini, Alberto Maria Segre, Philip M. Polgreen

Published: May 4, 2011 • <http://dx.doi.org/10.1371/journal.pone.0019467>

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[Abstract](#)[Introduction](#)[Methods](#)[Results](#)[Discussion](#)[Acknowledgments](#)[Author Contributions](#)[References](#)[Reader Comments \(1\)](#)[Media Coverage \(0\)](#)[Figures](#)

Abstract

Twitter is a free social networking and micro-blogging service that enables its millions of users to send and read each other's "tweets," or short, 140-character messages. The service has more than 190 million registered users and processes about 55 million tweets per day. Useful information about news and geopolitical events lies embedded in the Twitter stream, which embodies, in the aggregate, Twitter users' perspectives and reactions to current events. By virtue of sheer volume, content embedded in the Twitter stream may be useful for tracking or even forecasting behavior if it can be extracted in an efficient manner. In this study, we examine the use of information embedded in the Twitter stream to (1) track rapidly-evolving public sentiment with respect to H1N1 or swine flu, and (2) track and measure actual disease activity. We also show that Twitter can be used as a measure of public interest or concern about health-related events. Our results show that estimates of influenza-like illness derived from Twitter chatter accurately track reported disease levels.

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?

Subject Areas

Twitter	
Influenza	
H1N1	
Swine influenza	
Infectious disease s...	
Public and occupati...	
Vaccines	

Wikipedia Usage Estimates Prevalence of Influenza-Like Illness in the United States in Near Real-Time

David J. McIver*, John S. Brownstein

Boston Children's Hospital, Harvard Medical School, Boston, Massachusetts, United States of America

Abstract

Circulating levels of both seasonal and pandemic influenza require constant surveillance to ensure the health and safety of the population. While up-to-date information is critical, traditional surveillance systems can have data availability lags of up to two weeks. We introduce a novel method of estimating, in near-real time, the level of influenza-like illness (ILI) in the United States (US) by monitoring the rate of particular Wikipedia article views on a daily basis. We calculated the number of times certain influenza- or health-related Wikipedia articles were accessed each day between December 2007 and August 2013 and compared these data to official ILI activity levels provided by the Centers for Disease Control and Prevention (CDC). We developed a Poisson model that accurately estimates the level of ILI activity in the American population, up to two weeks ahead of the CDC, with an absolute average difference between the two estimates of just 0.27% over 294 weeks of data. Wikipedia-derived ILI models performed well through both abnormally high media coverage events (such as during the 2009 H1N1 pandemic) as well as unusually severe influenza seasons (such as the 2012–2013 influenza season). Wikipedia usage accurately estimated the week of peak ILI activity 17% more often than Google Flu Trends data and was often more accurate in its measure of ILI intensity. With further study, this method could potentially be implemented for continuous monitoring of ILI activity in the US and to provide support for traditional influenza surveillance tools.

Citation: McIver DJ, Brownstein JS (2014) Wikipedia Usage Estimates Prevalence of Influenza-Like Illness in the United States in Near Real-Time. PLoS Comput Biol 10(4): e1003581. doi:10.1371/journal.pcbi.1003581

Editor: Marcel Salathé, Pennsylvania State University, United States of America

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Competing Interests: The authors have declared that no competing interests exist.

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advanced search

OPEN ACCESS PEER-REVIEWED

RESEARCH ARTICLE

Monitoring Influenza Epidemics in China with Search Query from Baidu

Qingyu Yuan , Elaine O. Nsoesie , Benfu Lv, Geng Peng, Rumi Chunara, John S. Brownstein

Published: May 30, 2013 • <http://dx.doi.org/10.1371/journal.pone.0064323>

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Abstract

Introduction

Methods

Results

Discussion

Supporting Information

Author Contributions

References

Reader Comments (2)

Media Coverage (0)

Figures

Abstract

Several approaches have been proposed for near real-time detection and prediction of the spread of influenza. These include search query data for influenza-related terms, which has been explored as a tool for augmenting traditional surveillance methods. In this paper, we present a method that uses Internet search query data from Baidu to model and monitor influenza activity in China. The objectives of the study are to present a comprehensive technique for: (i) keyword selection, (ii) keyword filtering, (iii) index composition and (iv) modeling and detection of influenza activity in China. Sequential time-series for the selected composite keyword index is significantly correlated with Chinese influenza case data. In addition, one-month ahead prediction of influenza cases for the first eight months of 2012 has a mean absolute percent error less than 11%. To our knowledge, this is the first study on the use of search query data from Baidu in conjunction with this approach for estimation of influenza activity in China.



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Subject Areas

Influenza



Swine influenza



Influenza viruses



H1N1



China



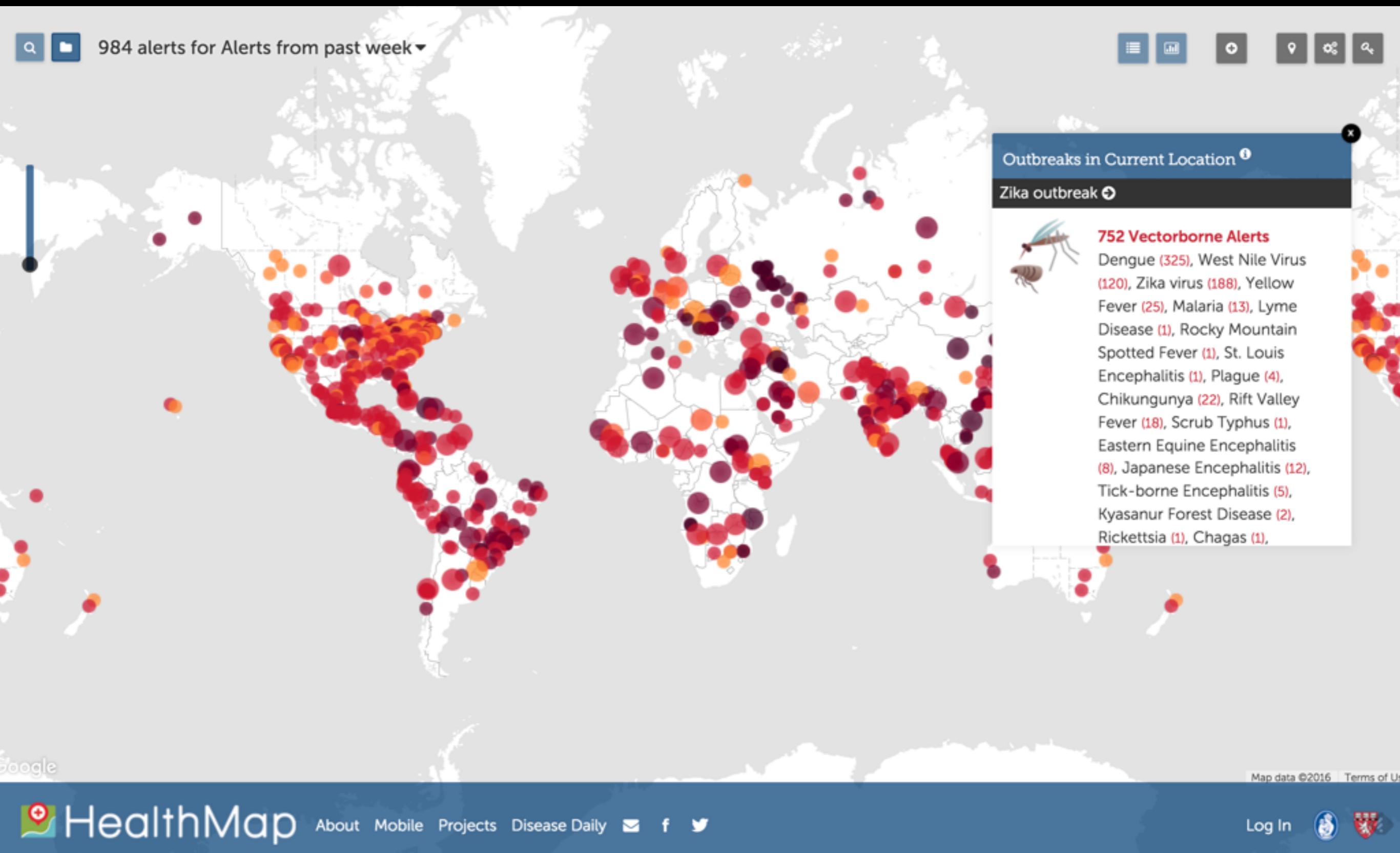
Influenza A virus



Infectious disease s...



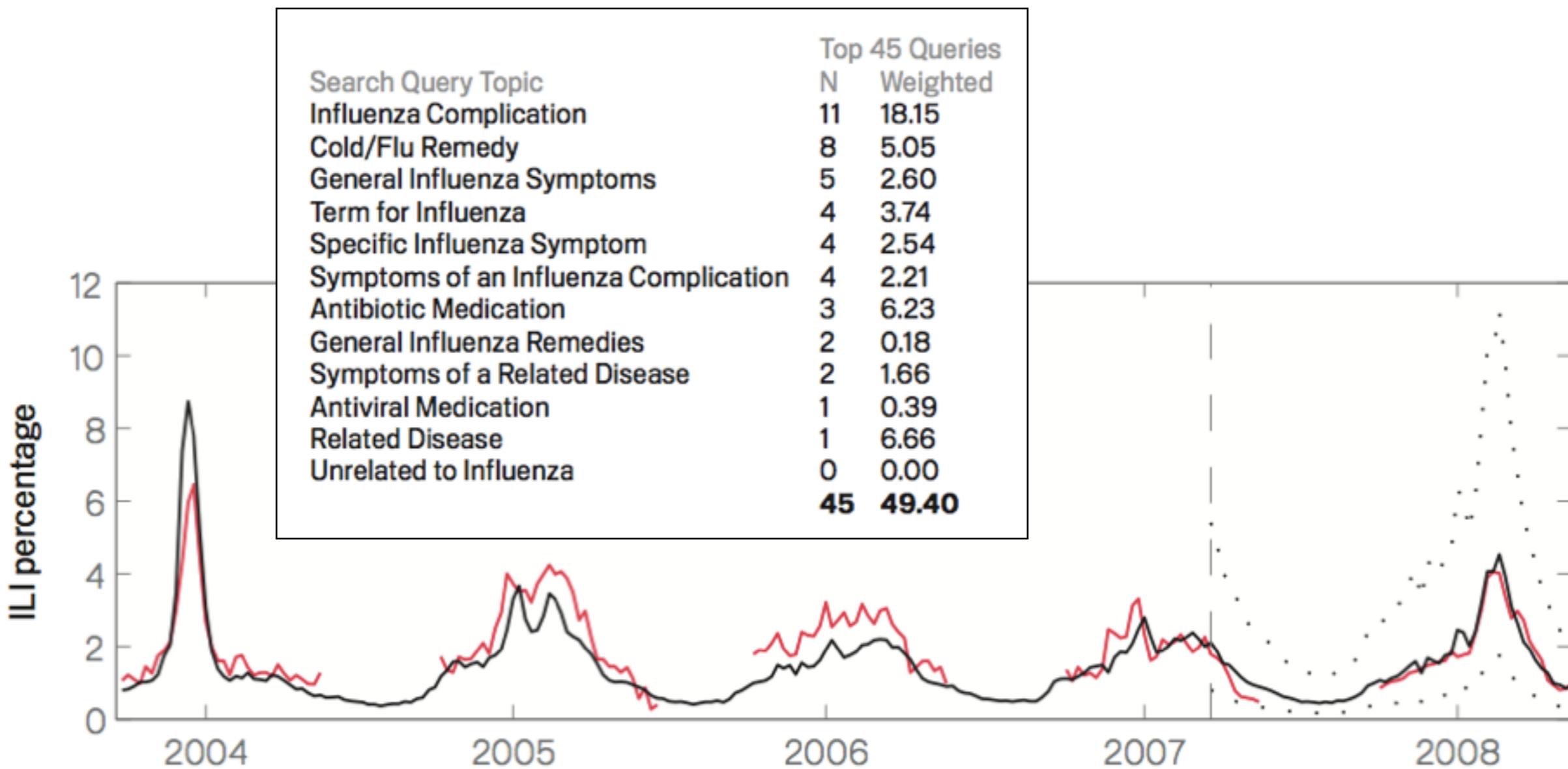
healthmap.org



Detecting influenza epidemics using search engine query data

Jeremy Ginsberg¹, Matthew H. Mohebbi¹, Rajan S. Patel¹, Lynnette Brammer²,
Mark S. Smolinski¹ & Larry Brilliant¹

¹Google Inc. ²Centers for Disease Control and Prevention



NATURE | NEWS

عربي

When Google got flu wrong

US outbreak foxes a leading web-based method for tracking seasonal flu.

Declan Butler

Science 14 March 2014:
 Vol. 343 no. 6176 pp. 1203–1205
 DOI: 10.1126/science.1248506

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POLICY FORUM

BIG DATA

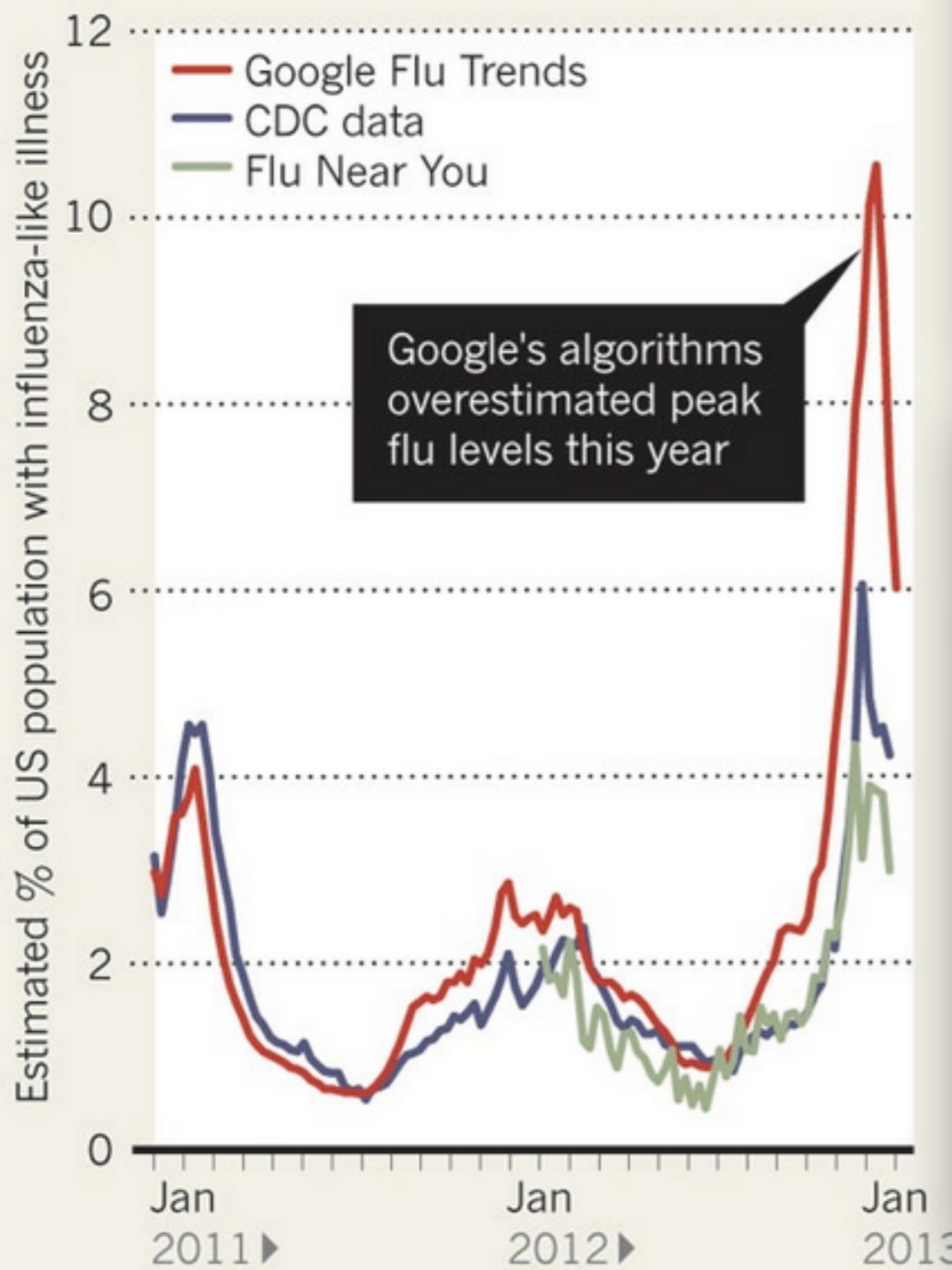
The Parable of Google Flu: Traps in Big Data Analysis

David Lazer^{1,2,*}, Ryan Kennedy^{1,3,4}, Gary King³, Alessandro Vespignani^{5,6,3}[+ Author Affiliations](#)[✉ Corresponding author. E-mail: d.lazer@neu.edu.](#)

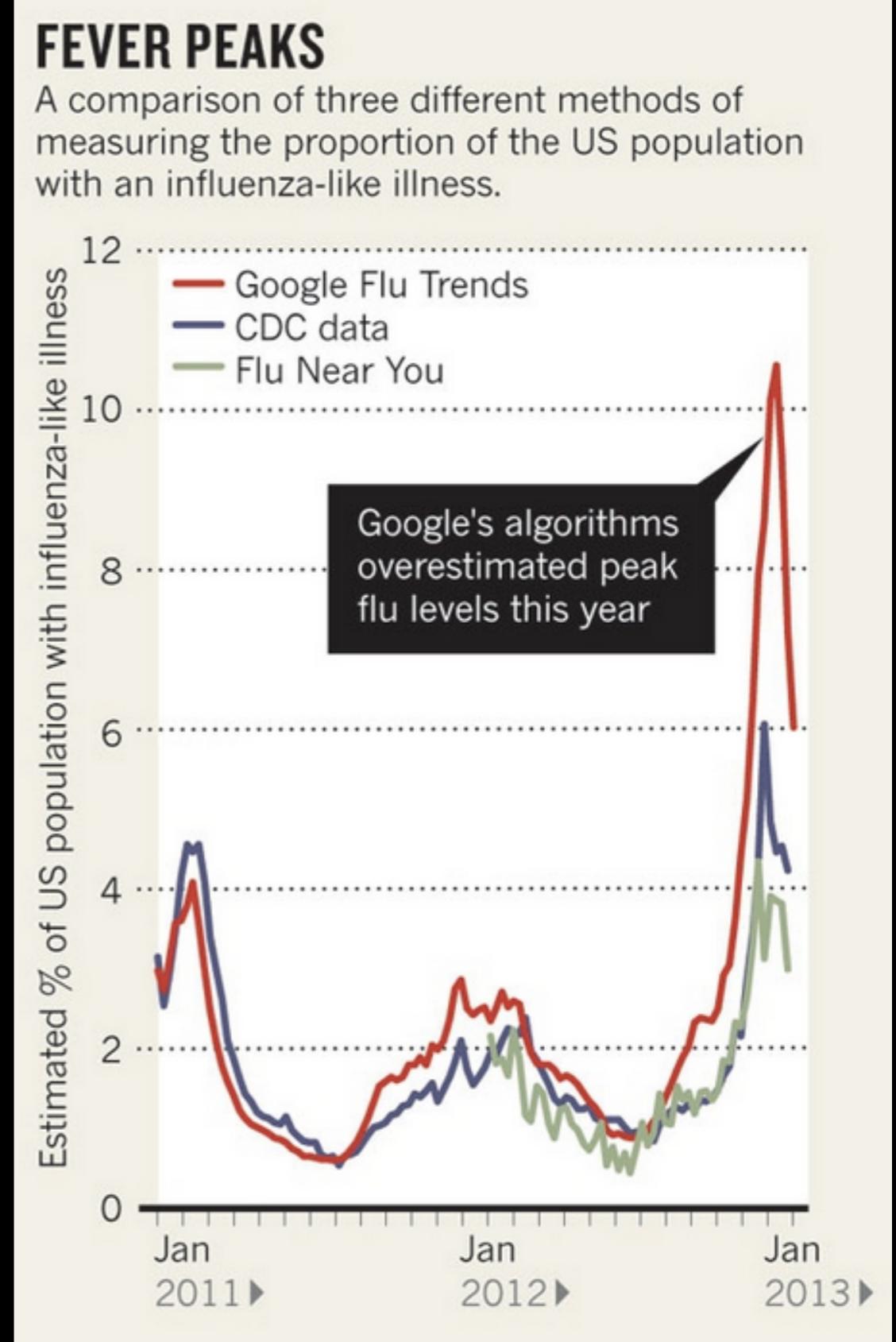
In February 2013, Google Flu Trends (GFT) made headlines but not for a reason that Google executives or the creators of the flu tracking system would have hoped. *Nature* reported GFT predicting more than double the proportion of doctor visits for influenza-like illness (ILI) compared with the US Centers for Disease Control and Prevention (CDC), which bases its estimates on surveillance from laboratories across the United States (1, 2). This happened despite the fact that GFT often predicted CDC reports. Given that GFT is often held up as an exemplary use of big data (3), what lessons can we draw from this error?

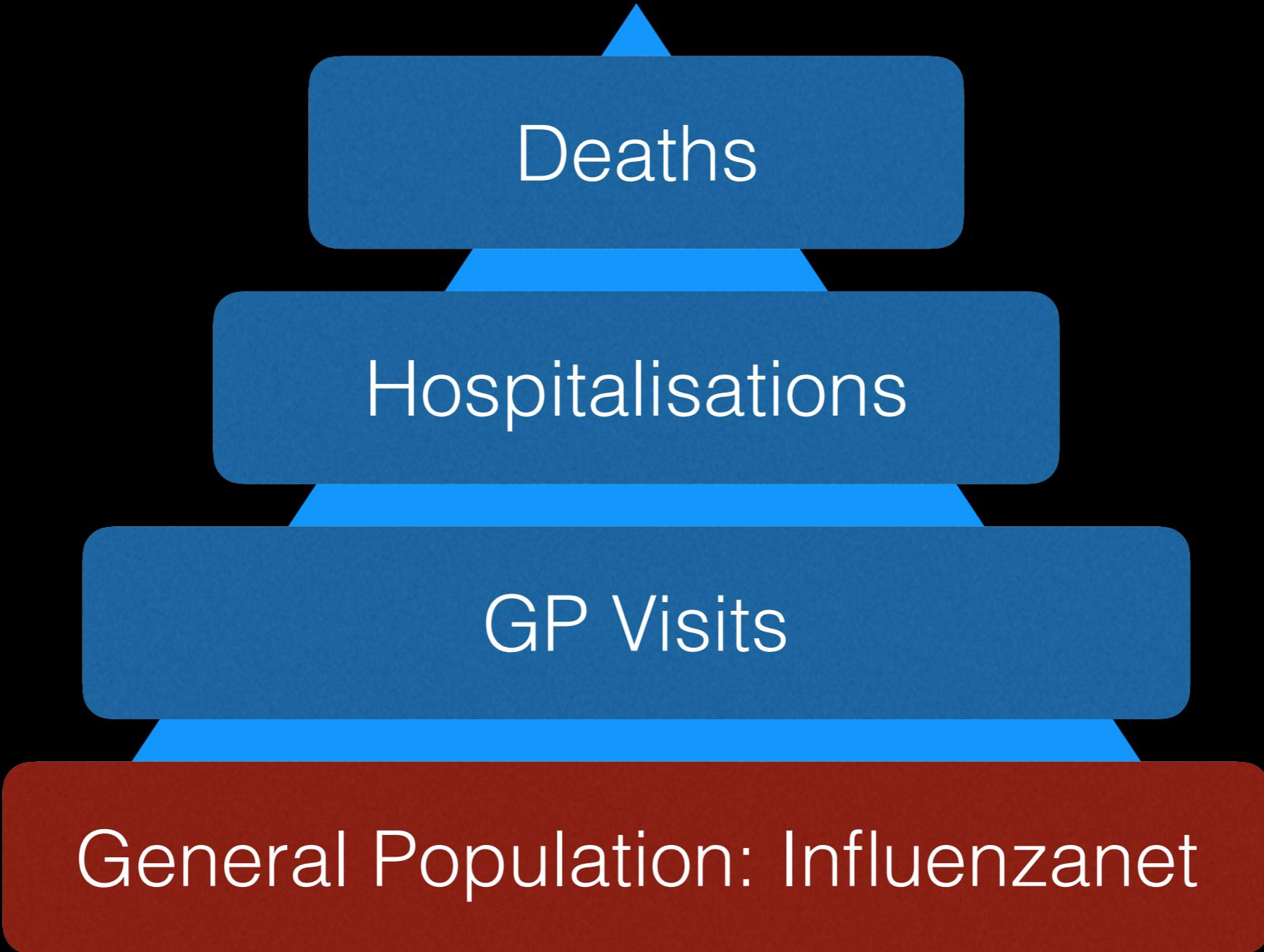
FEVER PEAKS

A comparison of three different methods of measuring the proportion of the US population with an influenza-like illness.



- Passive data sources don't describe who is well
- Low specificity





Deaths

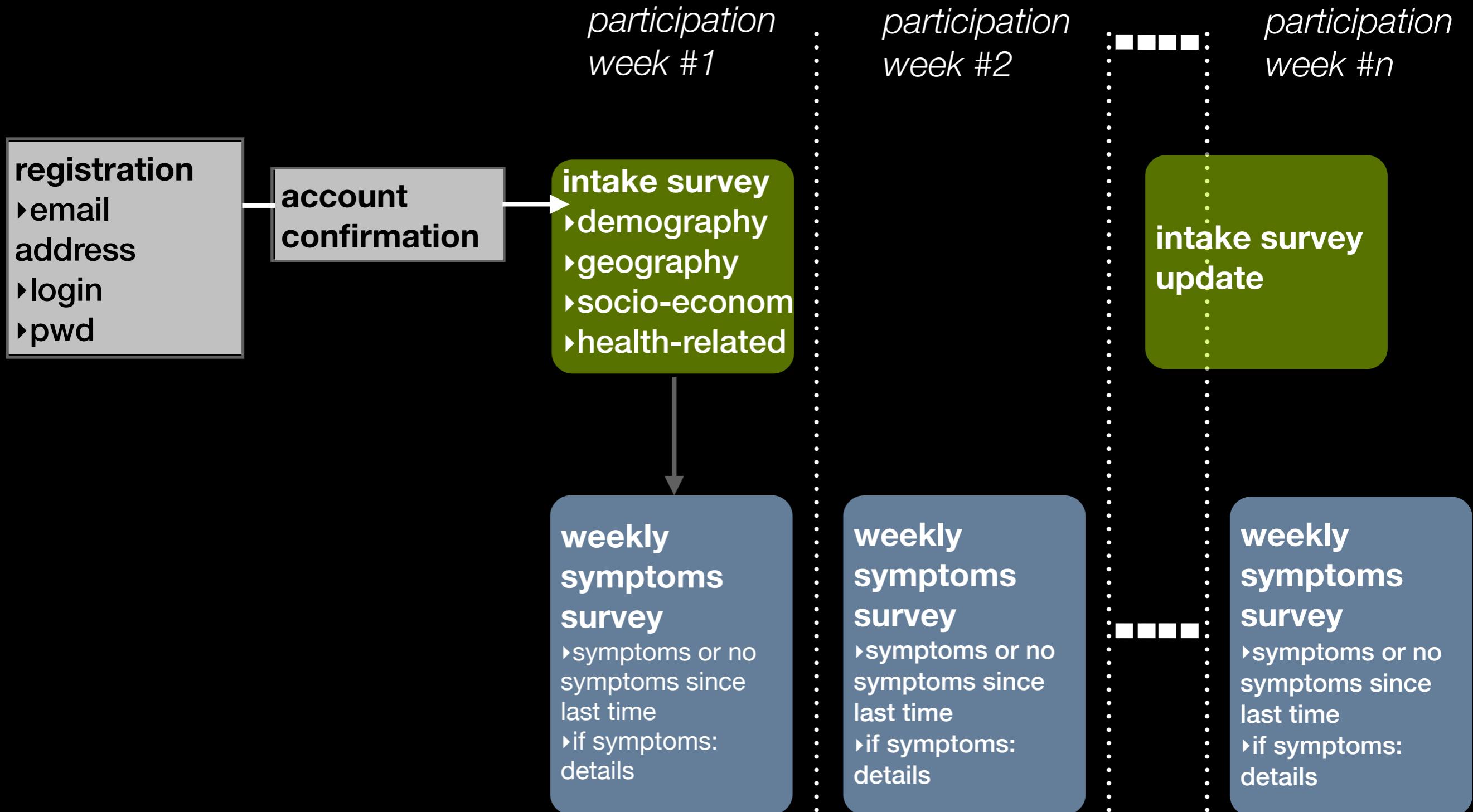
Hospitalisations

GP Visits

General Population: Influenzanet

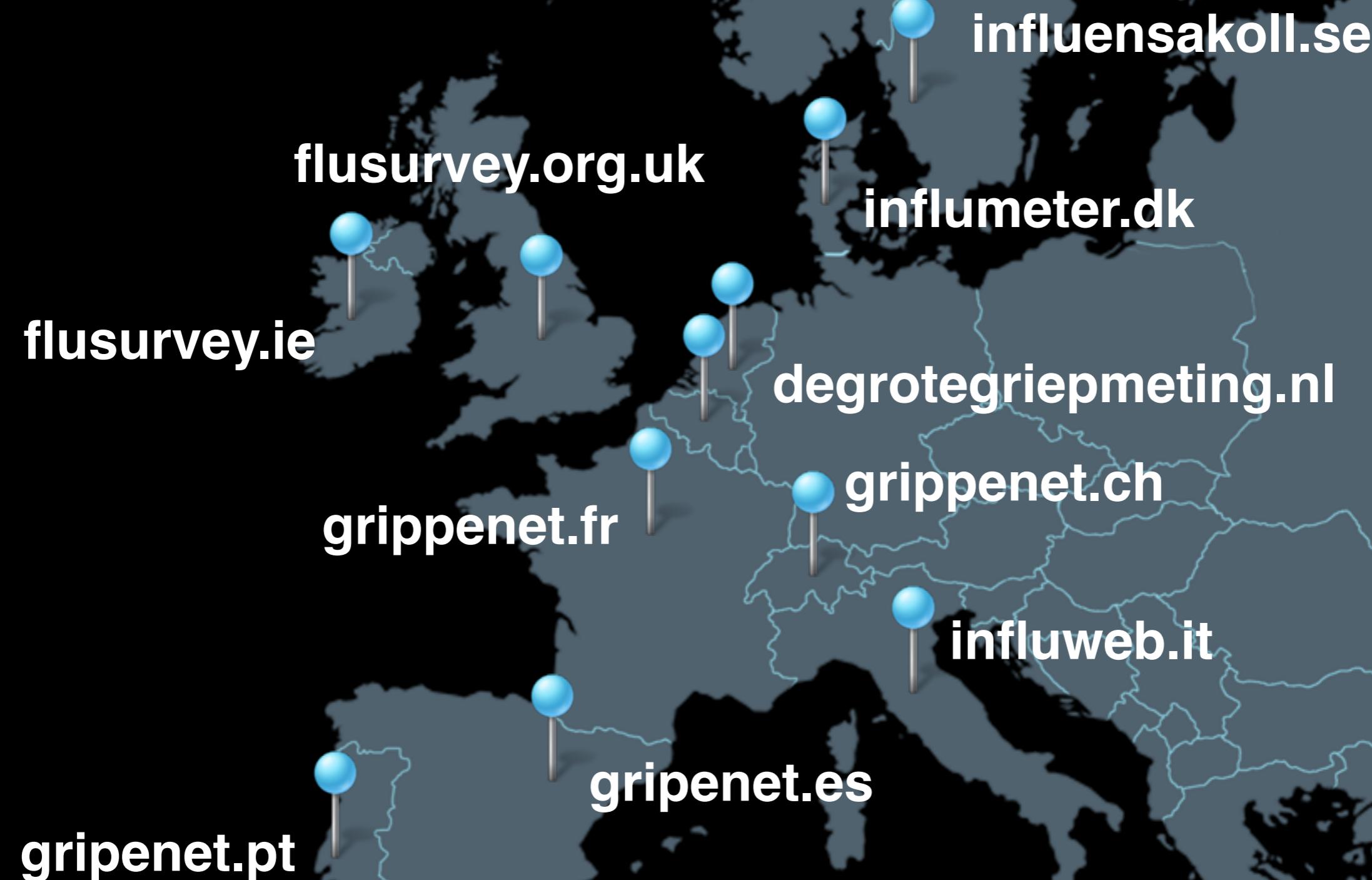


INFLUENZANET study design

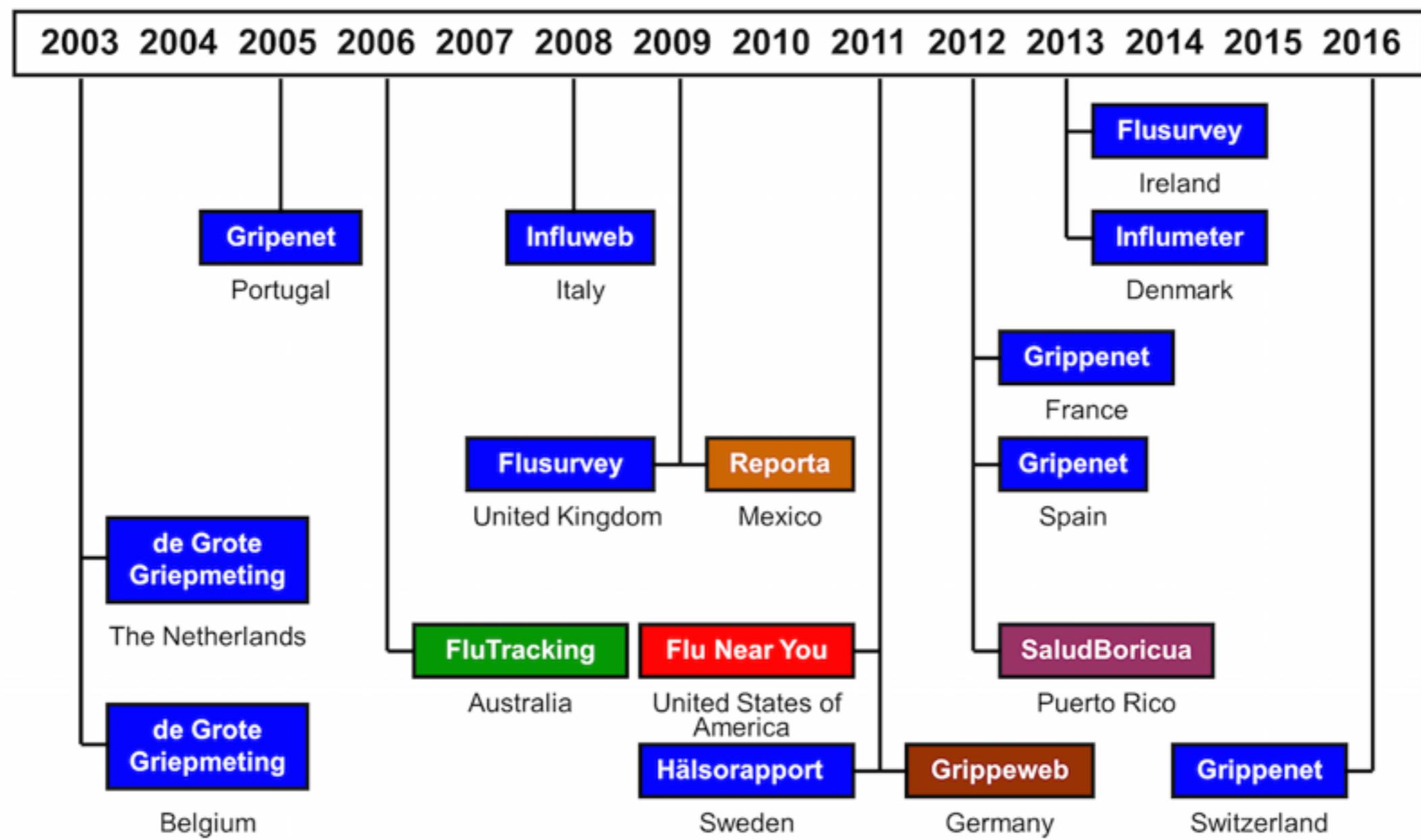


uniform data collection across countries

INFLUENZANET.INFO



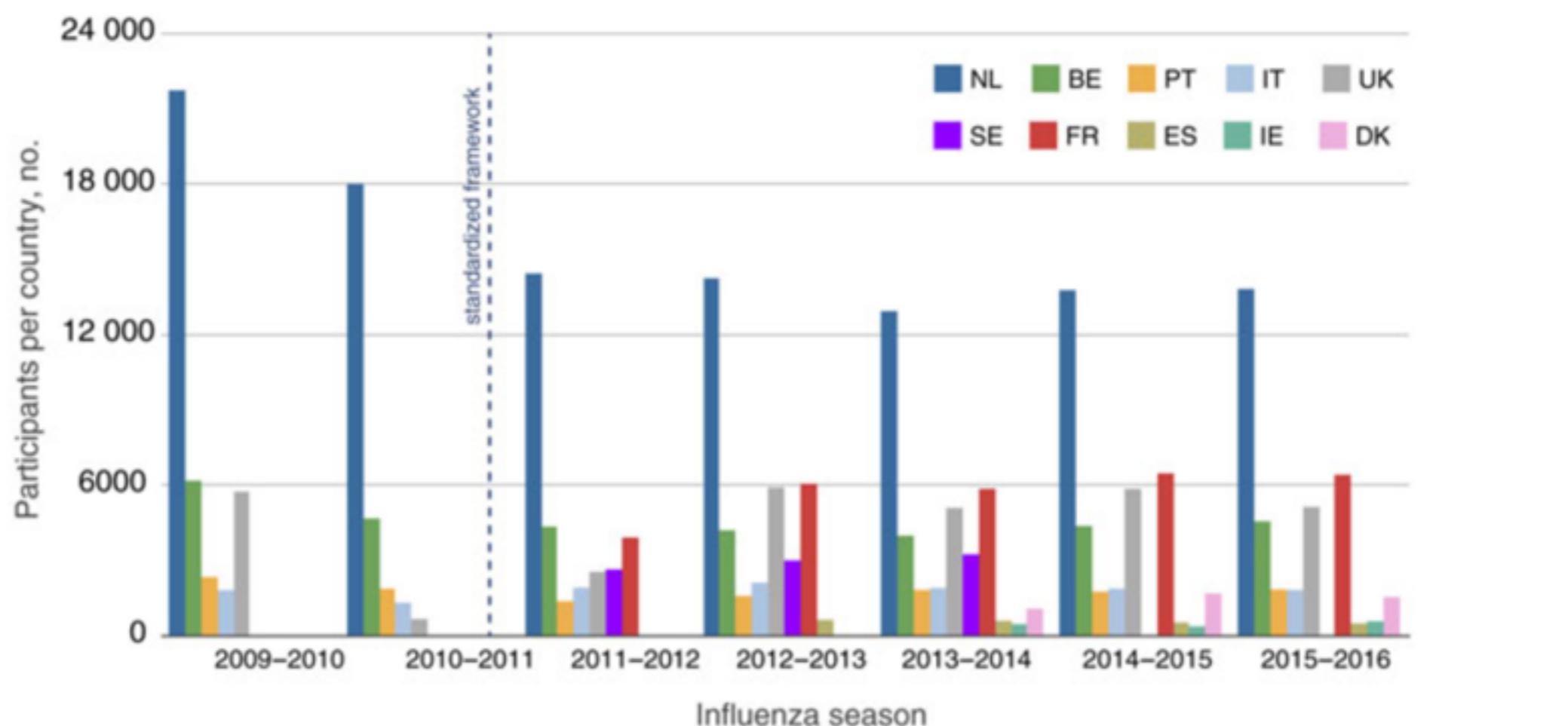
INFLUENZANET: a timeline



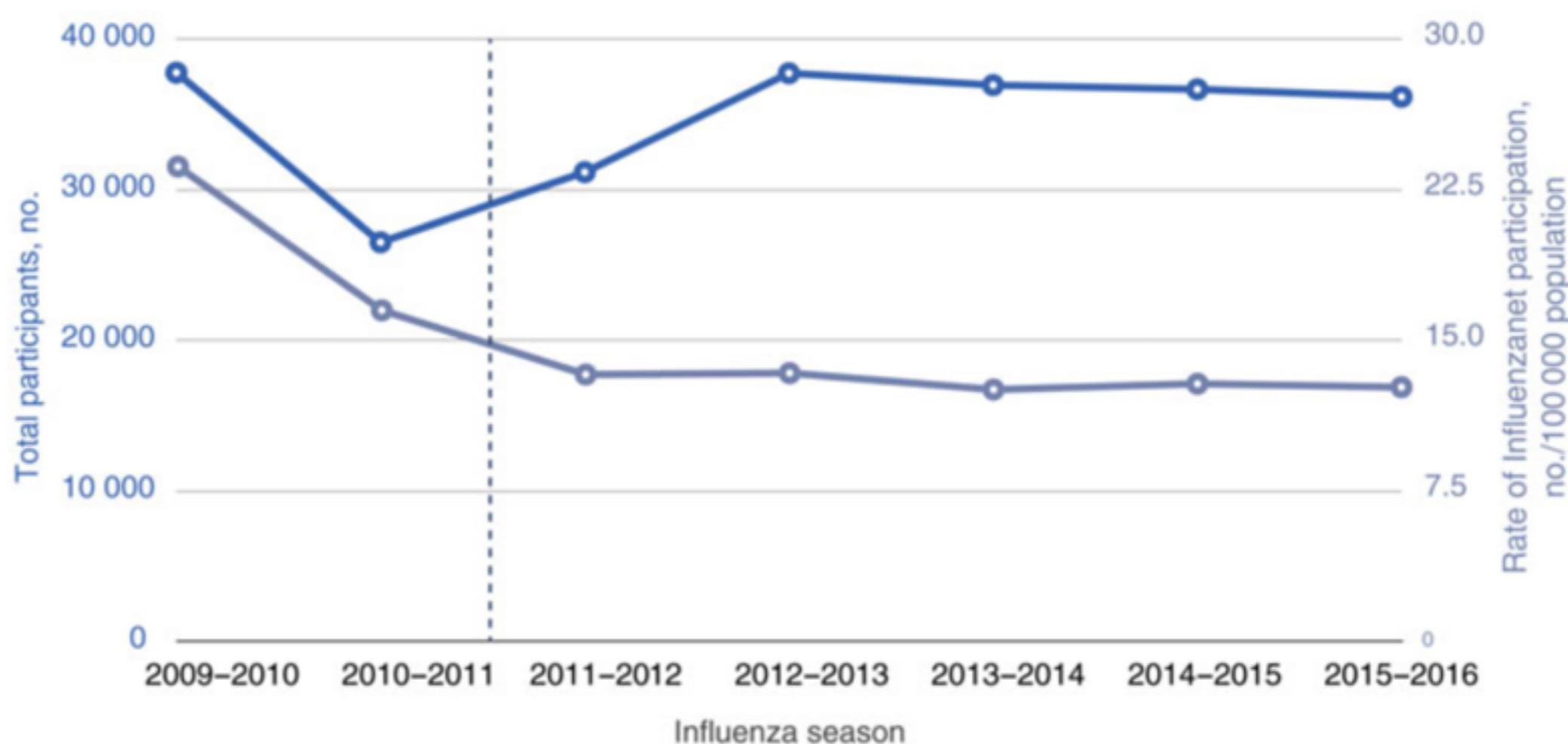
Koppeschaar CE, Colizza V, Guerrisi C, Turbelin C, Duggan J, Edmunds WJ, Kjelsø C, Mexia R, Moreno Y, Meloni S, Paolotti D, Perrotta D, van Straten E, Franco AO
Influenzanet: Citizens Among 10 Countries Collaborating to Monitor Influenza in Europe
JMIR Public Health Surveill 2017;3(3):e66

study design: key questions

- weekly individual symptoms (or lack thereof)
- vaccine coverage
- healthcare seeking
- drugs uptake (including antibiotics)
- perceived severity
- duration of the symptoms
- absenteeism



C. Guerrisi et al,
*Participatory
 Syndromic
 Surveillance of
 Influenza in Europe*
 Journal of Infectious
 Diseases (2016) 214
 (suppl 4): S386-
 S392.



INFLUENZANET.INFO

influweb.it - ISI Foundation & Istituto Superiore di Sanità, Italy

grippenet.fr - INSERM, France

gripenet.pt - INSA, Portugal

gripenet.es - University of Saragoza, Spain

flusurvey.net - Public Health England

influensakoll.se - Public Health Agency of Sweden

influmeter.dk - Staten Serum Institute, Denmark

fr.grippenet.ch, de.grippenet.ch - Global Health Institute, Geneva

flusurvey.ie - University of Galway

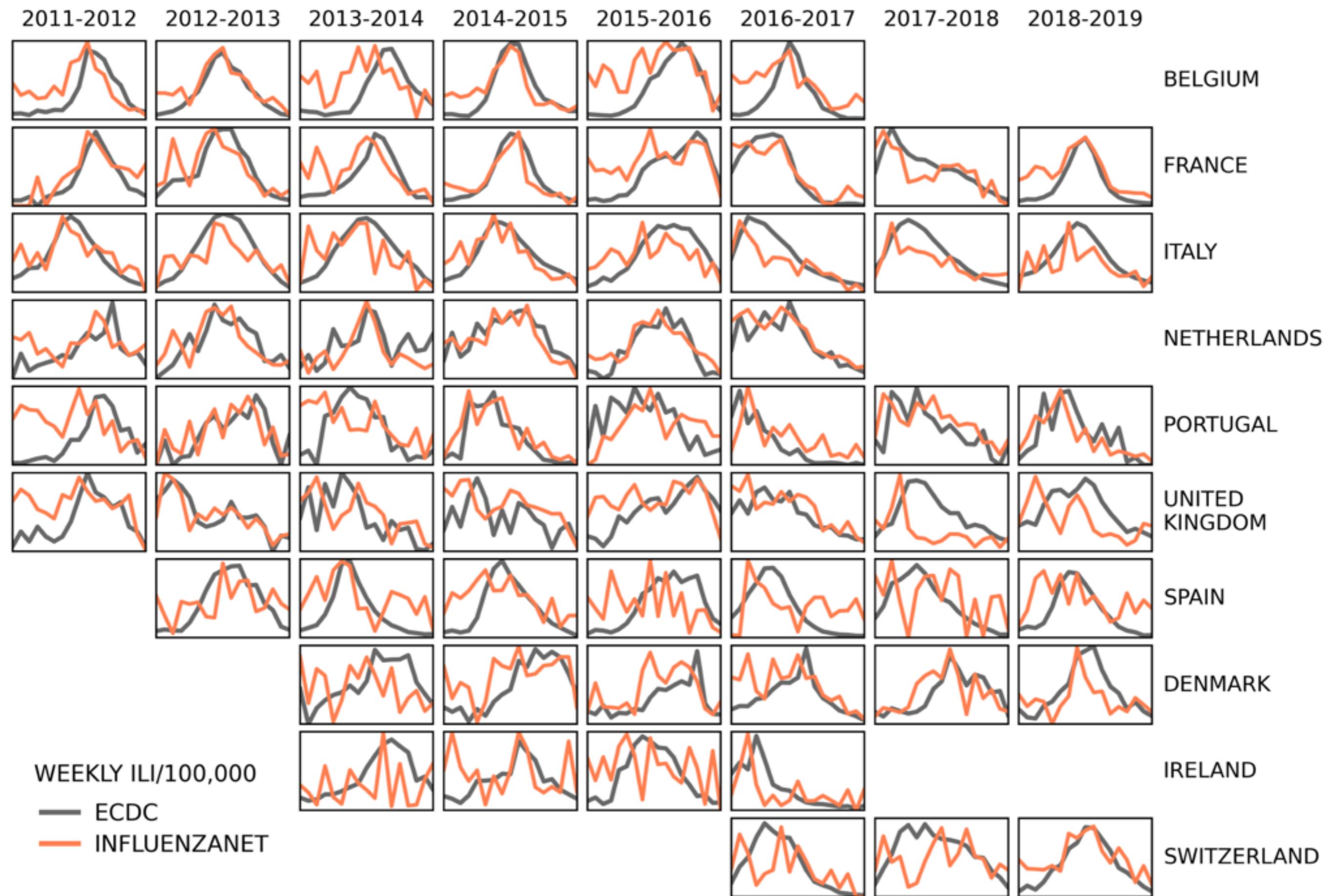
grippeweb.rki.de - Robert Koch Institute, Germany

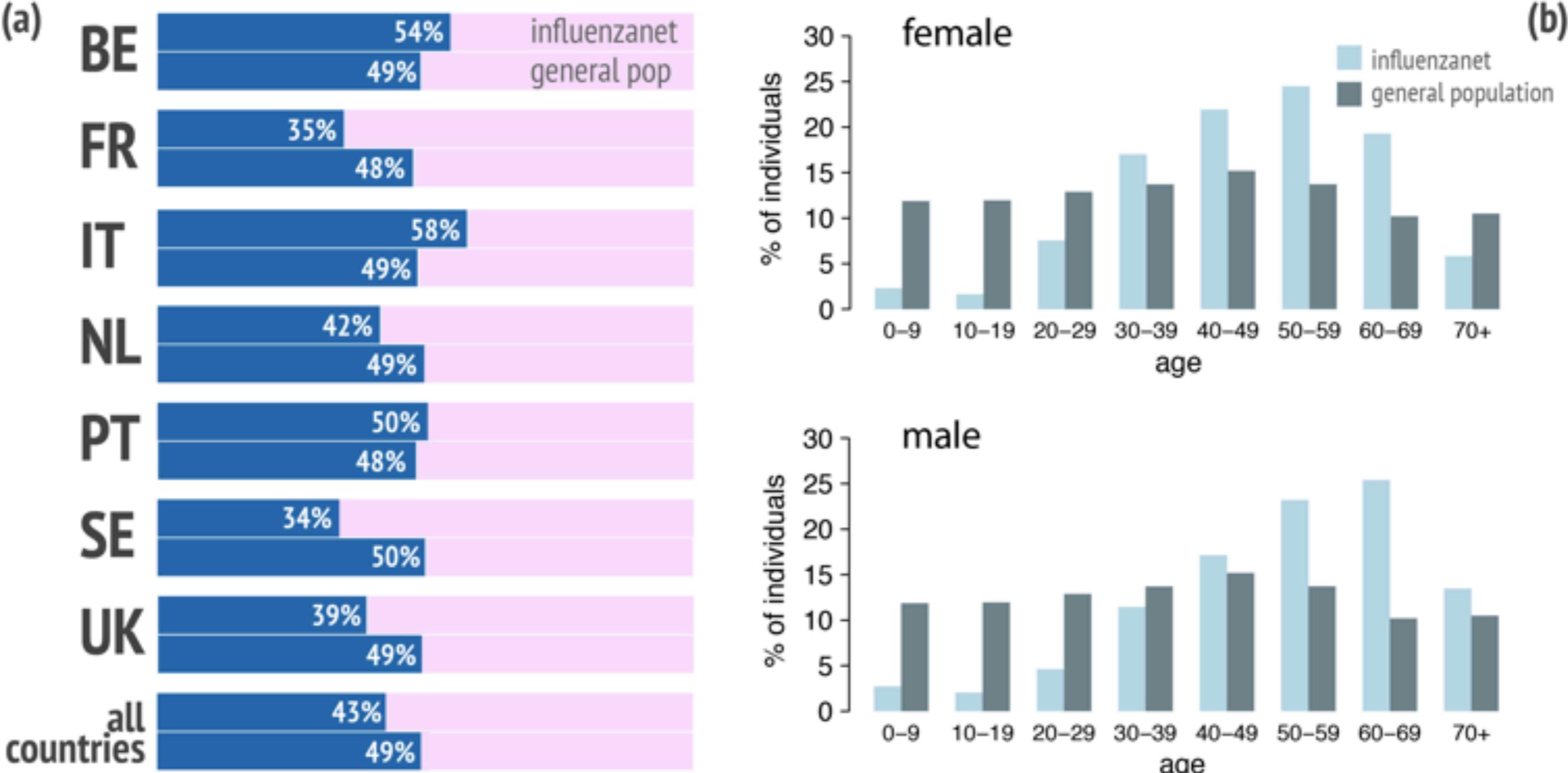


www.influensakoll.se
Kartlägger förekomst och spridning av influensa i Sverige



GrippeWeb





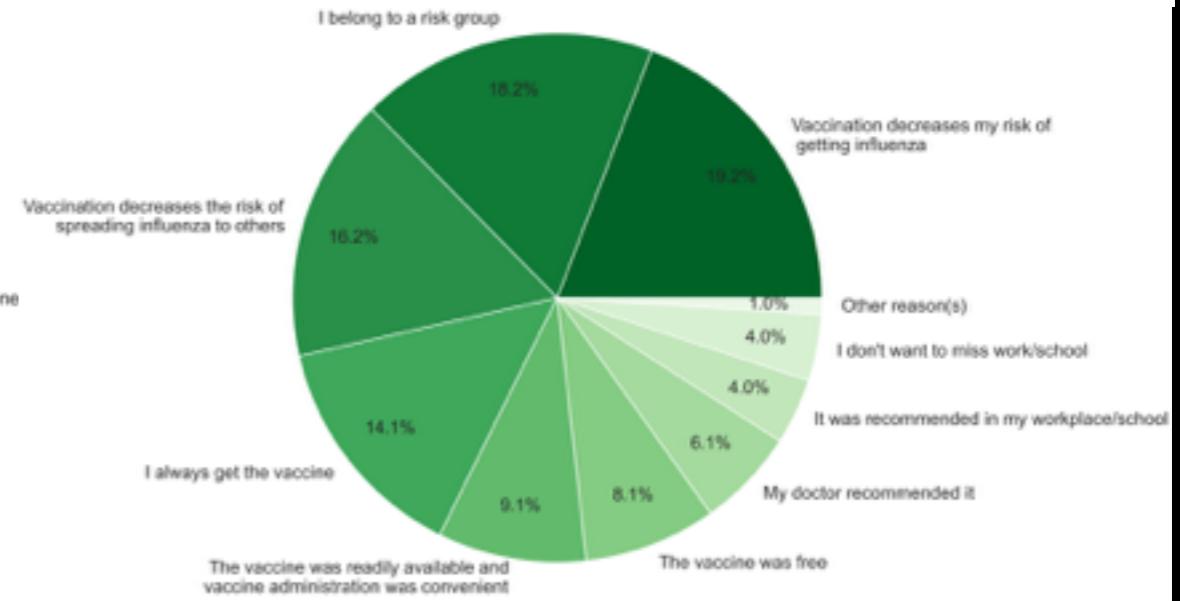
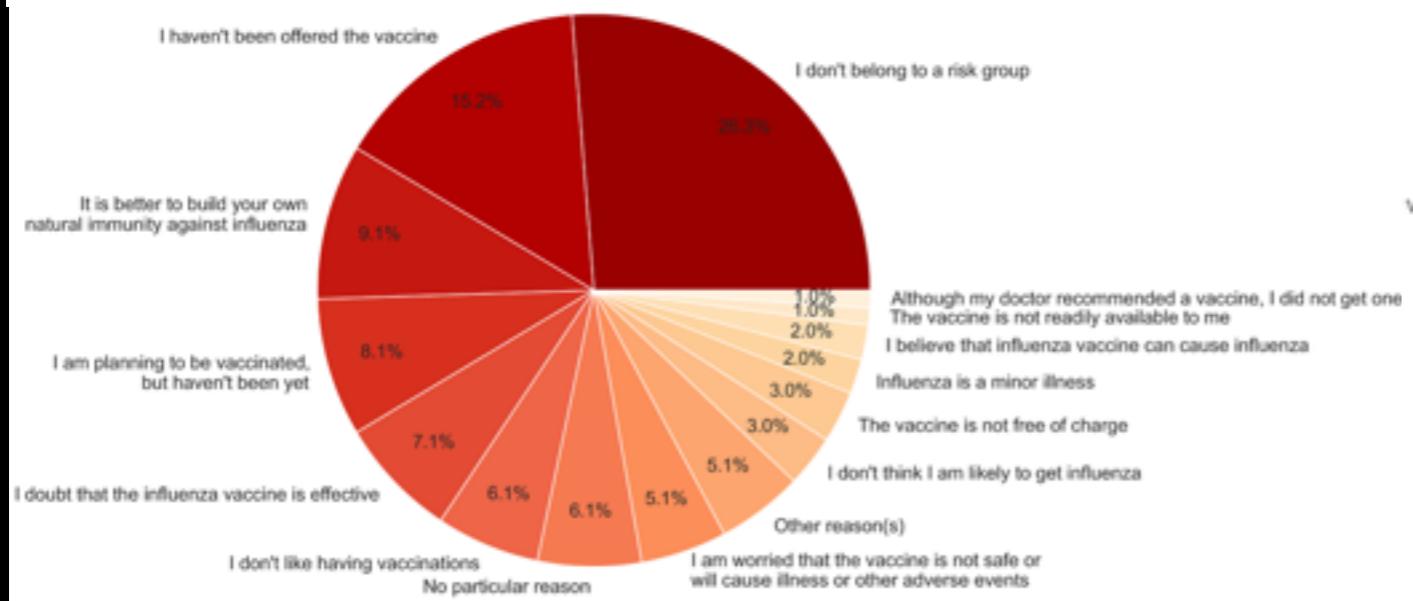
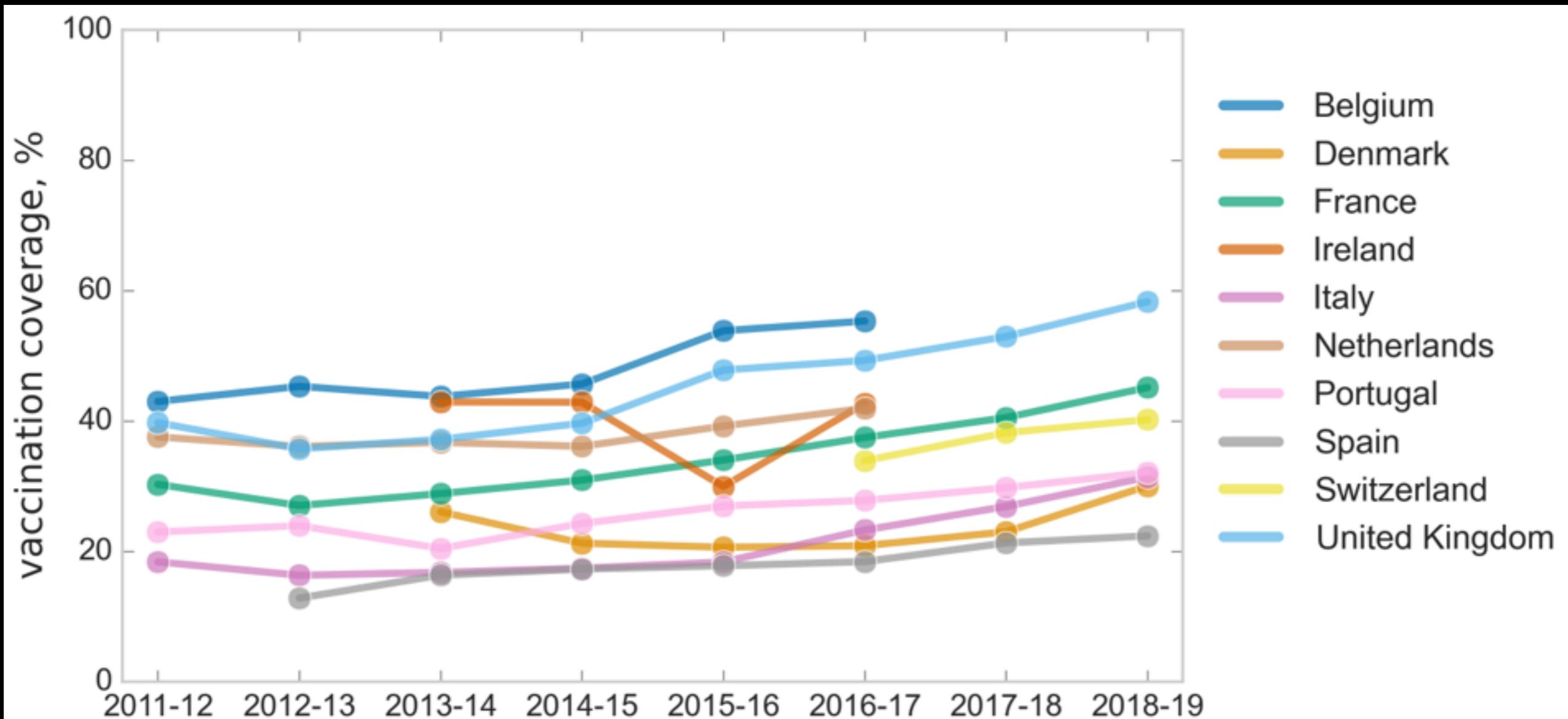
AGE / GENDER

P. Cantarelli et al.,

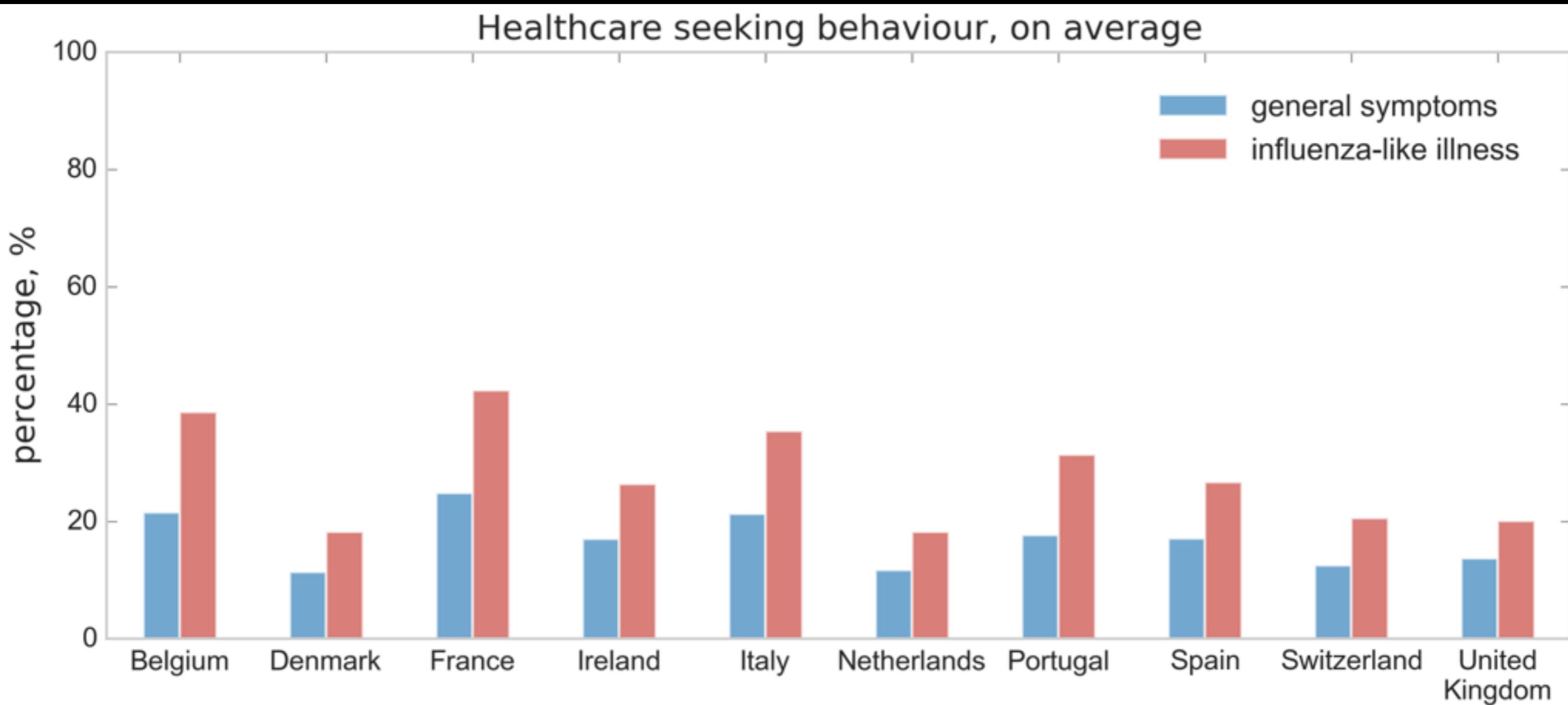
The representativeness of a European multi-center network for influenza-like-illness participatory surveillance

BMC Public Health 2014, 14:984

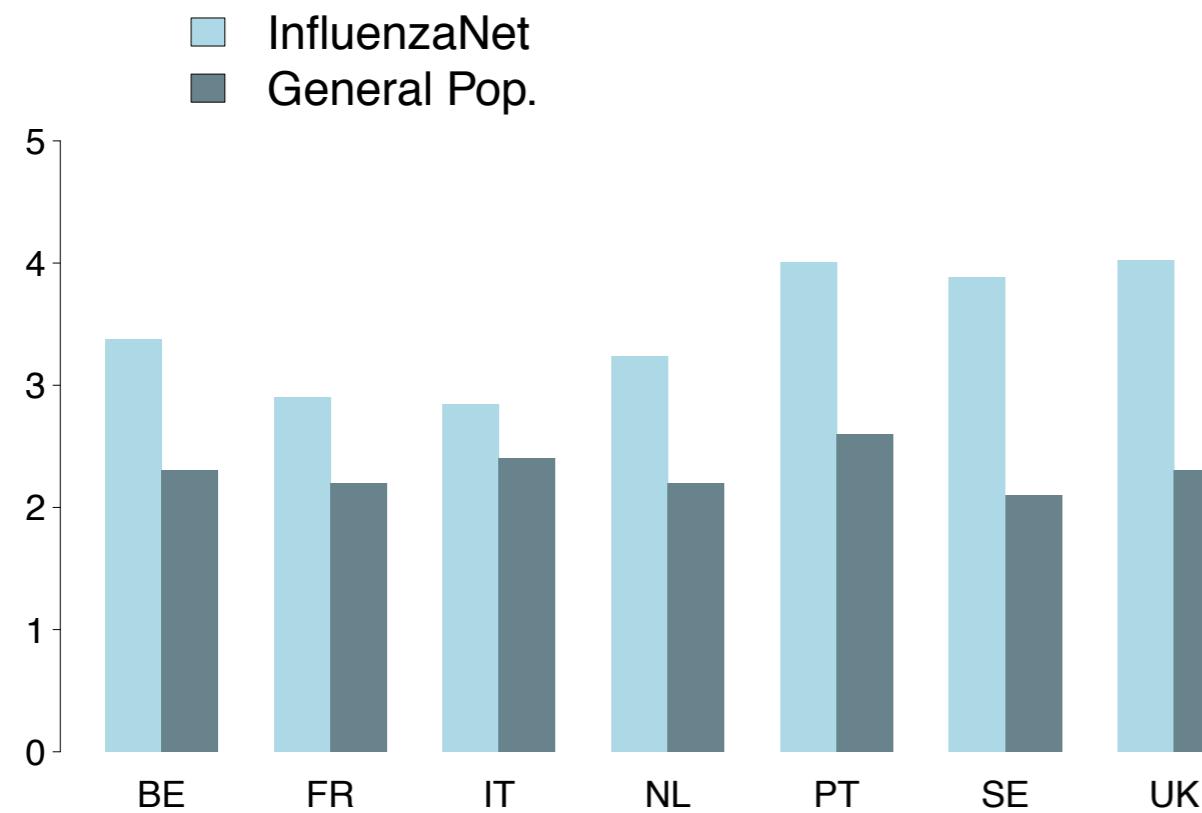
VACCINATION COVERAGE



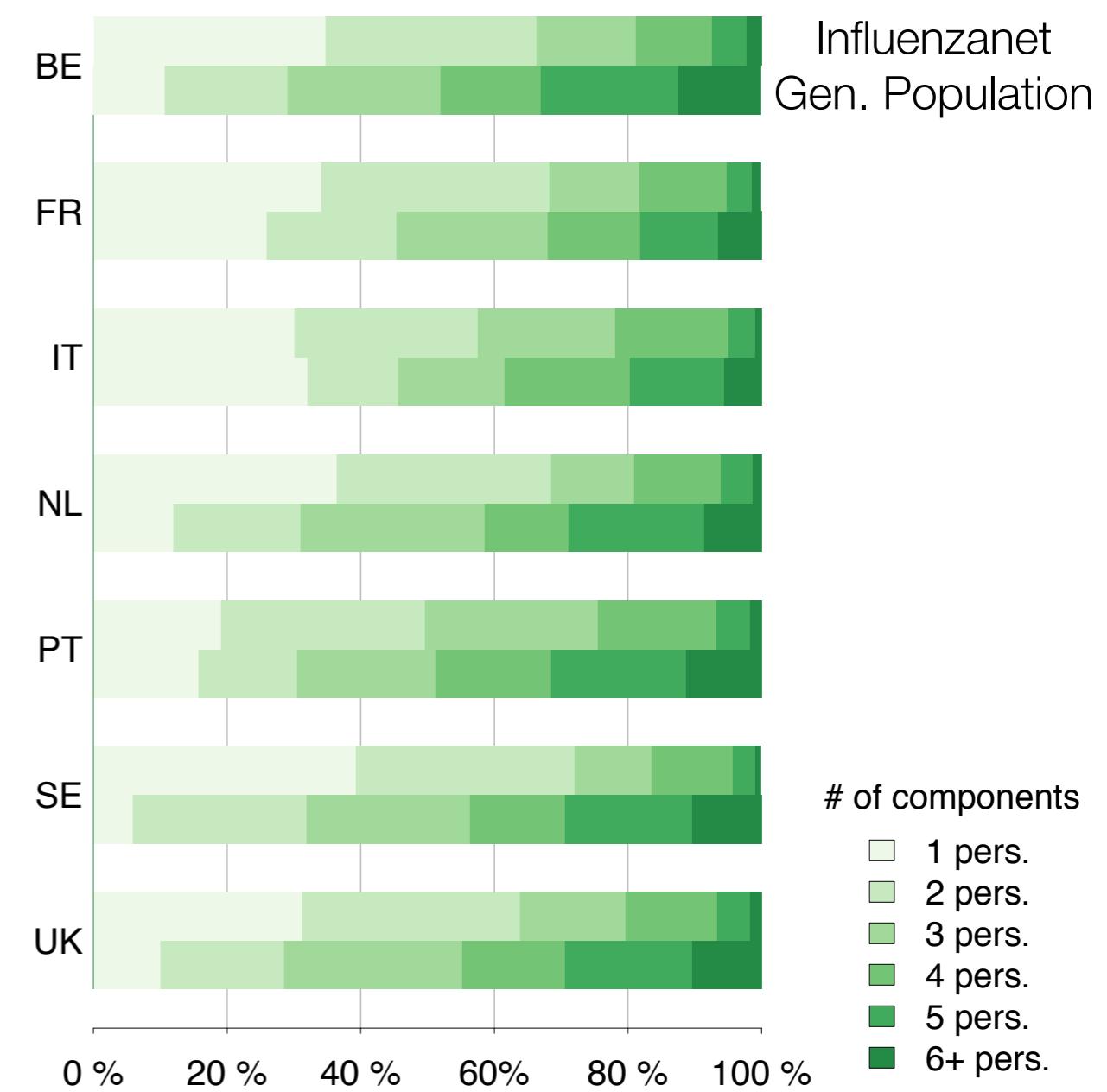
HEALTHCARE SEEKING



Average size of families



Distribution of Family Composition



SOCIO-ECONOMIC INDICATORS

P. Cantarelli et al.,

The representativeness of a European multi-center network for influenza-like-illness participatory surveillance

BMC Public Health 2014, 14:984

	OR	CI	P
Gender	1.52	1.33, 1.73	0.03
Groups of children	1.47	1.22, 1.77	0.03
Unvaccinated	1.94	1.66, 2.27	<0.001
18-24	0.95	0.64, 1.42	0.80
25-34	1.12	0.81, 1.55	0.50
35-44	1.42	1.04, 1.94	0.03
45-64	1.51	1.12, 2.03	<0.01
65+	0.95	0.65, 1.38	0.79
Living w child	1.10	0.95, 1.28	0.31
Smoking	1.32	1.07, 1.64	0.01
Risk factor	1.53	1.27, 1.83	<0.001
Public transport	0.91	0.82, 1.15	0.73

RISK FACTORS FOR ILI (UK)

Adler A et al.,
Incidence and risk factors for influenza-like-illness in the UK: online surveillance using Flusurvey
 BMC Infect Dis. 2014, 14:232

Countries	NL, BE, PT, IT ^a	UK ^b	FR ^b
Study period	2003–2013	2014–2015	2014–2015
Study participants per season, no., mean	24 666 ^c	2629	4475
Female sex	1.22 (1.17–1.28)	1.25 (1.14–1.36)	1.12 (1.02–1.23)
Vaccinated	0.80 (.71–.91) ^d	S ^e	0.87 (.78–.97)
Age, y			
NL, BE, PT, IT			
<18	1.59 (1.46–1.74)
18–49	Reference
50–64	0.82 (.78–.86)
≥65	0.46 (.41–.51)
UK, FR			
0–14	...	1.27 (1.03–1.50)	0.95 (.76–1.16)
15–44	...	Reference	Reference
45–64	...	0.99 (.89–1.11)	0.87 (.76–0.98)
≥65	...	0.82 (.71–.94)	0.68 (.58–.79)
Children in household (vs living alone)	1.31 (1.22–1.40)	S ^e	S ^e
Contact with groups ^f	ND	1.11 (1.01–1.21)	1.12 (1.01–1.23)
Smoker	1.16 (1.10–1.22)	S ^e	S ^e
Underlying health condition			
Asthma ^g	1.58 (1.47–1.69)
Diabetes	1.27 (1.15–1.41)
Heart	1.29 (1.13–1.47)
Kidney	1.23 (.80–1.90)
Immune	1.23 (1.02–1.49)
Any	...	S ^e	1.17 (1.05–1.30)
Having respiratory allergies	ND	1.14 (1.05–1.24)	1.19 (1.07–1.29)
Declaring often having ILI	ND	ND	1.31 (1.17–1.45)
Sports participation for >1 h/wk	0.95 (.90–1.00)	ND	ND
Having pets			
Dogs	1.15 (1.09–1.22)
Cats	1.07 (1.02–1.12)
Any	...	ND	1.17 (1.08–1.28)
Daily transportation method			
Bike/foot	0.95 (.90–1.00)	NS ^h	S ^e
Car	Reference
Public	0.97 (.89–1.05)

RISK FACTORS FOR ILI (multicountry)

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Devenez acteur de la surveillance de la grippe

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Objectifs

Estimer la fréquence de la grippe chez les femmes enceintes et le nombre de femmes vaccinées contre la grippe pendant leur grossesse en France métropolitaine.

G-GrippeNet est un projet de recherche fonctionnant sur le même principe que GrippeNet.fr, qui s'adresse spécifiquement aux **femmes enceintes**.

G-GrippeNet permet de participer activement à la surveillance de la grippe en France, de façon volontaire, anonyme et bénévole. Les données recueillies permettront de faire avancer la recherche sur la grippe afin de **mieux protéger les femmes enceintes et leurs bébés**.

Toutes les femmes enceintes résidant en France métropolitaine peuvent participer à G-GrippeNet, pour la durée qu'elles souhaitent. La participation à l'étude ne prend pas plus de 5 minutes par semaine. Le processus d'inscription est le même que pour Grippenet.fr, toutes les informations que vous trouverez sur ce site sont valables pour l'étude G-GrippeNet sauf mention contraire.

[Un commentaire, un problème ?](#)

5431 participants, 9004 comptes

Nom d'utilisateur

Mot de passe

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[Login ou mot de passe oublié ?](#)

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flusurvey



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i-sense
EPSRC IRC in Early Warning Sensing
Systems for Infectious Diseases

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Virological Swabbing

Virological Swabbing

This year at Flusurvey we're pleased to announce that we are piloting taking swab samples from a subset of the participants to ascertain whether, once experiencing symptoms, they actually do have the flu virus or not.



Self-Swabbing for Virological Confirmation of Influenza-Like Illness Among an Internet-Based Cohort in the UK During the 2014-2015 Flu Season: Pilot Study

C. Wenham, E. Gray, C. Keane, M. Donati, D. Paolotti, R. Pebody, E. Fragaszy, R. McKendry, J. Edmunds

J Med Internet Res 20,3 (2018)

Why have I been chosen to take part in the swabbing sample?

You have been invited to take part in an additional flusurvey sub-study. This year, in addition to collecting information provided by you online relating to your symptoms during the flu season, we are asking a small sample of people to undertake a swabbing

What is the definition of Influenza-like illness?

Sudden onset of symptoms

AND

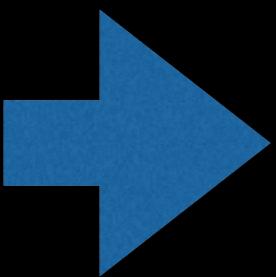
*at least one of the following four systemic symptoms:
Fever or feverishness, Malaise, Headache, Myalgia*

AND

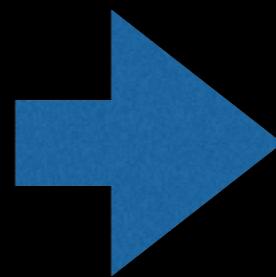
*at least one of the following three respiratory symptoms:
Cough, Sore throat, Shortness of breath*

“What if the observed symptoms are the result of a superposition of latent syndromes characterised by an unknown incidence and an unknown composition in terms of symptoms?”

Weekly
Symptoms
Survey



1. Fever
2. Chills
3. Runny/blocked nose
4. Sneezing
5. Sore throat
6. Cough
7. Shortness of breath
8. Headache
9. Muscle/joint pain
10. Chest pain
11. Feeling tired (malaise)
12. Loss of appetite
13. Phlegm
14. Watery, bloodshot eyes
15. Nausea
16. Vomiting
17. Diarrhoea
18. Stomachache
19. Sudden Onset



time
series of daily
symptoms
counts

boolean variables

$$\mathbf{X} = [x_{ij}]$$

matrix whose elements contains the occurrence of symptom j on day i

Latent Syndromes detection

- it is reasonable to expect that a specific combination of symptoms reported by a user is the symptomatic expression of one or more illnesses, i.e. syndromes, experienced by the user.
- In accordance with this consideration, we postulate that the time series x_{ij} of observed symptoms counts are the result of a linear mixing process driven by K unknown sources, corresponding to the latent syndromes we want to detect.

$$x_{ij} = \sum_{k \in \{1, \dots, K\}} w_{ik} h_{kj} + e_{ij}.$$

Latent Syndromes detection

The mixing equations can be expressed in matrix notation:

$$\mathbf{X} = \mathbf{W}\mathbf{H} + \mathbf{E}$$

$$\mathbf{W} = [w_{ik}], \mathbf{H} = [h_{kj}], \mathbf{E} = [e_{ij}]$$

In this notation, the problem of detecting the unknown K latent sources can then be formulated as a matrix decomposition problem

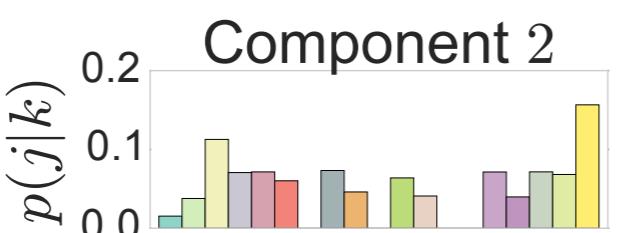
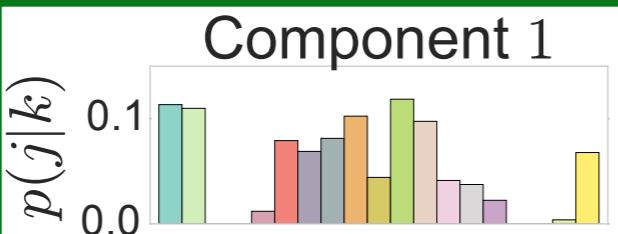
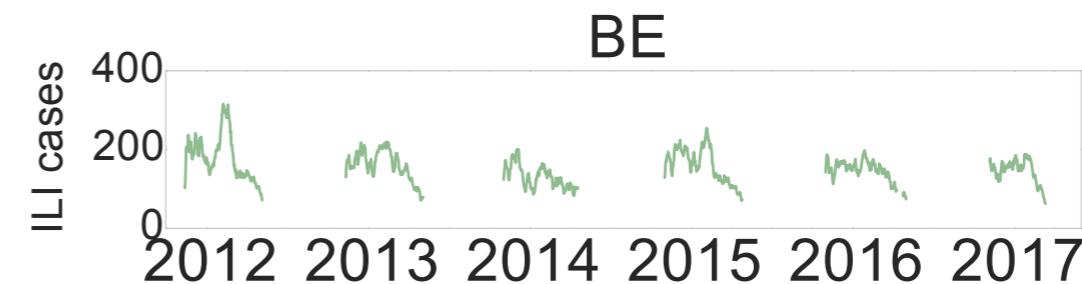
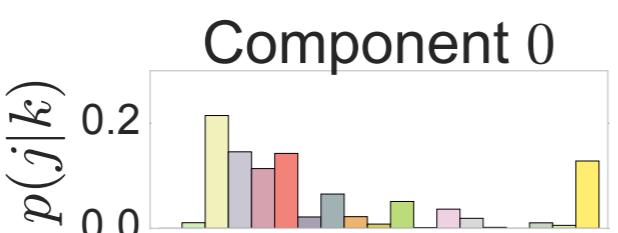
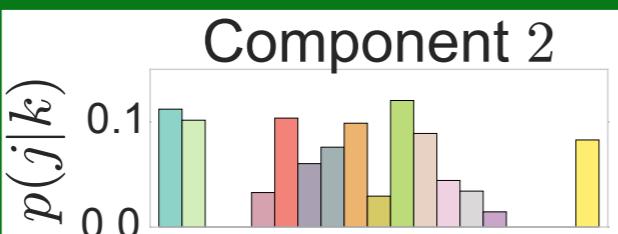
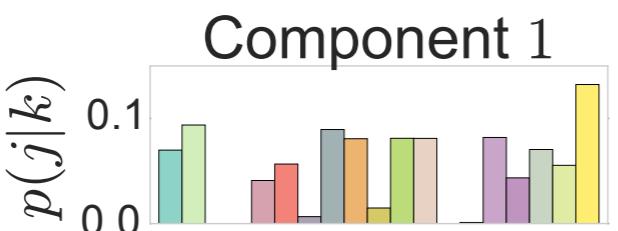
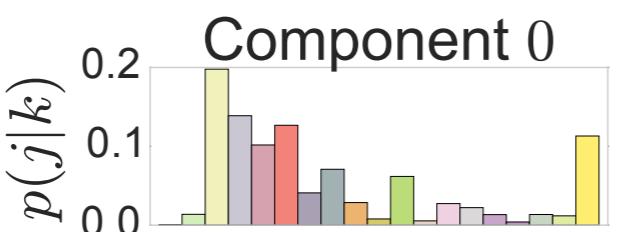
Non-negative matrix factorization

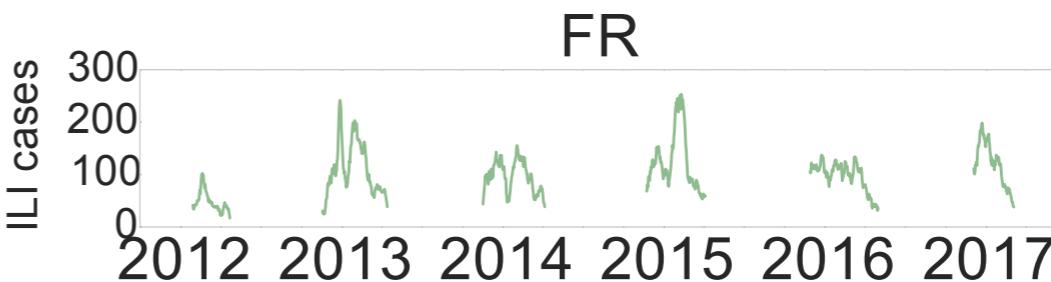
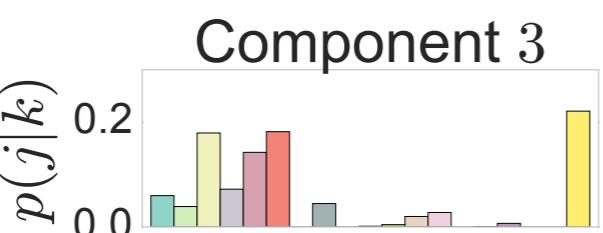
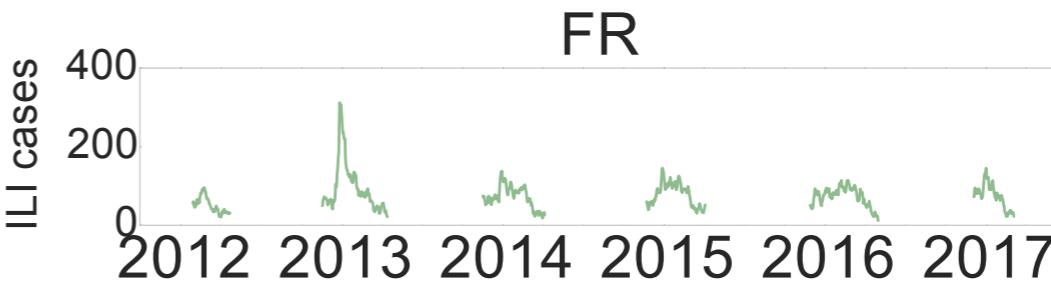
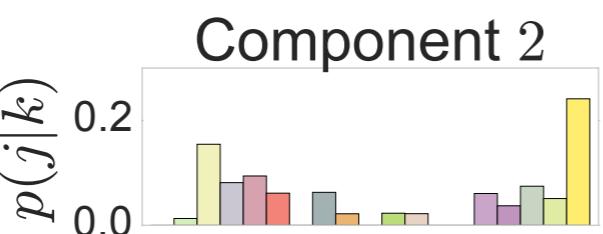
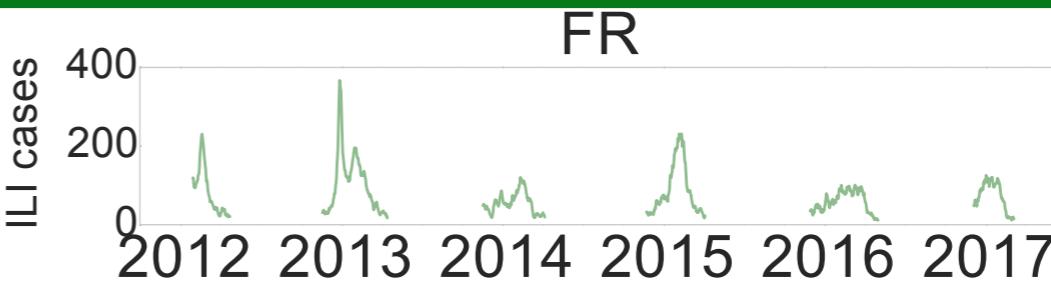
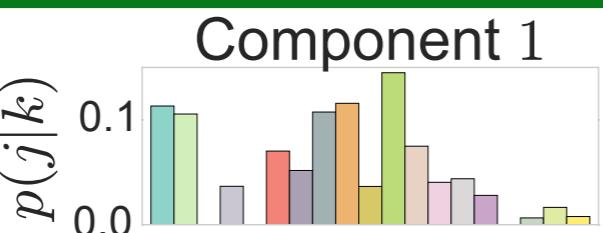
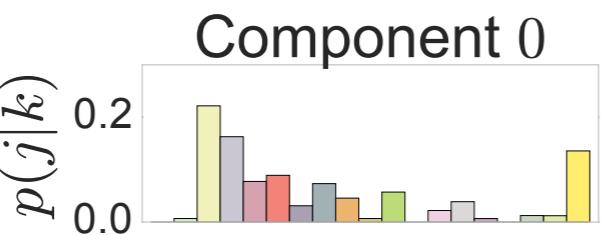
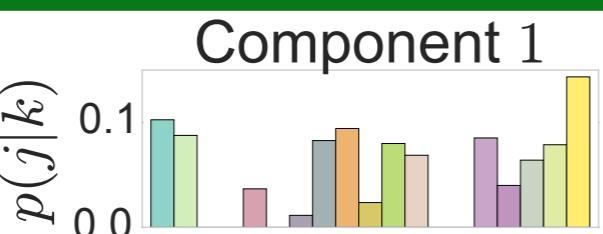
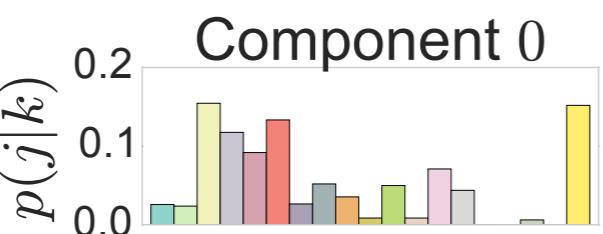
As a result of the probabilistic formulation, the total number of symptoms counts N will be proportionally split among K latent components according to $p(k)$. $p(j|k)$ describes each component in terms of the expected proportion of symptoms. According to this formulation, the total number of counts associated to a latent component k in day i will be given by:

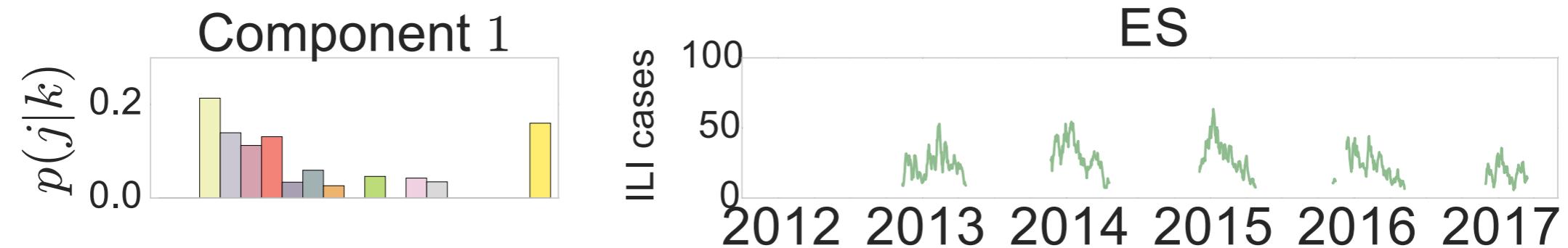
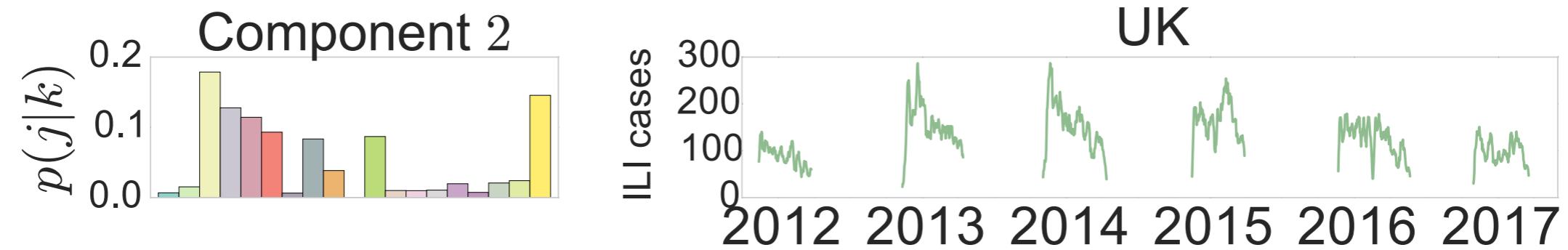
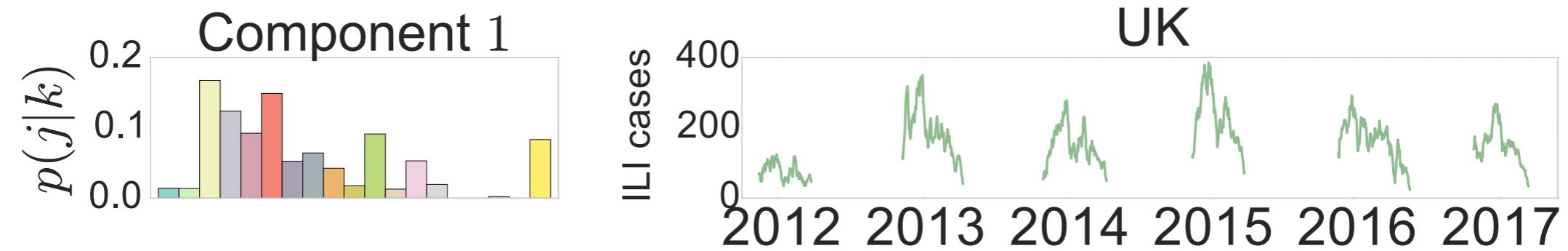
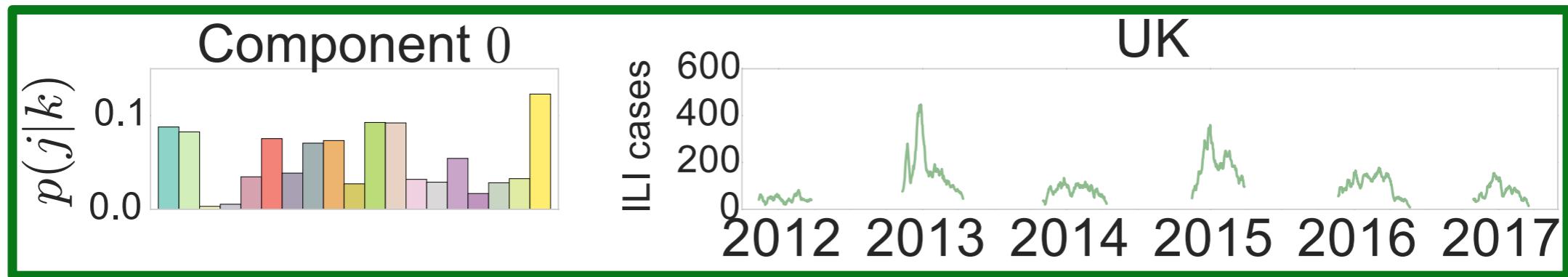
$$y_{ik} = N p(i, k) = N p(k) p(i|k)$$

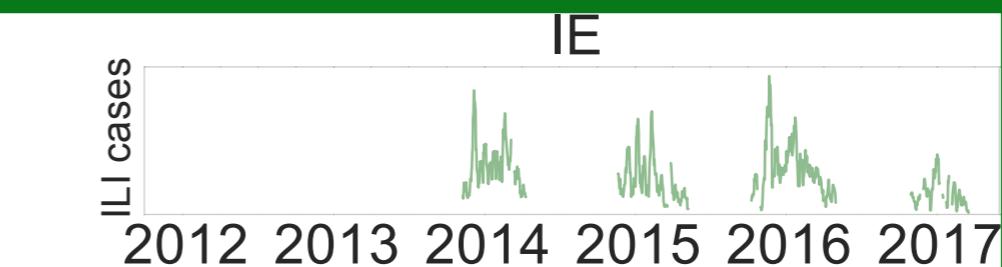
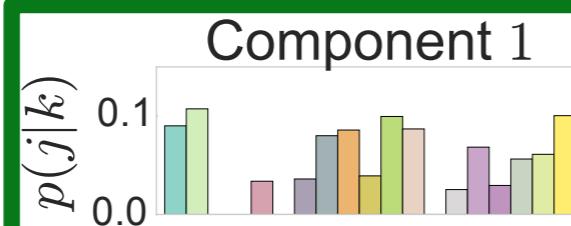
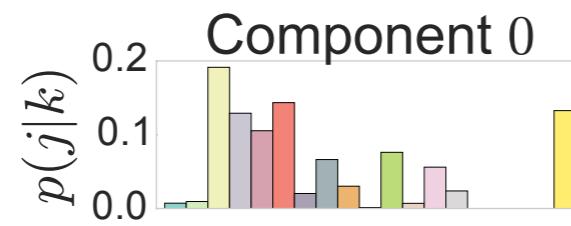
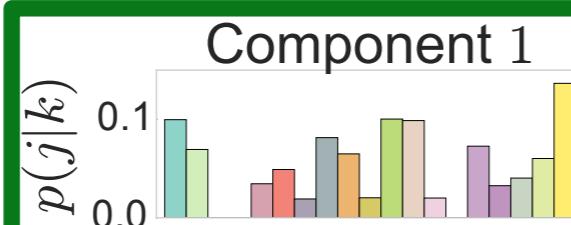
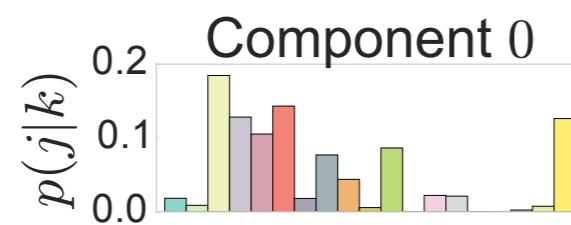
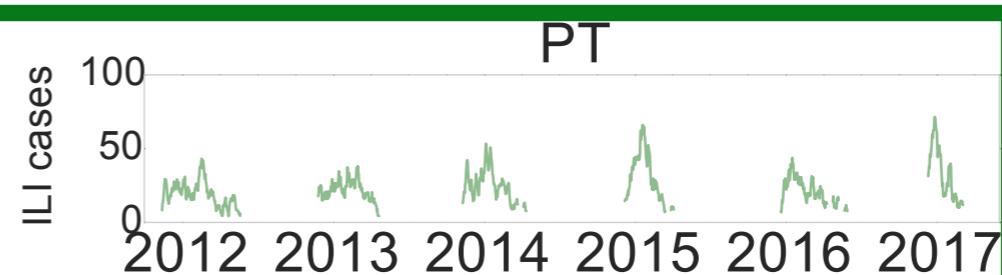
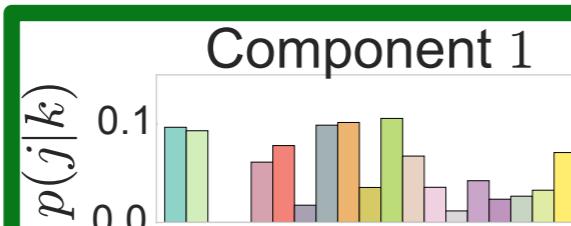
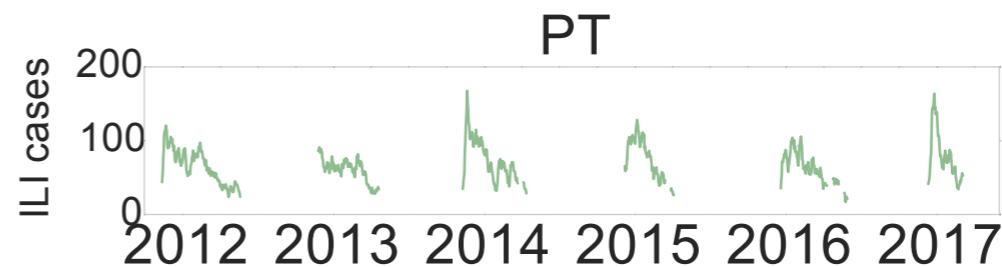
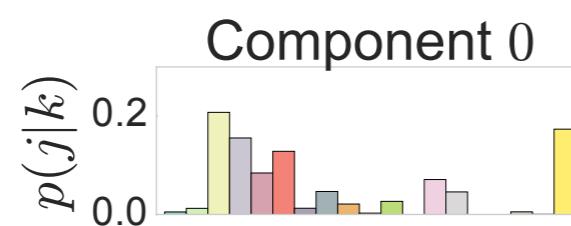
Back to data

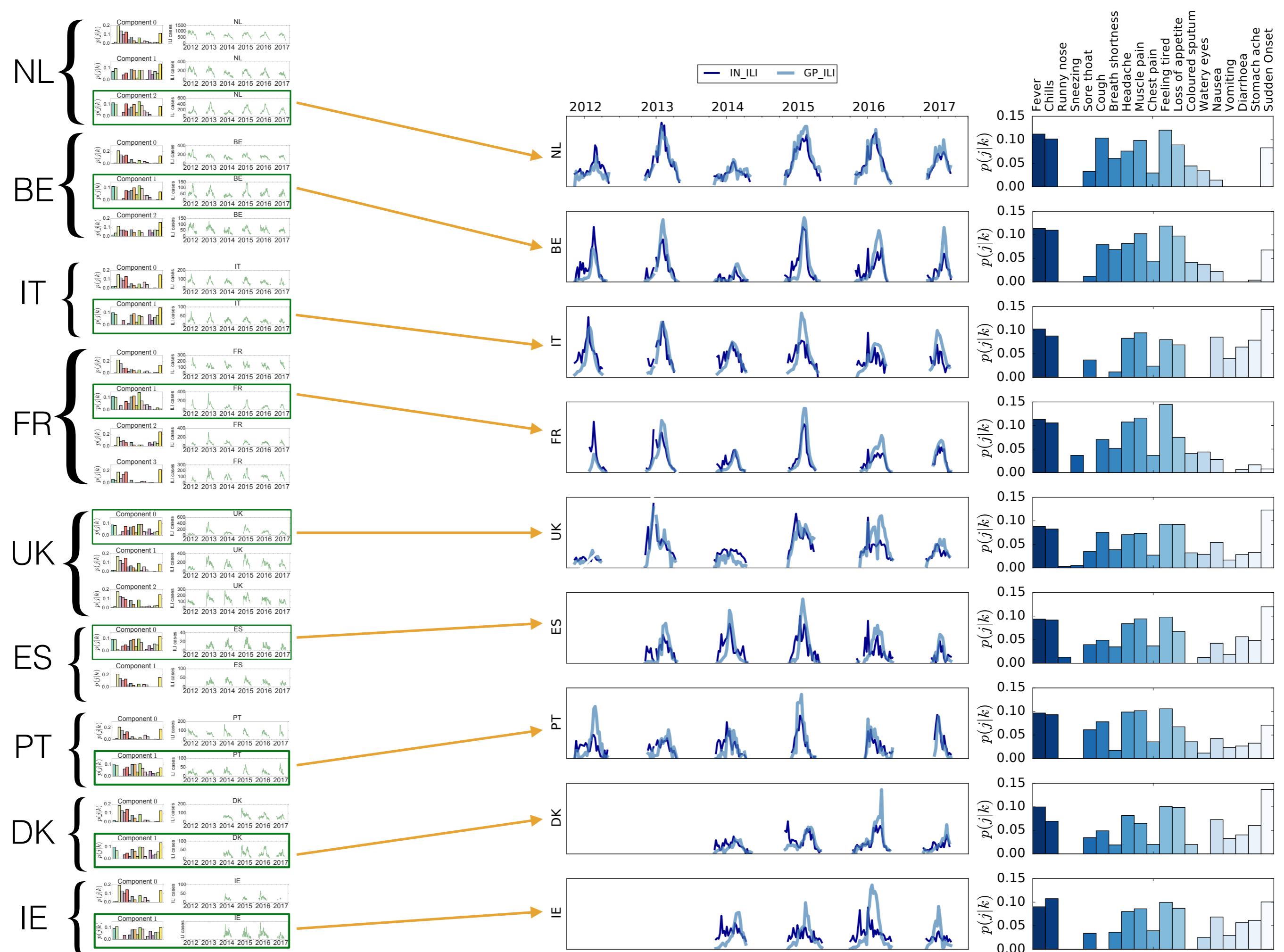
	NL	BE	IT	FR	UK	ES	PT	DK	IE
Number of seasons	6	6	6	6	6	5	6	4	4
Average participants per season	13,450	4,209	1,830	5,757	4,676	526	1,663	1,391	406
Average # surveys per season	206,987	67,420	17,807	68,567	45,543	5,894	17,852	22,782	3,220
Average % of surveys with symptoms	20%	16%	19%	20%	29%	22%	17%	18%	25%
Average of surveys per participant per season	15	16	9	12	9	11	10	16	8









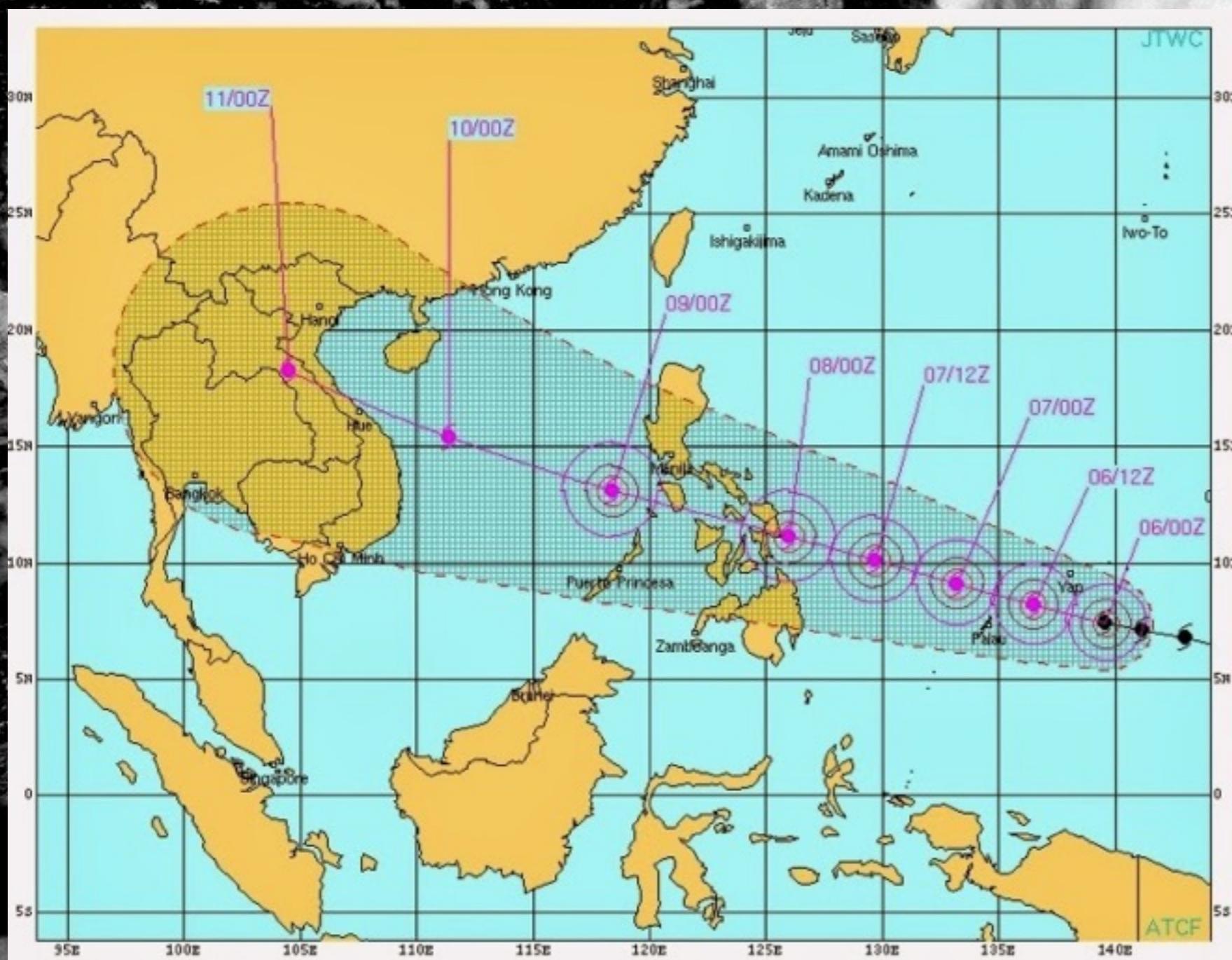


	NL	BE	IT	FR	UK	ES	PT	DK	IE
(i) Correlation between IN_ILI_ECDC and IN_ILI for the seasons 2011-2017	0.91	0.92	0.86	0.83	0.92	0.86	0.84	0.90	0.82
(ii) Correlation between IN_ILI and GP_ILI for the seasons 2011-2017	0.88	0.80	0.69	0.79	0.74	0.65	0.66	0.71	0.38
(iii) Correlation between IN_ILI_ECDC and GP_ILI for the seasons 2011-2017	0.79	0.72	0.80	0.86	0.75	0.67	0.63	0.68	0.23
(iv) Correlation between IN_ILI prediction and GP_ILI for the season 2016-2017	0.85	0.82	0.69	0.80	0.60	0.84	0.80	0.76	0.60
(v) Correlation between IN_ILI_ECDC and IN_ILI for the season 2016-2017	0.85	0.82	0.86	0.93	0.67	0.59	0.88	0.80	0.71

Unsupervised extraction of epidemic syndromes from participatory influenza surveillance self-reported symptoms, K. Kalimeri, M. Delfino, C. Cattuto, D. Perrotta, V. Colizza, C. Guerrisi, C. Turbelin, J. Duggan, J. Edmunds, C. Obi, R. Pebody, A. O Franco, Y. Moreno, S. Meloni, C. Koppeschaar, C. Kjelsø, R. Mexia, D. Paolotti, PLOS Computational Biology 15 / 4 (2019)

What's next - I?

- Virological confirmation is needed to estimate more accurately the scaling factor
- extension of the method to other countries and syndromes
- Assess the validity of the methods for detection of new emerging diseases



Forecasting seasonal influenza

Baseline

Model: forecasting influenza through sentinel doctors reports

$$y_{w+k} = \alpha_1 \tilde{y}_{w-1} + \alpha_2 \tilde{y}_{w-2} + \alpha_3 \tilde{y}_{w-3}$$

y_w denotes the ILI incidence value reported by the sentinel doctors at week w ($k=0$ nowcasting, $k>1$ forecasting)

Participatory Surveillance + sentinel doctors reports

$$y_{w+k} = \sum_{i=1}^3 \alpha_i \tilde{y}_{w-i} + \gamma_0 \tilde{z}_w$$

Linear Autoregressive exogenous model:

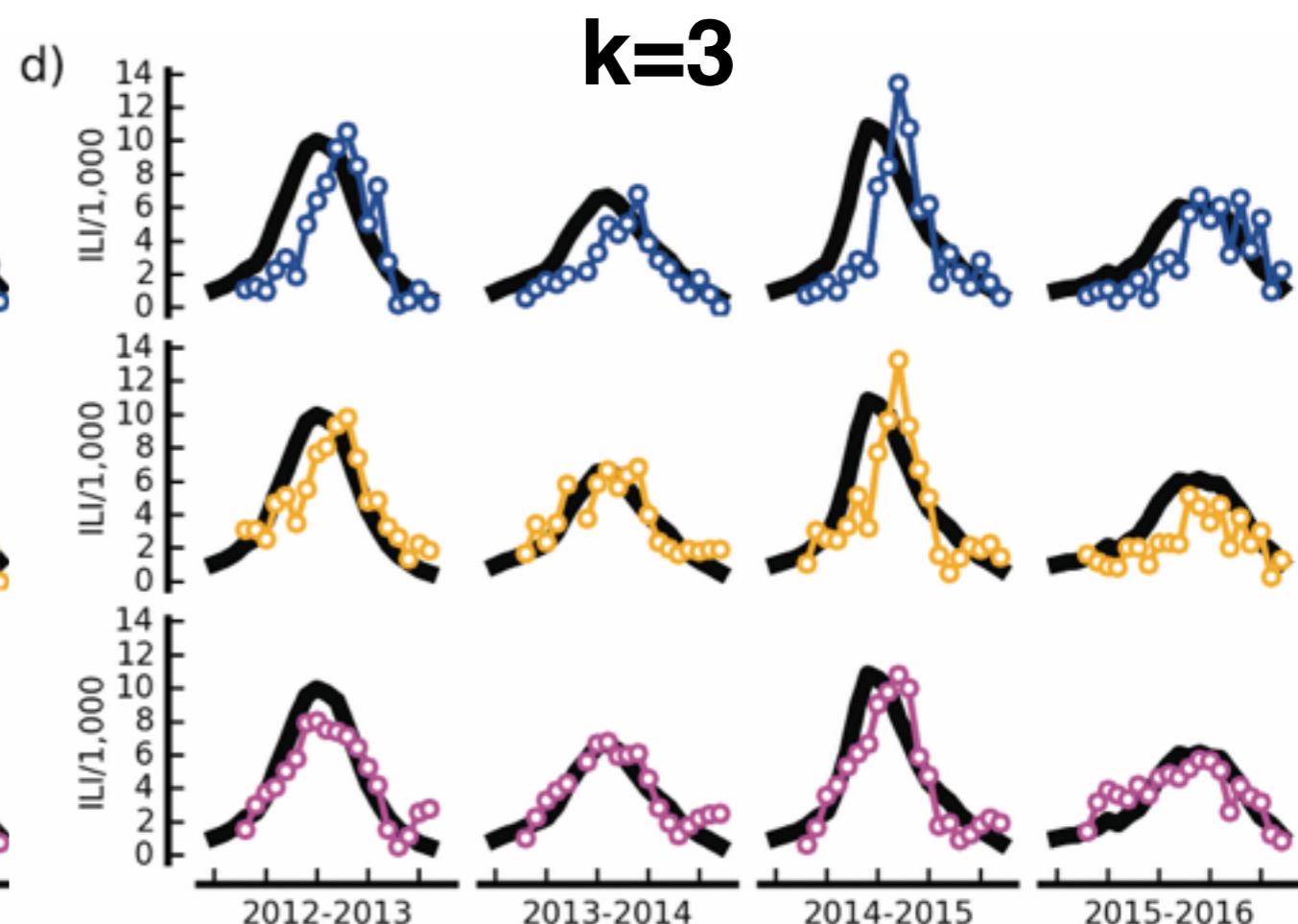
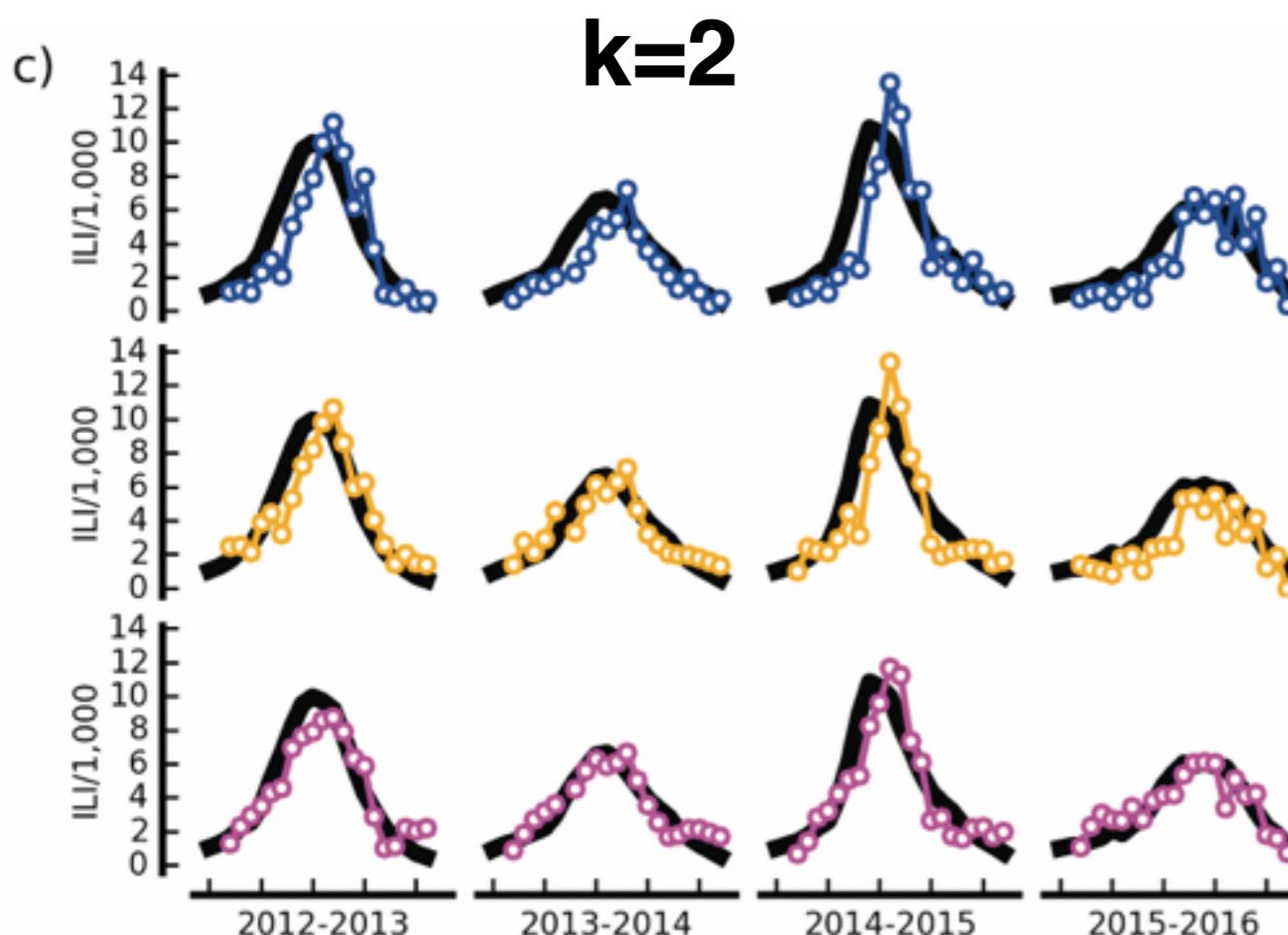
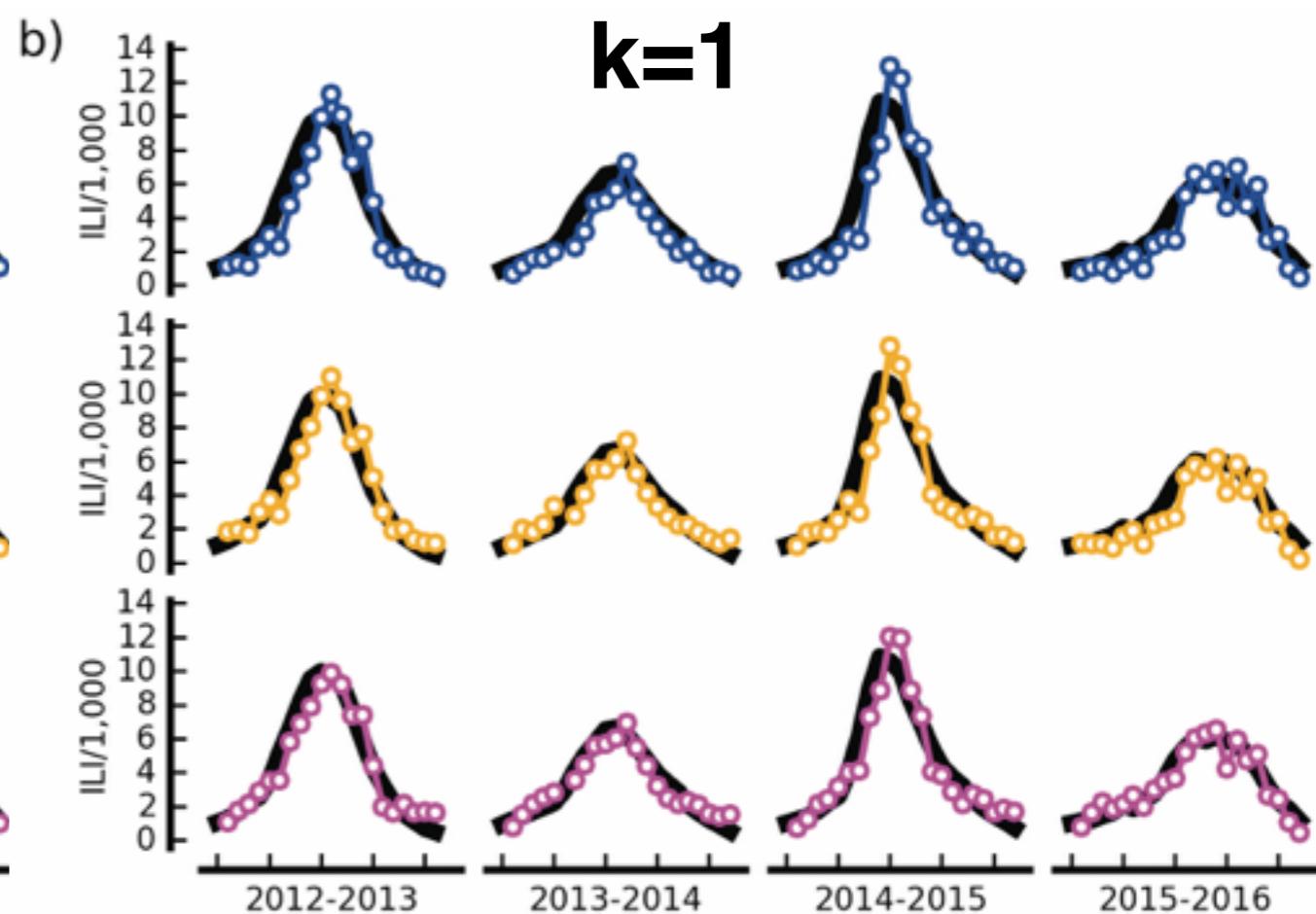
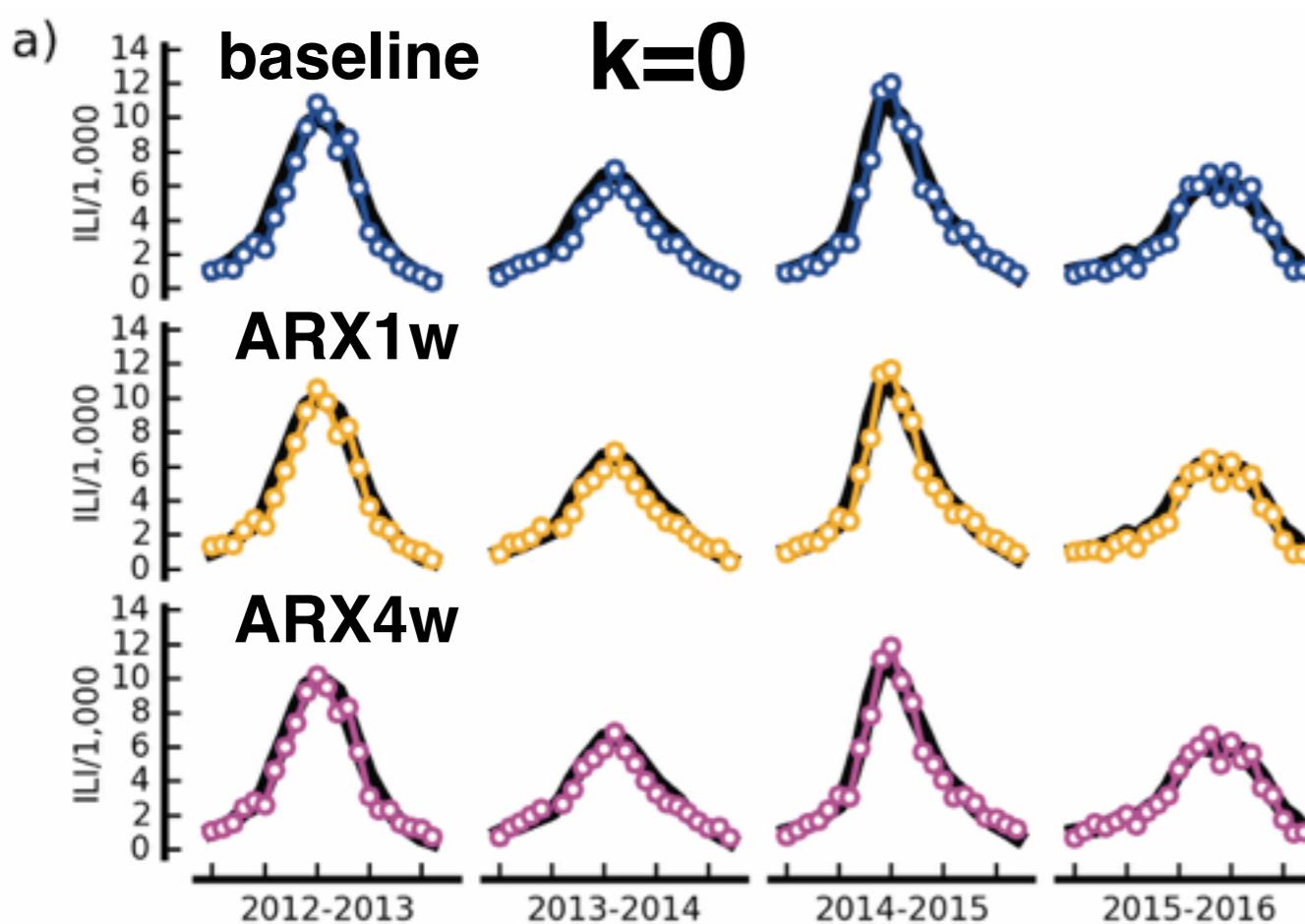
ARX1w, ARX4w

$$y_{w+k} = \sum_{i=1}^3 \alpha_i \tilde{y}_{w-i} + \sum_{j=0}^3 \gamma_j \tilde{z}_{w-j}$$

INFLUENZANET in Italy: influweb.it

Participation to Influweb during the five influenza seasons under study.

Season	Number of registered users	Average number of weekly active users	Total number of symptoms surveys
2011-2012	2,270	1,085	14,681
2012-2013	3,057	1,165	15,789
2013-2014	3,513	1,244	18,471
2014-2015	3,753	1,290	20,224
2015-2016	4,053	1,224	20,823



Similarity metrics and peak analysis of the forecasting models with respect to the ground truth.

The best performing model per metric is bold faced.

	Model	Similarity Metrics			Peak Analysis		
		CORR	MAE (ILI/1,000)	RMSE (ILI/1,000)	MAPE (%)	weeks lag [Min, Max]	PE (%) Mean [Min, Max]
$k = 0$	baseline	0.981	0.45	0.5955	12.93	[0, 1]	8.6 [4.8, 10.9]
	ARX_{1w}	0.983	0.42	0.5419	12.95	[0, 1]	5.5 [3.5, 7.7]
	ARX_{4w}	0.984	0.39	0.5052	13.13	[0, 1]	5.6 [1.9, 9.0]
$k = 1$	baseline	0.934	0.83	1.1174	22.39	[1, 2]	13.7 [8.9, 19.3]
	ARX_{1w}	0.942	0.73	0.9635	24.31	[1, 2]	9.3 [0.8, 18.1]
	ARX_{4w}	0.957	0.63	0.7952	25.63	[0, 1]	5.5 [0.7, 10.6]
$k = 2$	baseline	0.835	1.29	1.7497	33.79	[2, 3]	13.8 [7.7, 24.3]
	ARX_{1w}	0.864	1.05	1.4488	32.72	[2, 3]	11.7 [6.5, 23.0]
	ARX_{4w}	0.918	0.84	1.0885	34.62	[0, 2]	5.2 [0.3, 12.0]
$k = 3$	baseline	0.677	1.80	2.4287	45.23	[0, 3]	9.7 [2.3, 23.2]
	ARX_{1w}	0.751	1.42	1.9404	43.23	[0, 3]	10.7 [1.7, 22.1]
	ARX_{4w}	0.879	1.02	1.3058	42.93	[0, 3]	7.2 [0.6, 19.7]

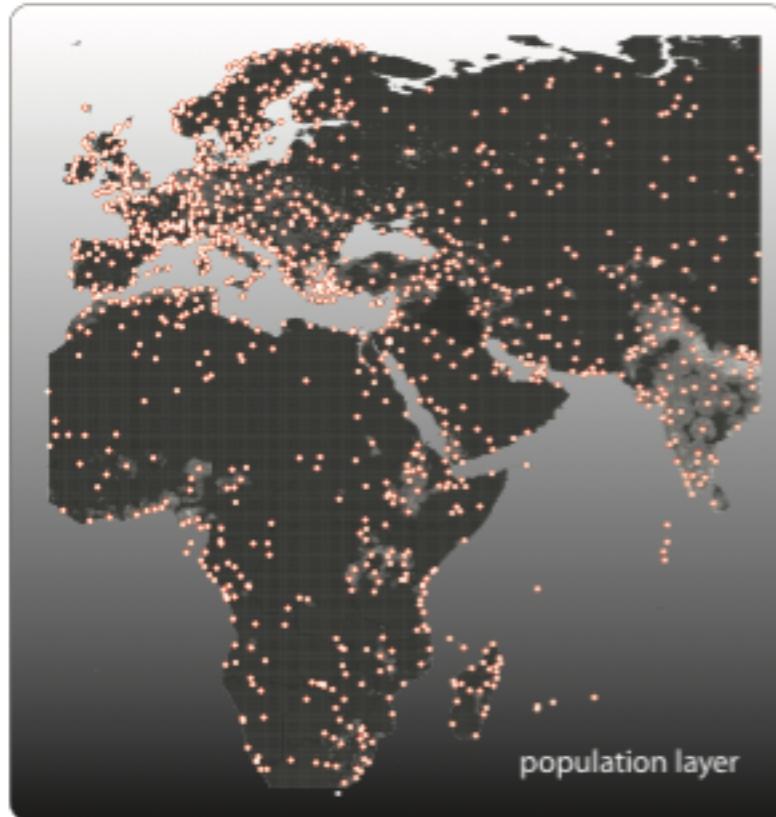
MAE = Mean Absolute Error

RMSE = Root Mean Square Error

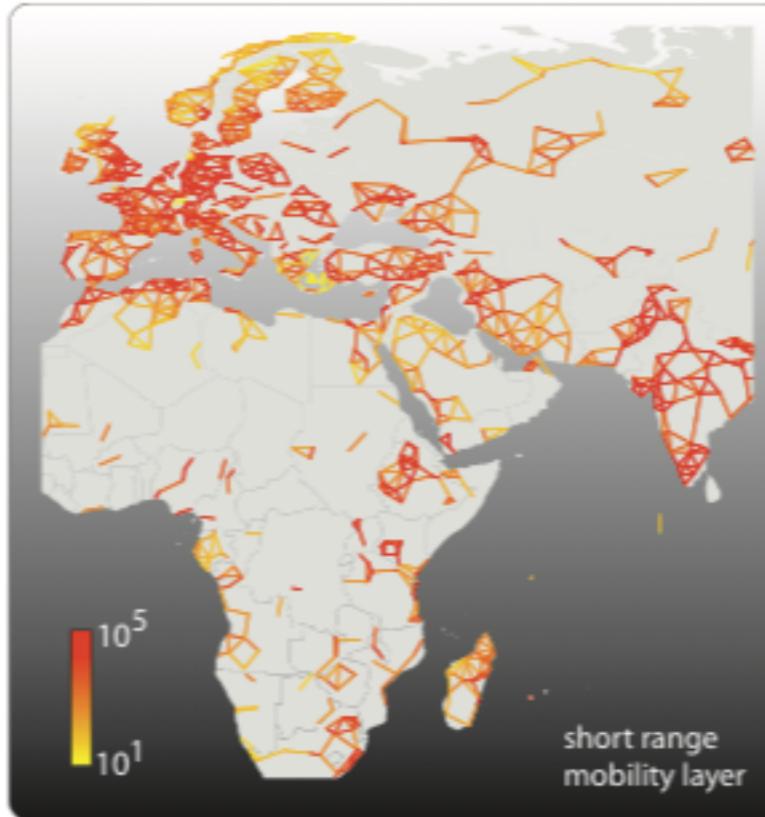
MAPE = Mean Absolute Percent Error

proceedings of WWW2017: D. Perrotta, M. Tizzoni, D. Paolotti, “Using Participatory Web-based Surveillance Data to Improve Seasonal Influenza Forecasting in Italy”

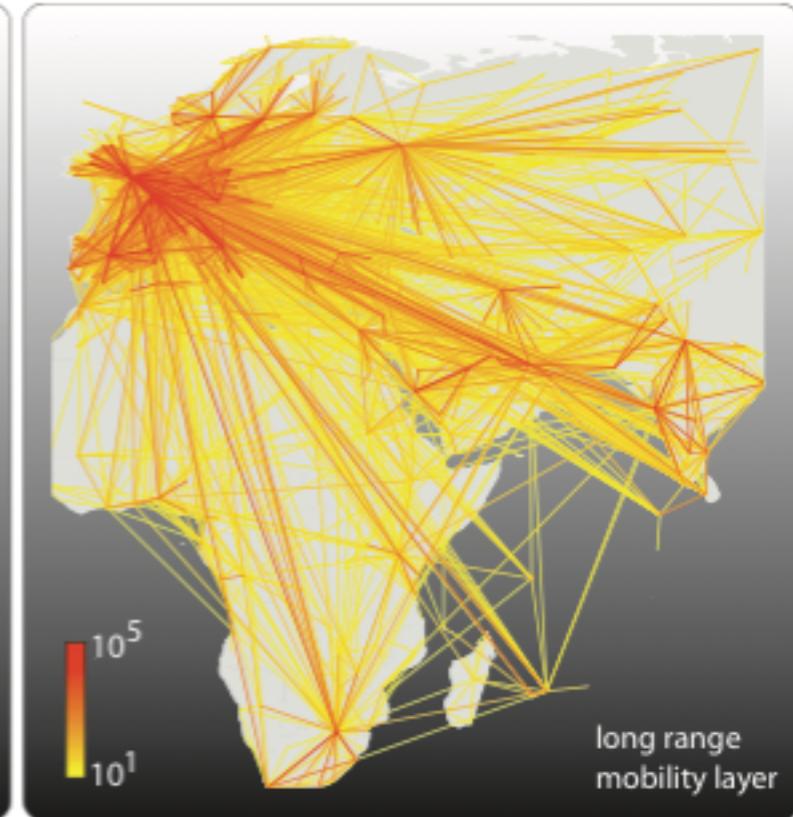
THE MODEL: GLEAM



Population
layer



Commuting

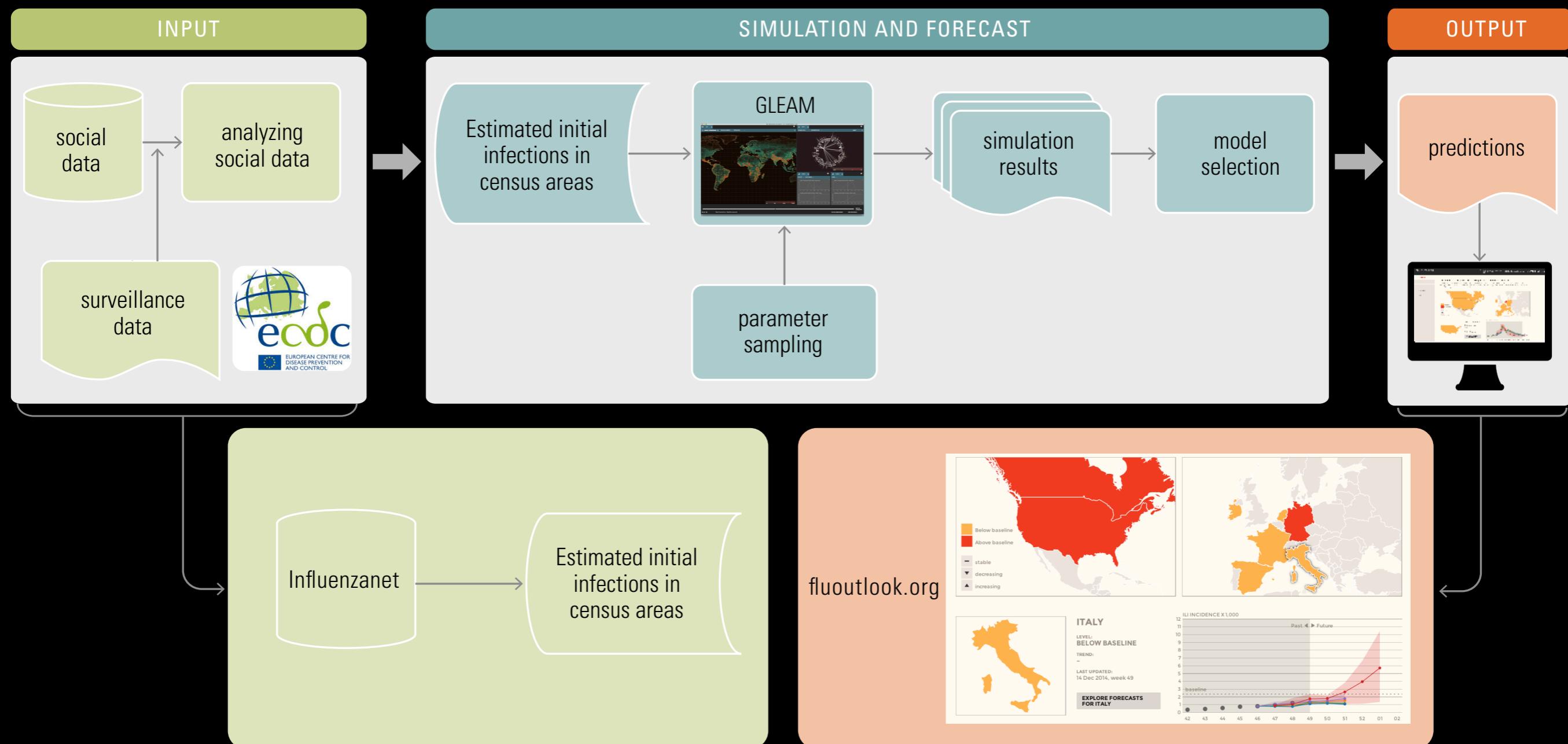


Air travel

Human mobility layers

Geographic scale

Forecasting seasonal influenza: fluoutlook.org



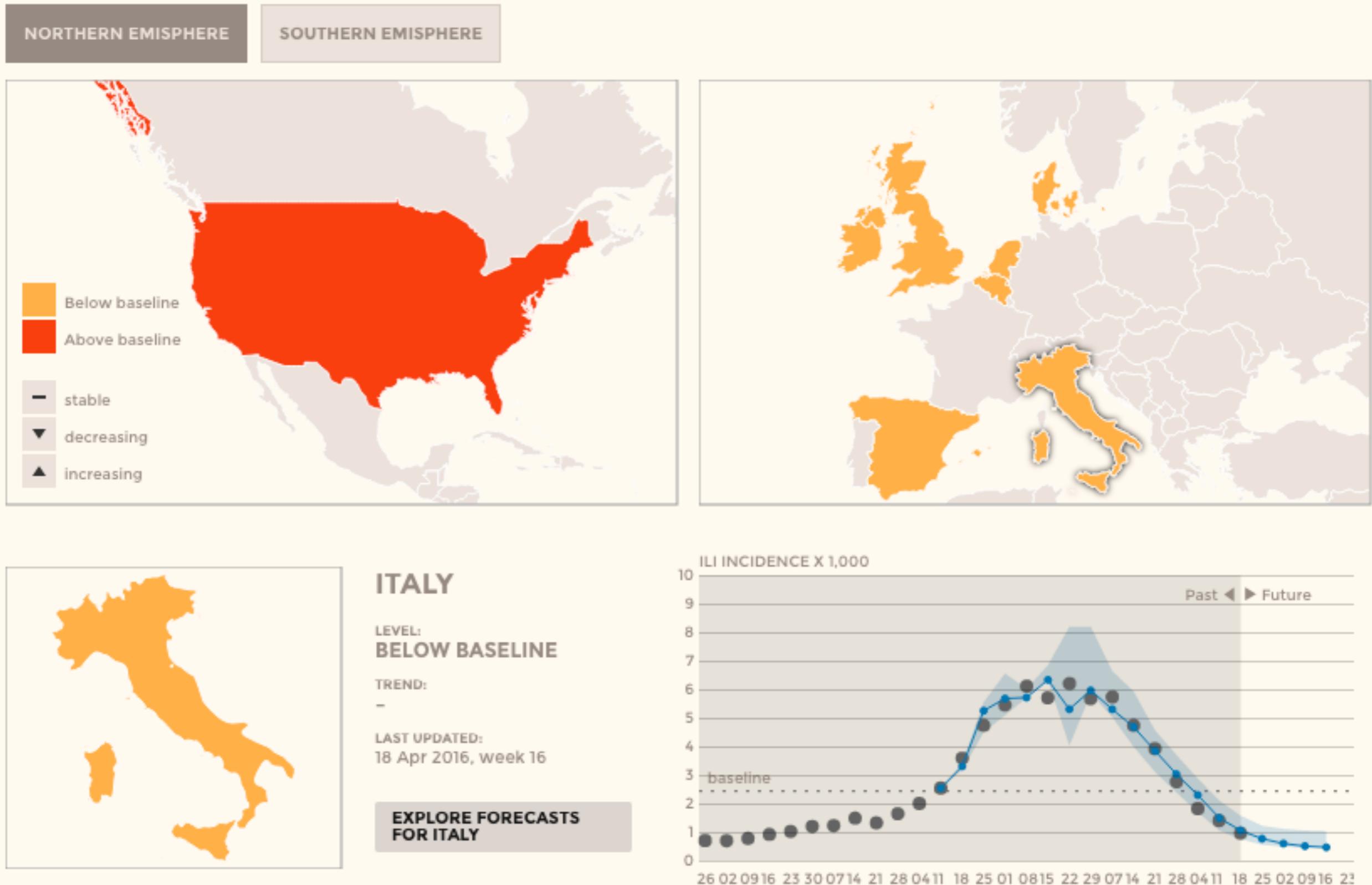
Social Data Mining and Seasonal Influenza Forecasts: The FluOutlook Platform

Q. Zhang, C. Gioannini, D. Paolotti, N. Perra, D. Perrotta, M. Quaggiotto, M. Tizzoni, A. Vespignani

Lecture Notes in Computer Science 9286 (2015) p.237-240

Fluoutlook is a web platform for the exploration of influenza forecasts

It provides a visual interface to numerical forecasts of the current influenza season in North America and Europe through maps and charts. Activity data and forecasts are updated weekly, based on the reports of the official influenza surveillance systems in each country. [Tell me more.](#)



Conclusions - 1

In summary, Influenzanet is a well-established standardized participatory surveillance system for ILI in Europe, covering more than one third of the 28 EU member states. Its strength lies in:

- (1) the **standardized technological and epidemiological framework** for a coherent surveillance across countries and the centralized fashion that characterize the technological deployment (except for France and Sweden, all the platforms have been deployed and are now managed by ISI Foundation in Turin, Italy).
- (2) the ability to timely **monitor ILI in the general population**, including individuals who do not seek medical assistance;
- (3) its **sensitivity** in detecting substantial changes in population health earlier than GP sentinel networks;
- (4) its potential **scalability** to large numbers with rather **limited costs**;
- (5) its **flexibility** in exploring different ILI definitions;
- (6) the detailed profile data allowing **individual-level epidemiological** analyses generally not possible in standard systems;
- (7) its potential extension to **other diseases**.

What's next - II?

- Participatory surveillance in developing countries
- Behavioural changes
- Other health-related issues: pregnant women and vaccination, chronic conditions, vaccine hesitancy
- Validation of passive web data: linking behavioural data from participatory systems with on-line data

ECDC/WHO bulletin: indicators

- Weekly number of active users
- Weekly incidence (with various ILI case definition)
- % of ILI cases in contact with healthcare
- cumulative % of vaccinated by key target group

Conclusions - 2

- Its limitations are mainly due to the self-selected sample, potential misreporting, and lack of validation by a physician or by virological testing. However, the agreement found with GP incidence trends suggests that these limitations have little effect once results are adjusted for lack of representativeness.
- Main challenges remain the baseline maintenance resources to sustain the system in the long run and the recruitment and retention of participants. While the identification of sociocultural determinants for participation will provide additional insights, the strong willingness for engagement found in most countries' participants confirms the feasibility of the approach.
- Moreover, the platform represents a crucial channel for communication with the public, to inform and increase awareness, an increasingly important aspect after the 2009 pandemic.

INFLUENZANET.INFO

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influweb

grippenet.fr

gripe^{net}①

flusurvey ①

www.influensakoll.se
Kartlägger förekomst och spridning av influensa i Sverige

influmeter.dk
Kortlægger forekomst og spredning af influenza i Danmark.

flusurvey.ie

grippe^{net}
GrippeWeb

- *C. Gioannini*
- *D. Perrotta*
- *M. Quaggiotto*
- *L. Rossi*
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- *M. Delfino*
- *K. Kalimeri*
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LABORATORY FOR THE MODELING OF BIOLOGICAL
AND SOCIO-TECHNICAL SYSTEMS



Northeastern



Thank you!

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