Public Health Surveillance

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"The continuous, systematic collection, analysis, and interpretation of healthrelated data needed for the planning, implementation, and evaluation of public health practice."

The goal is to tie disease activity to geographic location.

Serves as an early warning system for impending public health emergencies.

Documents the impact of an intervention, or tracks progress towards specified goals, such as mitigation or containment.

Monitors and clarifies the epidemiology of health problems, informing public health policy and strategies.



Syndromic means "factors related to disease," *e.g.*, symptoms.

Syndromic surveillance samples and collects disease-related data to impute the speed, size and tempo of an outbreak. When does it start? How fast does it grow? How long does it last?

As an example, when tracking influenza, we track %ILI, or "percent influenzalike illness," defined as the proportion of outpatients seen with "fever of 37.8C or higher combined with a cough and/or sore throat but no other apparent cause."

Other, non syndromic, surveillance systems might track factors on the periphery of real disease data, such as people's perceptions of disease, or other behaviors (e.g., hand hygiene, cell-phone social distancing measures) that are indirectly related to disease.



In 2008, Google flu trends used Google search term frequency and (geographic) origin to track %ILI by geographic area.

Flu trends used IP address location from the search to determine geographic area, anonymizing individual user queries by aggregating up.

They reported surprising accuracy values, and seemed to "lead" CDC's ability to estimate onset and peak of the curve by about two weeks.

Unfortunately, in 2013 Google flu trends spectacularly underperformed. Why?



Goodle flu trends overfits the data.

People who search for flu aren't really good at diagnosing flu; "cough" and "fever" are not specific to flu, and searches for individual flu-like symptoms may not meet the definition of ILI.

Correlation is not causality; some accused Google flu trends of simply predicting "winter."

Searches for flu related issue closely track news about the flu; later, Google corrected for prevalence of flu in news.



During the 2009 H1N1 swine flu epidemic (originating in Mexico and US), we quickly assembled a system to collect all tweets that contained what we determined were influenza-related words.

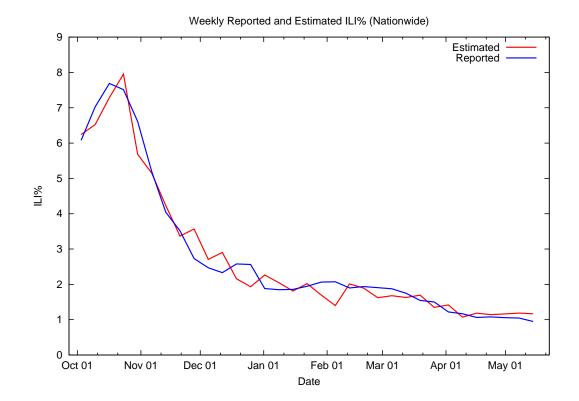
A total of 686,812 H1N1-related tweets out of 607,576,369 total tweets in a 5-week period starting 4/28/09.

Each tweet contains a geolocation code: sometimes, the mobile device reports GPS coordinates; other times, we used the user's self-declared home location.

We were hoping that tweet analysis would provide an inexpensive way to determine not only levels of anxiety and concern but also to gauge response to news and official public health messaging.



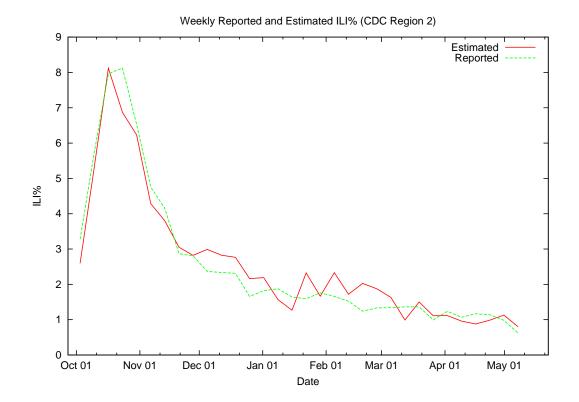
Example: Using Twitter to Monitor Public Health Concerns



Overall performance.



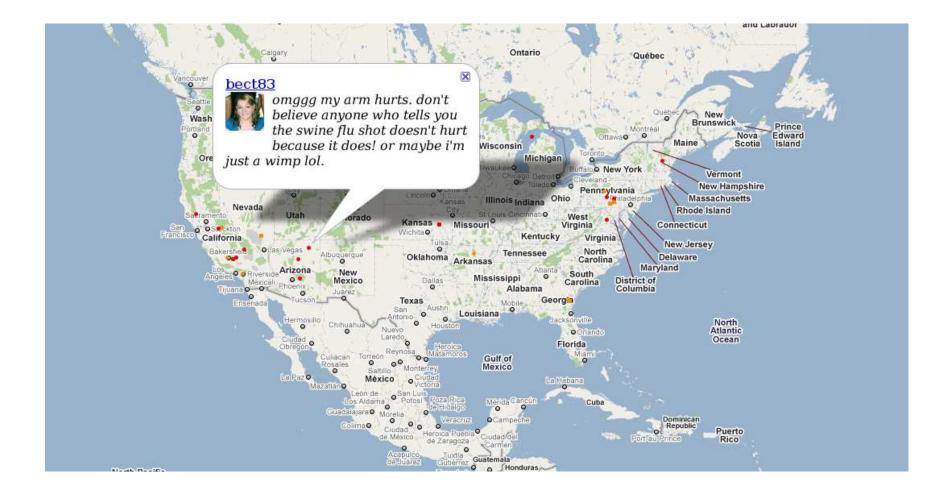
Example: Using Twitter to Monitor Public Health Concerns



Out-of-sample validation: predicting region 2 ILI from other region's tweets.



Live Twitter Feed for H1N1 Swine Flu





The reemergence of syphilis at the end of the 1990's is likely due to the advent of internet-mediated anonymous sexual encounters, most typically within the MSM community.

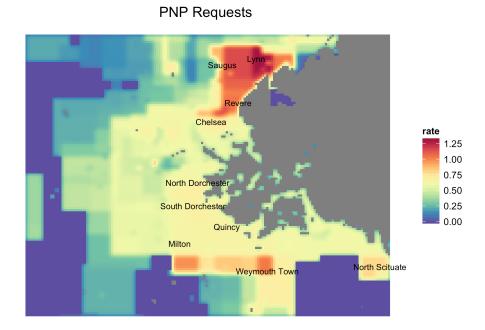
This community accounts for about 2% of the population, but 63% of new HIV cases, which are not uniformly distributed among racial and ethnic MSM subpopulations.

Understanding the online hookup market is a critical step to understanding STD disparities and designing effective public health interventions.

In 2008-10, we collected 136M Craigslist personals ads to understand the geographic footprints and sexual behaviors of anonymous casual sex-seeking individuals in order to understand their impact on STD transmission.



Example: Craigslist, Internet-Mediated Anonymous Sex, and STDs



In collaboration with the Boston Department of Public Health, we were able to use the Craigslist corpus to build an accurate picture of risky sexual behaviors at very high geographic resolution.

http://vimeo.com/48687015



These projects, as well as our work on instrumenting HCWs and monitoring hand-hygiene compliance, can be seen as surveillance efforts, but they are not really syndromic surveillance.

To some degree, they all exhibit the same problems as Google Flu Trends (overfitting, correlation/causality, etc.).

The paper we discuss today is different; maybe less sexy from a big data perspective, but it relies on professionals making judgments about actual symptoms, rather than indirect measurements of behaviors or effects of disease.



IDPH runs an ILI surveillence network using volunteer sentinel providers.

These volunteer sentinel providers report their weekly patient visits and the number of patients with ILI.

Because they are not paid for their efforts, these providers represent a convenience sample of pediatricians, internists, family practice physicians, group practices and hospital clinics.

Not everyone volunteers; there are some costs, which are largely supported by the providers, as well as IDPH.

The assigned paper addresses a very specific issue: how can we get the best information while minimizing cost?



Because the network is all volunteer, the 22 existing 2009 surveillance sites were not placed in geographically optimal locations.

Are the sites truly representative of the extend of disease? Do they capture sufficient activity? How does one choose the "best" sites?

As usual, what emerges is a location optimization problem, where a utility function needs to be formulated to meet some defined surveillance goal.

One view is to treat this as a facilities location problem, where we maximize the population "covered" within a certain geographic radius from each sentinel location.



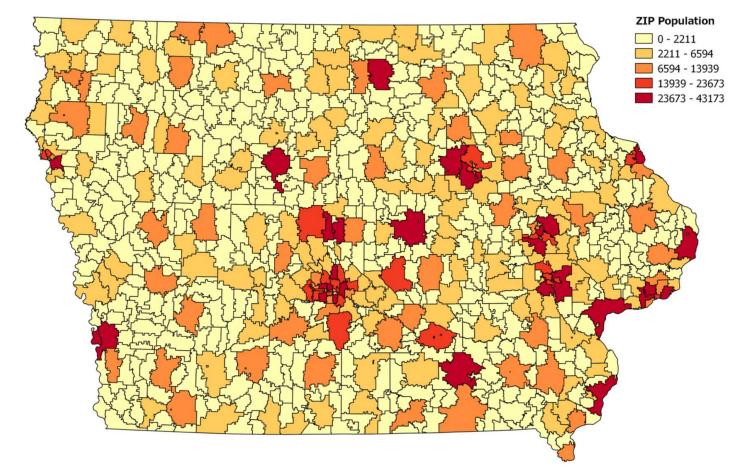


Figure 1 Map showing population distribution in Iowa by ZIP code. All 935 ZIP codes in Iowa are shaded by population. Darker colors indicate larger populations while lighter colors indicate smaller populations.



minimize:
$$\sum_{i=1}^{N} s_i$$

subject to:
$$\sum_{j=1}^{M} w_j \prod_{i=1}^{N} (1 - s_i y_{ij}) \le \tau$$

where:

N is the number of candidate surveillance sites

M is the number of population quanta (*e.g.*, ZCTA) that cover the state

 w_j is the *weight* of population quanta j

 s_i is the *selection variable* (with value 1 if site *i* is selected and 0 otherwise)

 y_{ij} is the *coverage variable* (with value 1 if population quanta *j* is serviced by candidate surveillance site *i* and 0 otherwise), and

 τ is the maximum number of people we're willing to leave uncovered.

Note that the problem is NP-hard; in practice, we used a greedy approximation algorithm to solve it.



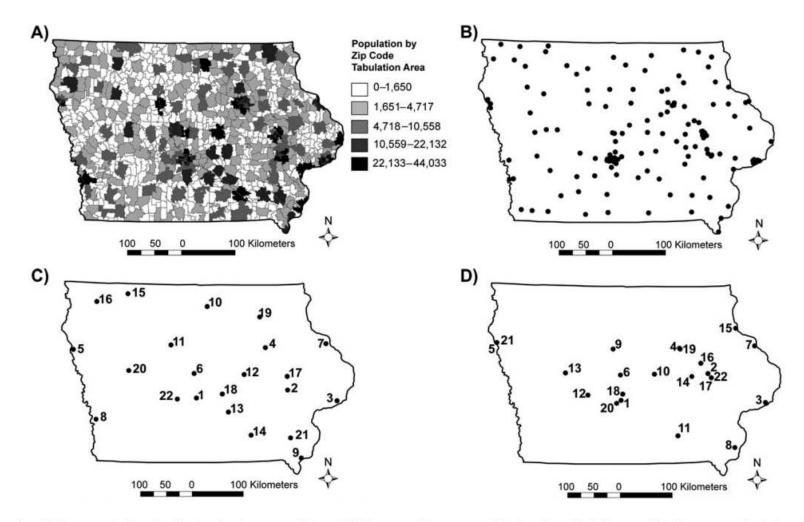


Figure 1. A) The population distribution for the state of Iowa. B) The 143 different possible locations for influenza-like illness sentinel sites. C) The 22 influenza-like illness sentinel locations (based on the 143 sentinel locations) chosen by the authors' maximal coverage model. The numbers represent the order in which the sites were chosen by the model. D) The 22 Iowa Department of Public Health influenza-like illness sentinel locations for the 2006–2007 influenza season. The numbers represent the order in which the existing sites should have been chosen to maximize coverage.



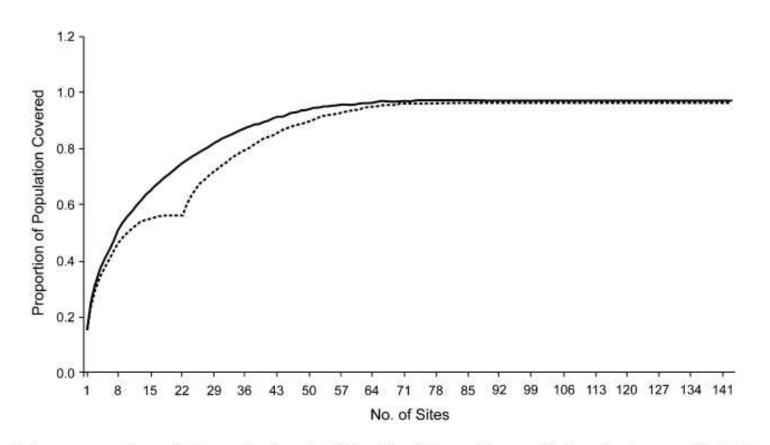


Figure 2. Population coverage (proportion) as a function of additional hospital surveillance units for a fixed, prespecified distance of 20 miles (32.2 km). The upper, solid curve represents the proportion of population coverage as each new surveillance site is added if the sites are selected de novo; the lower, dashed curve represents the proportion of population coverage when new surveillance sites are added incrementally to the 22 original 2006–2007 lowa Department of Public Health-influenza-like illness surveillance locations.



There are a number of underlying assumptions. First, influenza does not respect political borders. There is much more activity in Council Bluffs than what can be accounted for by Council Bluffs alone.

Second, we assume travel and mixing are uniform by distance, and not by time (can you find I80 on the map?).

Third, assumes all individuals are equal, and at equal risk for disease: is "maximal coverage" even the right utility function?

Fourth, assumes each sentinel site has unlimited capacity for, *e.g.*, testing, and therefore doesn't ever miss a case regardless of how busy they might be.



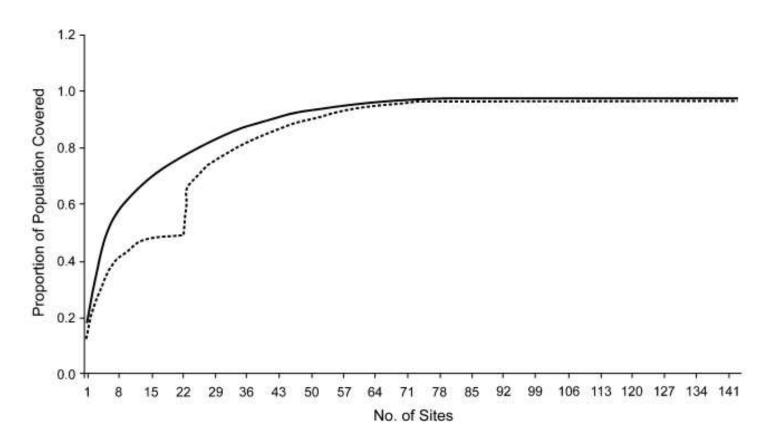


Figure 4. Population coverage (proportion) as a function of additional health care provider surveillance units, for a fixed, prespecified distance of 20 miles (32.2 km), considering the population adjacent to Iowa's border. The upper, solid curve represents the proportion of population coverage as each new surveillance site is added if the sites are selected by the algorithm; the lower, dashed curve represents the proportion of population coverage starting from the 22 currently selected surveillance sites.

University of Iowa COMP COL computational epidemiology research

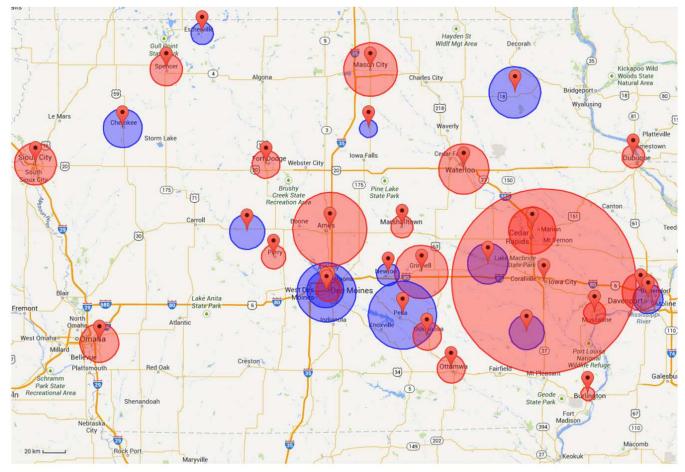


Figure 2 Map comparing 19 existing sites against 19 sites chosen using MCM-NC. The 19 existing sites (blue circles) and 19 sites calculated using the MCM-NC (red circles) are shown together. Circles around each marker indicate the average driving distance between patient homes and provider location in the Medicaid dataset. The large red circle around lowa City represents the University of lowa Hospitals and Clinics; the average driving distance is 45.47 miles (73.18 kilometers). The MCM-NC tends to choose sites in the more densely populated regions in lowa. These sites often contain more reputable hospitals and clinics, and as a result, many people are willing to drive further distances to be seen at these locations. The existing network neglects certain populous regions of lowa (such as Council Bluffs near Omaha) while potentially over-covering other regions (such as Des Moines). Although the Medicaid dataset is used to display average driving distances in this figure, recall that only population data are used to select sites for a network.



A subsequent paper addresses a few more of these issues and poses a slightly different question: if designing a network, how many sites do we need?

Fairchild, Polgreen, Foster, Rushton and Segre (2013), "How Many Suffice? A Computational Framework for Sizing Sentinel Surveillance Networks," in *International Journal of Health Geographics* 12:56



The paper describes a *capacitated maximal coverage* that limits each site to a certain capacity, allowing multiple sites to be co-located if capacity requires it.

It compares both capacitated and non-capacitated MCMs with an alternative kmedian model that minimizes the sum of the distances from each individual to their nearest site.

It uses the Huff metric, a probabilistic geographic measure that captures the chance that a case will be "noticed" in a sentinel site's catchment area to evaluate performance of the network (and not simply maximal coverage).

Finally, it uses simulation to optimize the size of the surveillance network for both intensity and timing of outbreaks based on eight years of Medicaid data and the Huff metric.

