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Social network analysis and modeling of cellphone-based syndromic surveillance data for Ebola in Sierra Leone

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ABSTRACT

Objective: To explore and visualize the connectivity of suspected Ebola cases and surveillance callers who used cellphone technology in Moyamba District in Sierra Leone for Ebola surveillance, and to examine the demographic differences and characteristics of Ebola surveillance callers who make more calls as well as those callers who are more likely to make at least one positive Ebola call.

Methods: Surveillance data for 393 suspected Ebola cases (192 males, 201 females) were collected from October 23, 2014 to June 28, 2015 using cellphone technology. UCINET and NetDraw software were used to explore and visualize the social connectivity between callers and suspected Ebola cases. Poisson and logistic regression analyses were used to do multivariable analysis.

Results: The entire social network was comprised of 393 ties and 745 nodes. Women (*AOR* = 0.33, 95% *CI* [0.14, 0.81]) were associated with decreased odds of making at least one positive Ebola surveillance call compared to men. Women (*IR* = 0.63, 95% *CI* [0.49, 0.82]) were also associated with making fewer Ebola surveillance calls compared to men.

Conclusion: Social network visualization can analyze syndromic surveillance data for Ebola collected by cellphone technology with unique insights.

1. Introduction

Ebola virus disease (EVD) was first identified in 1976 in Zaire but there has been more cases of the disease in other countries in recent time [1]. EVD is characterised by febrile illnesses and is naturally transmitted by animal and vector hosts [2]. The disease is caused by one of the four distinct hemorrhagic fever viruses (HFVs) family: *Flaviviradae, Arenaviradae, Bunyaviradae, and Filoviradae*. *Ebola virus* of the family *Filoviradae* which was responsible for the 2014 West African Ebola outbreak has 5 different virus strains: Sudan virus, Tai Forest virus, Reston virus, Ebola virus, and Bundibugyo virus. The 2014 outbreak is the largest so far and recorded more than 20000 cases and 10000 deaths [3].

Sierra Leone was one of the countries greatly affected by the 2014 West Africa Ebola outbreak [4]. Sierra Leone's first Ebola case was confirmed on 27th May 2014. Prior to 2014 Sierra Leone's preparedness for an EVD outbreak was lacking. The country's EVD response involved the formation of an Ebola technical task force that was responsible for EVD surveillance, case identification, case tracking and monitoring.

Contact tracing, Ebola case identification, treatment, management, and the effective response to both patients and the community have been shown to be effective for EVD surveillance in the past [5]. Correct coordination between Ebola case isolation and treatment, contact tracing and follow-up of contact for 21 days after exposure was very effective in controlling the spread of Ebola outbreak in West Africa in 2014 [6]. Sierra Leone embarked on using cellphone technology to assist with its EVD surveillance in 2014.

Cellphone technology has revolutionized disease surveillance by serving as a channel through which people reveal their public health concerns, locations, and movements. In Haiti, public health experts successfully predicted the spatial evolution of the 2010 cholera outbreak using cellphone calls and SMS messages obtained from more than 2 million mobile phone SIM cards [7].

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Following the 2014 Ebola outbreak, researchers in West Africa called for the use of routine syndromic surveillance systems that rely on data supplied via cellphones. Researchers also recommended the use of cellphones for relaying surveillance data from communities affected by an EVD outbreak to peripheral health centers [8]. In Guinea, smartphones were used to communicate real-time surveillance data for contact tracing and case identification during the 2014 West Africa EVD outbreak [9]. Given the surge in global cellphone usage and the increasing popularity of cellphone-based epidemiological surveillance more research is needed to explore its efficiency and community uptake. Furthermore, few studies have taken advantage of the network-like data generated from cellphone-based surveillance in which callers and cases are interconnected through a web of calls.

This study used social network analysis to evaluate Ebola surveillance data collected in Sierra Leone by cellphone technology. This study specifically used network analysis to explore and visualize the connectivity between suspected Ebola cases and callers who used cellphone technology for Ebola surveillance; to examine demographic differences in surveillance callers who used the cellphone surveillance system; and to determine the characteristics of efficient Ebola surveillance callers (i.e., Ebola surveillance callers whose calls were subsequently confirmed to be Ebola positive).

2. Material and methods

2.1. Study setting and participants

Surveillance data were collected from Moyamba District in southern Sierra Leone from October 2014 to June 2015 using cellphones. The district is approximately 6 902 sq/km with a population of 258 506 [10]. In 2014 the population of Sierra Leone was estimated to be 6 315 627 of which approximately 4 215 000 (70%) had cellphones [11]. The Moyamba District Health Manager Team (MDHMT) Ebola Taskforce, in collaboration with Action Contra la famine (ACF), started cellphone-based syndromic surveillance for Ebola in October 2014. Specifically, community members in Moyamba District were encouraged to call the MDHMT Ebola surveillance hotline to report suspected Ebola cases (dead or alive). Community members who called the center provided their names, telephone number, and the village in which they were residing. The caller also provided the name of the person that they suspected to have Ebola and the person's sex, age and the village of the person that they suspected to have Ebola. The caller could call about any person, regardless of their age, gender, or ethnicity. Within 24 h of receiving a call, the MDHMT Ebola Taskforce would dispatch a Community Health Officer and Ebola contact tracers to the location of the person suspected to have Ebola and transfer him/her to an Ebola treatment center for diagnosis. If the suspected Ebola case was deceased his/her body was transferred to the local mortuary for safe burial. Deceased persons however were not tested for Ebola.

This study is an analysis of Ebola surveillance data collected by the MDHMT Ebola Taskforce from October 2014 through May 2015. The dataset for analysis included callers' names, their telephone numbers, and their village of residence, as well as the cases' name, sex, and location (village), whether the person was sick or dead at the time the surveillance call was made, and if there were sick or dead people at the residence of the suspected Ebola case at the time the call was made to the MDHMT call center. The dataset also included the Ebola lab result for suspected Ebola cases. There were 353 surveillance callers and 393 suspected Ebola cases, including one caller who subsequently became a suspected Ebola case. By cross-referencing names and demographic characteristics of callers and cases, a whole network covering the entire surveillance period was constructed and the connectivity of callers and cases was determined. UCINET software [12] was used to analyze the network of callers and suspected Ebola cases. Specifically, UCINET was used to compute degree centrality. NetDraw [12] was used to visualize the network components in order to depict the social ties among callers and suspected Ebola cases. The analysis was also used to estimate the number of calls made by each caller; this estimate served as the outcome in the multivariate analysis.

2.2. Analysis

SAS 9.2 version [13] was used for descriptive and summary statistical analysis of the characteristics of the callers and suspected Ebola cases. Poisson and logistic regressions were used to determine the factors associated with the number of Ebola surveillance calls made and the likelihood of making at least one positive Ebola surveillance call respectively. Specifically, Poisson regression was used to determine the gender difference in the number of Ebola surveillance calls made controlling in the model for the week in which the surveillance call was made, and the Ebola prevalence of the village in which calls were made. Logistic regression was used to determine the gender difference associated with making at least one positive Ebola surveillance call, controlling in the model for the week in which Ebola surveillance calls were made, whether the person for whom the Ebola surveillance call was made was sick or dead, and the Ebola prevalence of the village in which calls were made.

2.3. Ethics and privacy

The University of Kentucky Institutional Review Board reviewed the protocol for the secondary analyses described in this capstone and determined that it met federal criteria to be exempt.

3. Results

3.1. Descriptive characteristics of suspected Ebola cases

Surveillance data for 393 suspected Ebola cases (192 males, 201 females) were collected from October 23, 2014 to June 28, 2015 using cellphone technology. The descriptive characteristics of suspected Ebola cases, callers, status of Ebola suspected cases, lab results and type of Ebola surveillance call made are presented in Table 1. The average age of the suspected Ebola cases was 23.5 years (standard deviation = 29.5). Three hundred and twenty-four (82.4%) of the suspected Ebola cases were reported sick at the time data was collected while 69 (17.6%) were deceased. Two hundred and twenty (68%) of the sick suspected Ebola cases were females while 104 (32%) were males. Twenty-