



# Predicting COVID-19 using lioness optimization algorithm and graph convolution network

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## Abstract

In this paper, a graph convolution network prediction model based on the lioness optimization algorithm (LsOA-GCN) is proposed to predict the cumulative number of confirmed COVID-19 cases in 17 regions of Hubei Province from March 23 to March 29, 2020, according to the transmission characteristics of COVID-19. On the one hand, Spearman correlation analysis with delay days and LsOA are used to capture the dynamic changes of feature information to obtain the temporal features. On the other hand, the graph convolutional network is used to capture the topological structure of the city network, so as to obtain spatial information and finally realize the prediction task. Then, we evaluate this model through performance evaluation indicators and statistical test methods and compare the results of LsOA-GCN with 10 representative prediction methods in the current epidemic prediction study. The experimental results show that the LsOA-GCN prediction model is significantly better than other prediction methods in all indicators and can successfully capture spatio-temporal information from feature data, thereby achieving accurate prediction of epidemic trends in different regions of Hubei Province.

**Keywords** COVID-19 · Epidemic prediction · Lioness optimization algorithm · Graph convolutional network

## 1 Introduction

Corona Virus Disease 2019 (COVID-19), which emerged in late 2019 and broke out in early 2020, is one of the most severe public health emergencies since the founding of the People's Republic of China. As of 18:00 on November 30, 2021, China has reported a total of 127, 859 confirmed cases and a total of 5697 deaths. Currently, there are still 2913 confirmed cases; a total of 262, 505, 360 confirmed cases and a total of 5, 227, 122 deaths have been reported globally, and there are presently 20, 202, 074 confirmed

cases. Figures 1 and 2 show the distribution of cumulative confirmed cases in China and globally, respectively.<sup>1</sup>

Due to its long duration and wide spread, infectious diseases not only seriously threaten the life safety of people, but also affect people's mental health for a long time. Improper handling can easily cause social chaos. Especially in recent years, sudden epidemic infectious diseases have continued to appear in the world. For example, SARS in 2002 quickly spread to 32 countries and regions in a short time. As of August 16, 2003, the global death toll reached 919, with a case fatality rate of nearly 11%. In 2009, the H1N1 broke out in the United States on a large scale and spread to 214 countries and regions, causing nearly 200, 000 deaths. In addition, there are Ebola virus, HIV, dengue, and so on.

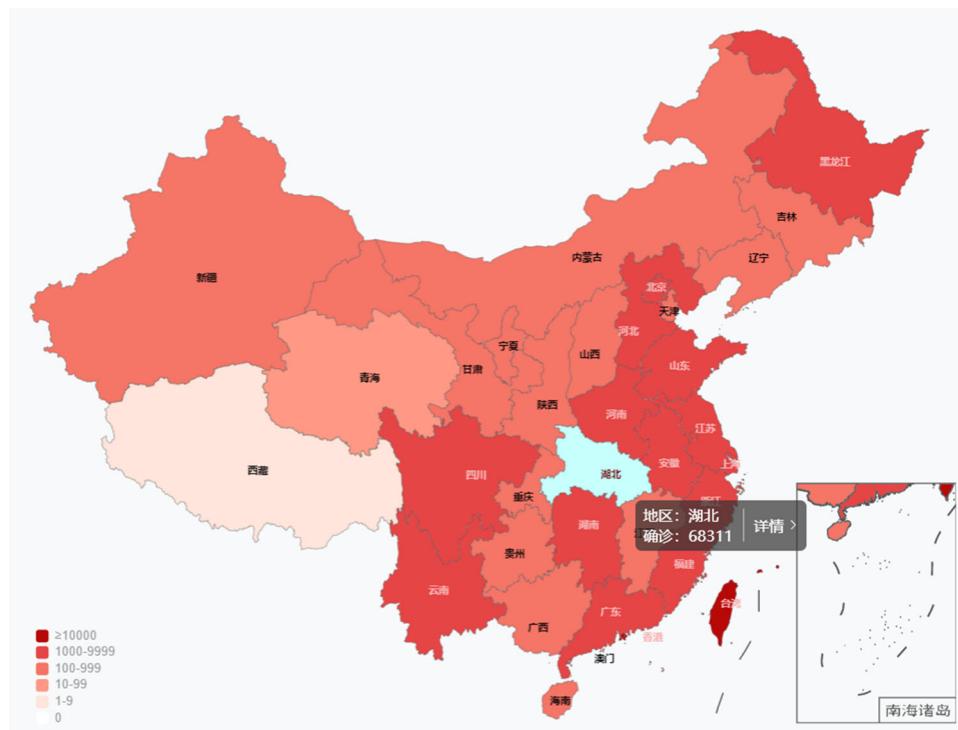
At present, the number of confirmed cases of COVID-19 is still increasing. How to monitor and predict sudden epidemics in a timely and effective manner, and eliminate the crisis in the bud, plays an essential role in the prevention and control of the epidemic. Given these reasons,

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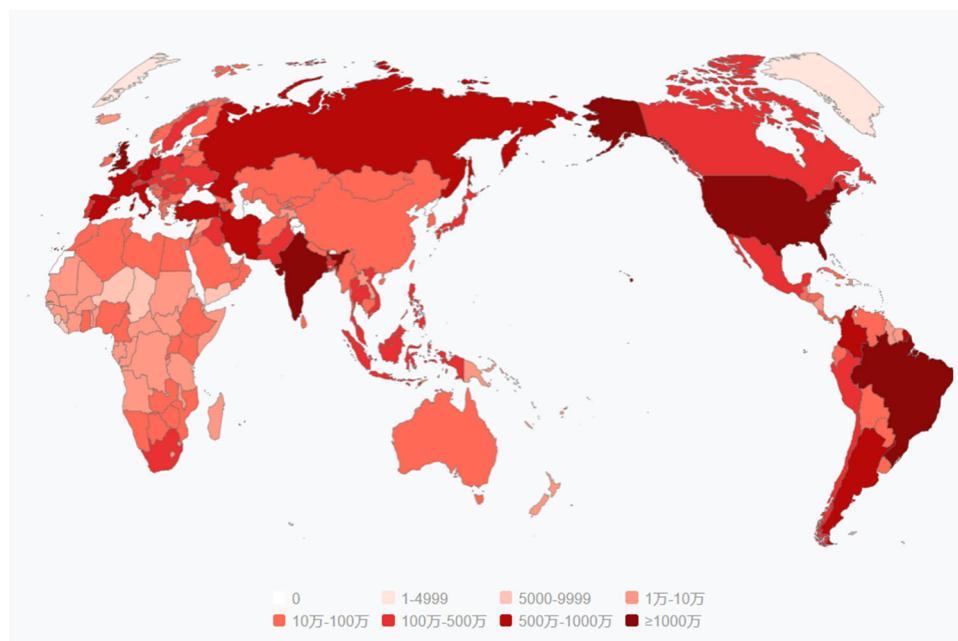
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<sup>1</sup> Source of the picture: [https://voice.baidu.com/act/newpneumonia/newpneumonia/?from=osari\\_aladin\\_banner](https://voice.baidu.com/act/newpneumonia/newpneumonia/?from=osari_aladin_banner).

**Fig. 1** The cumulative number of confirmed cases in China, including cured and deaths



**Fig. 2** The cumulative number of confirmed cases globally, including cured and deaths



domestic and foreign scholars have carried out a series of studies on the prediction of infectious diseases. To sum up, these methods can be divided into the following categories:

(1) Time series prediction model. The traditional time series prediction model is an effective tool for disease prediction, which mainly realizes the analysis of the

spreading law and development trend of the epidemic by fitting epidemic data. Zhang et al. (2016) decomposed the monthly time series data of 11 class C notifiable diseases in China from 2009 to 2014 into a seasonal index representing the seasonal law of infectious diseases and a linear regression model describing the long-term trend of

infectious diseases. And based on this, an autoregressive integrated moving average (ARIMA) model was established to predict the incidence of these infectious diseases. Wang et al. (2019) selected the monthly incidence data of brucellosis from 2004 to 2013 in Jinzhou city, Liaoning Province, China. They analyzed the epidemiological characteristics and prevalence situation by using the autoregressive integrated moving average (ARIMA) model and constructed a prediction model to predict the incidence of brucellosis from 2005 to 2014. Qi et al. (2020) collected the monthly incidence data of Hemorrhagic Fever with Renal Syndrome (HFRS) in Shandong Province from 2009 to 2018. And they used the seasonal autoregressive fractionally integrated moving average (SARFIMA) model to predict the incidence of HFRS. The model evaluation results showed that the SARFIMA model could better fit the dynamic changes of the monthly incidence of HFRS in Shandong Province, and the excellent degree of fitting and predictive ability. Singh et al. (2020) used an autoregressive integrated moving average (ARIMA) model with covariates to fit daily confirmed COVID-19 cases in Malaysia from 22 January to 31 March 2020. Then, they verified this prediction model with confirmed cases from 1 to 17 April 2020. Finally, the daily confirmed cases in Malaysia from 18 April to 1 May 2020 were predicted with satisfaction. Aslan et al. (2022) developed a SEIQR-type deterministic model, which mainly used a system of ordinary differential equations (ODEs) to analyze the dynamics of the epidemic in Hubei Province, and achieved accurate prediction of the local epidemic situation through social distancing rate and confirmed case rate.

(2) Infectious diseases model. In the field of epidemic transmission research, scholars have put forward many models of infectious diseases that can be used to predict the number of infected people, among which the representative ones are: SIR, SIRS, SEIR models, etc. The main idea of this type of model is to divide the population within the epidemic range of infectious diseases into susceptible, exposed, infectious, recovered, etc., and establish and solve differential equations based on the transformation relationship between different groups. Guo et al. (2016) analyzed the transmission data of the Ebola virus by introducing the Jacobian matrix to solve the differential equation of the SIR model and constructed a comprehensive compartment model to fit and predict the transmission trend of the Ebola epidemic. Kuperman et al. (2001) used the SIRS model to study the transmission rules of infectious diseases on the small world network and found that when network nodes change from a recovered state to a susceptible state, infectious diseases will exist in the net-

work for a long time, but the infection rate is low. Ding et al. (2004) used the SIJR model to analyze SARS epidemic data in Hong Kong, Singapore, and Canada and obtained essential parameters such as the epidemic transmission rate, and then realized the prediction of SARS transmission by adjusting the parameters in this model. Merler et al. (2011) and Wu et al. (2020) both established the SEIR model and included “exposed” into the simulated population to simulate the development trend of H1N1 in Europe and the COVID-19 in major cities of China, respectively. In addition, there are still many scholars who extend the infectious disease model by adding or adjusting the internal structure (Wang et al. 2022a, b; Yi et al. 2021; Liao et al. 2020; Cao et al. 2020; Bi et al. 2021). Liu et al. (2020) proposed a susceptible-asymptomatic-infected-removed (SAIR) model based on the social network to describe the spread of COVID-19 and analyzed the real epidemic data in Wuhan and finally selected an appropriate method to control the spread of COVID-19. In the classical SIR model, the transmission speed is a constant. However, in reality, the transmission speed will change through various interventions such as personal protection, community lockdowns, and urban control. Therefore, Jia et al. (2020) added a transmission modifier to the SIR model, established a dynamic extended SIR model (eSIR) for infectious diseases, which can dynamically change the transmission speed and predict the epidemic trend of COVID-19 in Italy. In view of the fact that the SEIR model does not consider the influence caused by random factors such as population migration and flow between regions, Yang et al. (2020) introduced input/output flows on the basis of the SEIR model, making the modified SEIR (M-SEIR) model more suitable for the actual population change when it is used to predict the urban epidemic. Wang et al. (2022a, b) introduced the complex network model and the intelligent agent on the basis of M-SEIR model, and proposed a city-level structured prediction and evaluation (CSPE) model for COVID-19 epidemic. The model has effectively deduced the epidemic situation in Hubei Province. For the COVID-19 pandemic, Manuel et al. (2020) also proposed a susceptible-exposed-symptomatic-infectious-asymptomatic infectious-recovered (SEIAR) model considering population and disease mortality based on the SEIR model, which aims to reduce the number of infectors by assessing the isolation situation.

(3) Machine learning prediction method. In recent years, many machine learning algorithms have also begun to be used for the prediction of infectious diseases. Wang et al. (2017) used the annual county-level human brucellosis (HB) cases and data of 37 environmental variables in Inner

Mongolia, China, to build an artificial neural network (ANN) model, which was used to determine essential predictors of human brucellosis and predict the annual county-level cases of human brucellosis. Chen et al. (2018) introduced different predictors for different diseases, countries, and prediction windows, established different LASSO prediction models, and analyzed the importance of different predictors under various conditions by evaluating the prediction performance of these models. Research results showed that different predictors have different effects in different situations, and short-term forecasts usually perform better than long-term forecasts. Hasan (2020) proposed a hybrid model combining ensemble empirical mode decomposition (EEMD) and artificial neural network (ANN). This model first used EEMD to decompose the epidemic time series data and then sent these decomposed IMF components into the neural network model for training. Eventually, the trained neural network was used to predict the trend of COVID-19. Fang et al. (2020) established a random forest model based on meteorological factors to predict the incidence of infectious diarrhea in Jiangsu Province, China. The prediction model better fitted the dynamic characteristics of the disease epidemic. Gumaei et al. (2021) used gradient boosting regression (GBR) to build a training model to predict the total number of daily confirmed COVID-19 cases around the world. Xie et al. (2021) proposed a nonlinear time-varying transmission rate model based on support vector regression in order to overcome the shortcomings of a single model for insufficient extraction of effective information and low prediction accuracy, and applied it to long-term prediction of COVID-19 epidemic in Hubei Province. Smith and Alvarez (2021) took COVID-19 patients in Wuhan, Hubei, China, as the research object and applied a series of machine learning models to analyze the factors affecting the death of COVID-19 patients. They finally obtained the conclusion that the Shapley value is helpful to determine the death factors of COVID-19.

(4) Deep learning prediction method. In recent years, deep learning methods have performed well in fields such as data mining, analysis, and prediction. They have provided new technologies and techniques for studying epidemic transmission. Chae et al. (2018) used search query data, social media big data, and weather data to sort out the factors that affect the occurrence of infectious diseases to build OLS, ARIMA, DNN, and LSTM prediction models and compare their prediction performance. The study found that the prediction model using deep learning is more suitable for predicting the trend of infectious diseases. Wang et al. (2020) used the LSTM network and rolling update mechanism to continuously input new prediction

results into the next iteration model for training, and finally made a long-term prediction of epidemic trends in Russia, Peru, and Iran. In this article, the authors also introduced the Diffusion Index (DI) to evaluate the effectiveness of preventive measures such as social isolation and lockdown. Hao et al. (2020) used Elman neural network, LSTM and SVM to predict the cumulative confirmed cases, deaths, and cured cases in Wuhan, Hubei Province, and used SVM with fuzzy granulation to predict the confirmed new cases, new deaths and new cured cases. Finally, they analyzed the applicability of different methods. Rauf et al. (2021) proposed to use long short-term memory (LSTM) network, Recurrent Neural Network (RNN), and Gate Recurrent Unit (GRU), while considering time variables and data nonlinearity, to evaluate each model feature to predict the number of confirmed COVID-19 cases in the Asia-Pacific region in the next 10 days. Chen et al. (2021) developed a data-driven workflow to extract, process, and develop deep learning methods, and built multivariate long and short-term memory (LSTM) models with different architectures to predict the COVID-19 epidemic, thereby supplementing existing modeling methods.

In addition, when predicting the incidence trend of specific infectious diseases, not all features should be included in the prediction model. Sometimes, you can get better prediction results by deleting some unimportant features. Meta-heuristic algorithm is a common feature selection method, which has been widely concerned by scholars (Jia et al. 2019; Turabieh et al. 2021; Samieyan et al. 2022). Depending on the source of inspiration, meta-heuristic algorithms can be divided into the four categories: (1) biological evolutionary algorithms that simulate the process and mechanism of biological evolution in nature. Among these algorithms, the most widely used is genetic algorithm (GA)(Holland 1992) with the basic operations of selection, crossover and mutation; in addition, there are differential evolution (DE), evolutionary strategy (ES), etc.; (2) algorithms inspired by physical phenomena and laws in nature, such as multiverse optimizer (MVO) inspired by three concepts in cosmology: white hole, black hole and wormhole, simulating the transfer process of matter in the universe from white hole to black hole through the wormhole (Mirjalili et al. 2015); gravitational search algorithm (GSA) based on Newton's law of universal gravitation (Rashedi et al. 2009); lightning search algorithm (LSA) based on the lightning mechanism (Shafeef et al. 2015), etc.; (3) algorithms that simulate various human behaviors in society, such as teaching–learning-based optimization (TLBO) that simulates the two stages of “teaching” and “learning” in the learning process to improve the ability of each individual(Rao et al. 2011);

heap-based optimizer (HBO) that uses the heap data structure to map corporate rank hierarchy to simulate three human behaviors(Askari et al. 2020), etc.; (4) swarm intelligence algorithms that simulate the social behavior of animal groups in nature, such as particle swarm optimization (PSO) designed to simulate the predatory behavior of birds(Eberhart and Kennedy 1995); whale optimization algorithm (WOA) proposed by imitating the hunting behavior of humpback whales(Mirjalili and Lewis 2016); chimp optimization algorithm (ChOA) for modeling hunting behavior of chimps according to differences in intelligence and sexual motivation(Khishe and Mosavi 2020); In addition, there are recently proposed marine predators algorithm (MPA)(Faramarzi et al. 2020), k-means clustering optimizer (KO)(Minh et al. 2022), etc. The principles of these algorithms are different, and their abilities to solve different types of problems are also different. As the “no free lunch (NFL)” theorem states, no algorithm can solve all types of optimization problems with equal simplicity and efficiency (Wolpert and Macready 1997). Therefore, we think it is necessary to develop new optimization algorithms to better achieve the purpose of feature selection.

In conclusion, many scholars have made great efforts in infectious disease prediction including COVID-19 prediction, and the results have been fruitful. However, by analyzing the current situation of infectious disease prediction research, we found that the existing models still have the following problems:

- (1) Some models are constructed based on the historical data of a single region, ignoring the spatial correlation of infectious disease transmission, which affects the improvement of the model prediction accuracy;
- (2) Some prediction models only use directly related data, such as the number of confirmed cases and the number of isolated people, which makes the prediction results unable to timely reflect the influence of other factors (such as social environment and government response policies) in the process of infectious disease transmission, and finally makes the prediction accuracy difficult to be satisfied;
- (3) Some models do not consider the time lag effect of eclipse period on transmission. Virus infection can have a certain eclipse period (the time from contact with the source of transmission to the onset of symptoms), and even asymptomatic patients can appear. Ignoring the time lag will also affect the prediction performance;
- (4) Finally, when there are many feature data input into the model, there may be some irrelevant features. If these features are forcibly incorporated into the prediction

model without selection, it will not only increase the computation, but also affect the prediction performance of the prediction model.

In order to solve the above problems, this paper proposed a graph convolution network prediction model based on the lioness optimization algorithm (LsOA-GCN) according to the transmission characteristics of the COVID-19. The main contributions are as follows:

- (1) A novel population-based lioness optimization algorithm was proposed to filter the feature data with time delay (Baidu index), and based on this, a feature matrix for epidemic prediction was constructed;
- (2) A COVID-19 prediction model, LsOA-GCN, which comprehensively uses the lioness optimization algorithm and graph convolutional network (GCN) (Kipf and Welling 2017), was proposed;
- (3) Experiments were conducted on epidemic data from 17 regions in Hubei Province, China, and compared with 10 prediction methods, which verified the superiority of the LsOA-GCN prediction model.

The rest of this paper is composed as follows. The second part introduces the LSOA-GCN prediction model in detail, including problem definition, adjacency matrix construction, data source and model construction, etc. The third part compares and analyzes the performance of different prediction methods through model evaluation indicators and statistical test methods. The last part summarizes the proposed prediction model.

## 2 LsOA-GCN model

### 2.1 Problem definition

(1) City network graph: $G$ . In this paper,  $G = (V, E)$  is used to describe the topological structure of Hubei Province's city network. We regard each region in Hubei Province as a node in the network graph, where,  $V = \{v_1, v_2, \dots, v_n\}$  represents the set of city nodes in Hubei Province,  $n$  is the total number of nodes, and  $E$  represents the set of edges between cities.

(2) Adjacency matrix:  $A^{n \times n}$ . The spatial information between cities is mapped into the city network graph  $G$ , and the degree of correlation between city nodes is represented by edges. This paper intends to use the migration relationship graph to reflect the degree of mutual influence between various regions in Hubei Province, which is represented by  $G = (V, A)$ , where  $A \in R^{n \times n}$  is the adjacency matrix.

(3) Feature matrix:  $X^{n \times t \times d}$  We use the Baidu Search Index value of keywords related to the epidemic as the attribute features of the nodes in the network, which was represented by  $X \in R^{n \times t \times d}$ , where  $t$  is the number of days, and  $d$  is the dimension of the node attribute features. For example,  $X_t \in R^{n \times i}$  represents the Baidu Search Index value of  $i$  keywords in  $n$  regions on the  $t$ -th day.

Therefore, the epidemic prediction problem solved in this paper can be expressed as: under the premise of a given city network topology graph  $G$  and the feature matrix  $X$ , we planned to predict the cumulative number of confirmed cases in each region of Hubei province in the future through graph convolution network. The formula is as follows:

$$[Y_{t+1}, \dots, Y_{t+T}] = GCN(G; (X_{t-s}, \dots, X_{t-1}, X_t)) \quad (1)$$

where  $s$  is the length of the historical time series,  $T$  is the length of the future time series that needs to be predicted,  $Y_{t+1}$  represents the cumulative number of confirmed cases in each region of Hubei Province on the  $(t + 1)$ -th day, and  $X_t$  represents the feature vector of each region on day  $t$ .

## 2.2 Construction of adjacency matrix

Since the outbreak of the COVID-19 has the characteristic of “person-to-person transmission”, this paper assumes that the large-scale population movement during the epidemic will have a certain impact on the spread of the epidemic. And COVID-19 has a certain eclipse period. During the eclipse period, some patients have only mild respiratory symptoms, such as fever, fatigue, dry cough, etc., but they are also infectious, and there are even asymptomatic patients. Based on this, this paper collected the population migration index values of 17 regions in Hubei province from January 10, 2020 to March 15, 2020 (a total of 66 days) and performed a time-delayed Spearman correlation analysis with the cumulative confirmed cases (from January 24, 2020 to March 29, 2020) in each region to construct an adjacency matrix.

$$A_{ij} = ave |(spearman(x_i, y_{ji}))| \quad i, j \in [1, 17] \quad (2)$$

where the Spearman correlation coefficient

$$\rho = \frac{\sum_{i,j} (x_i - \bar{x})(y_{ji} - \bar{y})}{\sqrt{\sum_i (x_i - \bar{x})^2 \sum_{i,j} (y_{ji} - \bar{y})^2}}, \quad x_i \text{ represents the cumulative}$$

number of confirmed cases in region  $i$ , and  $y_{ji}$  represents the population migration index from region  $j$  to region  $i$ .

## 2.3 Data source

The epidemic data in this paper come from the daily epidemic notification provided by the National Health Commission ([http://www.nhc.gov.cn/xcs/yqtb/list\\_gzbd.shtml](http://www.nhc.gov.cn/xcs/yqtb/list_gzbd.shtml)) and the COVID-19 epidemic situation in Hubei Province provided by the Hubei Health Committee (<https://wjw.hubei.gov.cn/>) to collect the cumulative number of confirmed COVID-19 cases in 17 regions of Hubei Province from January 24, 2020, to March 29, 2020.

The migration data used to construct the adjacency matrix in this paper come from Baidu Maps Huiyan (<http://qianxi.baidu.com/>) developed by Baidu, and we collected and processed the migration scale index of 17 regions in Hubei Province from January 10, 2020 to March 15, 2020. The keyword data used to construct the feature matrix is also derived from the Baidu Search Index (<https://index.baidu.com/v2/index.html#/>) provided by Baidu. Set the location parameter as “cities in Hubei Province”, the time parameter as “2020-01-10 ~ 2020-03-15”, and the source parameter as “PC+ Mobile”.

## 2.4 Model construction

### 2.4.1 Overall overview

In this section, we give a comprehensive overview of how to use the LsOA-GCN model for epidemic prediction. The LsOA-GCN model is mainly composed of two parts. One is the construction of the feature matrix for capturing temporal information, the other is the graph convolution operation for capturing spatial information. First, the keyword data was sorted out through Spearman correlation analysis with a time lag, and the initial feature matrix was constructed to capture the time series information between the current number of confirmed cases and the Baidu Search Index at different periods in the past. Then LsOA was used to further filter the feature matrix to improve the accuracy of the model and avoid unnecessary feature interference. Finally, the feature matrix was used as input to obtain the spatial features of the epidemic-related data through the graph convolution network, to get the prediction results.

### 2.4.2 Capture temporal information

In an information society, with the continuous innovation and development of PC and mobile terminals, the Internet has become the primary and critical way for people to

collect information from the outside world. Netizens can use various information search platforms to gather the information they want to know and obtain real-time data on the network. According to the 47th “Statistical Reports on Internet Development in China” released by the China Internet Network Information Center (CNNIC), as of December 2020, there have been 989 million Chinese netizens, an increase of 85.4 million compared with March 2020. The Internet has become an essential channel for the public to obtain information. Among the many search platforms, Baidu is the most popular information search platform, with nearly 90% of users in China, accounting for a significant market share. Among them, Baidu Index is a data sharing platform based on massive netizens’ behavior data in Baidu. Through the Baidu Index, we can study keyword search trends, gain insights into the interests and needs of netizens, monitor public opinion trends, and locate audience features. In recent years, Baidu Search Index has been used to predict some epidemic diseases, such as dengue (Li et al. 2017), HIV/AIDS (Huang et al. 2020) and Brucellosis (Zhao et al. 2020a). Therefore, it is scientific and feasible to use Baidu Search Index as a feature to predict the COVID-19.

During the epidemic, there is a complicated time series correlation between the current number of confirmed cases and the Baidu Search Index at different periods in the past. Therefore, obtaining the temporal features is also a key issue in epidemic prediction. The first step to obtain time information is the selection of keywords. The selection methods include empirical word selection, technical word selection (Ginsberg et al. 2009), and range word selection (Alessa and Faezipour 2018). The keyword selection used to construct the feature matrix this time mainly adopted the range word selection method, supplemented by the empirical word selection method. Firstly, the selection range of keywords should be roughly determined according to the features of the epidemic, transmission routes, protective measures, etc. Then, relevant words can be recommended through the demand graph function in Baidu Index, while reducing the workload, as far as possible to ensure the completeness of the epidemic-related vocabulary. Based on these, we set up a keyword database from the four dimensions of prevention, symptoms, common words and other related aspects of COVID-19, and initially selected 40 specific keywords, namely “vaccine”, “hand washing”, “mask”, “disinfection”, “nucleic acid”, “flu”, “diarrhea”, “dry cough”, “pneumonia”, “cough”, “fever”, “dyspnea”, “high temperature”, “stuffy nose”, “asymptomatic infection”, “close contact”, “suspected case”, “aerosol transmission”, “infection”, “novel coronavirus”, “NCP”, “SARS-CoV-2”, “COVID-19”, “novel

coronary pneumonia”, “novel coronavirus pneumonia”, “virus pneumonia”, “coronavirus”, “novel”, “epidemic”, “Center for Disease Control and Prevention”, “Red Cross Society”, “Spring Festival”, “Health Commission”, “SARS”, “Zhong Nanshan”, “children”, “medical”, “government”, “Wuhan” and “lockdown of the city”.

The specific construction steps of the feature matrix are as follows:

- (1) Establish a keyword database from the 4 dimensions mentioned above, and initially screen out 40 keywords;

- (2) Time-delayed Spearman correlation analysis is performed between Baidu Index (search index, information index) of 40 keywords and the cumulative number of COVID-19 cases;

- (3) Further screen the keyword database and select keyword data with a correlation coefficient greater than 0.5;

- (4) Collect keyword data according to the delay days corresponding to the maximum correlation coefficient of each keyword in the thesaurus, and construct a feature matrix based on this;

- (5) When dealing with practical problems, not all features must be included in the prediction model, and sometimes deleting some unimportant features may improve the performance of the model. Feature selection is a common data dimensionality reduction method, which has been widely concerned by scholars. They regard feature selection as a 0–1 problem and usually choose the meta-heuristic algorithm to solve (Guo et al. 2022; Jia et al. 2022; Sha et al. 2020). Inspired by this, we used the swarm intelligence optimization algorithm to further filter the feature matrix. Its main purpose is to delete the feature data that is not related to the problem in the original feature matrix, reduce the feature dimension and reduce the running time. This point will be explained in detail in the next part.

#### 2.4.3 Filter feature matrix

In recent years, various intelligent optimization algorithms have emerged one after another, such as the classic Particle Swarm Optimization (Eberhart and Kennedy 1995), Whale Optimization Algorithm (Mirjalili and Lewis 2016), and the newly developed Marine Predator Algorithm (Faramarzi et al. 2020), etc. However, when dealing with practical problems, these algorithms will converge prematurely and easily fall into the risk of local optimum, making it difficult to solve all practical application problems (Faramarzi et al. 2020).

Therefore, we proposed a population-based meta-heuristic optimization algorithm, Lioness Optimization

Algorithm (LsOA), to further filter the feature matrix to avoid the interference of unnecessary features. The lion-inspired optimization algorithm appeared in 2012: The Lion's Algorithm (2012) (Rajakumar 2012), which mainly simulates the three behaviors of lion mating, territory takeover, and territory defense, to find optimal solution; The second is Lion pride optimizer (2022) (Wang et al. 2012), which is developed around the mating behavior of lions, and on this basis, three evolutionary strategies are proposed. Survival of the fittest is its main idea; The third chapter is the Lion Optimization Algorithm (2016) (Yazdani and Jolai 2016), which not only draws on the mating and competition behaviors of the lions, but also introduces hunting, migration, patrol and other mechanisms. It can be seen that although the evolution process of the above algorithms incorporate the natural behaviors of lions hunting and mating, they fail to consider some of the core optimization mechanisms in the lion hunting process, resulting in their shortcomings such as slow convergence speed and difficulty in finding global extreme value. Aiming at the shortcomings of the above algorithms, this paper proposed LsOA. Because the optimization process of the algorithm is closer to the hunting behavior of the lions, the lioness is the main performer of the lion hunting. Inspired by this, this algorithm mainly focuses on the hunting behavior of lionesses, and comprehensively considers the two mechanisms of team hunting and elite hunting. During the period, Levy flight and Brownian motion are also introduced. While retaining the strong global exploration ability of the original lion pride algorithms, the local mining ability of the algorithm is improved, which can solve various mathematical optimization problems more effectively, which is completely different from the previous algorithms.

In LsOA, the prey is the global optimal solution, and the current optimal solution is considered to be lioness A or the top lioness. The lions mentioned are all candidate solutions, and they are all lionesses by default. LsOA is mainly composed of the following six mechanisms.

#### (1) Track prey.

In the hunting process, the lions tend to spread out into a fan shape to surround the prey in the center, and the group of lions in the “center” of the fan shape will consciously adjust their positions according to the positions of other lions and prey. The mathematical model of this behavior is shown in Eqs. (3) and (4), where Eq. (3) defines the step length and direction of the current lion group to the target position, and Eq. (4) defines the position of the lion group after the update.

$$\vec{D} = \left| \vec{C} \cdot \overrightarrow{\text{Prey}} - \vec{X}(t) \right| \quad (3)$$

$$\vec{X}(t+1) = \overrightarrow{\text{Prey}} - \vec{A} \cdot \vec{D} \quad (4)$$

where  $t$  is the current iteration number,  $\vec{X}(t)$  is the position vector of the lioness,  $\vec{D}$  is the distance between the current lioness and the prey, represents the product,  $\overrightarrow{\text{Prey}}$  is the position vector of the prey,  $\vec{A}$  and  $\vec{C}$  are coefficient vectors, and their calculation formulas are as follows:

$$\vec{A} = 2\vec{a} \cdot \vec{r}_1 - \vec{a} \quad (5)$$

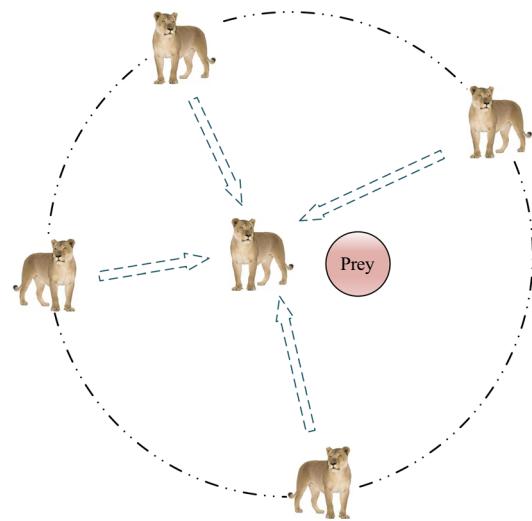
$$\vec{C} = 2\vec{r}_2 \quad (6)$$

where  $\vec{a} = 2 - \frac{2\text{Iter}}{\text{Max\_iter}} \in [0, 2]$ , to simulate approaching the prey, as the iteration progresses, the value of  $\vec{a}$  decreases linearly from 2 to 0, and  $\vec{A} \in [-\vec{a}, \vec{a}]$ . If  $|\vec{A}| < 1$ , it will attack the prey and focus on exploitation at this time; if  $|\vec{A}| > 1$ , it will force to stay away from the prey and focus on exploration at this time.  $\vec{C} \in [0, 2]$  is a random value, which represents the weight value of the influence of the position of the current optimal solution on the updated position.  $\vec{r}_1, \vec{r}_2$  are both random vectors between  $[0, 1]$ .

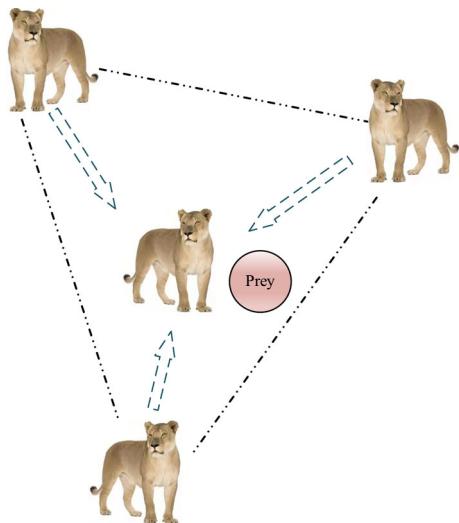
#### (2) Capture prey.

When capturing prey, we believe that the location of the target prey should be near the “central circle” of the hunting team. And this center circle is composed of the four best candidate solutions in the population (lioness A, lioness B, lioness C, and lioness D) and the average of their four. The mathematical model is as follows:

$$\begin{aligned} \vec{D}_A &= \left| \vec{C}_1 \cdot \vec{X}_A - \vec{X} \right|, \vec{D}_B = \left| \vec{C}_2 \cdot \vec{X}_B - \vec{X} \right| \\ \vec{D}_C &= \left| \vec{C}_3 \cdot \vec{X}_C - \vec{X} \right|, \vec{D}_D = \left| \vec{C}_4 \cdot \vec{X}_D - \vec{X} \right| \end{aligned} \quad (7)$$



**Fig. 3** The construction process of the “central circle”



**Fig. 4** The construction process of the elite matrix

$$\vec{X}_a = \vec{X}_A - \vec{A}_1 \cdot (\vec{D}_A), \vec{X}_b = \vec{X}_B - \vec{A}_2 \cdot (\vec{D}_B) \quad (8)$$

$$\vec{X}_c = \vec{X}_C - \vec{A}_3 \cdot (\vec{D}_C), \vec{X}_d = \vec{X}_D - \vec{A}_4 \cdot (\vec{D}_D)$$

$$\vec{X}_{ave} = \frac{\vec{X}_a + \vec{X}_b + \vec{X}_c + \vec{X}_d}{4} \quad (9)$$

where the calculation of  $A_i$ ,  $C_i(i=1, 2, 3, 4)$  are shown in Eq. (5) and Eq. (6),  $\vec{X}_A, \vec{X}_B, \vec{X}_C, \vec{X}_D$  are the positions of lioness A, lioness B, lioness C and lioness D, respectively,  $\vec{X}_{ave}$  is the average of  $\vec{X}_a, \vec{X}_b, \vec{X}_c, \vec{X}_d$ , and  $\vec{X}$  is the position of the current prey.

### (3) “Central Circle” and Elite Matrix.

#### “Central Circle”:

According to Eqs. (8) and (9), the positions of the lions at the center of the hunting team can be determined. It is updated based on the positions of the 4 lionesses closest to the prey in the current population, which are  $\vec{X}_a, \vec{X}_b, \vec{X}_c, \vec{X}_d$ , respectively, and the last candidate is the average  $\vec{X}_{ave}$  of these four candidates. These five positions constitute the hunting team’s “Center circle”:

$$E\_Team = [\vec{X}_a; \vec{X}_b; \vec{X}_c; \vec{X}_d; \vec{X}_{ave}] \quad (10)$$

Figure 3 shows the construction process of the “central circle”. Since the position of the target prey in the search space is not known a priori, this article assumes that the position of the target prey may be located near any of these five positions, and the probability of the five candidates being selected is the same, which is 0.2.

#### Elite Matrix:

The elite matrix is designed based on the elite hunting

mechanism. At this time, the population evolves into some agents of the elite, through which agents are used to continuously test the direction and position of the elite, and finally achieve the goal of capturing the prey.

In designing the elite hunting mechanism, this paper did not use a single lioness to carry out optimization iterations, but took the elite hunting mechanism as inspiration, combined with the swarm intelligence mechanism to design: that is, according to the results of the population iteration, the position of the elite is determined on the basis of the lioness of a specific size, and then the optimization iteration is carried out under the guidance of the group search mechanism.

Figure 4 shows the construction process of the elite matrix. In order to avoid the risk of falling into a local optimum caused by relying only on a single elite position, here we draw on the principle of triangular stability, and replace the common method for determining the position of the elite from the common last-round best to a joint decision by the top three agent positions (lioness A, lioness B, and lioness C) in the last-round fitness value. The calculation formula is as follows:

$$\text{Top\_lioness\_pos} = \frac{\vec{X}_A + \vec{X}_B + \vec{X}_C}{3} \quad (11)$$

where  $\vec{X}_A, \vec{X}_B, \vec{X}_C$  are the position vectors of lioness A, lioness B and lioness C, respectively. After the position of the top lioness is determined, copy it  $n$  times to build the elite matrix:

$$\text{Elite\_lioness} = \begin{bmatrix} X_{1,1}^I & X_{1,2}^I & \cdots & X_{1,d}^I \\ X_{2,1}^I & X_{2,2}^I & \cdots & X_{2,d}^I \\ \vdots & \vdots & \ddots & \vdots \\ \vdots & \vdots & \ddots & \vdots \\ X_{n,1}^I & X_{n,2}^I & \cdots & X_{n,d}^I \end{bmatrix}_{n \times d} \quad (12)$$

where  $\overrightarrow{X^I}$  is the position vector of the top lioness in the lion group.

### (4) The march strategy.

With the increasing number of optimization iterations, the risk of the lions falling into local optimization is increasing. In order to enable the lions to quickly explore a new area, this paper designed the march strategy. The implication is that when  $r2 \leq M$ , unifies all dimensions of the predator into a unique value (march). Otherwise, each dimension will randomly select a value in the  $E\_Team$  to be updated (random). The formulas are as follows:

$$\text{Prey}(i,j) = E\_Team(\text{randi}(\text{size}(E\_Team, 1)), :), \text{IF}(r2 > M) \quad (13)$$

$$\text{Prey}(i,:) = E\_Team(\text{randi}(\text{size}(E\_Team, 1)), :), \text{IF}(r2 \leq M) \quad (14)$$

where  $\text{randi}()$  is used to generate pseudo-random integers;  $\text{size}()$  is used to return the size of the vector.  $M$  indicates the degree of influence of the march strategy on the process. The prey here is a matrix with the same dimension as the elite matrix.

#### (5) Phase-focused mechanism.

In order to avoid premature convergence of LsOA and in reference to the behavioral process of lion hunting, the whole iteration process was divided into three stages in this paper, namely the early iteration, the middle iteration, and the late iteration (Faramarzi et al. 2020). And different search mechanisms were used in each stage.

##### ① the early iteration.

While  $\text{Iter} < \frac{1}{3} \text{Max\_iter}$

$$\begin{aligned} \overrightarrow{\text{step}}_i &= \overrightarrow{\text{RB}} \otimes (\overrightarrow{\text{Elite\_lioness}}_i - \overrightarrow{\text{RB}} \otimes \overrightarrow{\text{Prey}}_i) i = 1, \dots, n \\ \overrightarrow{\text{Prey}}_i &= \overrightarrow{\text{Prey}}_i + P \cdot \vec{R} \otimes \overrightarrow{\text{step}}_i \end{aligned} \quad (15)$$

where  $\text{Iter}$  is the current iteration number and  $\text{Max\_iter}$  is the maximum iteration number.  $\overrightarrow{\text{RB}}$  is a vector of random numbers representing Brownian motion, the symbol  $\otimes$  means multiplication term by term.  $P = 0.5$ ,  $R$  is a random vector in  $[0, 1]$ .

##### ② the middle iteration.

While  $\frac{1}{3} \text{Max\_iter} < \text{Iter} < \frac{2}{3} \text{Max\_iter}$ .

$$\begin{aligned} \overrightarrow{\text{step}}_i &= \overrightarrow{\text{RB}} \otimes \left( \overrightarrow{\text{RB}} \otimes \overrightarrow{\text{Elite\_lioness}}_i - \overrightarrow{\text{Prey}}_i \right) i = n/2, \dots, n \\ \overrightarrow{\text{Prey}}_i &= \overrightarrow{\text{Elite\_lioness}}_i + P \cdot \text{CF} \otimes \overrightarrow{\text{step}}_i \end{aligned} \quad (17)$$

where  $\text{CF} = \left(1 - \frac{\text{Iter}}{\text{Max\_Iter}}\right)^{\left(2 \frac{\text{Iter}}{\text{Max\_Iter}}\right)}$ , it is regarded as an adaptive parameter that controls the step length of the lion's movement.  $\overrightarrow{\text{RB}} \otimes \overrightarrow{\text{Elite\_lioness}}_i$  simulates the behavior of a predator (lioness).

##### ③ the late iteration.

While  $\text{Iter} > \frac{2}{3} \text{Max\_iter}$

$$\begin{aligned} \overrightarrow{\text{step}}_i &= \overrightarrow{\text{RL}} \otimes \left( \overrightarrow{\text{RL}} \otimes \overrightarrow{\text{Elite\_lioness}}_i - \overrightarrow{\text{Prey}}_i \right) i = 1, \dots, n \\ \overrightarrow{\text{Prey}}_i &= \overrightarrow{\text{Elite\_lioness}}_i + P \cdot \text{CF} \otimes \overrightarrow{\text{step}}_i \end{aligned} \quad (18)$$

where  $\overrightarrow{\text{RL}} \otimes \overrightarrow{\text{Elite\_lioness}}_i$  simulates the behavior of a predator (lioness).

#### (6) The impact of FADs.

FADs (Filmalter et al. 2011) is called “fish aggregating devices”, which is a kind of raft-like structure, usually made of some waste products. This kind of device is easy to attract and gather a large number of fish, making it difficult for them to escape. There are similar facilities (regions) on the African savannah, such as wetlands or swamps. When lions hunt here, they often get caught in them and cannot get out.

Given this point, this paper introduced FADs into the algorithm. FADs is considered to be local optima, and the effect is that lions are trapped in these areas. The FADs effect is expressed mathematically as:

$$\overrightarrow{\text{Prey}}_i = \begin{cases} \overrightarrow{\text{Prey}}_i + \text{CF} [\vec{X}_{\min} + \vec{R} \otimes (\vec{X}_{\max} - \vec{X}_{\min})] \otimes \text{Ma} & \text{if } r \leq \text{FADs} \\ \overrightarrow{\text{Prey}}_i + [\text{FADs}(1 - r) + r] (\overrightarrow{\text{Prey}}_{r1} - \overrightarrow{\text{Prey}}_{r2}) & \text{if } r > \text{FADs} \end{cases} \quad (19)$$

For prey:

$$\begin{aligned} \overrightarrow{\text{step}}_i &= \overrightarrow{\text{RL}} \otimes \left( \overrightarrow{\text{Elite\_lioness}}_i - \overrightarrow{\text{RL}} \otimes \overrightarrow{\text{Prey}}_i \right) i = 1, \dots, n/2 \\ \overrightarrow{\text{Prey}}_i &= \overrightarrow{\text{Prey}}_i + P \cdot \vec{R} \otimes \overrightarrow{\text{step}}_i \end{aligned} \quad (16)$$

where  $\overrightarrow{\text{RL}}$  is the random number vector representing Levy's flight and  $\overrightarrow{\text{RL}} \otimes \overrightarrow{\text{Prey}}_i$  simulates the behavior of prey. For lions, this study assumed:

where  $\text{FADs} = 0.2$ , which means the probability that FADs will affect the optimization process.  $r$  is a uniform random number in  $[0, 1]$ .  $r1, r2$  represents the random index of the prey matrix.  $\vec{X}_{\max}, \vec{X}_{\min}$  are the maximum and minimum values of each dimension, respectively.  $\text{Ma}$  is a binary vector containing only 0 and 1. First, generate a random vector in  $[0, 1]$ , if it is less than 0.2, it will be regarded as 0; otherwise, it will be regarded as 1. The pseudo-code of LsOA is as follows:

---

```

1 Initialize search agents(Prey) population  $i = 1, \dots, n$ 
2 Assign free parameters:  $FADs = 0.2; P = 0.5; Q = 0.5; M = 0.9$ 
3 While  $Iter < Max\_iter$ 
4   Calculate the fitness of each search agent
5    $\vec{X}_A$  = the best search agent
6    $\vec{X}_B$  = the second best search agent
7    $\vec{X}_C$  = the third best search agent
8    $\vec{X}_D$  = the fourth best search agent
9   Update  $CF, \vec{a}, RL$  and  $RB$ 
10  If ( $q \leq Q$ ) % Team hunting
11    For each search agent
12      Update  $\vec{A}$  and  $\vec{C}$  by the Eq. (5) and Eq. (6)
13      Use  $\vec{A}$  and  $\vec{C}$  to calculate  $\vec{D}$ 
14      Calculate  $\vec{X}_a, \vec{X}_b, \vec{X}_c$  and  $\vec{X}_d$  by the Eq. (8)
15       $\vec{X}_{ave} = \frac{\vec{X}_a + \vec{X}_b + \vec{X}_c + \vec{X}_d}{4}$ 
16      Construct the “center circle”:  $E\_Team = \{\vec{X}_a, \vec{X}_b, \vec{X}_c, \vec{X}_d, \vec{X}_{ave}\}$ 
17      If ( $r_2 < M$ )
18        Update the position of the current search agent by the Eq.(13)
19      else if ( $r_2 \geq M$ )
20        Update the position of the current search agent by the Eq.(14)
21      end if
22    End for
23  Else if ( $q \leq Q$ ) % Elite hunting
24    Top_lioness_pos =  $\frac{\vec{X}_A + \vec{X}_B + \vec{X}_C}{3}$ 
25    Construct the Elite matrix and accomplish memory saving
26    For each search agent
27      If  $Iter < Max\_iter/3$ 
28        Update the position of the current search agent by the Eq.(15)
29      else if  $Max\_iter/3 < Iter < 2 \cdot Max\_iter/3$ 
30        For the first half of the populations ( $i=1, \dots, n/2$ )
31          Update the position of the current search agent by the Eq.(16)
32        End for
33        For the other half of the populations ( $i=n/2, \dots, n$ )
34          Update the position of the current search agent by the Eq.(17)
35        End for
36      else if  $Iter > 2 \cdot Max\_iter/3$ 
37        Update the position of the current search agent by the Eq.(18)
38      end if
39    End for
40  End if
41  Update  $Top\_lioness\_pos$  if there is a better solution
42  Applying FADs effect and update the position of the current search agent by the Eq.(19)
43   $Iter = Iter + 1$ 
44 End while
45 Return  $Top\_lioness\_pos$ 

```

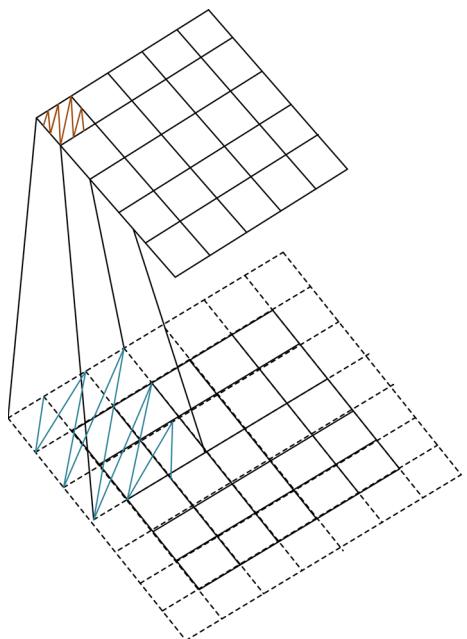
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Note: The value of each dimension is within the range of [0, 1]. If the value is greater than 0.5, it is regarded as 1; otherwise, it is regarded as 0. If the value in a certain dimension is 1, it means that the corresponding feature vector in this dimension is selected, otherwise, it will not be selected. The calculation formula of the fitness value is  $\text{fitness} = \text{GCN}(G; X)$ ,  $G$  is the city network graph,  $X$  is the feature matrix, and the output result is the value of Root Mean Squared Error (RMSE).

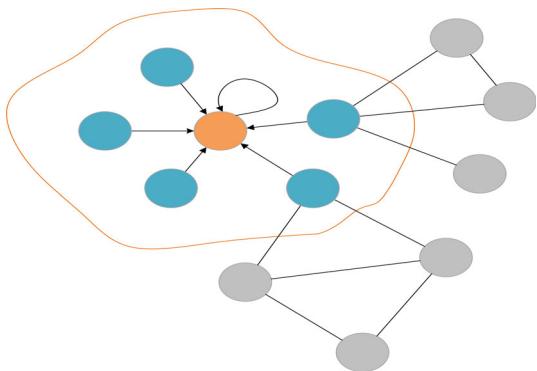
#### 2.4.4 Capture spatial information

At the beginning of the outbreak, it was the Spring Festival travel season. Most people are taking chances, and their awareness of epidemic prevention is weak. The large-scale population movement has caused the epidemic to spread quickly across the country in a short time. Since the outbreak of the COVID-19 has the feature of “person-to-person transmission”, when forecasting the number of confirmed COVID-19 cases in a certain region, it is necessary to consider not only the number of confirmed cases in the region in the past period of time, but also the inflow of confirmed cases from the surrounding regions to the region. How to characterize the interrelationships between complex spaces has become another vital issue in epidemic prediction.

The conventional convolutional neural network (convolutional neural network, CNN) can get local spatial information, but it can only apply to Euclidean space, such as images, regular grids, and so on. The principle is to get



**Fig. 5** A schematic diagram of common CNN two-dimensional convolution



**Fig. 6** Capture spatial features by obtaining topological relationships between nodes

the spatial relationship features of the image through the convolution kernel. The typical CNN two-dimensional convolution is shown in Fig. 5. However, the city network is not a two-dimensional grid, but in the form of a graph, which means that CNN cannot reflect the complex topology of the city network and cannot precisely obtain spatial information. In order to be able to extend convolution to topological graphs of non-European data structures, GCN is born. It can extract features from many irregular data structures in real life, typical of which is graph structure, or topological structure.

Give a set of graph data, which contains  $N$  nodes, each node has its own features, we used the nodes with  $D$ -dimensional features to construct a feature matrix  $X \in R^{N \times D}$ , and then used the mutual relationship between the nodes to construct the adjacency matrix  $A \in R^{N \times N}$ . Finally, input the feature matrix  $X$  and the adjacency matrix  $A$  into the network model. GCN is also a neural network layer, and its layer-to-layer propagation mode is as follows:

$$X^{(l+1)} = \sigma(\tilde{D}^{-\frac{1}{2}} \tilde{A} \tilde{D}^{-\frac{1}{2}} X^{(l)} W^{(l)}) \quad (20)$$

where  $X^{(l)}$  represents the feature matrix of the  $l$ -th layer,  $W^{(l)}$  represents the learnable parameters of the  $l$ -th layer.  $\tilde{A} = A + I_N$ ,  $A$  is the adjacency matrix, which is used to reflect the network topology in the graph data.  $I_N$  is the identity matrix, which is used to extract the features of its own node.  $\tilde{D} = \sum \tilde{A}_{ij}$  is the degree matrix, and the degree matrix is used to measure the importance of neighbor nodes.  $\sigma$  is a nonlinear activation function.

It can be seen from the above formula that the GCN model is equivalent to a feature extractor, which acts on the nodes of the graph, and gets together its first-order neighbors according to different weights, thereby capturing the spatial features between nodes. The specific operation is shown in Fig. 6. Suppose the orange node is the central node, and the blue nodes are the first-order neighbor node

of the central node. According to the network topology, the central node aggregates the features of itself and its first-order neighbor nodes, and finally obtains spatial information from the feature data.

The LsOA-GCN model used in this paper constructed a GCN model by superimposing two convolutional layers. A two-layer GCN model can be expressed as:

$$X^{(l+2)} = \text{Relu}(\hat{A}\text{Relu}(\hat{A}X^{(l)}W^{(l)})W^{(l+1)}) \quad (21)$$

where  $X^{(l)}$  represents the feature matrix of the  $l$ -th layer;  $\text{Relu}()$  is the activation function;  $\hat{A} = \tilde{D}^{-\frac{1}{2}}\tilde{A}\tilde{D}^{-\frac{1}{2}}$ ,  $\tilde{A} = A + I_N$ ,  $A$  is the adjacency matrix, and  $\tilde{D} = \sum \tilde{A}_{ij}$  is the degree matrix;  $W^{(l)}W^{(l+1)}$  represent the learnable parameters of layer  $l$  and layer  $l + 1$ , respectively.

#### 2.4.5 Loss function

The loss function is the most basic and critical element of the prediction model in the training process. Its purpose is to minimize the error between the actual value and the predicted value of cumulative confirmed cases. The loss function we defined in the LsOA-GCN model was the mean square loss function. The specific formula is as follows:

$$\text{loss} = \frac{\sum_{t=1}^T (Y_t - \hat{Y}_t)^2}{T} \quad (22)$$

where  $Y_t$  and  $\hat{Y}_t$ , respectively, represent the actual and predicted the cumulative number of confirmed cases on day  $t$ .

#### 2.4.6 LsOA-GCN model

In order to capture the temporal and spatial information of the feature data simultaneously, we used a graph convolutional network prediction model based on the lioness optimization algorithm (LsOA-GCN). The overall process of LsOA-GCN is shown in Fig. 7. The right part is the flow chart of this prediction model, and the left part is the feature screening process.

In short, the LSOA-GCN model can handle complex temporal dependence and spatial correlation. On the one hand, Spearman correlation analysis with delay days and LsOA are used to capture the dynamic changes of feature information to get the temporal features. On the other hand, we obtain the structure of the city network by graph convolutional network, to get spatial information and finally realize the prediction task.

## 3 Experiments

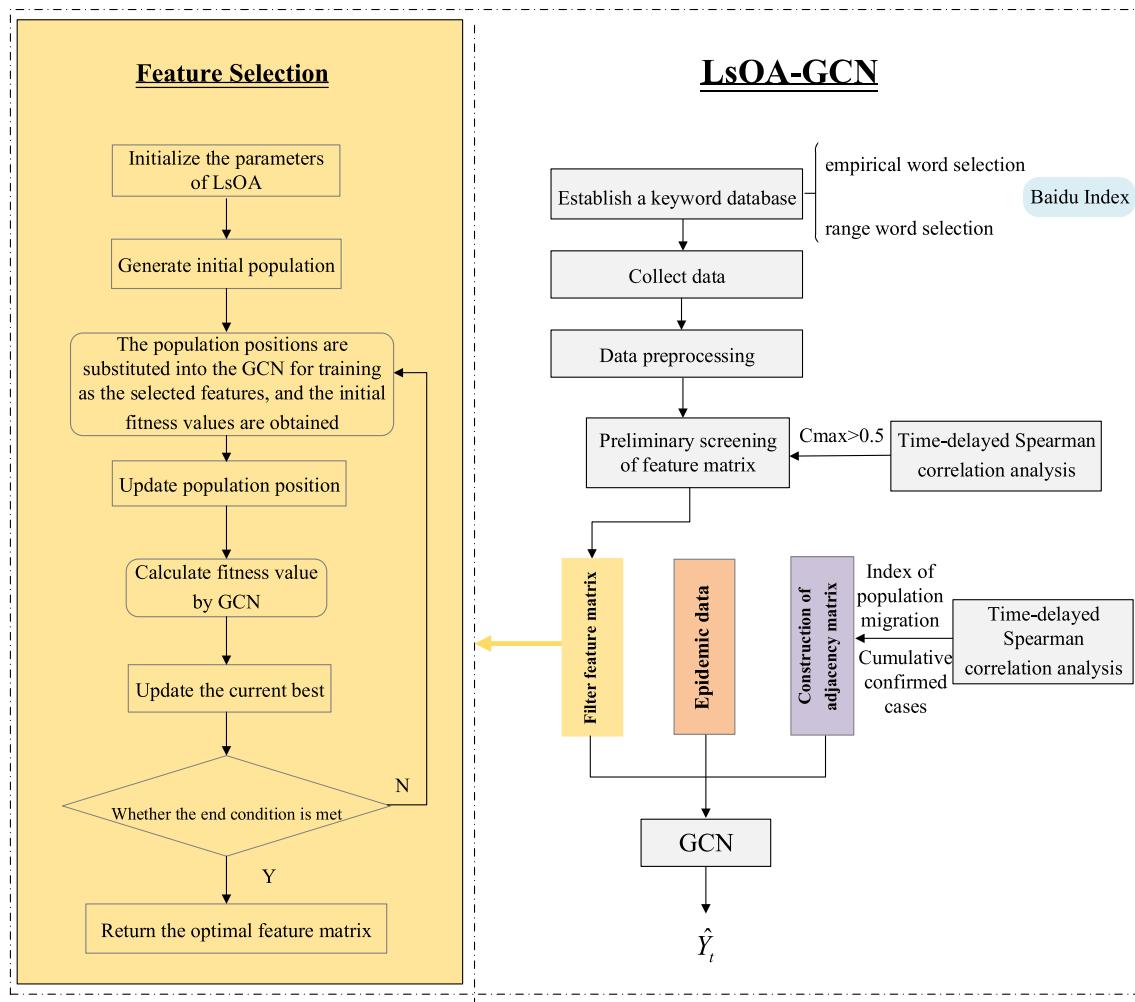
### 3.1 Data set

To verify the performance of the LsOA-GCN model, we predicted the number of confirmed cases of the epidemic for 66 days in 17 regions of Hubei Province. Due to proper epidemic control measures, the number of newly confirmed cases in Hubei Province was almost zero in the later period. Therefore, the prediction target of the confirmed epidemic data set in Hubei Province was set as the cumulative number of daily confirmed cases.

In the experiment, we selected 17 regions in Hubei Province as the primary research areas, namely Wuhan, Ezhou, Enshi, Huanggang, Huangshi, Jingmen, Jingzhou, Qianjiang, Shennongjia, Shiyan, Suizhou, Tianmen, Xiantao, Xianning, Xiangyang, Xiaogan and Yichang. The experimental data was mainly composed of three parts: The first was the migration data from January 10, 2020 to March 15, 2020. The 17\*17 adjacency matrix was constructed mainly based on the correlation between the migration scale index and the cumulative number of confirmed cases; The second was the keyword data from January 10, 2020 to March 15, 2020 (the date range varies according to the number of delayed days), which was used to construct a 17\*66\*44 feature matrix; The third was the epidemic data from January 24, 2020, to March 29, 2020. The daily cumulative number of confirmed cases in each region of Hubei Province was used as the final target value to form a 17\*66 target value matrix. During the period, all experimental data used were normalized, and according to the ratio of 9:1, the data set was divided into the training set and the test set. The length of the historical time series was set to 1, and finally the cumulative number of confirmed cases in the next day was predicted. The specific division is shown in Fig. 8.

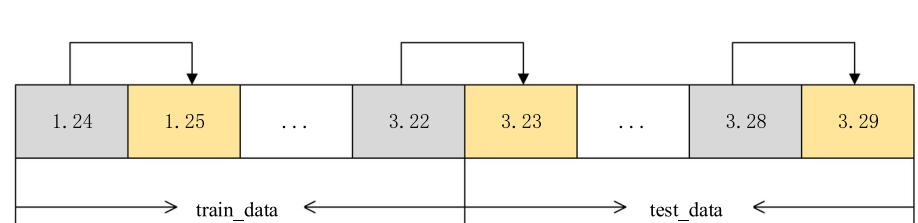
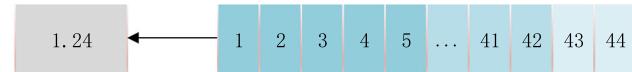
### 3.2 Evaluation indicator

At present, there are many indicators used to evaluate regression prediction models (Zhao et al. 2020b). RMSE, mean absolute error (MAE), mean absolute percentage error (MAPE), accuracy (Acc), coefficient of determination (R2), and explained variance score (Var) were selected as evaluation indicators in this paper to evaluate the difference between the real value  $Y_t$  and the predicted value  $\hat{Y}_t$  of the cumulative number of confirmed cases. RMSE, MAE, and MAPE are mainly used to measure the difference between the predicted value and the real value. The smaller the three indicators are, the smaller the difference between the predicted value and the actual value is, and the better the performance of the prediction model is. Accuracy is



**Fig. 7** LsOA-GCN model structure

**Fig. 8** Division of data set



mainly used to test the prediction accuracy of the model. R2 and Var are mainly used to measure the fitting degree between the predicted value and the actual value. The larger the three indicators are, the better the prediction effect is. The specific calculation formulas of these six indicators are as follows:

(1) RMSE:

$$RMSE = \sqrt{\frac{1}{T} \sum_{t=1}^T (Y_t - \hat{Y}_t)^2} \quad (23)$$

(2) MAE:

$$MAE = \frac{1}{T} \sum_{t=1}^T |Y_t - \hat{Y}_t| \quad (24)$$

(3) MAPE:

$$MAPE = \frac{100\%}{T} \times \sum_{t=1}^T \left| \frac{Y_t - \hat{Y}_t}{Y_t} \right| \quad (25)$$

(4) Acc:

$$Accuracy = 1 - \frac{\| Y - \hat{Y} \|_F}{\| Y \|_F} \quad (26)$$

(5) R2:

$$R^2 = 1 - \frac{\sum_{t=1}^T (Y_t - \hat{Y}_t)^2}{\sum_{t=1}^T (Y_t - \bar{Y})^2} \quad (27)$$

(6) Var:

$$\text{var} = 1 - \frac{\text{Var}\{Y - \hat{Y}\}}{\text{Var}\{Y\}} \quad (28)$$

where  $T$  is the predicted days;  $\| \cdot \|_F$  is the Frobenius norm that refers to as F-norm, which is a matrix norm that can be used to compare the error between the real matrix and the estimated matrix;  $\text{Var}\{\cdot\}$  is the variance calculation.

### 3.3 Correlation analysis between Baidu Index and the cumulative number of confirmed cases of COVID-19

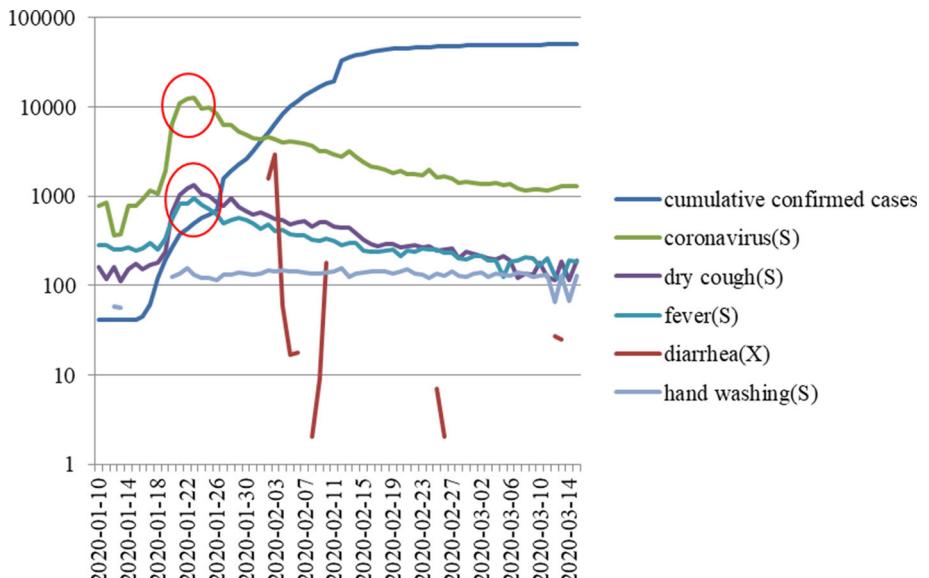
Since Baidu Index includes search index and information index, 40 keywords constitute 80 feature vectors. The search index shows the degree of Internet users' attention to keyword search and the continuous changes. It is based

on the data of Internet users' search volume on Baidu, with keywords as statistical objects, scientifically analyzes and calculates the weight of the search frequency of each keyword in Baidu. The information index shows the degree of attention and coverage of specific keywords and continuous changes in news information on the Internet. It is based on Baidu's intelligent distribution and recommended content data and is obtained by the weighted summation of the number of netizens' reading, commenting, forwarding, like, dislike and other behaviors. We first preprocessed the data, deleted the search index or information index of keywords with all zeros, and finally contained only 61 feature vectors.

Figure 9 shows the trend comparison between the five randomly selected feature vectors and the cumulative number of confirmed cases, from January 10, 2020, to March 15, 2020. The five feature vectors are coronavirus (S), dry cough (S), fever (S), diarrhea (X), and hand washing (S), where (S) represents the search index of the keyword and (X) describes the information index of the keyword. Since the ordinate used a logarithmic scale with a base of 10 and negative or zero values could not be drawn correctly on the logarithmic graph, keywords with the Baidu Index value of 0 would appear truncated and had no value in the graph.

As can be seen from Fig. 9, the change of Baidu Index values of individual keywords is earlier than the change of cumulative confirmed cases of COVID-19. This showed that, based on certain keywords, we could predict the epidemic trend in the future to a certain extent, and there was an apparent lag between the Baidu Index of keywords and the cumulative number of confirmed cases. Because of this, we assumed that the Baidu Index could be used as a predictive indicator in a period of time before the

**Fig. 9** Trend comparison between the five randomly selected feature vectors and cumulative number of confirmed cases



diagnosis, that is, the feature vector, and then the number of confirmed cases could be predicted.

Based on the above analysis, in this part, we mainly analyzed the correlation between the daily cumulative number of confirmed cases of the epidemic and the Baidu Index. Due to the eclipse period of COVID-19, there are even asymptomatic infections. And as the main feature vector of the prediction model, the Baidu Index needs to be at least a certain number of days in advance to have practical application value. Therefore, the Baidu Index values of keywords 0 to 14 days in advance (epidemic data was relatively lagging by 0 to 14 days) were successively used for correlation analysis with epidemic data.

Taking Wuhan as an example, we performed Spearman correlation analysis on the Baidu Search Index (search index, information index) of keywords with time delay (0–14d) and the cumulative number of confirmed cases in Wuhan. The results are shown in Table 1, and the correlation analysis results of other 16 regions are shown in Appendix Tables 13–28.

Table 1 shows the correlation coefficient value of each keyword with a delay of 0–14 days and the cumulative number of confirmed cases, where the correlation coefficient value corresponding to the optimal delay days has been shown in bold. For example, for novel coronary pneumonia (S), we found that the 14-day delay has the strongest explanatory power. For fever (S), the most explanatory delay time is 3 days. In this experiment, we believed that it is most appropriate to use the keyword data corresponding to the delay days with the strongest explanatory power, and used it for subsequent prediction tasks. In order to highlight the role of feature vectors and improve prediction performance, based on comprehensive consideration of 17 regions in Hubei Province, we only screened out the keyword data with the maximum correlation coefficient greater than 0.5, a total of 44, as shown in Table 2. Then, according to the delay days corresponding to the maximum correlation coefficient, the keyword data was moved, filled, and sorted out to obtain the final feature vectors.

After screening out 44 feature vectors, in order to further determine the importance of correlation analysis with delay days for epidemic prediction, we, respectively, predicted the sorted feature data and unsorted feature data on the GCN model, and the prediction results are shown in Tables 3 and 4, where Table 3 is a comparison of the overall prediction results of Hubei Province, and Table 4 is a comparison of the prediction results of 17 regions in Hubei Province. It was worth noting that due to proper epidemic control measures, the number of newly confirmed cases in various regions of Hubei Province in the later period was almost zero, and the daily cumulative number of confirmed cases was almost unchanged. Therefore, according to the

calculation formulas of evaluation indicators, when comparing the prediction results of various regions, we only used the four indicators of RMSE, MAE, MAPE, and Acc. It can be clearly seen from Tables 3 and 4 that the sorted feature data are better than the unsorted feature data in predicting effect, and it is better than the unsorted feature data in all indicators. This result proves that the use of correlation analysis with delay days to sort the feature data can improve the performance of the prediction model.

### 3.4 Performance analysis of lioness optimization algorithm on benchmark functions

After determining the optimal delay days for each feature, we then considered using lioness optimization algorithm to solve the feature selection problem, whose main task is to delete redundant and irrelevant features from the original feature data. In chapter 2.4.3, we introduced the lioness optimization algorithm. In this part, we mainly tested the performance of the proposed algorithm.

In this experiment, we tested the algorithm through 23 commonly used benchmark functions. The benchmark functions mainly include four categories: (1) unimodal functions (F1–F7), which have the unique optimal solution and are mainly used to test the exploitation performance of the algorithm. (2) Multimodal functions (F8–F13), with multiple local optimal solutions, are mainly used to test the exploration ability of the algorithm. (3) Multimodal functions with fixed dimension (F14–F23), which has low and fixed dimensions. The optimization algorithms involved in the comparison included classical particle swarm optimization (PSO), grey wolf optimizer(GWO) (Mirjalili et al. 2014), whale optimization algorithm(WOA), teaching–learning-based optimization(TLBO), and newly developed heap-based optimizer(HBO), marine predators algorithm(MPA) and K-means clustering algorithm(KO). The parameter settings of various algorithms and more details can be found in the corresponding references. To provide reliable comparison results, for all test functions, the algorithms independently ran 30 times in the same environment, and the number of function evaluations was 15,000. Therefore, under the same other conditions, the differences between algorithms are caused by different performance. Table 5 shows the test results of different algorithms on common benchmark functions.

It can be seen from Table 5 that for unimodal functions (F1–F7), the average values obtained by LsOA on F5 and F6 were far superior to other algorithms. On F7, LsOA also ranked first with a slight advantage. On F1–F4, compared with KO, LsOA performed poorly, but it was also superior to other algorithms by absolute advantages, showing strong competitiveness. For multimodal functions (F8–F23), LsOA reached global optimization on F9, F11, and F15–

**Table 1** Correlation analysis between keywords and the cumulative number of confirmed cases–Wuhan

Keywords	Delay days														
	14	13	12	11	10	9	8	7	6	5	4	3	2	1	0
Center for Disease Control and Prevention(S)	0.1247	0.1787	0.2346	0.2929	0.3559	0.4239	0.4956	0.5729	0.6464	0.7330	0.7933	<b>0.7954</b>	0.7911	0.7824	0.7743
Red Cross Society(S)	0.0283	0.0172	0.0596	0.1123	0.1690	0.2237	0.2814	0.3437	0.4134	0.4897	0.5693	0.6542	0.7477	0.8416	<b>0.8720</b>
Spring Festival(S)	0.8880	0.8911	0.8946	0.8986	<b>0.9009</b>	0.8994	0.8926	0.8860	0.8812	0.8788	0.8749	0.8701	0.8647	0.8581	0.8502
Health Commission(S)	0.0678	0.1150	0.1691	0.2274	0.2875	0.3560	0.4242	0.4935	0.5700	0.6539	0.7429	0.7821	<b>0.7844</b>	0.7733	0.7651
Vaccine(S)	0.1460	0.1085	0.0679	0.0239	0.0231	0.0749	0.1362	0.1963	0.2563	0.3217	0.3915	0.4672	0.4932	0.5363	<b>0.5418</b>
SARS(S)	0.4405	0.5084	0.5448	0.6189	0.6970	0.7842	0.8693	0.9261	0.9524	0.9817	0.9895	<b>0.9902</b>	0.9899	0.9894	0.9887
Hand washing(S)	0.2298	0.2060	0.1690	0.1284	0.0911	0.0426	0.0071	0.0598	0.1158	0.1774	0.2407	0.2931	<b>0.3013</b>	0.2609	0.2885
Children(S)	<b>0.6136</b>	0.6043	0.6105	0.6064	0.6061	0.6011	0.6014	0.5873	0.5707	0.5475	0.5355	0.5267	0.5175	0.6036	0.5960
Mask(S)	0.1552	0.2114	0.2701	0.3329	0.4003	0.4692	0.5413	0.6218	0.7057	0.7974	<b>0.8964</b>	0.8913	0.8851	0.8786	0.8714
Coronavirus(S)	0.2614	0.3194	0.3820	0.4483	0.5207	0.5978	0.6791	0.7663	0.8588	0.9574	0.9889	<b>0.9998</b>	0.9893	0.9888	0.9881
Novel(S)	0.1858	0.2432	0.3022	0.3663	0.4346	0.5055	0.5812	0.6635	0.7504	0.8434	0.9445	0.9667	<b>0.9670</b>	0.9655	0.9635
Disinfection(S)	0.0703	0.1208	0.1701	0.2234	0.2868	0.3465	0.4113	0.4833	0.5615	0.6373	0.7211	<b>0.7317</b>	0.7175	0.7026	0.6851
Lockdown of the city(S)	0.0240	0.0235	0.0742	0.1232	0.1733	0.2286	0.2912	0.3597	0.4327	0.5085	0.5891	0.6770	0.7717	<b>0.8739</b>	0.8665
Wuhan(S)	0.2114	0.2670	0.3326	0.3995	0.4659	0.5390	0.6134	0.6947	0.7869	0.8841	0.9433	<b>0.9455</b>	0.9434	0.9418	0.9384
Zhong Nanshan(S)	0.1781	0.2334	0.2904	0.3521	0.4205	0.4932	0.5689	0.6489	0.7341	0.8259	<b>0.9263</b>	0.9233	0.9192	0.9145	0.9095
epidemic(S)	<b>0.9191</b>	0.9160	0.9090	0.9039	0.8984	0.8924	0.8883	0.8845	0.8779	0.8714	0.8651	0.8571	0.8487	0.8399	0.8305
Flu(S)	0.4343	0.4723	0.5102	0.5521	0.5953	0.6490	0.7170	0.7802	0.8435	0.9111	0.9678	<b>0.9683</b>	0.9678	0.9665	0.9646
Diarrhea(S)	0.0656	0.1116	0.1652	0.2184	0.2790	0.3432	0.4081	0.4791	0.5538	0.6371	0.7232	0.8191	0.9184	0.9448	<b>0.9464</b>
Dry cough(S)	0.3505	0.3969	0.4594	0.5134	0.5879	0.6501	0.7144	0.7849	0.8594	0.9367	0.9860	<b>0.9993</b>	0.9888	0.9883	0.9876
Ncp(s)	<b>0.5801</b>	0.5725	0.5644	0.5561	0.5357	0.5233	0.5042	0.4817	0.4578	0.4324	0.4036	0.3730	0.3455	0.3103	0.2721
novel Coronary pneumonia(S)	<b>0.9480</b>	0.9439	0.9383	0.9373	0.9371	0.9346	0.9326	0.9303	0.9247	0.9206	0.9170	0.9114	0.9062	0.9009	0.8951
COVID-19(S)	<b>0.7171</b>	0.7119	0.7045	0.6986	0.6863	0.6765	0.6684	0.6559	0.6447	0.6330	0.6178	0.6023	0.5860	0.5676	0.5474
Novel coronavirus pneumonia(S)	0.0032	0.0408	0.0900	0.1460	0.2025	0.2663	0.3286	0.3944	0.4657	0.5457	0.6286	<b>0.7207</b>	0.7192	0.7111	
Asymptomatic infection(S)	<b>0.5199</b>	0.5022	0.4800	0.4601	0.4378	0.4142	0.3834	0.3512	0.3215	0.2901	0.2560	0.2189	0.1770	0.1314	0.0820
Aerosol transmission(S)	<b>0.4709</b>	0.4491	0.4250	0.4014	0.3768	0.3486	0.3181	0.2867	0.2514	0.2147	0.1752	0.1320	0.0849	0.0345	0.0202
Pneumonia(S)	0.1877	0.2412	0.3021	0.3647	0.4326	0.5065	0.5831	0.6624	0.7497	0.8428	0.9425	<b>0.9525</b>	0.9524	0.9501	0.9472
Cough(S)	0.4075	0.4556	0.4821	0.5211	0.5778	0.6139	0.6796	0.7160	0.7874	0.8513	0.8985	<b>0.9050</b>	0.9027	0.8981	0.8926
Fever(S)	0.7782	0.7958	0.8099	0.8295	0.8477	0.8644	0.8989	0.9202	0.9354	0.9597	0.9717	<b>0.9718</b>	0.9705	0.9689	0.9670
dyspnea(S)	0.2216	0.2765	0.3356	0.4018	0.4715	0.5451	0.6238	0.7083	0.7809	0.8764	0.9698	<b>0.9724</b>	0.9715	0.9704	0.9686
high temperature(S)	0.3187	0.3761	0.4367	0.5013	0.5578	0.6306	0.7048	0.7879	0.8781	0.9480	<b>0.9695</b>	0.9691	0.9674	0.9656	0.9635
novel coronavirus(S)	0.1182	0.1676	0.2229	0.2823	0.3448	0.4140	0.4871	0.5632	0.6446	0.7322	0.8260	0.9278	0.9549	<b>0.9650</b>	0.9636
SARS-CoV-2(S)	<b>0.7241</b>	0.7163	0.7067	0.7000	0.6957	0.6893	0.6792	0.6683	0.6540	0.6413	0.6280	0.6137	0.5978	0.5805	0.5616
Spring Festival(X)	0.7765	0.7760	0.7778	0.7833	<b>0.7858</b>	0.7786	0.7703	0.7597	0.7528	0.7427	0.7364	0.7262	0.7105	0.6941	0.6762
Red Cross Society(X)	0.1147	0.0777	0.0379	0.0030	0.0468	0.0968	0.1471	0.2017	0.2591	0.3080	0.3584	0.4287	0.5046	<b>0.5567</b>	0.5470

**Table 1** (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
government(X)	0.1745	0.1704	0.1351	0.0981	0.0884	0.0974	0.1565	0.2103	0.2334	0.2439
Health Commission(X)	0.1732	0.2229	0.2187	0.2570	0.3143	0.3699	0.3919	0.4613	0.5331	0.6098
medical(X)	0.1602	0.1231	0.0859	0.0501	0.0135	0.0306	0.0792	0.1337	0.1894	0.2473
vaccine(X)	0.1378	0.1077	0.0765	0.0441	0.0028	0.0442	0.0953	0.1484	0.2012	0.2579
SARS(X)	0.3243	0.3812	0.4357	0.4917	0.5574	0.5790	0.5926	0.6194	0.6805	0.7358
children(X)	0.3235	0.3048	0.2971	0.3468	0.4064	0.4691	0.4541	0.4354	0.4128	0.4341
mask(X)	0.1042	0.0666	0.0301	0.0150	0.0609	0.1053	0.1575	0.2168	0.2742	0.3381
infection(X)	0.3778	0.4038	0.4311	0.4598	0.4899	0.5216	0.5550	0.5901	0.6270	0.6659
coronavirus(X)	<b>0.5493</b>	0.5341	0.5183	0.5029	0.4854	0.4656	0.4433	0.4168	0.3938	0.3696
disinfection(X)	0.6499	0.6513	0.6526	0.6508	0.6505	0.6964	0.6981	0.6997	0.7018	0.7022
nucleic acid(X)	<b>0.4535</b>	0.4280	0.4065	0.3823	0.3555	0.3253	0.2918	0.2558	0.2199	0.1806
lockdown of the city(X)	<b>0.6956</b>	0.6932	0.6902	0.6872	0.6858	0.6853	0.6844	0.6812	0.6778	0.6742
Wuhan(X)	0.0945	0.0912	0.1337	0.1827	0.2412	0.3006	0.3612	0.4063	0.4712	0.5192
Zhong Nanshan(X)	0.0065	0.0335	0.0766	0.1237	0.1759	0.2338	0.2898	0.3537	0.4185	0.4877
epidemic(X)	0.2695	0.2373	0.2054	0.1743	0.1386	0.0993	0.0572	0.0069	0.0407	0.0898
flu(X)	<b>0.4739</b>	0.4532	0.4316	0.4130	0.3924	0.3645	0.3330	0.3015	0.2694	0.2353
diarrhea(X)	0.0572	0.0468	0.0358	0.0243	0.0023	0.0183	0.0300	0.0498	0.0654	0.0818
stuffy nose(X)	0.0938	0.1273	0.1524	0.1789	0.1974	0.2250	0.2478	0.2884	0.3185	0.3471
dry cough(X)	0.1158	0.0810	0.0439	0.0048	0.0403	0.0873	0.1414	0.1983	0.2547	0.3149
NCP(X)	<b>0.2867</b>	0.2724	0.2662	0.2596	0.2475	0.2283	0.2122	0.1936	0.1769	0.1592
close contact(X)	<b>0.7132</b>	0.7060	0.6949	0.6858	0.6782	0.6702	0.6597	0.6444	0.6300	0.6163
suspected case(X)	<b>0.4649</b>	0.4406	0.4211	0.3985	0.3715	0.3440	0.3136	0.2746	0.2405	0.2043
pneumonia(X)	<b>0.4079</b>	0.3867	0.3628	0.3407	0.3154	0.2871	0.2562	0.2203	0.1859	0.1511
cough(X)	0.4278	0.4759	0.5282	0.5770	0.5706	0.5538	0.5509	0.5581	0.5833	0.5958
dyspnea(X)	0.1956	0.1643	0.1255	0.0834	0.0386	0.0009	0.0116	0.0539	0.0519	0.0503
virus pneumonia(X)	<b>0.7062</b>	0.7032	0.6900	0.6832	0.6807	0.6787	0.6726	0.6653	0.6805	0.6733
novel coronavirus(X)	<b>0.6044</b>	0.5898	0.5741	0.5595	0.5424	0.5229	0.5009	0.4750	0.4518	0.4277

Optimal values are shown in bold in the table

**Table 2** The final keyword list of COVID-19

Category	Keyword
prevention	mask(S), mask(X), disinfection(X), nucleic acid(X)
symptoms	flu(S), diarrhea(S), dry cough(S), asymptomatic infection(S), pneumonia(S), cough(S), fever(S), dyspnea(S), high temperature(S), infection(X), stuffy nose(X), close contact(X), suspected case(X), pneumonia(X), cough(X)
common words	coronavirus(S), novel(S), epidemic(S), NCP(S), novel coronary pneumonia(S), COVID-19(S), novel coronavirus(S), coronavirus(X), virus pneumonia(X), novel coronavirus(X)
other related aspects of COVID-19	Red Cross Society(S), Spring Festival(S), Health Commission(S), SARS(S), lockdown of the city(S), Wuhan(S), Zhong Nanshan(S), Spring Festival(X), Red Cross Society(X), Health Commission(X), SARS(X), children(X), lockdown of the city(X), Wuhan(X), Zhong Nanshan(X)

The Baidu Index of the keyword here included search index and information index, where (S) describes the search index of the keyword, and (X) represents the information index of the keyword

F23. On F8, LsOA also had a weak advantage over other algorithms and achieved good results. For F10 and F14, LsOA tied for first with some algorithms and obtained the same results. On F12 and F13, LsOA showed an absolute advantage.

We comprehensively considered the test results of these benchmark functions. The last row in Table 5 is the final ranking result. The overall performance of LsOA on these 23 commonly used benchmark functions was excellent, with an average ranking of 1.17, ranking first in the comprehensive ranking, which was in turn superior to KO, MPA, TLBO, HBO, GWO, WOA, and PSO. The above experiments prove the superior performance of LsOA, which also shows that we can use this algorithm to screen the prediction features of COVID-19.

### 3.5 Setting training parameters and comparison methods

#### 3.5.1 Training parameters

The parameters used in the GCN training model included the length of history\future time series, feature dimension, learning rate, training epoch, batch size, number of hidden layer nodes, optimizer, loss function, and so on. The processor of the experimental hardware environment was Intel (R) Pentium (R) CPU N3700 @ 1.60 GHz 1.60 GHz, a server with 4 GB of memory, and the software environment consisted of Python 3.6.12 and TensorFlow 1.14.0.

Among the training parameters, the number of hidden layer nodes is a very important parameter in the GCN model. Different numbers of hidden layer nodes will greatly affect the prediction accuracy. Too many nodes will cause over-fitting, and too few nodes will cause the model to be inaccurate. Therefore, in this part, we conducted a sensitivity analysis on the number of hidden layer nodes and conducted experiments with different numbers of

hidden layer nodes, to analyze the performance of the model and finally determine the optimal number of hidden layer nodes by comparing the prediction results.

In this experiment, we determined the number of hidden layer nodes from {8, 16, 32, 64, 80, 100, 128, 144, 160}, conducted experiments on the premise that the number of training is 3000, and analyzed the change of prediction accuracy. Finally, the results of sensitivity analysis for different numbers of hidden layer nodes are shown in Fig. 10, where the abscissa represents the number of hidden layer nodes and the ordinate represents the corresponding value of each indicator.

Figure 10 shows the changes of various index values under different numbers of hidden layer nodes when the number of training times is 3000, where Fig. 10a and c, respectively, shows the changes of RMSE, MAE, and MAPE as the number of hidden layer nodes increases. It can be seen from these figures that when the number of hidden layer nodes is 16, the corresponding RMSE and MAE are the smallest, and MAPE is the smallest when the number of hidden layer nodes is 32, indicating that the difference between the predicted value and the actual value in the corresponding state is also the smallest. Figure 10b shows the changes of Accuracy and Var as the number of hidden layer nodes increases. It can be concluded from Fig. 10b that when the number of hidden layer nodes is 16, these three indicators all reach the maximum value, indicating that the model fitting effect is the best at this time. In summary, we could conclude that when the training times are set to 3000, the model has the best effect when the number of hidden layer nodes is set to 16.

We can also see from these figures that as the number of hidden layer nodes increases, the prediction accuracy first decreases, then increases, and then decreases. This is mainly because when the hidden layer node is larger than a certain level, the model complexity and calculation difficulty will greatly increase, thereby reducing the prediction

**Table 3** Comparison of prediction results in Hubei Province

Data set	Indicator	Data (sorted)	Data (unsorted)
Hubei Province	RMSE	322.277	1030.61
	MAE	74.0729	308.891
	MAPE	0.02144	0.08120
	Acc	0.97360	0.91558
	R2	0.99922	0.99202
	Var	0.99925	0.99274

accuracy. Therefore, according to the results of sensitivity analysis, we also set the number of hidden layer nodes as 16 in the subsequent experiment of LsOA-GCN. Finally, the parameters used for the LsOA-GCN training model we obtained are shown in Table 6.

### 3.5.2 Comparison methods

This part mainly compared several representative time series prediction methods in current epidemic prediction research, including multiple linear regression, support vector regression, backpropagation neural network, kernel ridge regression, Gaussian process regression, stochastic gradient descent regression, GAT, LSTM, and T-GCN models. The nine comparison methods are described as follows:

(1) Multiple Linear Regression (Yuchi et al. 2019). When performing regression analysis, we call regression with two or more independent variables under linear correlation conditions as multiple linear regression (MLR). This method uses the optimal combination of multiple independent variables to predict or estimate the dependent variable together. Because the relationship between multiple independent variables is integrated, it is usually a better method than linear regression, which is more effective and more realistic than using only one independent variable for prediction or estimation. In the comparison experiment, the parameters used in this method were all default values.

(2) Support Vector Regression (Brereton and Lloyd 2010). Support vector machine (SVM) is a class of generalized linear classifiers that perform binary classification of data in a supervised learning manner. Support vector regression (SVR) is an application model of support vector machines in regression problems. The support vector machine regression model has many variants based on different loss functions. The core idea is to find a separating hyperplane (hypersurface) that minimizes the expected risk. In the comparison experiment, the kernel type used in this method was 'poly', the penalty parameter C = 1.1, the kernel coefficient gamma of 'poly' was set as 'auto', the degree of the polynomial kernel function ('poly') was 3, epsilon = 0.1, the independent term in the kernel function coef0 = 1.0, and the rest of the parameters used default values.

**Table 4** Comparison of prediction results of 17 regions in Hubei Province

Data		Data (sorted))				Data (unsorted)			
Number	Region	RMSE	MAE	MAPE	Acc	RMSE	MAE	MAPE	Acc
0	Wuhan	1318.24	881.730	0.01763	0.97364	4219.93	3823.86	0.07647	0.91561
1	Ezhou	38.6784	26.1887	0.01879	0.97225	118.111	106.713	0.07655	0.91527
2	Enshi	6.72733	4.83126	0.01917	0.97330	21.2020	19.3225	0.07668	0.91586
3	Huanggang	83.8352	60.6461	0.02086	0.97116	248.487	226.466	0.07790	0.91452
4	Huangshi	27.9862	18.5732	0.01830	0.97243	86.9989	78.6410	0.07748	0.91429
5	Jingmen	24.3426	16.3154	0.01758	0.97377	79.0874	72.2794	0.07789	0.91478
6	Jingzhou	46.1738	31.7411	0.02009	0.97078	137.985	125.721	0.07957	0.91267
7	Qianjiang	5.51831	3.60669	0.01822	0.97213	16.7239	15.2088	0.07681	0.91554
8	Shennongjia	0.27478	0.20933	0.01903	0.97502	1.04942	0.97174	0.08834	0.90460
9	Shiyan	19.7104	13.8861	0.02066	0.97067	58.0828	53.0211	0.07890	0.91357
10	Suizhou	37.5744	25.8064	0.01974	0.97125	111.841	101.834	0.07791	0.91443
11	Tianmen	16.5381	12.1464	0.02449	0.96666	45.3164	41.8090	0.08429	0.90864
12	Xiantao	16.1987	10.9648	0.01907	0.97183	48.2514	43.9107	0.07637	0.91608
13	Xianning	47.6550	43.2417	0.05172	0.94300	103.582	98.5077	0.11783	0.87610
14	Xiangyang	32.7713	22.5675	0.01921	0.97211	101.452	92.3979	0.07864	0.91366
15	Xiaogan	96.6878	67.4415	0.01917	0.97252	302.343	275.598	0.07834	0.91406
16	Yichang	27.8529	19.3436	0.02078	0.97008	81.9497	74.8775	0.08043	0.91198

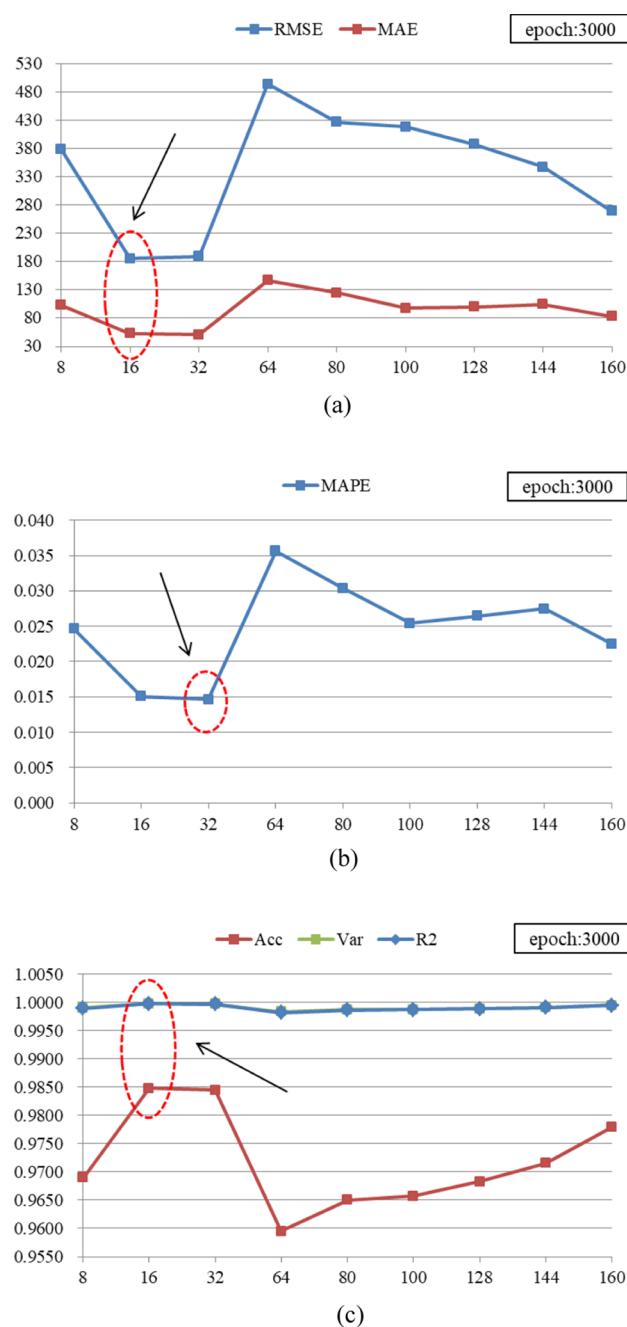
**Table 5** Test results of different algorithms on common benchmark functions

Function	LsOA	PSO	GWO	WOA	TLBO	HBO	MPA	KO
F1	Ave	3.47E-167	2.49E + 00	1.47E-27	2.40E-74	1.42E-38	6.09E-07	1.16E-09
	Rank	2	8	5	3	4	7	6
	Std	0.00E + 00	9.09E-01	2.82E-27	9.81E-74	1.11E-38	7.66E-07	9.16E-10
F2	Ave	4.19E-84	4.40E + 00	9.96E-17	2.35E-49	8.92E-20	8.15E-06	6.23E-06
	Rank	2	8	5	3	4	7	6
	Std	2.30E-83	1.20E + 00	7.23E-17	1.18E-48	4.22E-20	1.06E-05	3.33E-06
F3	Ave	5.23E-112	1.98E + 02	1.21E-05	3.95E + 04	5.99E-07	2.51E + 04	1.21E + 00
	Rank	2	6	4	8	3	7	5
	Std	1.87E-111	6.53E + 01	2.20E-05	1.23E + 04	5.80E-07	7.02E + 03	1.31E + 00
F4	Ave	1.46E-84	1.96E + 00	1.05E-06	4.55E + 01	4.38E-16	1.28E + 01	3.55E-04
	Rank	2	6	4	8	3	7	5
	Std	8.00E-84	1.93E-01	1.37E-06	2.88E + 01	2.63E-16	4.96E + 00	1.15E-04
F5	Ave	<b>5.59E-17</b>	1.18E + 03	2.72E + 01	2.79E + 01	2.57E + 01	1.08E + 02	2.68E + 01
	Rank	1	8	4	6	2	7	3
	Std	3.06E-16	7.58E + 02	7.77E-01	5.31E-01	4.78E-01	5.51E + 01	5.09E-01
F6	Ave	<b>1.28E-17</b>	2.20E + 00	7.89E-01	3.82E-01	3.01E-03	6.92E-07	1.02E-01
	Rank	1	8	7	6	3	2	4
	Std	2.79E-17	9.58E-01	4.22E-01	2.34E-01	5.29E-03	1.23E-06	9.23E-02
F7	Ave	<b>1.40E-04</b>	1.32E + 01	1.75E-03	3.87E-03	2.41E-03	3.52E-02	2.85E-03
	Rank	1	8	3	6	4	7	5
	Std	1.23E-04	1.11E + 01	8.78E-04	3.91E-03	8.77E-04	1.12E-02	1.36E-03
F8	Ave	<b>-1.26E + 04</b>	-6.25E + 03	-6.25E + 03	-1.02E + 04	-6.22E + 03	-1.16E + 04	-8.36E + 03
	Rank	1	7	6	3	8	2	4
	Std	1.10E-09	1.53E + 03	7.54E + 02	1.91E + 03	1.23E + 03	3.44E + 02	4.28E + 02
F9	Ave	<b>0.00E + 00</b>	1.58E + 02	2.78E + 00	<b>0.00E + 00</b>	1.90E + 01	1.51E + 01	<b>0.00E + 00</b>
	Rank	1	6	3	1	5	4	2
	Std	0.00E + 00	3.02E + 01	3.88E + 00	0.00E + 00	9.94E + 00	5.52E + 00	2.97E-03
F10	Ave	<b>8.88E-16</b>	2.71E + 00	1.07E-13	3.73E-15	5.03E-15	3.12E-02	6.47E-06
	Rank	1	7	4	2	3	6	5
	Std	0.00E + 00	3.84E-01	1.90E-14	2.36E-15	1.35E-15	1.70E-01	2.65E-06
F11	Ave	<b>0.00E + 00</b>	1.17E-01	5.84E-03	1.89E-02	<b>0.00E + 00</b>	1.73E-03	7.11E-09
	Rank	1	6	4	5	1	3	2
	Std	4.34E-13	6.47E-02	1.91E-02	2.26E-02	4.13E-04	1.89E-02	3.22E-03
F12	Ave	<b>6.05E-13</b>	5.14E-01	6.70E-01	4.85E-01	1.38E-01	2.56E-03	1.30E-01
	Rank	1	7	8	6	4	2	3
	Std	2.90E-12	1.94E-01	2.47E-01	2.38E-01	1.18E-01	8.50E-03	7.24E-02
F14	Ave	<b>0.99800</b>	3.19702	3.97097	3.74390	<b>0.99800</b>	<b>0.99800</b>	1.19346
	Rank	1	3	5	4	1	1	2

**Table 5** (continued)

Function		LsoA	PSO	GWO	WOA	TlBO	HBO	MPA	KO
F15	Std	2.52E-16	2.81E+00	3.91E+00	3.74E+00	4.12E-17	4.19E-16	0.54228	
	Ave	<b>0.00031</b>	0.00092	0.00713	0.00070	0.00118	0.00079	<b>0.00031</b>	0.00056
	Rank	1	5	7	3	6	4	1	2
F16	Std	2.43E-13	1.29E-04	9.52E-03	3.62E-04	3.64E-03	1.93E-04	2.30E-06	1.91E-04
	Ave	<b>-1.03163</b>	-1.03162						
	Rank	1	1	1	1	1	1	1	2
F17	Std	6.10E-12	4.88E-16	1.81E-05	3.03E-09	6.58E-16	6.39E-16	3.32E-14	7.93E-06
	Ave	<b>0.39789</b>	0.39790	0.39790	<b>0.39789</b>	<b>0.39789</b>	<b>0.39789</b>	<b>0.39789</b>	<b>0.39789</b>
	Rank	1	1	2	2	1	1	1	1
F18	Std	2.10E-12	0.00E+00	5.68E-05	2.26E-05	0.00E+00	0.00E+00	2.09E-12	4.59E-06
	Ave	<b>3.00000</b>	<b>3.00000</b>	3.00005	3.00010	<b>3.00000</b>	<b>3.00000</b>	<b>3.00000</b>	3.00018
	Rank	1	1	2	3	1	1	1	4
F19	Std	5.83E-14	5.34E-15	4.62E-05	2.38E-04	1.39E-15	1.38E-15	2.54E-14	2.09E-04
	Ave	<b>-3.86278</b>	<b>-3.86278</b>	-3.86149	-3.85491	<b>-3.86278</b>	<b>-3.86278</b>	<b>-3.86278</b>	-3.86210
	Rank	1	1	3	4	1	1	1	2
F20	Std	5.48E-12	1.89E-15	2.66E-03	1.11E-02	2.71E-15	2.71E-15	1.74E-12	8.32E-04
	Ave	<b>-3.32200</b>	-3.27047	-3.25757	-3.18350	-3.30784	<b>-3.32200</b>	<b>-3.32200</b>	-3.22099
	Rank	1	3	4	6	2	1	1	5
F21	Std	1.22E-09	5.99E-02	7.96E-02	1.79E-01	3.67E-02	5.95E-15	3.50E-10	6.66E-02
	Ave	<b>-10.1532</b>	-7.30681	-9.47737	-7.38607	-10.1493	-9.31835	<b>-10.1532</b>	-10.1422
	Rank	1	7	4	6	2	5	1	3
F22	Std	9.96E-08	3.21E+00	1.75E+00	3.01E+00	1.66E-02	2.21E+00	2.59E-09	1.89E-02
	Ave	<b>-10.4029</b>	-8.21827	-9.97077	-7.77028	-10.4028	-10.1733	<b>-10.4029</b>	-10.3930
	Rank	1	6	5	7	2	4	1	3
F23	Std	2.44E-08	3.23E+00	1.67E+00	2.86E+00	9.84E-04	1.26E+00	2.18E-09	9.95E-03
	Ave	<b>-10.5364</b>	-9.03874	-10.3560	-7.57344	-10.3551	-10.1484	<b>-10.5364</b>	-10.3476
	Rank	1	6	2	7	3	5	1	4
	Std	4.86E-08	2.78E+00	9.79E-01	3.33E+00	9.92E-01	1.48E+00	2.87E-09	9.86E-01
	Sum rank	27	127	99	105	66	88	64	63
	Mean rank	1.17	5.52	4.30	4.57	2.87	3.83	2.78	2.74
	Final rank	1	8	6	7	4	5	3	2

Optimal values are shown in bold in the table



**Fig. 10** Sensitivity analysis of hidden layer nodes

(3) BP neural network (Wang et al. 2015). BP neural network (backpropagation neural network, BPNN) is a multilayer feedforward network trained according to error backpropagation. Its basic idea is the gradient descent method, which uses gradient search technology to minimize the mean square error between the real output value and the expected output value of the network. It is the most widely used neural network. The BPNN used in the comparison experiment was a three-layer network structure, and the node number of input

layer, hidden layer, and output layer was 16, 32, and 1, respectively. The activation function was relu, loss was the mean square loss function, the optimizer used Adam algorithm, the training epoch was set to 3000, and the batch size was 7.

(4) Kernel Ridge Regression (Wu and Zhao 2020). The kernel ridge regression (KRR) algorithm introduces the kernel method based on the ridge regression algorithm. The independent variable space is mapped to the high-dimensional feature space through the kernel function, and then, the ridge regression method is used to analyze and process the data in the high-dimensional feature space. In other words, kernel ridge regression is a kind of regression algorithm, and its essence is to use ridge regression to make predictions. Ridge regression is linear regression, and it fits a straight line. When facing a nonlinear situation, you can map the independent variable space to the high-dimensional feature space by adding a kernel function, and then perform linear regression in the high-dimensional feature space. For the KRR model used in the comparison experiment, alpha = 1.0, the kernel used a linear kernel, the regularization strength coefficient was set to 0.9, and the rest of the parameters were default values.

(5) Gaussian Process Regression (Arthur et al. 2020). Gaussian process (GP) is a commonly used supervised learning method, which aims to solve regression problems and probabilistic classification problems. Gaussian process regression (GPR) implements the Gaussian process model in the case of regression. It is a nonparametric model that uses Gaussian process priors to perform regression analysis on data. In the comparison experiment, the kernel object used in this method was kernel = DotProduct() + WhiteKernel(), the generator used to initialize the center was random\_state = 0, and the rest of the parameters adopted default values.

(6) Stochastic Gradient Descent Regression (Ighalo et al. 2020). Stochastic gradient descent (SGD) is a simple but highly efficient method, which is mainly used for discriminant learning of linear classifiers under convex loss functions, such as (linear) support vector machines and logistic regression. Stochastic gradient descent regression supports different loss functions and penalties to fit linear regression models. The SGD model used in the comparison experiment, the maximum number of iterations was set as 1000, the stopping criterion tol was set as ‘0.001’, and the rest of the parameters adopted default values.

(7) GAT (Velikovi et al. 2017). Graph attention network (GAT), which aggregates neighbor nodes through the attention mechanism and realizes the adaptive allocation of different neighbor weights. This is different from GCN. The weights of different neighbors in GCN are

**Table 6** Parameters of training model

Training parameters	Mode or value
History time series/d	1
Future time series/d	1
Training set /d	59
Test set /d	7
Feature dimension	44
Learning rate	0.001
Training epoch	3000
Batch size	7
Hidden layer nodes	16
Network model	GCN
Optimizer	Adam
Number of graph convolutional layers	2
Loss function	ReLU
Feature optimization algorithm	LsOA
Population size	30
Max iteration	100

fixed, and they all come from the normalized Laplacian matrix. GAT greatly improves the expressive ability of the graph neural network model. The GAT model used in the comparison experiment was a double-headed three-layer network structure, the number of nodes in the three layers was 45, 16, and 1, respectively. The activation function was LeakyReLU, loss was the mean square loss function, the optimizer used the Adam algorithm, the default learning rate was 0.001, the training epoch was set to 3000, and the batch size was 7.

(8) LSTM (Hochreiter and Schmidhuber 1997). Long short-term memory (LSTM) is a time-cyclic neural network, which is specially designed to solve the long-term dependency problem of general recurrent neural network (RNN). The core concepts of LSTM lie in the cell state and the “gate” structure. The cell state is equivalent to the path of information transmission, so that information can be passed down in the serial connection, which can be regarded as the “memory” of the network. The addition and removal of information is achieved through the “door” structure, which learns which information should be saved or forgotten during the training process. The LSTM model used in the comparison experiment was a three-layer network structure. The number of nodes in the three layers was 45, 200, and 1, respectively. The optimizer used Adam algorithm, the initial learning rate was 0.005, the gradient threshold was set to 1, and after 125 cycles, the learning rate was reduced by multiplying by 0.1, the training epoch was 3000, and the batch size was 7.

(9) T-GCN (Zhao et al. 2020b). Temporal graph convolutional network (T-GCN) is proposed for traffic flow prediction and is mainly composed of two parts. One part is to use the graph convolutional network (GCN) to learn the connection relationship of roads in the traffic network graph, to model the spatial correlation of traffic temporal and spatial sequence data. The other part is to use gate recurrent unit (GRU) to obtain the time dependence of data. The T-GCN model used in the comparison experiment was composed of GCN and GRU. The optimizer used Adam algorithm, the initial learning rate was 0.001, the number of hidden layer nodes was 8, the activation function of the graph convolutional layer was sigmoid, the activation function of the GRU layer was tanh, the loss function was  $loss = ||Y_t - \hat{Y}_t|| + \lambda L_{reg}$ , the training epoch was 3000, and the batch size was 7.

### 3.6 Performance evaluation

In this section, in order to verify the effectiveness of the prediction model, we used the cumulative number of confirmed cases of COVID-19 from January 24, 2020, to March 29, 2020 (66 days in total), in 17 regions of Hubei Province and the Baidu Index of adjusted keywords as the data source, used the data in the data set before March 23, 2020, as the training set, and used the LsOA-GCN prediction model to predict the cumulative number of confirmed cases in the 7 days from March 23, 2020 to March 29, 2020. During training, the input data  $X_i$  is the keyword data on the  $i$ -th day, where  $X_i = [x^1, x^2, \dots, x^j]$ ,  $1 \leq i \leq 66$ ,  $1 \leq j \leq 44$ , and the output value  $Y_{i+1}$  is the cumulative number of confirmed cases on the  $(i+1)$ -th day. Finally, the prediction results were compared with the results of 10 representative prediction methods in current epidemic prediction research, including multiple linear regression, support vector regression, backpropagation neural network, kernel ridge regression, Gaussian process regression, stochastic gradient descent regression, GAT, LSTM, T-GCN and original GCN model. In order to visually test the prediction accuracy of each model, we compared the predicted value of each model with the actual value through evaluation indicators and obtained the RMSE, MAE, MAPE, Acc, R2, and Var of each model. The final results are shown in Tables 7 and 8, where Table 7 shows the comprehensive prediction results of Hubei Province and Table 8 shows the respective prediction results of 17 regions in Hubei Province.

It can be seen from the comprehensive prediction results in Table 7 that compared with the other 10 methods, the RMSE, MAE, and MAPE values of the cumulative number of confirmed cases predicted by the LsOA-GCN model are

**Table 7** Experimental results of LsOA-GCN and other methods–Hubei Province

Data	Metric	Methods	LsOA-GCN	GCN	LSTM	GAT	T-GCN	BPNN	GPR	KRR	MLR	SGD	SVR
Hubei Province	RMSE	<b>14.6572</b>	322.277	1047.15	2729.48	392.285	1459.15	3001.42	986.865	1250.69	1486.09	780.661	
	MAE	<b>5.40047</b>	74.0729	259.864	740.637	308.649	327.823	690.042	312.087	328.534	494.022	222.975	
	MAPE	<b>0.00350</b>	0.02144	0.03701	0.22437	0.55861	0.04046	0.06150	0.13179	0.05606	0.15712	0.04660	
	Acc	<b>0.99880</b>	0.97360	0.91422	0.77642	0.96787	0.88047	0.75414	0.91916	0.89755	0.87827	0.93605	
	R2	<b>1.00000</b>	0.99922	0.99176	0.94404	0.99884	0.98401	0.93233	0.99268	0.98825	0.98341	0.99542	
	Var	<b>1.00000</b>	0.99925	0.99219	0.94621	0.99887	0.98473	0.93541	0.99336	0.98863	0.98484	0.99553	

Optimal values are shown in bold in the table

the smallest, which are 14.65572, 5.4005, and 0.0035, respectively. It shows that the cumulative number of confirmed cases predicted by this method is closer to the actual value, and the error is smaller. For the three types of indicators, Acc, R2, and Var, the values are all close to 1, indicating that the prediction accuracy of the prediction model is high and it shows an absolute advantage. From the regional prediction results in Table 8, we could also see that when predicting the cumulative number of confirmed cases in Xianning, LsOA-GCN ranked third with only a slight difference in the two indicators of MAE and MAPE. In addition, for the three types of indicators, RMSE, MAE and MAPE, LsOA-GCN has an absolute advantage over other forecasting methods. For Acc, in these 17 regions, the value of Acc is within the range of 0.9914 to 0.9990. The closer Acc is to 1, the stronger the fitting degree between the actual value and the predicted value is, the higher the prediction accuracy is, and the more accurate the prediction method is. In addition, we also found that the same prediction model was used for 17 regions in Hubei Province, but the prediction results would be different to some extent. For example, when using LsOA-GCN for prediction, the RMSE value for Wuhan is 50.6662, Ezhou is 1.4078, and Enshi is 0.2504, which shows that the same prediction method has different prediction effects for different regions.

The convergence curve of Fig. 11 also proves the influence of using LsOA for feature screening on the prediction results, where the ordinate is the value of RMSE, and the abscissa is the number of iterations. It can be seen from Fig. 11 that at the beginning of the iteration, the RMSE value drops sharply. As the iteration progresses, the RMSE value gradually decreases and finally stabilizes. The final feature matrix was a 17\*66\*3 matrix, and its keyword features were dyspnea (S), novel coronavirus pneumonia (S) and stuffy nose (X). Among them, the two symptoms of dyspnea (S) and stuffy nose (X) are not unique to COVID-19, but they are easy to recognize and attract attention. For the general population, this is an indicator that may be infected with COVID-19. In addition, novel coronavirus pneumonia (S) is a commonly used term in the epidemic, indicating that as the epidemic continues to spread, people have further information needs for the basic knowledge of COVID-19. Based on the above analysis, we could believe that these three keywords can reflect the prevalence of COVID-19 to a certain extent and have certain predictive value.

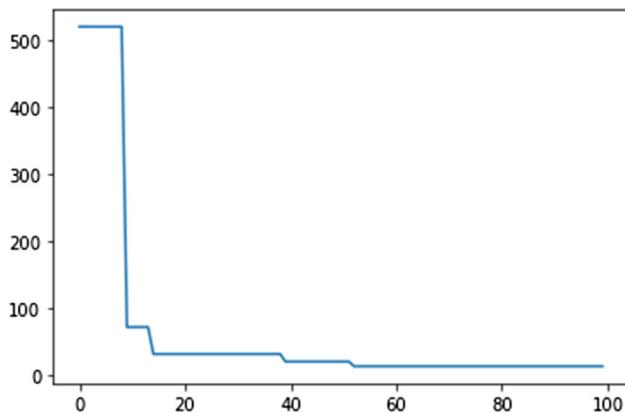
In general, for these six types of evaluation indicators, the LsOA-GCN prediction model has shown superior performance in Hubei Province or in the 17 regions of Hubei and is the best prediction method.

**Table 8** Experimental results of LsOA-GCN and other methods–17 regions in Hubei

Data	Metric	LsOA-GCN	GCN	LSTM	GAT	T-GCN	BPNN	GPR	KRR	MLR	SGD	SVR
Wuhan	RMSE	<b>50.6662</b>	1318.24	4310.42	11,134.0	787.196	6013.73	12,371.2	3992.33	5146.77	6062.66	3204.49
	MAE	<b>41.5418</b>	881.730	3832.77	8734.63	655.635	5071.99	10,897.2	3036.36	4731.97	5738.31	2964.49
	MAPE	<b>0.00083</b>	0.01763	0.07665	0.17467	0.01311	0.10143	0.21792	0.06072	0.09463	0.11475	0.05928
	Acc	<b>0.99899</b>	0.97364	0.91380	0.77735	0.98426	0.87974	0.75260	0.92016	0.89708	0.87876	0.93592
	RMSE	<b>1.40783</b>	38.6784	43.1421	480.575	500.479	58.7762	128.618	151.441	137.801	221.969	57.7294
	MAE	<b>1.17987</b>	26.1887	34.2657	343.917	492.940	48.0810	105.159	122.883	115.187	211.224	51.6682
Ezhou	MAPE	<b>0.00085</b>	0.01879	0.02458	0.24671	0.35362	0.03449	0.07544	0.08815	0.08263	0.15152	0.03706
	Acc	<b>0.99899</b>	0.97225	0.96905	0.65525	0.64098	0.95784	0.90773	0.89136	0.90115	0.84077	0.95839
	RMSE	<b>0.25039</b>	6.72733	9.18368	39.6688	258.790	7.19169	10.6373	55.7457	10.8859	47.8082	14.0554
	MAE	<b>0.20125</b>	4.83126	8.06020	33.0289	204.116	5.77993	8.63636	52.0468	8.79418	46.4684	13.3372
	MAPE	<b>0.00080</b>	0.01917	0.03198	0.13107	0.80999	0.02294	0.03427	0.20654	0.03490	0.18440	0.05293
	Acc	<b>0.99901</b>	0.97330	0.96356	0.84258	-0.02694	0.97146	0.95779	0.77879	0.95680	0.81029	0.94422
Enshi	RMSE	<b>3.22323</b>	83.8352	114.250	786.510	287.178	90.6541	188.220	488.267	198.876	474.418	209.138
	MAE	<b>2.51078</b>	60.6461	96.6943	607.871	271.633	85.5400	160.247	483.764	168.108	461.406	191.306
	MAPE	<b>0.00086</b>	0.02086	0.03326	0.20911	0.09344	0.02943	0.05512	0.16641	0.05783	0.15872	0.06581
	Acc	<b>0.99889</b>	0.97116	0.96070	0.72944	0.90121	0.96882	0.93525	0.83204	0.93159	0.83680	0.92806
	RMSE	<b>1.01808</b>	27.9862	23.0499	299.370	160.273	30.0910	69.7950	126.014	67.6836	147.611	60.1148
	MAE	<b>0.85338</b>	18.5732	19.3720	231.617	131.782	27.0843	41.0881	109.056	39.0791	139.819	53.9503
Huangshi	MAPE	<b>0.00084</b>	0.01830	0.01909	0.22819	0.12983	0.02668	0.04048	0.10744	0.03850	0.13775	0.05315
	Acc	<b>0.99900</b>	0.97243	0.97729	0.70505	0.84210	0.97035	0.93124	0.87585	0.93332	0.85457	0.94077
	RMSE	<b>0.98352</b>	24.3426	24.3273	322.172	513.461	7.52676	64.2156	124.861	65.6214	123.718	62.6998
	MAE	<b>0.79078</b>	16.3154	21.5688	252.195	511.540	6.41007	55.2451	109.424	56.6386	112.163	60.6048
	MAPE	<b>0.00085</b>	0.01758	0.02324	0.27176	0.55123	0.00691	0.05953	0.11791	0.06103	0.12086	0.06531
	Acc	<b>0.99894</b>	0.97377	0.97379	0.65283	0.44670	0.99189	0.93080	0.86545	0.92929	0.86668	0.93244
Jingmen	RMSE	<b>1.68224</b>	46.1738	38.4515	297.973	486.231	45.3420	45.5418	160.072	47.9307	229.225	104.973
	MAE	<b>1.36306</b>	31.7411	48.5009	250.867	463.086	32.2055	39.8533	147.699	41.4051	218.332	91.0327
	MAPE	<b>0.00086</b>	0.02009	0.03070	0.15878	0.29309	0.02038	0.02522	0.09348	0.02621	0.13818	0.05762
	Acc	<b>0.99894</b>	0.97078	0.96301	0.81141	0.69226	0.97130	0.97118	0.8869	0.96966	0.85492	0.93356
	RMSE	<b>0.20008</b>	5.51831	9.17594	88.2524	500.587	17.6562	13.2697	39.1590	12.8428	45.1208	8.19654
	MAE	<b>0.16711</b>	3.60669	6.96984	62.9022	480.283	16.0132	10.6344	34.1029	10.5190	42.8084	7.39665
Qianjiang	MAPE	<b>0.00084</b>	0.01822	0.03520	0.31769	2.42567	0.08087	0.93298	0.17224	0.05313	0.21620	0.03736
	Acc	<b>0.99899</b>	0.97213	0.95366	0.55428	-1.52822	0.91083	0.05371	0.80223	0.93514	0.77212	0.95860
	RMSE	<b>0.01086</b>	0.27478	0.73771	1.60061	32.6534	0.75230	1.36115	2.38661	1.39890	2.05334	0.46899
	MAE	<b>0.00902</b>	0.20933	0.66089	1.25474	27.5576	0.57620	0.96659	2.06045	0.97459	1.95007	0.34058
	MAPE	<b>0.00082</b>	0.01903	0.06008	0.11407	2.50524	0.05238	0.08787	0.18731	0.08860	0.17728	0.03096
	Acc	<b>0.99901</b>	0.97502	0.93294	0.85449	-1.96849	0.93161	0.87626	0.78304	0.87283	0.81333	0.95736

Table 8 (continued)

Data	Metric	LsOA-GCN	GCN	LSTM	GAT	T-GCN	BPNN	GPR	KRR	MLR	SGD	SVR
Shiyan	RMSE	<b>0.71915</b>	19.7104	26.1772	157.081	274.934	37.1574	50.4028	104.800	51.0943	109.373	20.1123
	MAE	<b>0.57114</b>	13.8861	19.2230	131.183	214.508	29.4689	44.9231	98.6214	45.5895	105.590	17.4851
	MAPE	<b>0.00085</b>	0.02066	0.02861	0.19521	0.31921	0.04385	0.06685	0.14676	0.06784	0.15713	0.02602
	Acc	<b>0.99893</b>	0.97067	0.96105	0.76625	0.5987	0.94471	0.92500	0.84405	0.92397	0.83724	0.97007
	RMSE	<b>1.39058</b>	37.5744	51.8354	645.079	468.449	28.2146	98.4990	263.211	96.7216	257.949	49.9525
	MAE	<b>1.12534</b>	25.8064	43.1949	388.660	398.648	23.9810	88.3543	246.807	85.9936	249.472	45.7563
Suizhou	MAPE	<b>0.00086</b>	0.01974	0.03305	0.29737	0.30501	0.01835	0.06760	0.18883	0.06579	0.19087	0.03501
	Acc	<b>0.99894</b>	0.97125	0.96034	0.50644	0.64158	0.97841	0.92464	0.79861	0.92600	0.80264	0.96178
	RMSE	<b>4.26481</b>	16.5381	17.3045	148.241	269.851	20.8620	39.0279	49.2429	51.5244	80.0063	21.3729
	MAE	<b>4.24731</b>	12.1464	14.1336	117.714	222.573	19.5843	28.7870	45.1768	37.9261	76.1957	15.5669
	MAPE	<b>0.00856</b>	0.02449	0.02850	0.23733	0.44873	0.03948	0.05804	0.09108	0.07646	0.15362	0.03138
	Acc	<b>0.99140</b>	0.96666	0.96511	0.70113	0.45595	0.95794	0.92131	0.90072	0.89612	0.83870	0.95691
Tianmen	RMSE	<b>0.60402</b>	16.1987	32.0901	231.922	186.895	37.8016	41.9303	89.9615	42.6721	118.372	21.3962
	MAE	<b>0.48564</b>	10.9648	25.1861	174.238	167.300	35.1417	33.5490	88.2869	34.1701	116.414	20.2326
	MAPE	<b>0.00084</b>	0.01907	0.04380	0.30302	0.29096	0.06112	0.05855	0.15354	0.05943	0.20246	0.03519
	Acc	<b>0.99895</b>	0.97183	0.94419	0.59666	0.67497	0.93426	0.92708	0.84355	0.92579	0.79414	0.96279
	RMSE	<b>32.1149</b>	47.6550	62.4732	535.267	152.921	51.3795	32.2676	113.252	35.3993	167.479	66.8367
	MAE	<b>32.1073</b>	43.2417	58.9477	360.307	122.046	46.4659	<b>24.5145</b>	104.926	27.0204	157.611	60.9347
Xiantao	MAPE	<b>0.03841</b>	0.05172	0.07051	0.43099	0.14599	0.05558	<b>0.02932</b>	0.12551	0.03232	0.18853	0.07289
	Acc	<b>0.96159</b>	0.94300	0.92527	0.35973	0.81708	0.93854	0.96140	0.86453	0.95766	0.79967	0.92005
	RMSE	<b>1.23615</b>	32.7713	25.3289	262.735	427.916	35.6762	64.3560	131.025	65.7992	141.774	98.3506
	MAE	<b>0.98278</b>	22.5675	21.0696	195.030	419.414	30.1233	58.5119	120.568	59.9261	131.521	97.2247
	MAPE	<b>0.00084</b>	0.01921	0.01793	0.16598	0.35695	0.02564	0.04980	0.10261	0.05100	0.11193	0.08274
	Acc	<b>0.99895</b>	0.97211	0.97844	0.77640	0.63582	0.96964	0.94523	0.88849	0.94400	0.87934	0.91630
Xiangyang	RMSE	<b>3.59745</b>	96.6878	175.942	790.340	69.5158	47.2554	105.199	397.664	90.0374	503.097	82.9717
	MAE	<b>2.88957</b>	67.4415	136.026	538.574	55.8627	42.1431	97.4514	391.336	86.0844	463.377	72.2803
	MAPE	<b>0.00082</b>	0.01917	0.03867	0.15309	0.01588	0.01198	0.02770	0.11124	0.02447	0.13172	0.02055
	Acc	<b>0.99898</b>	0.97252	0.94999	0.77534	0.98024	0.98657	0.97010	0.88696	0.97441	0.85699	0.97642
	RMSE	<b>0.97179</b>	27.8529	36.2150	214.007	452.742	70.5527	48.6621	125.226	50.7516	140.394	29.1905
	MAE	<b>0.78178</b>	19.3436	31.0428	166.834	408.107	52.4005	35.5812	112.358	35.6877	125.705	26.9711
Xiaogan	MAPE	<b>0.00084</b>	0.02078	0.03334	0.17920	0.43835	0.05628	0.03822	0.12069	0.03833	0.13502	0.02897
	Acc	<b>0.99896</b>	0.97008	0.96110	0.77013	0.51370	0.92422	0.94773	0.86549	0.94549	0.84920	0.96865



**Fig. 11** Convergence curve

### 3.7 Statistical test

Before statistical analysis, Friedman was used to calculate the average ranking of various prediction methods on different indicators. The results are shown in Table 9. According to Friedman mean rank, LsOA-GCN performed best when considering 17 regions in Hubei Province, ranking first under the four indicators of RMSE, MAE, MAPE, and Acc. Among them, for RMSE, the average ranking of LsOA-GCN was 1.00, which was superior to GCN, LSTM, BPNN, SVR, GPR, MLR, KRR, SGD, T-GCN, GAT, respectively, by 182%, 324%, 329%, and 376%, 482%, 512%, 729%, 800%, 853%, 912%; for MAE, the average ranking of LsOA-GCN was 1.12, which was superior to GCN, LSTM, BPNN, SVR, GPR, MLR, KRR, SGD, T-GCN, GAT, respectively, by 121%, 284%, 295%, 363%, 405%, 426%, 637%, 721%, 753%, 800%; for MAPE, the average ranking of LsOA-GCN was 1.12, which was superior to GCN, LSTM, BPNN, SVR, GPR, MLR, KRR, SGD, T-GCN, GAT, respectively, by 121%, 284%, 289%, 363%, 426%, 426%, 632%, 716%, 753%, 795%; for Acc, the average ranking of LsOA-GCN was 11.00, which was superior to GCN, LSTM, BPNN, SVR, GPR, MLR, KRR, SGD, T-GCN, GAT, respectively, by 17%, 29%, 29%, 34%, 46%, 47%, 66%, 72%, 78%, 82%. These results once again verify the superior performance of the LsOA-GCN prediction method. The calculation method is as follows:

$$\text{percentage} = \frac{\text{rank}_i - \text{rank}_{\text{LsOA-GCN}}}{\text{rank}_{\text{LsOA-GCN}}} \quad (29)$$

where  $\text{rank}_i$  represents the ranking of the prediction method  $i$ ,  $i$  is the prediction method used for comparison, and  $\text{rank}_{\text{LsOA-GCN}}$  is the ranking of LsOA-GCN.

When comparing methods, it is often necessary to perform statistical tests on experimental results. This part used some statistical test methods for comparing methods. First,

**Table 9** Friedman test ranking results

Friedman mean rank	RMSE	MAE	MAPE	Acc
LsOA-GCN	<b>1.00</b>	<b>1.12</b>	<b>1.12</b>	<b>11.00</b>
GCN	2.82	2.47	2.47	9.18
LSTM	4.24	4.29	4.29	7.76
GAT	10.12	10.06	10.00	1.94
T-GCN	9.53	9.53	9.53	2.47
BPNN	4.29	4.41	4.35	7.76
GPR	5.82	5.65	5.88	5.94
KRR	8.29	8.24	8.18	3.76
MLR	6.12	5.88	5.88	5.88
SGD	9.00	9.18	9.12	3.06
SVR	4.76	5.18	5.18	7.24

The three types of indicators, RMSE, MAE, and MAPE, should be as small as possible, and close to 0. Acc should be as large as possible, and close to 1

a nonparametric Friedman test was used to determine whether there are significant differences in the performance of all methods under each indicator. In order to make a reliable comparison, we considered 11 methods and tested them on 4 different indicators for 17 regions, where the indicator used in the first group was RMSE, the second group was MAE, the third group was MAPE, and the fourth group was Acc. First, the Friedman test requires the average ranking to be calculated first. Then, the Friedman test should consider the critical value obtained at the significance level ( $\alpha = 0.05, 0.1$ ) and compare the critical value with Friedman's statistical results to determine whether there is evidence that the null hypothesis is false. For the tests of these four indicators, we found that the results rejected the null hypothesis, which indicates that the performance of each group of methods has significant differences.

After the difference was determined, a "post hoc test" was needed to determine which methods have statistical differences in performance. For this purpose, we used the Bonferroni–Dunn test (Zar 2010). This test compared the proposed method, LsOA-GCN, with the other 10 methods; that is, the average ranking difference of each method was compared with the critical difference (CD). If the difference is greater than the critical difference, it means that the method with a good average ranking is statistically superior to the method with a bad average ranking; otherwise, there is no statistical difference between the two. The calculation formula of the critical difference is as follows:

$$CD = q_{\alpha} \sqrt{\frac{k(k+1)}{6N}} \quad (30)$$

where  $k$  is the number of methods used for comparison,  $N$  is the number of data sets, and the commonly used value of

$q_\alpha$  can be obtained by looking up the table. See Appendix B for the table.

The results of the Bonferroni–Dunn test are shown in Fig. 12. In order to facilitate observation and comparison, here we only used three indicators: RMSE, MAE, and MAPE. And the average rankings of 11 methods on these 3 indicators are all reflected in the bar chart. The value corresponding to the horizontal line is equivalent to the threshold, that is, the ranking of the compared method plus the value of CD. This part defined two thresholds at the significance level of 0.05 and 0.1, respectively. Each group used a different color to identify (the first group was blue, the second group was red, the third group was green), and the threshold was also distinguished by different colors.

It is worth noting that the method used for comparison (LsOA-GCN) can outperform those whose average ranking is higher than the threshold line (i.e., the height of the bar exceeds its corresponding line). It can be seen from Fig. 12 that in these three groups, LsOA-GCN is better than all methods, with an average ranking of 1.00, 1.12, and 1.12, respectively. In Group 1, LsOA-GCN is significantly better than the other 7 methods except GCN, LSTM and BPNN at the significance level of 0.05 and 0.1, respectively. In Group 2 and Group 3, LsOA-GCN is also significantly better than these 7 methods of GAT, T-GCN, GPR, KRR, MLR, SGD, and SVR at these two significant levels.

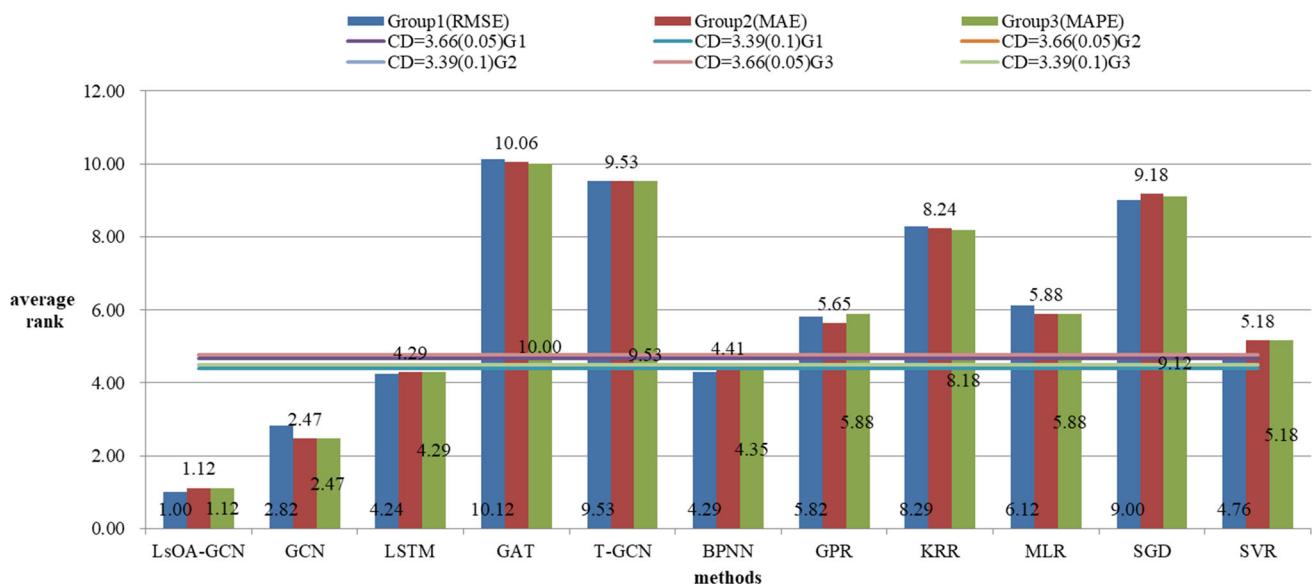
In addition to the Bonferroni–Dunn test, this study also considered Holm's method (Holm, 1979). This step: calculate the  $p$  value, sort the  $p$  value, and compare the  $p$  value with  $\alpha/i$ . If  $p < \alpha/i$ , reject the null hypothesis, that is, the difference is significant, where  $\alpha$  is the significance level and  $i$  is the method number. Here, we also only considered

the three indicators: RMSE, MAE, and MAPE. The test results of these three groups are shown in Tables 10, 11 and 12, respectively. It can be seen that for these three indicators, LsOA-GCN is significantly better than other methods at the significance level of 0.05 and 0.1, respectively. The above test results once again verify the superiority of the LsOA-GCN model.

## 4 Conclusion

Aiming at the spreading characteristics of the COVID-19, this paper proposed a graph convolutional network prediction model based on the lioness optimization algorithm (LsOA-GCN). The LsOA-GCN model is mainly composed of two parts. One is the construction of the feature matrix for capturing temporal information, and the other is the graph convolution operation for capturing spatial information. First, the keyword data were sorted out through Spearman correlation analysis with a time lag, and the initial feature matrix was constructed to capture the time series information between the current number of confirmed cases and the Baidu Search Index at different periods in the past. Then LsOA was used to further filter the feature matrix to improve the accuracy of the model and avoid unnecessary feature interference. Finally, the feature matrix was used as input to obtain the spatial features of the epidemic-related data through the graph convolution network, to get the prediction results.

We used the cumulative confirmed case data of 17 regions in Hubei Province from January 24, 2020 to March 29, 2020 to test the effectiveness of the prediction model,



**Fig. 12** Bonferroni–Dunn test for different methods and benchmark groups with  $\alpha = 0.05$  and  $\alpha = 0.1$

**Table 10** Holm's method test results of the first group

LsOA-GCN vs	Rank	z-value	p-value	a/i(0.05)	a/i(0.1)
GCN	2.82	-3.621	0.00029	0.005	0.01
LSTM	4.24	-3.621	0.00029	0.00556	0.01111
GAT	10.12	-3.621	0.00029	0.00625	0.0125
T-GCN	9.53	-3.621	0.00029	0.00714	0.01429
BPNN	4.29	-3.621	0.00029	0.00833	0.01667
GPR	5.82	-3.621	0.00029	0.01	0.02
KRR	8.29	-3.621	0.00029	0.0125	0.025
MLR	6.12	-3.621	0.00029	0.01667	0.03333
SGD	9.00	-3.621	0.00029	0.025	0.05
SVR	4.76	-3.621	0.00029	0.05	0.1

**Table 11** Holm's method test results of the second group

LsOA-GCN vs	Rank	z-value	p-value	a/i(0.05)	a/i(0.1)
GCN	2.47	-3.621	0.00029	0.005	0.01
LSTM	4.29	-3.621	0.00029	0.00556	0.01111
GAT	10.06	-3.621	0.00029	0.00625	0.0125
T-GCN	9.53	-3.621	0.00029	0.00714	0.01429
BPNN	4.41	-3.621	0.00029	0.00833	0.01667
KRR	8.24	-3.621	0.00029	0.01	0.02
SGD	9.18	-3.621	0.00029	0.0125	0.025
SVR	5.18	-3.621	0.00029	0.01667	0.03333
GPR	5.65	-3.527	0.00042	0.025	0.05
MLR	5.88	-3.527	0.00042	0.05	0.1

and used multiple indicators to evaluate the prediction model, namely RMSE, MAE, MAPE, Accuracy, Var and R2. And these results were compared with 10 typical prediction methods, including multiple linear regression, support vector regression, backpropagation neural network, kernel ridge regression, Gaussian process regression, stochastic gradient descent regression, GAT, LSTM and T-GCN. By analyzing the experimental results, we can draw the following conclusions:

- (1) Time-delay analysis can improve the accuracy of epidemic prediction. Firstly, by observing the trend graph of the features and the number of confirmed cases (Fig. 9), it can be found that the number of confirmed cases has a certain time lag relative to the search index. Then, the optimal lag days corresponding to each feature were determined by the correlation analysis (Table 1). Finally, we sorted out the feature data and found that the prediction results

**Table 12** Holm's method test results of the third group

LsOA-GCN vs	Rank	z-value	p-value	a/i(0.05)	a/i(0.1)
GCN	2.47	-3.621	0.00029	0.005	0.01
LSTM	4.29	-3.621	0.00029	0.00556	0.01111
GAT	10.00	-3.621	0.00029	0.00625	0.0125
T-GCN	9.53	-3.621	0.00029	0.00714	0.01429
BPNN	4.35	-3.621	0.00029	0.00833	0.01667
KRR	8.18	-3.621	0.00029	0.01	0.02
SGD	9.12	-3.527	0.00029	0.0125	0.025
SVR	5.18	-3.527	0.00029	0.01667	0.03333
GPR	5.88	-3.621	0.00035	0.025	0.05
MLR	5.88	-3.621	0.00035	0.05	0.1

based on the time-delay feature data were significantly better than the original feature data (Tables 3 and 4).

- (2) GCN can capture spatial information from feature data, which is helpful for prediction models to improve prediction accuracy. According to the characteristics of the epidemic transmission, this paper used the population migration index to simulate the population mobility during the epidemic, and constructed an adjacency matrix based on this. Finally, GCN was used to establish the spatial connection of various regions in Hubei Province, and to realize the prediction of the COVID-19 in various regions of Hubei Province. Through experiments, we found that compared with other prediction models, the GCN and LsOA-GCN models achieved better results in various indicators (Tables 7 and 8).
- (3) There is also a significant influence between the selection of prediction features and the prediction performance of the model. Compared with the single GCN model, LsOA-GCN performed better in all indicators. It can be seen that feature selection is helpful to improve the prediction accuracy of the model.

In the future, we will consider using the LsOA-GCN model for financial market forecasting, traffic forecasting, and environmental pollution forecasting, etc., to determine that this prediction model can also have excellent performance in other fields.

## Appendix A

See Tables 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28.

**Table 13** Correlation analysis between keywords and the cumulative number of confirmed cases—Enshi

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Center for Disease Control and Prevention(S)	<b>0.3237</b>	0.2770	0.2437	0.2224	0.1610	0.1188	0.1063	0.1163	0.0809	0.0589
Red Cross Society(S)	0.1399	0.0867	0.0537	0.0108	0.0334	0.0960	0.1552	0.2167	0.2669	0.3201
Spring Festival(S)	0.7663	0.7712	0.7875	0.7949	0.7854	0.7765	0.7962	<b>0.8015</b>	0.7757	0.7762
Health Commission(S)	0.2105	0.1429	0.0856	0.0532	0.0088	0.0330	0.1001	0.1813	0.2524	0.3022
vaccine(S)	0.3071	0.2645	0.2108	0.1769	0.1539	0.1238	0.0634	0.0331	0.0157	0.0834
SARS(S)	0.1259	0.1911	0.2494	0.3237	0.4018	0.4849	0.5659	0.6570	0.7322	0.8226
hand washing(S)	0.2000	<b>0.2014</b>	0.1964	0.1444	0.1514	0.1490	0.1142	0.1191	0.1178	0.1100
children(S)	0.6004	0.6214	0.6559	0.6562	0.7348	0.7266	0.7277	<b>0.7373</b>	0.7185	0.7226
mask(S)	0.0208	0.0847	0.1458	0.1917	0.2589	0.3310	0.4201	0.4952	0.5788	0.6689
coronavirus(S)	0.1105	0.1764	0.2462	0.3150	0.3892	0.4586	0.5426	0.6337	0.7304	0.8266
novel(S)	0.1115	0.0568	0.0054	0.0367	0.1020	0.1569	0.2075	0.2778	0.3638	0.4575
disinfection(S)	0.0209	0.0126	0.0572	0.0615	0.0897	0.1562	0.2167	0.2663	0.2958	0.3195
lockdown of the city(S)	0.0488	0.0028	0.0574	0.0935	0.1261	0.1683	0.2369	0.2852	0.3230	0.3804
Wuhan(S)	0.1004	0.1595	0.2221	0.2898	0.3520	0.4186	0.4960	0.5864	0.6698	0.7573
Zhong Nanshan(S)	0.0094	0.0497	0.1132	0.1730	0.2276	0.2998	0.3869	0.4652	0.5348	0.6108
epidemic(S)	0.8986	0.8896	<b>0.8989</b>	0.8976	0.8986	0.8924	0.8557	0.8434	0.8359	0.8362
flu(S)	0.1138	0.1488	0.2081	0.2690	0.3367	0.3863	0.4329	0.5283	0.6038	0.7011
diarrhea(S)	0.0489	0.0013	0.0413	0.0870	0.1532	0.2012	0.2099	0.2562	0.3260	0.3925
dry cough(S)	0.0980	0.1534	0.2158	0.2839	0.3501	0.3957	0.4620	0.5332	0.6147	0.7045
NCPS(S)	<b>0.6958</b>	0.6789	0.6468	0.6117	0.5859	0.5689	0.5436	0.5157	0.4965	0.4673
novel coronary pneumonia(S)	0.9213	0.9189	0.9204	<b>0.9220</b>	0.9195	0.9219	0.9153	0.9155	0.9216	0.9210
COVID-19(S)	<b>0.7571</b>	0.7346	0.7228	0.7040	0.6897	0.6868	0.6532	0.6181	0.6106	0.5890
novel coronavirus pneumonia(S)	0.0949	0.0396	0.0179	0.0827	0.1513	0.2250	0.3033	0.3795	0.4580	0.5335
asymptomatic infection(S)	<b>0.5755</b>	0.5617	0.5457	0.5085	0.4689	0.4577	0.4375	0.4303	0.3861	0.3714
aerosol transmission(S)	<b>0.3040</b>	0.2676	0.2470	0.2115	0.2107	0.1769	0.1350	0.0918	0.0529	0.0106
pneumonia(S)	0.0406	0.1065	0.1721	0.2390	0.3068	0.3781	0.4587	0.5505	0.6380	0.7277
cough(S)	0.1876	0.1763	0.1308	0.0739	0.0104	0.0456	0.0840	0.1221	0.1658	0.2248
fever(S)	0.1696	0.1498	0.1726	0.2230	0.2277	0.3341	0.4036	0.4415	0.4922	0.5547
dyspnea(S)	0.2037	0.2300	0.2489	0.2780	0.3380	0.3938	0.4485	0.4939	0.5738	0.6096
high temperature(S)	0.1508	0.2116	0.2538	0.3130	0.3517	0.4087	0.4949	0.5348	0.5703	0.5842
novel coronavirus(X)	0.0633	0.0044	0.0716	0.1291	0.1955	0.2654	0.3437	0.4327	0.5199	0.6039
SARS-CoV-2(S)	0.3584	0.3674	<b>0.3767</b>	0.3400	0.3505	0.3212	0.3329	0.2725	0.2802	0.3066
Spring Festival(X)	0.8245	0.8293	0.8361	0.8405	0.8498	<b>0.8519</b>	0.8234	0.8250	0.8339	0.8216
Red Cross Society(X)	0.2608	0.1989	0.1430	0.0956	0.0515	0.0168	0.0063	0.0644	0.1592	0.2471
government(X)	0.2272	0.2154	0.2048	0.1831	0.1578	0.1304	0.2128	0.2992	0.3046	0.2948

Table 13 (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Health Commission(X)	0.0115	0.0852	0.0866	0.1236	0.1801	0.2616	0.3120	0.3981	0.4769	0.5501
medical(X)	0.1648	0.1071	0.0632	0.0307	0.0030	0.0479	0.0750	0.1436	0.1928	0.2449
vaccine(X)	0.2126	0.1872	0.1723	0.1473	0.1208	0.0751	0.0441	0.0004	0.0479	0.1020
SARS(X)	0.0709	0.1464	0.2098	0.2727	0.3383	0.3876	0.4290	0.5170	0.5970	0.6744
children(X)	0.5049	0.4898	0.5237	0.5064	0.5366	0.5671	0.5990	0.5832	0.5656	0.5984
mask(X)	0.0837	0.0245	0.0004	0.0336	0.0738	0.0969	0.1338	0.2130	0.2787	0.3318
infection(X)	0.3101	0.3351	0.3469	0.3617	0.3832	0.4057	0.4317	0.4564	0.4810	0.5068
coronavirus(X)	<b>0.6522</b>	0.6120	0.5973	0.5751	0.5537	0.5444	0.5554	0.5130	0.4623	0.4347
disinfection(X)	0.4771	0.4875	0.4984	0.5097	0.5216	0.5340	0.5469	0.5605	0.5747	0.5897
nucleic acid(X)	<b>0.5088</b>	0.4744	0.4670	0.4350	0.4056	0.4147	0.4210	0.3634	0.2985	0.2519
lockdown of the city(X)	0.5063	0.5199	0.5396	0.5512	0.5603	0.5691	0.5785	0.5908	0.5824	0.5974
Wuhan(X)	0.0173	0.0265	0.0831	0.1615	0.2280	0.2789	0.3194	0.3952	0.4650	0.5405
Zhong Nanshan(X)	0.1785	0.1177	0.0839	0.0544	0.0005	0.0723	0.1071	0.1709	0.2299	0.2987
epidemic(X)	<b>0.5325</b>	0.4863	0.4533	0.4208	0.3815	0.3573	0.3554	0.3022	0.2364	0.1925
flu(X)	<b>0.4801</b>	0.4305	0.4019	0.3980	0.4113	0.4049	0.3944	0.3688	0.3359	0.3030
diarrhea(X)	0.0599	0.0558	0.0482	0.0402	0.0095	0.0023	0.0147	0.0184	0.0108	0.0029
stuffy nose(X)	0.0522	0.0081	0.0445	0.0667	0.0963	0.1350	0.1517	0.1926	0.2229	0.2283
dry cough(X)	0.2052	0.1602	0.1053	0.0431	0.0232	0.0863	0.0967	0.1626	0.2411	0.3265
NCP(X)	<b>0.2270</b>	0.1987	0.1489	0.1337	0.1284	0.1504	0.1738	0.1137	0.0431	0.0154
close contact(X)	<b>0.8081</b>	0.7922	0.7893	0.7727	0.7497	0.7511	0.7850	0.7556	0.7172	0.6975
suspected case(X)	<b>0.5266</b>	0.4850	0.4632	0.4259	0.3915	0.3464	0.3534	0.3217	0.2582	0.2127
pneumonia(X)	<b>0.5749</b>	0.5312	0.5144	0.4860	0.4571	0.4410	0.4447	0.3957	0.3382	0.3007
cough(X)	0.3457	0.4151	0.4588	0.4968	0.4867	0.4583	0.4679	0.4962	0.5309	0.5967
dyspnea(X)	0.1800	0.1277	0.0982	0.0612	0.0303	0.0334	0.0465	0.0060	0.0047	0.0002
virus pneumonia(X)	0.5401	0.5466	0.5183	0.5236	0.5360	0.5790	0.5719	0.6141	0.6469	0.6569
novel coronavirus(X)	<b>0.7072</b>	0.6667	0.6511	0.6290	0.6073	0.5949	0.6239	0.5868	0.5229	0.4892

Optimal values are shown in bold in the table

**Table 14** Correlation analysis between keywords and the cumulative number of confirmed cases—Ezhou

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Center for Disease Control and Prevention(S)	0.0168	0.0281	0.0481	0.0660	0.0792	0.1027	0.1166	0.1398	0.1826	0.2229
Red Cross Society(S)	0.0167	0.0030	0.0472	0.0987	0.1421	0.1985	0.2038	0.2480	0.2961	0.3389
Spring Festival(S)	0.7893	0.7941	0.8063	0.8224	<b>0.8229</b>	0.8186	0.8105	0.8139	0.7999	0.7852
Health Commission(S)	0.1318	0.0787	0.0236	0.0205	0.0715	0.1166	0.1856	0.2420	0.2873	0.3491
vaccine(S)	0.1600	0.1449	0.1154	0.0732	0.0282	0.0183	0.0469	0.1039	0.1295	0.1887
SARS(S)	0.2443	0.3114	0.3677	0.4376	0.5175	0.5707	0.6497	0.7397	0.8167	<b>0.8964</b>
hand washing(S)	<b>0.0994</b>	0.0860	0.0927	0.0916	0.0837	0.0806	0.0774	0.0740	0.0357	0.0236
children(S)	0.2256	0.2237	0.2212	0.2530	<b>0.2707</b>	0.2490	0.2133	0.1946	0.2306	0.2250
mask(S)	0.0306	0.0836	0.1487	0.1995	0.2490	0.3239	0.3955	0.4616	0.5295	0.6040
coronavirus(S)	0.1599	0.2137	0.2711	0.3356	0.3957	0.4592	0.5248	0.5994	0.6810	0.7786
novel(S)	0.0689	0.1003	0.1486	0.2103	0.2664	0.3126	0.3782	0.4432	0.5101	0.5675
disinfection(S)	0.1245	0.1410	0.1480	0.1951	0.2198	0.2254	0.2413	0.2651	0.2860	0.3033
lockdown of the city(S)	0.1007	0.0800	0.0507	0.0259	0.0140	0.0568	0.1102	0.1515	0.1825	0.2367
Wuhan(S)	0.1335	0.1900	0.2554	0.3198	0.3847	0.4597	0.5432	0.6262	0.7146	0.8123
Zhong Nanshan(S)	0.0960	0.1555	0.2224	0.2742	0.3294	0.4029	0.4747	0.5457	0.6185	0.7187
epidemic(S)	0.9057	<b>0.9162</b>	0.9144	0.9055	0.8977	0.8986	0.8860	0.8739	0.8660	0.8608
flu(S)	0.1302	0.1784	0.2393	0.2999	0.3669	0.4260	0.4920	0.5649	0.6457	0.7453
diarrhea(S)	0.0943	0.1347	0.1780	0.2171	0.2679	0.3225	0.3813	0.4375	0.4859	0.5406
dry cough(S)	0.2608	0.3084	0.3449	0.3954	0.4422	0.4994	0.5480	0.6100	0.6757	0.7368
NCPS(S)	<b>0.5195</b>	0.4977	0.4718	0.4630	0.4472	0.4243	0.4027	0.3809	0.3497	0.3304
novel coronary pneumonia(S)	<b>0.9234</b>	0.9183	0.9199	0.9188	0.9172	0.9149	0.9020	0.9032	0.8952	0.8836
COVID-19(S)	0.5771	<b>0.5842</b>	0.5753	0.5632	0.5429	0.5312	0.5344	0.5121	0.5007	0.5023
novel coronavirus pneumonia(S)	0.2868	0.2523	0.2264	0.1878	0.1524	0.1170	0.0647	0.0098	0.0503	0.1045
asymptomatic infection(S)	<b>0.4011</b>	0.3851	0.3605	0.3438	0.3217	0.2889	0.2632	0.2276	0.1960	0.1855
aerosol transmission(S)	<b>0.2472</b>	0.2243	0.1940	0.1739	0.1465	0.1360	0.1039	0.0734	0.0580	0.0319
pneumonia(S)	0.0894	0.1492	0.2122	0.2776	0.3417	0.4125	0.4923	0.5719	0.6609	0.7541
cough(S)	0.0266	0.0124	0.0745	0.1216	0.1616	0.2136	0.2691	0.3290	0.3878	0.4671
fever(S)	0.3552	0.3369	0.3733	0.4224	0.4120	0.4752	0.5236	0.5676	0.6176	0.6711
dyspnea(S)	0.1875	0.2445	0.2743	0.3002	0.3475	0.3904	0.4250	0.4776	0.5238	0.5677
high temperature(S)	0.2199	0.2490	0.2892	0.3294	0.3650	0.4139	0.4514	0.5242	0.5747	0.6221
novel coronavirus(S)	0.0634	0.1205	0.1831	0.2471	0.3135	0.3871	0.4648	0.5465	0.6335	0.7192
SARS-CoV-2(S)	0.2497	0.2479	0.2520	<b>0.2562</b>	0.2222	0.2196	0.2234	0.2272	0.1909	0.1768
Spring Festival(X)	0.7922	0.7826	0.7875	<b>0.7946</b>	0.7939	0.7811	0.7750	0.7695	0.7832	0.7806
Red Cross Society(X)	0.1895	0.1684	0.1338	0.0883	0.0267	0.0238	0.0673	0.1224	0.1799	0.2176
government(X)	0.3444	0.3551	0.3277	0.2777	0.2421	0.2455	0.3234	0.3860	<b>0.3861</b>	0.3791

**Table 14** (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Health Commission(X)	0.1056	0.1434	0.1680	0.2294	0.3045	0.3563	0.3885	0.4522	0.5326	0.6050
medical(X)	0.1187	0.0993	0.0648	0.0265	0.0129	0.0473	0.0859	0.1387	0.1825	0.2311
vaccine(X)	0.2192	0.2079	0.1917	0.1563	0.1107	0.0633	0.0168	0.0461	0.1000	0.1537
SARS(X)	0.1632	0.2022	0.2590	0.3199	0.3837	0.4460	0.4964	0.5511	0.6319	0.7090
children(X)	0.4266	0.4081	0.4320	0.4571	0.5035	0.5350	0.5617	0.5427	0.5760	0.6100
mask(X)	0.0876	0.0624	0.0214	0.0311	0.0893	0.1361	0.1851	0.2380	0.2979	0.3538
infection(X)	0.2880	0.3106	0.3191	0.3384	0.3600	0.3799	0.4007	0.4225	0.4453	0.4694
coronavirus(X)	0.5795	<b>0.5824</b>	0.5651	0.5358	0.5040	0.4834	0.4643	0.4207	0.3974	0.3682
disinfection(X)	0.4820	0.4927	0.5039	0.5156	0.5278	0.5406	0.5540	0.5680	0.5827	0.5981
nucleic acid(X)	<b>0.5021</b>	0.4973	0.4749	0.4322	0.3895	0.3627	0.3365	0.2950	0.2508	0.2158
lockdown of the city(X)	0.5894	0.6028	0.6037	0.6129	0.6260	0.6414	0.6547	0.6694	0.6872	0.7037
Wuhan(X)	0.0130	0.0329	0.0836	0.1344	0.1868	0.2429	0.3178	0.3881	0.4619	0.5294
Zhong Nanshan(X)	0.1143	0.0528	0.0995	0.0334	0.0920	0.1595	0.2271	0.2887	0.3528	0.4377
epidemic(X)	<b>0.4186</b>	0.4153	0.3898	0.3497	0.3085	0.2796	0.2500	0.2039	0.1526	0.1153
flu(X)	<b>0.4764</b>	0.4740	0.4652	0.4571	0.4329	0.3963	0.3596	0.3181	0.3015	0.2708
diarrhea(X)	0.0736	0.0854	0.0934	0.1018	0.1126	0.1282	0.1227	0.1348	0.1498	0.1609
stuffy nose(X)	0.1002	0.0848	0.0504	0.0092	0.0159	0.0400	0.0525	0.1051	0.1419	0.1893
dry cough(X)	0.1648	0.1506	0.1132	0.0645	0.0129	0.0289	0.0630	0.1169	0.1733	0.2295
NCP(X)	<b>0.2111</b>	0.2079	0.1700	0.1284	0.0982	0.0711	0.0571	0.0460	0.0380	0.0154
close contact(X)	<b>0.7652</b>	0.7549	0.7376	0.7168	0.6961	0.6830	0.6748	0.6604	0.6414	0.6139
suspected case(X)	<b>0.4923</b>	0.4622	0.4440	0.4081	0.3746	0.3471	0.3270	0.2931	0.2580	0.2210
pneumonia(X)	0.4894	<b>0.4918</b>	0.4717	0.4381	0.4025	0.3789	0.3578	0.3140	0.2703	0.2383
cough(X)	0.3218	0.3518	0.4024	0.4576	0.4575	0.4397	0.4285	0.4385	0.4650	0.4799
dyspnea(X)	0.2056	0.1868	0.1551	0.1258	0.0865	0.0584	0.0001	0.0496	0.0430	0.0381
virus pneumonia(X)	0.6049	0.5914	0.5974	0.6067	0.6213	0.6277	0.6076	0.6084	0.6244	0.6350
novel coronavirus(X)	<b>0.6216</b>	0.6207	0.6030	0.5746	0.5200	0.4953	0.4638	0.4382	0.4055	0.3696

Optimal values are shown in bold in the table

**Table 15** Correlation analysis between keywords and the cumulative number of confirmed cases—Huanggang

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Center for Disease Control and Prevention(S)	0.1299	0.0954	0.0454	0.0049	0.0485	0.1074	0.1605	0.2167	0.2956	0.3570
Red Cross Society(S)	0.2596	0.2178	0.1694	0.1167	0.0732	0.0214	0.0191	0.0801	0.1300	0.1864
Spring Festival(S)	0.6785	0.7261	0.7618	0.8213	0.8816	0.9108	<b>0.9190</b>	0.9111	0.9082	0.9051
Health Commission(S)	0.0180	0.0621	0.1082	0.1582	0.2167	0.2714	0.3320	0.3942	0.4809	0.5640
vaccine(S)	0.1731	0.1264	0.0973	0.0433	0.0008	0.0492	0.1206	0.1754	0.2547	0.3391
SARS(S)	0.1809	0.2464	0.3158	0.3883	0.4645	0.5473	0.6325	0.7258	0.8232	0.9240
hand washing(S)	0.1512	0.1119	0.0750	0.1122	0.0799	0.0377	0.0054	0.0353	0.0606	0.0820
children(S)	0.6428	0.6386	0.6288	0.6213	0.6516	0.6458	0.6562	<b>0.6630</b>	0.6500	0.6414
mask(S)	0.0872	0.1443	0.1903	0.2445	0.2955	0.3601	0.4270	0.5070	0.5915	0.6807
coronavirus(S)	0.1362	0.1987	0.2619	0.3319	0.4069	0.4853	0.5696	0.6577	0.7510	0.8520
novel(S)	0.0603	0.1217	0.1779	0.2338	0.2939	0.3535	0.4180	0.4857	0.5752	0.6535
disinfection(S)	0.1311	0.1056	0.0734	0.0296	0.0154	0.0772	0.1127	0.1548	0.2260	0.2679
lockdown of the city(S)	0.1662	0.1294	0.0764	0.0433	0.0100	0.0486	0.1124	0.1706	0.2289	0.2840
Wuhan(S)	0.1277	0.1865	0.2481	0.3161	0.3878	0.4609	0.5382	0.6240	0.7222	0.8185
Zhong Nanshan(S)	0.1051	0.1604	0.2313	0.2970	0.3687	0.4407	0.5174	0.6047	0.6856	0.7786
epidemic(S)	0.8731	0.8730	0.8784	<b>0.8812</b>	0.8699	0.8601	0.8397	0.8231	0.7989	0.7857
flu(S)	0.0954	0.1571	0.2088	0.2684	0.3370	0.4077	0.4794	0.5715	0.6641	0.7642
diarrhea(S)	0.1043	0.0621	0.0272	0.0146	0.0512	0.1152	0.1578	0.2364	0.2895	0.3523
dry cough(S)	0.1227	0.1683	0.2327	0.2983	0.3576	0.4284	0.5065	0.5858	0.6761	0.7646
NCPS(S)	<b>0.6990</b>	0.6771	0.6584	0.6313	0.6221	0.5939	0.5733	0.5477	0.5145	0.4840
novel coronary pneumonia(S)	0.9133	0.9179	0.9162	0.9179	0.9221	<b>0.9289</b>	0.9213	0.9221	0.9066	0.9021
COVID-19(S)	<b>0.7744</b>	0.7644	0.7677	0.7656	0.7652	0.7571	0.7348	0.7157	0.6901	0.6690
novel coronavirus pneumonia(S)	0.2566	0.2148	0.1735	0.1280	0.0813	0.0244	0.0254	0.0907	0.1547	0.2183
asymptomatic infection(S)	<b>0.6947</b>	0.6629	0.6477	0.6321	0.6175	0.5998	0.5832	0.5559	0.5269	0.4890
aerosol transmission(S)	<b>0.3725</b>	0.3480	0.3278	0.2961	0.2589	0.2183	0.1813	0.1464	0.1240	0.0963
pneumonia(S)	0.0720	0.1308	0.1912	0.2540	0.3236	0.3962	0.4763	0.5579	0.6524	0.7399
cough(S)	0.1527	0.2004	0.2663	0.3387	0.3256	0.3959	0.3778	0.4444	0.4285	0.5015
fever(S)	0.2157	0.2431	0.2679	0.3293	0.3754	0.4255	0.4871	0.5344	0.6138	0.6984
dyspnea(S)	0.0021	0.0340	0.0817	0.1446	0.2000	0.2386	0.3138	0.3638	0.4328	0.4920
high temperature(S)	0.0606	0.1303	0.1753	0.2349	0.2925	0.3601	0.4133	0.4957	0.5672	0.6590
novel coronavirus(S)	0.0249	0.0830	0.1417	0.2055	0.2750	0.3454	0.4239	0.5059	0.5945	0.6851
SARS-CoV-2(S)	<b>0.4572</b>	0.4306	0.4159	0.3922	0.3688	0.3557	0.3519	0.3229	0.3240	0.3196
Spring Festival(X)	0.7969	0.7969	0.8055	0.8007	<b>0.8126</b>	0.8122	0.8115	0.8051	0.8005	0.7885
Red Cross Society(X)	0.2368	0.2040	0.1686	0.1351	0.0941	0.0401	0.0298	0.0947	0.1661	0.2298
government(X)	0.2891	0.2645	0.2154	0.2003	0.1912	0.1644	0.2159	0.2857	0.3286	0.3331

Table 15 (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Health Commission(X)	0.0475	0.1061	0.1355	0.2039	0.2669	0.3367	0.3885	0.4571	0.5400	0.6073
medical(X)	0.1034	0.0720	0.0334	0.0127	0.0269	0.0654	0.1106	0.1602	0.2056	0.2571
vaccine(X)	0.1989	0.1813	0.1405	0.1274	0.0977	0.0521	0.0020	0.0682	0.1303	0.2020
SARS(X)	0.1688	0.2311	0.2930	0.3389	0.4096	0.4682	0.5344	0.6007	0.6835	0.7711
children(X)	0.4165	0.4024	0.4315	0.4660	0.5125	0.5488	0.5318	0.5075	0.5590	0.5325
mask(X)	0.0702	0.0503	0.0185	0.0035	0.0562	0.0980	0.1605	0.2131	0.2760	0.3227
infection(X)	0.3208	0.3333	0.3593	0.3757	0.3993	0.4219	0.4455	0.4702	0.4962	0.5235
coronavirus(X)	<b>0.5996</b>	0.5813	0.5509	0.5561	0.5331	0.5081	0.4760	0.4396	0.4042	0.3691
disinfection(X)	0.5890	0.5931	0.6069	0.6212	0.6362	0.6519	0.6684	0.6857	0.7038	0.7229
nucleic acid(X)	<b>0.4791</b>	0.4717	0.4594	0.4541	0.4318	0.3939	0.3475	0.2981	0.2708	0.2385
lockdown of the city(X)	0.5486	0.5642	0.5704	0.5630	0.5759	0.5835	0.5947	0.6119	0.6260	0.6423
Wuhan(X)	0.0333	0.0805	0.1408	0.1767	0.2323	0.2732	0.3351	0.3908	0.4743	0.5504
Zhong Nanshan(X)	0.0531	0.0148	0.0431	0.0725	0.1320	0.1840	0.2551	0.3346	0.4127	0.4900
epidemic(X)	0.2732	0.2378	0.1891	0.1734	0.1288	0.0774	0.0234	0.0363	0.0978	0.1558
flu(X)	<b>0.4246</b>	0.4149	0.4029	0.4064	0.3937	0.3861	0.3532	0.3059	0.2524	0.2139
diarrhea(X)	0.0514	0.0572	0.0141	0.0116	0.0215	0.0270	0.0348	0.0435	0.0523	0.0900
stuffy nose(X)	0.0195	0.0119	0.0506	0.0624	0.0849	0.1183	0.1460	0.1921	0.2289	0.2737
dry cough(X)	0.1687	0.1254	0.0693	0.0446	0.0087	0.0554	0.1168	0.1740	0.2355	0.2956
NCP(X)	<b>0.2803</b>	0.2563	0.2533	0.2459	0.2137	0.1814	0.1409	0.1001	0.0814	0.0626
close contact(X)	<b>0.7574</b>	0.7460	0.7422	0.7525	0.7441	0.7286	0.7079	0.6857	0.6629	0.6585
suspected case(X)	<b>0.4694</b>	0.4463	0.4112	0.4005	0.3815	0.3544	0.3241	0.2889	0.2624	0.2236
pneumonia(X)	<b>0.4100</b>	0.3842	0.3406	0.3334	0.2960	0.2670	0.2245	0.1800	0.1292	0.0878
cough(X)	0.3545	0.4022	0.4695	0.5374	0.5329	0.5268	0.5256	0.5437	0.5637	0.5688
dyspnea(X)	0.1702	0.1454	0.1197	0.1010	0.0727	0.0413	0.0006	0.0489	0.0792	0.1311
virus pneumonia(X)	0.5668	0.5621	0.5693	0.5604	0.6003	0.6119	0.6208	0.6166	0.6307	<b>0.6415</b>
novel coronavirus(X)	<b>0.6409</b>	0.6226	0.5955	0.5984	0.5748	0.5502	0.5184	0.4839	0.4516	0.4194

Optimal values are shown in bold in the table

**Table 16** Correlation analysis between keywords and the cumulative number of confirmed cases—Huangshi

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Center for Disease Control and Prevention(S)	0.1050	0.0698	0.0516	0.0218	0.0380	0.0991	0.1204	0.1440	0.2080	0.2512
Red Cross Society(S)	0.0065	0.0364	0.0821	0.1100	0.1501	0.2024	0.2651	0.3214	0.3983	0.4324
Spring Festival(S)	0.7827	0.8068	0.8202	0.8376	0.8369	<b>0.8392</b>	0.8288	0.8265	0.8051	0.7924
Health Commission(S)	0.0196	0.0324	0.0818	0.1407	0.1935	0.2617	0.3099	0.3767	0.4584	0.5435
vaccine(S)	0.1915	0.1630	0.1294	0.0875	0.0210	0.0290	0.0820	0.1428	0.2031	0.2546
SARS(S)	0.2036	0.2694	0.3389	0.4108	0.4692	0.5462	0.6278	0.7170	0.8034	0.8970
hand washing(S)	0.1611	0.1145	0.1215	0.0985	0.0785	0.0296	0.0062	0.0288	0.0234	0.0478
children(S)	0.3862	0.4047	0.4128	0.3945	0.4181	0.4983	0.5005	0.4799	0.4552	0.4913
mask(S)	0.0760	0.1313	0.1752	0.2438	0.3001	0.3775	0.4520	0.5470	0.6295	0.7093
coronavirus(S)	0.1506	0.2111	0.2771	0.3439	0.4187	0.4975	0.5782	0.6670	0.7628	0.8627
novel(S)	0.1032	0.1577	0.2004	0.2559	0.3174	0.3687	0.4263	0.5087	0.5671	0.6442
disinfection(S)	0.0119	0.0083	0.0213	0.0707	0.0938	0.1145	0.1607	0.2000	0.2510	0.2628
lockdown of the city(S)	0.1228	0.0713	0.0317	0.0226	0.0816	0.1358	0.1996	0.2588	0.3243	0.4051
Wuhan(S)	0.1394	0.2027	0.2652	0.3299	0.3949	0.4702	0.5535	0.6475	0.7399	0.8365
Zhong Nanshan(S)	0.1076	0.1783	0.2411	0.3021	0.3755	0.4587	0.5370	0.6162	0.6998	0.8057
epidemic(S)	<b>0.8768</b>	0.8693	0.8574	0.8479	0.8417	0.8407	0.8132	0.7871	0.7671	0.7603
flu(S)	0.1593	0.2087	0.2680	0.2935	0.3383	0.4015	0.4607	0.5370	0.6186	0.7110
diarrhea(S)	0.0300	0.0063	0.0141	0.0235	0.0765	0.1335	0.1912	0.2403	0.3088	0.3691
dry cough(S)	0.1627	0.2100	0.2596	0.3245	0.3916	0.4689	0.5292	0.5979	0.6672	0.7591
NCPS(S)	<b>0.5531</b>	0.5277	0.5011	0.4619	0.4441	0.4323	0.4144	0.4000	0.3882	0.3584
novel coronary pneumonia(S)	<b>0.9093</b>	0.9080	0.8967	0.8803	0.8744	0.8670	0.8674	0.8504	0.8444	0.8437
COVID-19(S)	<b>0.7714</b>	0.7529	0.7393	0.7245	0.7033	0.6874	0.6537	0.6289	0.6095	0.6044
novel coronavirus pneumonia(S)	<b>0.3518</b>	0.3295	0.3013	0.2620	0.2327	0.1879	0.1388	0.0919	0.0338	0.0160
asymptomatic infection(S)	<b>0.5724</b>	0.5702	0.5445	0.5231	0.4941	0.4688	0.4571	0.4349	0.3847	0.3482
aerosol transmission(S)	<b>0.3396</b>	0.3047	0.2708	0.2390	0.2396	0.2150	0.1871	0.1608	0.1385	0.1187
pneumonia(S)	0.0789	0.1365	0.1990	0.2651	0.3324	0.4023	0.4835	0.5684	0.6607	0.7489
cough(S)	0.0893	0.0558	0.0249	0.0379	0.0614	0.1119	0.0956	0.1589	0.2217	0.2648
fever(S)	0.2737	0.2749	0.2804	0.3076	0.2819	0.3567	0.4153	0.4905	0.5741	0.5789
dyspnea(S)	0.0262	0.0717	0.1125	0.1621	0.2093	0.2648	0.3212	0.3786	0.4362	0.4617
high temperature(S)	0.0293	0.0727	0.1247	0.1836	0.2322	0.2534	0.3267	0.3838	0.4431	0.4445
novel coronavirus(S)	0.0132	0.0737	0.1341	0.1928	0.2583	0.3276	0.4000	0.4859	0.5728	0.6570
SARS-CoV-2(S)	<b>0.3035</b>	0.2838	0.2455	0.2480	0.2490	0.2463	0.2397	0.2241	0.1939	0.2028
Spring Festival(X)	0.7885	0.7817	0.7860	<b>0.8009</b>	0.7955	0.7798	0.7703	0.7648	0.7765	0.7742
Red Cross Society(X)	0.2044	0.1803	0.1515	0.0971	0.0437	0.0185	0.0290	0.0919	0.1485	0.1847
government(X)	0.3155	0.3131	0.3021	0.2509	0.2212	0.2307	0.3161	<b>0.3770</b>	0.3687	0.3702

**Table 16** (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Health Commission(X)	0.0947	0.1324	0.1436	0.2079	0.2703	0.3168	0.3348	0.4052	0.4935	0.5546
medical(X)	0.1997	0.1636	0.1388	0.0944	0.0727	0.0521	0.0227	0.0398	0.1215	0.1826
vaccine(X)	0.2159	0.2053	0.1935	0.1598	0.1115	0.0578	0.0035	0.0757	0.1307	0.1828
SARS(X)	0.1453	0.1842	0.2370	0.2996	0.3518	0.4050	0.4471	0.5089	0.6002	0.6833
children(X)	0.3785	0.3583	0.3871	0.4194	0.4642	0.4968	0.5206	0.5489	0.5836	0.6137
mask(X)	0.0951	0.0681	0.0350	0.0108	0.0519	0.0781	0.1314	0.1960	0.2522	0.2926
infection(X)	0.2861	0.3080	0.3300	0.3611	0.3861	0.4075	0.4298	0.4532	0.4778	0.5037
coronavirus(X)	0.5883	<b>0.5892</b>	0.5784	0.5474	0.5263	0.5340	0.5127	0.4663	0.4398	0.4158
disinfection(X)	0.5571	0.5697	0.5829	0.5967	0.6111	0.6262	0.6420	0.6586	0.6759	0.6942
nucleic acid(X)	0.5163	<b>0.5255</b>	0.5032	0.4502	0.4114	0.3911	0.3762	0.3621	0.3238	0.3053
lockdown of the city(X)	0.5565	0.5736	0.6005	0.6125	0.6278	0.6432	0.6506	0.6633	0.6800	0.6975
Wuhan(X)	0.0349	0.0462	0.0978	0.1588	0.1982	0.2442	0.3123	0.3727	0.4481	0.5298
Zhong Nanshan(X)	0.1258	0.0668	0.0318	0.0189	0.0717	0.1192	0.1664	0.2356	0.3158	0.3904
epidemic(X)	<b>0.4479</b>	0.4437	0.4247	0.3772	0.3460	0.3442	0.3081	0.2560	0.2155	0.1778
flu(X)	<b>0.4803</b>	0.4692	0.4558	0.4392	0.4269	0.3997	0.3838	0.3497	0.3127	0.2720
diarrhea(X)	0.0052	0.0085	0.0040	0.0201	0.0516	0.0767	0.0699	0.0543	0.0944	0.0914
stuffy nose(X)	0.0810	0.0923	0.0965	0.1226	0.1563	0.1669	0.1735	0.2269	0.2909	0.3016
dry cough(X)	0.1775	0.1307	0.0917	0.0326	0.0160	0.0638	0.1118	0.1628	0.2205	0.2886
NCP(X)	<b>0.2382</b>	0.2340	0.1878	0.1373	0.1040	0.0786	0.0700	0.0729	0.0674	0.0486
close contact(X)	0.7599	<b>0.7777</b>	0.7594	0.7388	0.7126	0.6978	0.6968	0.6808	0.6562	0.6407
suspected case(X)	<b>0.5137</b>	0.4917	0.4793	0.4433	0.4115	0.4019	0.3895	0.3434	0.2917	0.2642
pneumonia(X)	<b>0.5000</b>	0.4971	0.4853	0.4444	0.4160	0.3974	0.3628	0.3186	0.2873	0.2532
cough(X)	0.2899	0.3306	0.3621	0.4054	0.3997	0.3818	0.3602	0.3637	0.3959	0.4062
dyspnea(X)	<b>0.2590</b>	0.2403	0.2172	0.1885	0.1543	0.1334	0.0921	0.0380	0.0178	0.0195
virus pneumonia(X)	0.5645	0.5506	0.5589	0.5755	0.5945	0.5891	0.5803	0.5953	0.6165	0.6211
novel coronavirus(X)	0.6364	<b>0.6372</b>	0.6233	0.5971	0.5771	0.5843	0.5797	0.5380	0.4991	0.4682

Optimal values are shown in bold in the table

**Table 17** Correlation analysis between keywords and the cumulative number of confirmed cases—Jingmen

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Center for Disease Control and Prevention(S)	0.0810	0.0358	0.0089	0.0491	0.1052	0.1376	0.1130	0.1705	0.2241	0.3044
Red Cross Society(S)	0.1650	0.1177	0.1049	0.0945	0.0542	0.0003	0.0569	0.1281	0.1959	0.2509
Spring Festival(S)	0.7382	0.7275	0.7654	<b>0.8046</b>	0.8019	0.7789	0.7696	0.7724	0.7589	0.7389
Health Commission(S)	0.1311	0.1009	0.0632	0.0205	0.0377	0.0909	0.1465	0.2017	0.2813	0.3214
vaccine(S)	<b>0.2932</b>	0.2736	0.2492	0.2325	0.2025	0.1878	0.1097	0.0307	0.0091	0.0569
SARS(S)	0.1871	0.2559	0.3288	0.4054	0.4788	0.5545	0.6396	0.7365	0.8353	0.9017
hand washing(S)	0.2357	0.2122	<b>0.2425</b>	0.1912	0.1974	0.1933	0.1593	0.1329	0.1795	0.1160
children(S)	0.3497	0.3204	0.3180	0.3451	0.3616	<b>0.3814</b>	0.3249	0.3392	0.3584	0.2760
mask(S)	0.0157	0.0576	0.1102	0.1731	0.2290	0.2733	0.3462	0.4260	0.5165	0.5985
coronavirus(S)	0.1316	0.1967	0.2619	0.3364	0.4064	0.4802	<b>0.5631</b>	0.6448	0.7343	0.8352
novel(S)	0.0643	0.1035	0.1700	0.2249	0.2467	0.2968	0.3791	0.4232	0.4920	0.5905
disinfection(S)	0.0267	0.0598	0.1195	0.1272	0.1324	0.1866	0.2268	0.2648	0.3308	0.3565
lockdown of the city(S)	0.1184	0.0638	0.0219	0.0065	0.0415	0.0686	0.1293	0.2022	0.2368	0.2851
Wuhan(S)	0.1145	0.1754	0.2310	0.2951	0.3579	0.4270	0.5020	0.5916	0.6797	0.7651
Zhong Nanshan(S)	0.0275	0.0873	0.1535	0.2215	0.2857	0.3653	0.4612	0.5674	0.6452	0.7228
epidemic(S)	<b>0.8886</b>	0.8803	0.8830	0.8848	0.8770	0.8616	0.8365	0.8133	0.7888	0.7946
flu(S)	0.1440	0.2040	0.2482	0.3166	0.3640	0.4169	<b>0.4777</b>	0.5638	0.6316	0.7294
diarrhea(S)	0.0095	0.0366	0.0880	0.1438	0.1987	0.2602	0.3242	0.3965	0.4554	0.5295
dry cough(S)	0.0475	0.1132	0.1762	0.2296	0.2913	0.3512	0.4507	0.5315	0.5945	0.6815
NCPS(S)	<b>0.5602</b>	0.5335	0.4999	0.4686	0.4349	0.3970	0.3856	0.3671	0.3679	0.3469
novel coronary pneumonia(S)	<b>0.9293</b>	0.9253	0.9214	0.9200	0.9221	0.9145	0.9080	0.8965	0.8941	0.8729
COVID-19(S)	<b>0.7525</b>	0.7477	0.7453	0.7447	0.7357	0.7446	0.7227	0.6882	0.6587	0.6383
novel coronavirus pneumonia(S)	0.3966	<b>0.4014</b>	0.3725	0.3347	0.3004	0.2650	0.2268	0.1832	0.1313	0.0750
asymptomatic infection(S)	<b>0.6621</b>	0.6553	0.6367	0.6187	0.6116	0.5851	0.5800	0.5646	0.5190	0.4554
aerosol transmission(S)	<b>0.2966</b>	0.2739	0.2485	0.2154	0.1844	0.1810	0.1613	0.1474	0.1423	0.1224
pneumonia(S)	0.0566	0.1172	0.1784	0.2355	0.3062	0.3725	0.4447	0.5226	0.6232	0.6980
cough(S)	0.0288	0.0422	0.0759	0.1724	0.2275	0.2652	0.2948	0.3336	0.3529	0.4516
fever(S)	0.0961	0.1462	0.2359	0.3018	0.2319	0.2647	0.3179	0.3446	0.3973	0.4488
dyspnea(S)	0.1235	0.1993	0.2358	0.3096	0.3510	0.4229	0.5010	0.5844	0.6345	0.6603
high temperature(S)	0.0656	0.1166	0.1429	0.2237	0.2363	0.2972	0.3021	0.3462	0.4521	0.5069
novel coronavirus(S)	0.0022	0.0642	0.1322	0.2029	0.2670	0.3345	0.4100	0.4854	0.5657	0.6460
SARS-CoV-2(S)	<b>0.4834</b>	0.4653	0.4426	0.4456	0.4557	0.4362	0.4094	0.3695	0.3318	0.3167
Spring Festival(X)	0.7888	0.7854	0.8104	0.7917	0.8026	0.7954	<b>0.8368</b>	0.7823	0.7684	0.7583
Red Cross Society(X)	0.1735	0.1511	0.0979	0.0782	0.0299	0.0194	0.1228	0.1366	0.2173	0.2755
government(X)	0.3107	0.3080	0.2909	0.2507	0.2623	0.2502	0.2584	0.3645	0.4012	0.3943

Table 17 (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Health Commission(X)	0.0136	0.0697	0.1018	0.1709	0.2393	0.2896	0.3815	0.4175	0.4961	0.5242
medical(X)	0.1145	0.0985	0.0683	0.0392	0.0002	0.0446	0.0914	0.1336	0.1892	0.2439
vaccine(X)	0.2461	0.2246	0.1783	0.1603	0.1379	0.1216	0.0192	0.0067	0.0469	0.1188
SARS(X)	0.1225	0.1809	0.2521	0.2930	0.3525	0.3821	0.4781	0.4998	0.5654	0.6637
children(X)	0.3833	0.3928	0.3889	0.3563	0.3923	0.4242	0.5024	0.5102	0.5115	0.5433
mask(X)	0.1209	0.1006	0.0709	0.0586	0.0145	0.0189	0.1158	0.1324	0.1776	0.2400
infection(X)	0.3281	0.2940	0.3182	0.3515	0.4023	0.3871	0.4480	0.4743	0.4691	0.5319
coronavirus(X)	<b>0.5842</b>	0.5546	0.5177	0.5162	0.5007	0.4896	0.4552	0.4210	0.4187	0.3904
disinfection(X)	0.4721	0.4836	0.4929	0.5017	0.5035	0.5150	0.5357	0.5474	0.5750	0.5873
nucleic acid(X)	0.4991	<b>0.5058</b>	0.4854	0.4643	0.4642	0.4239	0.3343	0.3388	0.3492	0.3106
lockdown of the city(X)	0.5523	0.5790	0.5948	0.6359	0.6184	0.6333	0.6184	0.6581	0.6678	0.6555
Wuhan(X)	0.0367	0.0840	0.1329	0.1601	0.2144	0.2720	0.3413	0.3951	0.4990	0.5969
Zhong Nanshan(X)	0.1293	0.0854	0.0288	0.0015	0.0428	0.0748	0.1912	0.2301	0.2855	0.3703
epidemic(X)	<b>0.4150</b>	0.3857	0.3547	0.3236	0.3166	0.2827	0.2331	0.1596	0.1437	0.1030
flu(X)	<b>0.4611</b>	0.4511	0.4264	0.4075	0.4054	0.4170	0.3977	0.3384	0.3104	0.2939
diarrhea(X)	0.0556	0.0828	0.0730	0.0828	<b>0.1072</b>	0.0937	0.0417	0.0749	0.0513	0.0424
stuffy nose(X)	0.1024	0.0776	0.0262	0.0070	0.0526	0.0643	0.1286	0.1443	0.1863	0.2602
dry cough(X)	0.1803	0.1468	0.0722	0.0490	0.0041	0.0233	0.1185	0.1224	0.1823	0.2316
NCP(X)	<b>0.1607</b>	0.1538	0.1322	0.1125	0.0684	0.0348	0.0059	0.0314	0.0354	0.0610
close contact(X)	<b>0.7448</b>	0.7287	0.7208	0.7292	0.7010	0.6636	0.6239	0.6039	0.6007	0.5981
suspected case(X)	<b>0.5152</b>	0.5058	0.4599	0.4061	0.4109	0.3573	0.2717	0.2664	0.2653	0.2032
pneumonia(X)	<b>0.4836</b>	0.4654	0.4026	0.4045	0.3822	0.3824	0.2911	0.2744	0.2437	0.1821
cough(X)	0.3059	0.3511	0.4283	0.4694	0.4519	0.4275	0.4570	0.4171	0.4566	0.4631
dyspnea(X)	0.2276	<b>0.2563</b>	0.2373	0.2002	0.1587	0.1559	0.0836	0.0907	0.0808	0.0746
virus pneumonia(X)	0.5483	0.5342	0.5454	0.5454	0.5847	0.6235	0.6586	0.6135	0.6333	0.6453
novel coronavirus(X)	<b>0.6553</b>	0.6241	0.5881	0.5543	0.5330	0.5099	0.4672	0.4594	0.4338	0.4632

Optimal values are shown in bold in the table

**Table 18** Correlation analysis between keywords and the cumulative number of confirmed cases—Jingzhou

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Center for Disease Control and Prevention(S)	0.0972	0.1061	0.0681	0.0318	0.0088	0.0574	0.1386	0.1825	0.2238	0.3029
Red Cross Society(S)	0.2385	0.1859	0.1325	0.1022	0.0587	0.0123	0.0263	0.0896	0.1312	0.2025
Spring Festival(S)	0.7395	0.7628	0.8038	0.8187	0.8308	0.8521	<b>0.8584</b>	0.8519	0.8377	0.8360
Health Commission(S)	0.0015	0.0553	0.1088	0.1626	0.2280	0.3054	0.3798	0.4382	0.5044	0.5967
vaccine(S)	0.2360	0.1913	0.1671	0.1381	0.0918	0.0302	0.0031	0.0692	0.1135	0.1952
SARS(S)	0.1707	0.2381	0.3100	0.3838	0.4603	0.5403	0.6282	0.7210	0.8187	0.9198
hand washing(S)	<b>0.2960</b>	0.2606	0.2289	0.1922	0.1663	0.1216	0.1063	0.0829	0.0431	0.0238
children(S)	0.4189	0.4044	0.3780	0.3959	0.4220	0.4341	<b>0.4634</b>	0.4264	0.4097	0.3769
mask(S)	0.1038	0.1553	0.2175	0.2696	0.3309	0.3864	0.4599	0.5435	0.6207	0.7087
coronavirus(S)	0.1322	0.1946	0.2619	0.3305	0.4052	0.4862	0.5694	0.6564	0.7499	0.8516
novel(S)	0.0306	0.0799	0.1339	0.1861	0.2457	0.3226	0.3924	0.4768	0.5639	0.6544
disinfection(S)	0.0327	0.0117	0.0326	0.0828	0.1153	0.1609	0.2313	0.2733	0.3038	0.3572
lockdown of the city(S)	0.1070	0.0541	0.0135	0.0384	0.0832	0.1504	0.1994	0.2543	0.3014	0.3702
Wuhan(S)	0.1249	0.1848	0.2483	0.3125	0.3890	0.4674	0.5450	0.6356	0.7268	0.8296
Zhong Nanshan(S)	0.0876	0.1445	0.2064	0.2723	0.3520	0.4253	0.5053	0.5834	0.6775	0.7670
epidemic(S)	0.8964	<b>0.9001</b>	0.8941	0.8937	0.8934	0.8758	0.8623	0.8458	0.8467	0.8325
flu(S)	0.1302	0.1704	0.1994	0.2684	0.3408	0.4031	0.4809	0.5702	0.6599	0.7511
diarrhea(S)	0.0430	0.0013	0.0400	0.0989	0.1714	0.2106	0.2700	0.3287	0.4087	0.4834
dry cough(S)	0.1946	0.2555	0.3192	0.3847	0.4333	0.5044	0.5135	0.6051	0.6948	0.7680
NCPS(S)	<b>0.7113</b>	0.6914	0.6802	0.6547	0.6319	0.6111	0.5834	0.5575	0.5302	0.4968
novel coronary pneumonia(S)	0.9258	<b>0.9330</b>	0.9280	0.9303	0.9525	0.9292	0.9277	0.9139	0.9127	0.8991
COVID-19(S)	<b>0.7992</b>	0.7946	0.7970	0.7834	0.7683	0.7577	0.7493	0.7218	0.7089	0.6887
novel coronavirus pneumonia(S)	0.2258	0.1876	0.1399	0.0928	0.0450	0.0119	0.0674	0.1235	0.1864	0.2606
asymptomatic infection(S)	<b>0.7379</b>	0.7172	0.6877	0.6628	0.6608	0.6373	0.6076	0.5893	0.5599	0.5449
aerosol transmission(S)	<b>0.5028</b>	0.4743	0.4489	0.4155	0.3787	0.3393	0.3092	0.2777	0.2351	0.1843
pneumonia(S)	0.0614	0.1206	0.1818	0.2475	0.3195	0.3965	0.4746	0.5589	0.6490	0.7449
cough(S)	0.1385	0.1038	0.0397	0.0071	0.0636	0.1137	0.1844	0.2439	0.3207	0.3734
fever(S)	0.2194	0.2036	0.2329	0.2734	0.3329	0.3930	0.4336	0.4538	0.5395	0.6131
dyspnea(S)	0.1462	0.2118	0.2682	0.2528	0.3123	0.3890	0.4484	0.5259	0.5904	0.6689
high temperature(S)	0.0004	0.0301	0.0953	0.1457	0.2080	0.2751	0.3535	0.4147	0.4620	0.5360
novel coronavirus(S)	0.0015	0.0609	0.1206	0.1832	0.2523	0.3283	0.4043	0.4868	0.5728	0.6615
SARS-CoV-2(S)	0.4581	<b>0.4716</b>	0.4476	0.4543	0.4195	0.4032	0.4066	0.3842	0.3769	0.3604
Spring Festival(X)	0.7844	0.7843	0.7896	<b>0.7950</b>	0.7828	0.7839	0.7852	0.7832	0.7790	0.7509
Red Cross Society(X)	0.1417	0.0963	0.0625	0.0121	0.0209	0.0800	0.1484	0.2326	0.3028	0.3416
government(X)	0.3433	0.3564	0.3140	0.2605	0.2408	0.2567	0.3073	0.3538	0.3768	0.4119

**Table 18** (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Health Commission(X)	0.0498	0.0990	0.1268	0.1906	0.2696	0.3365	0.3615	0.4424	0.5176	0.6061
medical(X)	0.1542	0.1247	0.0751	0.0185	0.0112	0.0613	0.1068	0.1505	0.1933	0.2436
vaccine(X)	0.2128	0.1889	0.1719	0.1292	0.1095	0.0861	0.0374	0.0260	0.0793	0.1275
SARS(X)	0.1083	0.1669	0.2182	0.2795	0.3242	0.3905	0.4388	0.5174	0.5934	0.6638
children(X)	0.3436	0.3556	0.4009	0.4241	0.4535	0.5130	0.4898	0.4672	0.4775	0.5151
mask(X)	0.0503	0.0152	0.0196	0.0602	0.1001	0.1599	0.2132	0.2725	0.3226	0.3821
infection(X)	0.3250	0.3409	0.3531	0.3786	0.4052	0.4287	0.4533	0.4791	0.5062	0.5346
coronavirus(X)	<b>0.6675</b>	0.6395	0.6190	0.5906	0.5516	0.5242	0.4871	0.4513	0.4146	0.3830
disinfection(X)	0.5670	0.5798	0.5931	0.6070	0.6216	0.6368	0.6528	0.6695	0.6871	0.7055
nucleic acid(X)	<b>0.5120</b>	0.4889	0.4719	0.4528	0.4107	0.3734	0.3293	0.2804	0.2427	0.2118
lockdown of the city(X)	0.5968	0.5402	0.5572	0.5661	0.5753	0.5881	0.6016	0.6158	0.6308	0.6467
Wuhan(X)	0.0275	0.0192	0.0774	0.1332	0.2065	0.2629	0.3240	0.3917	0.4651	0.5445
Zhong Nanshan(X)	0.1314	0.0988	0.0565	0.0064	0.0370	0.0935	0.1470	0.2152	0.2907	0.3650
epidemic(X)	<b>0.4958</b>	0.4684	0.4364	0.3966	0.3451	0.3156	0.2729	0.2160	0.1556	0.1171
flu(X)	<b>0.4355</b>	0.4239	0.4177	0.4100	0.4140	0.4077	0.3904	0.3502	0.2987	0.2720
diarrhea(X)	0.0194	0.0352	0.0478	0.0553	0.0725	0.0888	0.0859	0.0834	0.1014	0.1200
stuffy nose(X)	0.0882	0.0542	0.0226	0.0201	0.0390	0.0756	0.0987	0.1500	0.2044	0.2068
dry cough(X)	0.1696	0.1167	0.0610	0.0064	0.0235	0.0828	0.1466	0.2177	0.2799	0.3508
NCP(X)	0.2420	0.2145	0.1826	0.1740	0.1726	0.1323	0.0914	0.0547	0.0179	0.0037
close contact(X)	<b>0.7843</b>	0.7668	0.7541	0.7493	0.7622	0.7423	0.7196	0.6914	0.6853	0.6845
suspected case(X)	<b>0.5190</b>	0.4831	0.4643	0.4350	0.4234	0.3912	0.3487	0.3122	0.2854	0.2624
pneumonia(X)	<b>0.5164</b>	0.4889	0.4605	0.4241	0.4213	0.3953	0.3584	0.3023	0.2540	0.2229
cough(X)	0.3198	0.3666	0.4249	0.4901	0.4634	0.4467	0.4377	0.4575	0.4922	0.4981
dyspnea(X)	0.2021	0.1729	0.1514	0.1235	0.1004	0.0648	0.0298	0.0026	0.0019	0.0181
virus pneumonia(X)	0.5512	0.5490	0.5213	0.5438	0.5432	0.5762	0.5780	0.5895	0.6101	<b>0.6159</b>
novel coronavirus(X)	<b>0.6930</b>	0.6723	0.6544	0.6309	0.5930	0.5696	0.5423	0.5105	0.4663	0.4385

Optimal values are shown in bold in the table

**Table 19** Correlation analysis between keywords and the cumulative number of confirmed cases—Qianjiang

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Center for Disease Control and Prevention(S)	<b>0.0865</b>	0.0834	0.0791	0.0721	0.0665	0.0592	0.0512	0.0493	0.0408	0.0332
Red Cross Society(S)	0.0571	0.0089	0.0116	0.0550	0.0953	0.1427	0.1727	0.2203	0.2534	0.3135
Spring Festival(S)	0.6851	0.7290	0.7552	<b>0.7556</b>	0.7491	0.7371	0.7099	0.6997	0.6856	0.6662
Health Commission(S)	0.0200	0.0338	0.0734	0.1206	0.1490	0.2003	0.2506	0.3184	0.3835	0.4425
vaccine(S)	0.1707	0.1418	0.1083	0.0759	0.0440	0.0083	0.0471	0.1009	0.1233	0.1703
SARS(S)	0.1996	0.2650	0.3367	0.4112	0.4920	0.5657	0.6342	0.7201	0.8105	0.9011
hand washing(S)	<b>0.1386</b>	0.1345	0.1331	0.1316	0.1285	0.1299	0.1251	0.0817	0.0787	0.0773
children(S)	0.2831	0.2557	0.2543	0.2549	0.2555	0.2540	0.2530	0.2597	0.3052	0.3051
mask(S)	0.1035	0.0388	0.0051	0.0325	0.0935	0.1606	0.2107	0.2555	0.3339	0.3976
coronavirus(S)	0.0913	0.1585	0.2255	0.3017	0.3762	0.4572	0.5445	0.6344	0.7311	0.8320
novel(S)	0.0502	0.0692	0.1226	0.1632	0.2246	0.2557	0.3075	0.3769	0.4353	0.4971
disinfection(S)	0.0543	0.0532	0.0521	0.0509	0.0473	0.0506	0.0469	0.0454	0.0439	0.0423
lockdown of the city(S)	0.1248	0.1808	0.2035	0.2286	0.2539	0.2823	0.3154	0.3440	0.3794	0.4116
Wuhan(S)	0.0978	0.1711	0.2314	0.3044	0.3678	0.4415	0.5252	0.6106	0.7038	0.8074
Zhong Nanshan(S)	0.0677	0.1371	0.2028	0.2652	0.3420	0.4173	0.5065	0.5959	0.6903	0.7793
epidemic(S)	<b>0.8756</b>	0.8523	0.8361	0.8347	0.8268	0.8184	0.8197	0.8102	0.7845	0.7725
flu(S)	0.2278	0.2494	0.2934	0.3396	0.3839	0.4309	0.4707	0.4968	0.5668	0.6358
diarrhea(S)	0.1210	0.1807	0.2380	0.2981	0.3289	0.3648	0.4014	0.4603	0.5060	0.5636
dry cough(S)	0.2520	0.3006	0.3320	0.3647	0.4249	0.4602	0.5015	0.5603	0.6183	0.6658
NCPS(S)	<b>0.3920</b>	0.3593	0.3318	0.3047	0.3026	0.2705	0.2370	0.1975	0.1594	0.1216
novel coronary pneumonia(S)	0.8860	0.8911	0.8897	0.8964	0.8912	0.8873	0.8955	<b>0.8967</b>	0.8835	0.8721
COVID-19(S)	0.6170	<b>0.6299</b>	0.6208	0.6114	0.6237	0.6076	0.5979	0.5733	0.5852	0.5616
novel coronavirus pneumonia(S)	0.2969	0.2478	0.2058	0.1688	0.1177	0.0684	0.0176	0.0262	0.0939	0.1461
asymptomatic infection(S)	0.4758	<b>0.4832</b>	0.4557	0.4366	0.4051	0.4112	0.3887	0.3557	0.3340	0.3029
aerosol transmission(S)	<b>0.1829</b>	0.1501	0.1448	0.1375	0.1298	0.0890	0.0539	0.0158	0.0330	0.0491
pneumonia(S)	0.0120	0.0822	0.1569	0.2226	0.2887	0.3631	0.4367	0.5188	0.6128	0.7101
cough(S)	0.0434	0.0920	0.0913	0.1138	0.1085	0.1663	0.2178	0.2902	0.3475	0.3487
fever(S)	0.5531	0.5444	0.5461	0.5881	0.5682	0.5701	0.5530	0.5985	0.6171	0.6671
dyspnea(S)	0.1877	0.2046	0.2449	0.2668	0.2868	0.3088	0.3336	0.3593	0.3837	0.4120
high temperature(S)	0.3142	0.3377	0.3601	0.3888	0.4160	0.4429	0.4714	0.5088	0.5403	0.5740
novel coronavirus(S)	0.0824	0.0162	0.0473	0.1092	0.1792	0.2541	0.3321	0.4115	0.5004	0.5909
SARS-CoV-2(S)	0.2069	0.2111	0.2155	0.2201	0.2248	0.2298	0.2350	0.2404	0.2461	<b>0.2520</b>
Spring Festival(X)	0.7968	0.8059	0.8177	<b>0.8236</b>	0.8202	0.8218	0.8147	0.7944	0.8039	0.7963
Red Cross Society(X)	0.1527	0.0965	0.0366	0.0282	0.0820	0.1373	0.1996	0.2403	0.3150	0.3680
government(X)	0.2483	0.2515	0.2234	0.2048	0.2012	0.1856	0.2298	0.3182	0.3585	0.3560

Table 19 (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Health Commission(X)	0.0568	0.1310	0.1454	0.1886	0.2429	0.2983	0.3253	0.3751	0.4583	0.5447
medical(X)	0.1603	0.1030	0.0533	0.0011	0.0409	0.0866	0.1277	0.1595	0.2235	0.2753
vaccine(X)	0.2381	0.1906	0.1632	0.1357	0.1045	0.0766	0.0311	0.0102	0.0283	0.0796
SARS(X)	0.0823	0.1522	0.2217	0.2950	0.3701	0.4420	0.4991	0.5391	0.6275	0.7081
children(X)	0.4820	0.4601	0.4908	0.5139	0.5426	0.5739	0.6029	0.5858	0.6162	0.6533
mask(X)	0.1003	0.0409	0.0131	0.0528	0.0932	0.1360	0.1785	0.2161	0.2832	0.3475
infection(X)	0.3136	0.3308	0.3502	0.3689	0.3869	0.4072	0.4273	0.4496	0.4745	0.4991
coronavirus(X)	<b>0.6745</b>	0.6405	0.6105	0.5856	0.5546	0.5236	0.4929	0.4916	0.4595	0.4246
disinfection(X)	0.4219	0.4309	0.4403	0.4502	0.4604	0.4711	0.4823	0.4941	0.5063	0.5192
nucleic acid(X)	<b>0.5197</b>	0.4808	0.4470	0.4349	0.4050	0.3808	0.3676	0.3638	0.3152	0.2637
lockdown of the city(X)	0.4891	0.4968	0.5094	0.5197	0.5288	0.5400	0.5510	0.5650	0.5764	0.5909
Wuhan(X)	0.0180	0.0638	0.1233	0.1879	0.2711	0.3509	0.4197	0.4687	0.5501	0.6143
Zhong Nanshan(X)	0.1772	0.1141	0.0521	0.0018	0.0386	0.0966	0.1695	0.2076	0.2778	0.3437
epidemic(X)	<b>0.5792</b>	0.5358	0.4991	0.4648	0.4275	0.3887	0.3448	0.3366	0.2951	0.2451
flu(X)	<b>0.5131</b>	0.4752	0.4337	0.4086	0.3939	0.3827	0.3689	0.3624	0.3491	0.3273
diarrhea(X)	0.0097	0.0192	0.0253	0.0317	0.0383	0.0511	0.0526	0.0539	0.0681	0.0765
stuffy nose(X)	0.0610	0.0075	0.0623	0.1072	0.1078	0.1192	0.1110	0.1510	0.2063	0.1866
dry cough(X)	0.2420	0.2070	0.1620	0.1109	0.0493	0.0099	0.0681	0.1036	0.1760	0.2486
NCP(X)	<b>0.2050</b>	0.1687	0.1650	0.1589	0.1245	0.1213	0.1155	0.1094	0.0642	0.0219
close contact(X)	<b>0.7952</b>	0.7761	0.7588	0.7514	0.7284	0.7075	0.6986	0.7087	0.6866	0.6609
suspected case(X)	<b>0.5455</b>	0.5005	0.4627	0.4345	0.3899	0.3511	0.3119	0.2975	0.2582	0.2103
pneumonia(X)	<b>0.5972</b>	0.5558	0.5203	0.4894	0.4512	0.4141	0.3709	0.3653	0.3246	0.2760
cough(X)	0.2695	0.3375	0.4260	0.4826	0.4785	0.4720	0.4743	0.4575	0.4998	0.5304
dyspnea(X)	0.1710	0.1246	0.0800	0.0396	0.0005	0.0372	0.0557	0.0892	0.0767	0.0598
virus pneumonia(X)	0.5645	0.5924	0.5769	0.5692	0.5773	0.5903	0.6114	0.5919	0.6524	<b>0.6826</b>
novel coronavirus(X)	<b>0.7223</b>	0.6907	0.6625	0.6407	0.6114	0.5836	0.5546	0.5553	0.5213	0.4844

Optimal values are shown in bold in the table

**Table 20** Correlation analysis between keywords and the cumulative number of confirmed cases—Shennongjia

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Center for Disease Control and Prevention(S)	0.1026	0.1047	0.1068	0.1090	<b>0.1113</b>	0.0934	0.0913	0.0891	0.0868	0.0844
Red Cross Society(S)	0.2322	0.3017	0.2966	0.2913	0.2858	0.2801	0.2740	0.2677	0.2611	0.2542
Spring Festival(S)	0.3536	0.3402	0.3451	0.4074	0.3831	0.3849	0.4170	0.3819	0.4203	0.4209
Health Commission(S)	0	0.0038	0.0352	0.0319	0.0285	0.0250	0.0319	0.0327	0.0334	0.0388
vaccine(S)	0.0602	0.0570	0.0537	0.0503	0.0467	0.0429	0.0390	0.0349	0.0306	0.0835
SARS(S)	0.2742	0.3386	0.4105	0.4792	0.5391	0.5606	0.5823	0.6163	0.6554	0.6947
hand washing(S)	0.1026	0.1047	0.1068	0.1090	0.1113	<b>0.1137</b>	0.0913	0.0891	0.0868	0.0844
children(S)	0.1463	0.1492	0.1522	0.1554	0.1587	0.1621	0.1657	0.1694	0.1733	0.1774
mask(S)	0.1193	0.1835	0.2063	0.2239	0.2644	0.2603	0.2940	0.3721	0.3839	0.3940
coronavirus(S)	0.0061	0.0398	0.0490	0.1004	0.1512	0.2187	0.2741	0.3170	0.3872	0.4429
novel(S)	0.1463	0.1439	0.1413	0.1387	0.1360	0.1331	0.1302	0.1271	0.1919	0.1901
disinfection(S)	0.1026	0.1047	0.1068	0.1090	0.1113	0.1137	0.1161	0.1188	0.1215	0.1243
lockdown of the city(S)	0.2101	0.2066	0.2031	0.1994	0.2381	0.2800	0.3268	0.3256	0.3348	0.3492
Wuhan(S)	0.0310	0.0897	0.1360	0.1519	0.2254	0.2101	0.2401	0.3303	0.4176	0.4896
Zhong Nanshan(S)	0.0379	0.0035	0.0273	0.0854	0.1125	0.1831	0.2483	0.3137	0.4284	0.5262
epidemic(S)	0.7322	0.7497	0.7625	0.7578	0.7776	<b>0.7875</b>	0.7434	0.7014	0.7070	0.6819
flu(S)	0.1413	0.1348	0.2200	0.3020	0.3226	0.3442	0.3810	0.3863	0.3944	0.4446
diarrhea(S)	0.0000	0.0027	0.0054	0.0083	0.0113	0.0145	0.0177	0.0454	0.0433	0.0412
dry cough(S)	0.1026	0.1009	0.0992	0.0973	0.0954	0.0934	0.0913	0.0891	0.0868	0.0844
NCPS(S)	0.1463	0.1492	<b>0.1522</b>	0.0083	0.0113	0.1331	0.1302	0.1271	0.1238	0.1204
novel coronary pneumonia(S)	0.6436	0.6586	<b>0.6743</b>	0.6517	0.6297	0.6477	0.6666	0.6184	0.6387	0.6206
COVID-19(S)	0.1026	0.1047	0.1068	0.1090	0.1113	0.1137	0.1161	0.1188	0.1215	0.1243
novel coronavirus pneumonia(S)	0.2735	0.2284	0.1756	0.1066	0.0282	0.0348	0.0977	0.1338	0.1606	0.1547
asymptomatic infection(S)	0.1026	0.1009	0.0992	0.0973	0.0954	0.0934	0.0913	0.0891	0.0868	0.0844
aerosol transmission(S)	0.0620	0.0654	0.0556	0.0522	<b>0.1679</b>	0.1644	0.1608	0.1569	0.1529	0.1488
pneumonia(S)	0.2923	0.2285	0.1767	0.0912	0.0294	0.0060	0.0676	0.1590	0.2500	0.3554
cough(S)	0.1463	0.1492	0.1522	0.1554	0.1587	0.1621	<b>0.1657</b>	0.0212	0.0248	0.0285
fever(S)	0.0000	0.0027	0.0054	0.0083	<b>0.1360</b>	0.1331	0.1302	0.1271	0.1238	0.1204
dyspnea(S)	0.1026	0.1047	0.1068	0.1090	<b>0.1113</b>	0.0934	0.0913	0.0891	0.0868	0.0844
high temperature(S)	0.1026	0.1009	0.0992	0.0973	0.0954	0.0934	0.0913	0.0891	0.0868	0.0844
novel coronavirus(S)	0.1984	0.1864	0.1241	0.0698	0.0049	0.0870	0.1585	0.2494	0.3712	0.4850
SARS-CoV-2(S)	0.1026	0.1047	0.1068	0.1090	0.1113	0.1137	0.1161	0.1188	0.1215	0.1243
Spring Festival(X)	0.7494	0.7428	0.7666	0.7840	<b>0.7969</b>	0.7598	0.7808	0.7796	0.7910	0.7624
Red Cross Society(X)	0.1711	0.1287	0.0962	0.0608	0.0134	0.0175	0.0021	0.0404	0.1191	0.1924
government(X)	0.1518	0.1763	0.1476	0.0915	0.1130	0.1266	0.1808	0.2629	0.2665	0.3497

**Table 20** (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Health Commission(X)	0.0090	0.0661	0.0791	0.1241	0.1627	0.1532	0.1107	0.2173	0.3149	0.3750
medical(X)	0.1439	0.1300	0.1098	0.0602	0.0315	0.0360	0.0613	0.0268	0.0950	0.1699
<b>vaccine(X)</b>	<b>0.2743</b>	0.2419	0.2486	0.2320	0.2027	0.2085	0.2081	0.1454	0.0990	0.0507
SARS(X)	0.0049	0.0669	0.1307	0.1929	0.2576	0.3189	0.3828	0.4721	0.5747	0.6885
children(X)	0.1026	0.1009	0.0992	0.0973	0.0954	0.0934	0.0913	0.0891	0.0868	0.0844
mask(X)	0.1080	0.0687	0.0665	0.0405	0.0172	0.0257	0.0381	0.1254	0.2158	0.2672
infection(X)	0.1805	0.1776	0.1745	0.2199	0.2705	0.2692	0.2679	0.2786	0.2955	0.3051
coronavirus(X)	<b>0.7398</b>	0.6987	0.6773	0.6437	0.6093	0.6435	0.5881	0.5316	0.4806	0.4627
disinfection(X)	0.2367	0.2415	0.2465	0.2517	0.2572	0.2628	0.2687	0.2749	0.2814	0.2882
nucleic acid(X)	<b>0.4908</b>	0.4749	0.4323	0.3835	0.3317	0.3600	0.3831	0.3159	0.2489	0.2467
lockdown of the city(X)	0.4066	0.4153	0.4243	0.4337	0.4436	0.4539	0.4646	0.4759	0.4878	0.5002
Wuhan(X)	0.0727	0.0265	0.0301	0.0798	0.1226	0.1737	0.2104	0.2715	0.3734	0.4592
Zhong Nanshan(X)	0.3097	0.2751	0.2372	0.1825	0.1090	0.0388	0.0599	0.1477	0.2232	0.2730
epidemic(X)	<b>0.7167</b>	0.6761	0.6539	0.6365	0.6089	0.5617	0.5970	0.5394	0.4800	0.4506
flu(X)	<b>0.4929</b>	0.4328	0.3807	0.3705	0.3556	0.2907	0.2847	0.2219	0.1690	0.1297
diarrhea(X)	0.1026	0.1047	0.1068	<b>0.1090</b>	0.0954	0.0934	0.0913	0.0891	0.0868	0.0844
stuffy nose(X)	0.0479	0.0555	0.0121	0.0456	0.1074	0.1064	0.0963	0.1575	0.2063	0.2509
dry cough(X)	0.2304	0.1822	0.1105	0.0618	0.0027	0.0505	0.1155	0.1459	0.1948	0.2483
NCP(X)	0.2367	<b>0.2415</b>	0.1544	0.0596	0.0278	0.0228	0.0176	0.0121	0.0064	0.0004
close contact(X)	0.7662	0.7847	0.8040	<b>0.8243</b>	0.8073	0.7706	0.7934	0.7475	0.7094	0.6852
suspected case(X)	<b>0.5111</b>	0.4981	0.4831	0.4762	0.4322	0.4555	0.4020	0.3458	0.2933	0.2631
pneumonia(X)	<b>0.6758</b>	0.6332	0.6061	0.5780	0.5373	0.5720	0.5165	0.4564	0.4017	0.3881
cough(X)	0.2337	0.2972	0.3279	0.3971	0.3715	0.3591	0.3525	0.3845	0.4642	0.4588
dyspnea(X)	0.1066	0.1182	0.1303	0.1388	0.1503	0.1624	0.1734	0.1887	0.1762	0.2267
virus pneumonia(X)	0.4311	0.4250	0.3903	0.4239	0.4840	0.5374	0.5574	0.6127	0.6053	<b>0.6145</b>
novel coronavirus(X)	<b>0.7858</b>	0.7448	0.7218	0.6948	0.6622	0.6129	0.5572	0.4982	0.4512	0.4406

Optimal values are shown in bold in the table

**Table 21** Correlation analysis between keywords and the cumulative number of confirmed cases—Shiyan

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Center for Disease Control and Prevention(S)	0.0331	0.0891	0.1379	0.1930	0.2411	0.2705	0.2855	0.3349	0.4004	0.4398
Red Cross Society(S)	0.2750	0.2347	0.1968	0.1415	0.0810	0.0325	0.0234	0.0743	0.1508	0.2244
Spring Festival(S)	0.7856	0.8085	0.8338	0.8469	0.8594	0.8587	<b>0.8674</b>	0.8619	0.8646	0.8616
Health Commission(S)	0.0307	0.0232	0.0865	0.1435	0.2021	0.2776	0.3471	0.4302	0.5196	0.5935
vaccine(S)	0.1787	0.1727	0.1243	0.0879	0.0474	0.0122	0.0179	0.0705	0.1208	0.1969
SARS(S)	0.1753	0.2412	0.3124	0.3869	0.4644	0.5430	0.6294	0.7181	0.8195	0.9252
hand washing(S)	<b>0.1800</b>	0.1389	0.1069	0.1037	0.0723	0.0690	0.0229	0.0113	0.0343	0.0488
children(S)	0.3452	0.3485	0.3246	<b>0.3867</b>	0.3862	0.3791	0.3516	0.3067	0.2782	0.2467
mask(S)	0.0461	0.1127	0.1742	0.2164	0.2884	0.3547	0.4274	0.5042	0.5763	0.6609
coronavirus(S)	0.1209	0.1848	0.2483	0.3189	0.3876	0.4694	0.5560	0.6436	0.7386	0.8407
novel(S)	0.0363	0.0810	0.1271	0.1901	0.2497	0.3186	0.3920	0.4559	0.5204	0.6108
disinfection(S)	0.1104	0.1029	0.0949	0.0567	0.0001	0.0183	0.0570	0.0740	0.0969	0.1275
lockdown of the city(S)	0.0626	0.0061	0.0476	0.0892	0.1353	0.2013	0.2559	0.3043	0.3777	0.4416
Wuhan(S)	0.1263	0.1897	0.2517	0.3227	0.3976	0.4783	0.5632	0.6499	0.7429	0.8509
Zhong Nanshan(S)	0.0560	0.1190	0.1744	0.2311	0.3011	0.3892	0.4753	0.5532	0.6346	0.7203
epidemic(S)	<b>0.5496</b>	0.5098	0.4954	0.4655	0.4468	0.4183	0.3637	0.3234	0.2633	0.2182
flu(S)	0.2272	0.2479	0.3055	0.3643	0.3915	0.4677	0.5021	0.5939	0.6737	0.7661
diarrhea(S)	0.0470	0.0978	0.1492	0.1644	0.2280	0.2756	0.3328	0.3932	0.4533	0.5141
dry cough(S)	0.1856	0.2428	0.2903	0.3299	0.3972	0.4791	0.5604	0.6382	0.6938	0.7656
NCPS(S)	<b>0.6761</b>	0.6580	0.6376	0.6142	0.5874	0.5588	0.5407	0.5119	0.4854	0.4548
novel coronary pneumonia(S)	0.8885	<b>0.8941</b>	0.8901	0.8779	0.8744	0.8551	0.8410	0.8401	0.8364	0.8184
COVID-19(S)	<b>0.8130</b>	0.7973	0.7828	0.7803	0.7813	0.7608	0.7545	0.7333	0.7088	0.6973
novel coronavirus pneumonia(S)	<b>0.4067</b>	0.3911	0.3601	0.3216	0.2768	0.2394	0.2006	0.1603	0.1023	0.0631
asymptomatic infection(S)	<b>0.7259</b>	0.7039	0.6802	0.6673	0.6369	0.6022	0.5736	0.5340	0.5103	0.4772
aerosol transmission(S)	<b>0.3240</b>	0.3017	0.2725	0.2305	0.1943	0.1612	0.1167	0.0793	0.0441	0.0250
pneumonia(S)	0.0720	0.1327	0.1959	0.2604	0.3269	0.4019	0.4877	0.5753	0.6661	0.7599
cough(S)	0.0359	0.0608	0.0879	0.1496	0.2159	0.2782	0.3255	0.4100	0.4024	0.4057
fever(S)	0.3585	0.3722	0.4072	0.4510	0.4780	0.5444	0.5913	0.6164	0.5946	0.6389
dyspnea(S)	0.1255	0.1703	0.2333	0.2852	0.3300	0.3649	0.4290	0.4876	0.5551	0.6051
high temperature(S)	0.0764	0.1249	0.1857	0.2539	0.2874	0.3616	0.4221	0.4727	0.5305	0.6100
novel coronavirus(S)	0.0188	0.0397	0.0954	0.1735	0.2510	0.3215	0.3995	0.4811	0.5693	0.6627
SARS-CoV-2(S)	0.4787	<b>0.4916</b>	0.4758	0.4524	0.4261	0.4042	0.3827	0.3894	0.3652	0.3667
Spring Festival(X)	0.7795	0.7859	0.7831	0.7867	<b>0.7882</b>	0.7700	0.7754	0.7725	0.7640	0.7442
Red Cross Society(X)	0.1580	0.1008	0.0578	0.0191	0.0371	0.0973	0.1580	0.2301	0.3173	0.3526
government(X)	0.1193	0.1032	0.1121	0.1783	0.2180	<b>0.2291</b>	0.1667	0.1266	0.1092	0.0811

Table 21 (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Health Commission(X)	0.1786	0.2217	0.2318	0.2937	0.3748	0.3907	0.3887	0.4605	0.5506	0.6269
medical(X)	0.0060	0.0480	0.0755	0.1208	0.1671	0.2048	0.2290	0.2733	0.3246	0.3595
vaccine(X)	0.1112	0.0739	0.0387	0.0064	0.0472	0.0945	0.1293	0.1671	0.2271	0.2971
SARS(X)	0.1175	0.1804	0.2392	0.3053	0.3741	0.4165	0.4758	0.5302	0.6051	0.6867
children(X)	0.6002	0.5912	0.5762	0.6090	0.6435	0.6798	0.6669	0.6526	0.6585	0.6410
mask(X)	0.0102	0.0277	0.0682	0.1031	0.1459	0.1846	0.2488	0.3072	0.3744	0.4344
infection(X)	0.3403	0.3604	0.3813	0.4031	0.4259	0.4498	0.4749	0.5011	0.5287	0.5576
coronavirus(X)	<b>0.5882</b>	0.5638	0.5341	0.5030	0.4676	0.4687	0.4363	0.3936	0.3515	0.3086
disinfection(X)	0.4576	0.4673	0.4773	0.4878	0.4988	0.5721	0.5861	0.6009	0.6163	0.6326
nucleic acid(X)	<b>0.3942</b>	0.3783	0.3501	0.3293	0.3098	0.2639	0.2158	0.1606	0.0999	0.0594
lockdown of the city(X)	0.4971	0.5059	0.5152	0.5251	0.5554	0.5463	0.5578	0.5699	0.5827	0.5963
Wuhan(X)	0.1067	0.1395	0.1970	0.2683	0.3303	0.3768	0.4520	0.5057	0.5848	0.6569
Zhong Nanshan(X)	0.0239	0.0221	0.0634	0.1220	0.1898	0.2307	0.2979	0.3649	0.4449	0.5343
epidemic(X)	0.1559	0.1086	0.0692	0.0176	0.0403	0.0709	0.1268	0.1961	0.2646	0.3342
flu(X)	<b>0.3419</b>	0.3149	0.3038	0.2975	0.2790	0.2692	0.2494	0.2232	0.1842	0.1292
diarrhea(X)	0.0815	0.0974	0.1068	0.1166	0.1268	0.1374	0.1486	0.1603	0.1725	0.1853
stuffy nose(X)	0.1920	0.2169	0.2262	0.2241	0.2243	0.2242	0.2276	0.2720	0.3101	0.3226
dry cough(X)	0.1349	0.1077	0.0578	0.0003	0.0574	0.0863	0.1484	0.2145	0.2822	0.3369
NCP(X)	<b>0.1891</b>	0.1547	0.1584	0.1556	0.1509	0.1435	0.0920	0.0488	0.0081	0.0320
close contact(X)	<b>0.7903</b>	0.7823	0.7610	0.7496	0.7445	0.7297	0.7093	0.6844	0.6566	0.6426
suspected case(X)	<b>0.4277</b>	0.4017	0.3662	0.3356	0.2999	0.2842	0.2508	0.2001	0.1559	0.1066
pneumonia(X)	0.1746	0.1335	0.0831	0.0286	0.0321	0.0633	0.1210	0.1950	0.2663	0.3408
cough(X)	0.6119	0.6375	0.6639	0.6921	0.6925	0.6643	0.6619	0.6751	0.6875	0.7024
dyspnea(X)	0.0657	0.0274	0.0099	0.0545	0.0909	0.1042	0.1303	0.1583	0.1396	0.1095
virus pneumonia(X)	0.5793	0.5869	0.5720	0.5777	0.6113	0.6111	0.6516	0.6720	0.7001	<b>0.7208</b>
novel coronavirus(X)	<b>0.6381</b>	0.6109	0.5864	0.5633	0.5308	0.5324	0.5033	0.4638	0.4251	0.3856

Optimal values are shown in bold in the table

**Table 22** Correlation analysis between keywords and the cumulative number of confirmed cases—Suizhou

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Center for Disease Control and Prevention(S)	0.0234	0.0065	0.0544	0.1111	0.1690	0.2241	0.2908	0.3233	0.3937	0.4328
Red Cross Society(S)	0.1276	0.0963	0.0423	0.0064	0.0581	0.1154	0.1709	0.1988	0.2564	0.3324
Spring Festival(S)	0.5994	0.6145	0.6554	0.6691	<b>0.6719</b>	0.6636	0.6428	0.6126	0.6119	0.5988
Health Commission(S)	0.1771	0.1124	0.0456	0.0231	0.0803	0.1549	0.2120	0.2842	0.3695	0.4442
vaccine(S)	0.2365	0.2008	0.1544	0.1163	0.0801	0.0484	0.0153	0.0193	0.0623	0.0934
SARS(S)	0.1331	0.2013	0.2727	0.3487	0.4285	0.5129	0.6023	0.6961	0.7809	0.8615
hand washing(S)	0.0815	0.0908	0.1005	0.1107	0.1213	0.1324	0.1439	0.1561	0.1688	0.1821
children(S)	0.4765	0.5094	0.4824	0.4871	0.4796	0.4427	0.4822	0.4968	0.5260	0.5317
mask(S)	0.0092	0.0349	0.1039	0.1619	0.2080	0.2723	0.3446	0.4060	0.4736	0.5418
coronavirus(S)	0.0902	0.1530	0.2237	0.2999	0.3771	0.4558	0.5391	0.6248	0.7148	0.8177
novel(S)	0.0154	0.0603	0.1227	0.1739	0.2406	0.2793	0.3558	0.4133	0.4742	0.5524
disinfection(S)	0.0643	0.0149	0.006	0.0522	0.1005	0.1426	0.1661	0.1909	0.2172	0.2450
lockdown of the city(S)	0.0351	0.0152	0.0372	0.0604	0.1117	0.1661	0.1975	0.2308	0.2787	0.3173
Wuhan(S)	0.0799	0.1428	0.2194	0.2936	0.3704	0.4486	0.5228	0.6045	0.6978	0.8012
Zhong Nanshan(S)	0.0340	0.1022	0.1743	0.2474	0.3199	0.3899	0.4707	0.5600	0.6490	0.7400
epidemic(S)	0.8918	<b>0.8980</b>	0.8756	0.8681	0.8707	0.8709	0.8708	0.8736	0.8798	0.8649
flu(S)	0.0004	0.0582	0.1227	0.1983	0.2755	0.3332	0.3966	0.4788	0.5523	0.6457
diarrhea(S)	0.0849	0.0525	0.0118	0.0668	0.1217	0.1661	0.2129	0.2624	0.3461	0.4042
dry cough(S)	0.1317	0.1864	0.2443	0.3119	0.3510	0.4133	0.4393	0.4871	0.5702	0.6573
NCPS(S)	<b>0.5722</b>	0.5443	0.5164	0.4869	0.4582	0.4290	0.3954	0.3668	0.3306	0.2957
novel coronary pneumonia(S)	0.8897	0.8942	0.8904	0.8929	<b>0.8949</b>	0.8921	0.8817	0.8632	0.8526	0.8427
COVID-19(S)	<b>0.6989</b>	0.6770	0.6695	0.6449	0.6290	0.6059	0.5877	0.5753	0.5623	0.5689
novel coronavirus pneumonia(S)	<b>0.4262</b>	0.3841	0.3416	0.2982	0.2718	0.2253	0.1850	0.1435	0.0952	0.0467
asymptomatic infection(S)	<b>0.6322</b>	0.6204	0.6049	0.5764	0.5391	0.5053	0.4744	0.4331	0.3840	0.3486
aerosol transmission(S)	<b>0.2804</b>	0.2466	0.2083	0.1680	0.1312	0.1197	0.1076	0.0739	0.0251	0.0132
pneumonia(S)	0.0147	0.0745	0.1477	0.2191	0.2915	0.3697	0.4394	0.5164	0.6031	0.6967
cough(S)	0.1429	0.0848	0.0323	0.0164	0.0655	0.1307	0.1949	0.2773	0.3426	0.4185
fever(S)	0.1906	0.2208	0.2218	0.2661	0.3197	0.3829	0.4263	0.4724	0.5216	0.5532
dyspnea(S)	0.1094	0.1653	0.2203	0.2501	0.2981	0.3645	0.4031	0.4440	0.5105	0.5585
high temperature(S)	0.1954	0.2376	0.2642	0.3167	0.3647	0.4150	0.4752	0.5152	0.5575	0.6023
novel coronavirus(S)	0.0750	0.0198	0.0454	0.1194	0.1848	0.2586	0.3398	0.4209	0.5035	0.6002
SARS-CoV-2(S)	0.2926	0.2986	<b>0.3049</b>	0.2635	0.2677	0.2721	0.2304	0.2324	0.1848	0.1837
Spring Festival(X)	0.7865	0.7714	0.7835	0.7945	<b>0.7969</b>	0.7902	0.7922	0.7841	0.7673	0.7648
Red Cross Society(X)	0.2236	0.2067	0.1535	0.0928	0.0302	0.0254	0.0760	0.1380	0.1805	0.2490
government(X)	0.2938	0.3135	0.2910	0.2557	0.2565	0.2542	0.3038	0.3565	0.3897	<b>0.4053</b>

**Table 22** (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Health Commission(X)	0.0424	0.1068	0.1429	0.2116	0.2635	0.3265	0.3618	0.4493	0.5091	0.5993
medical(X)	0.2156	0.1593	0.0975	0.0376	0.0233	0.0721	0.1206	0.1704	0.2442	0.3140
vaccine(X)	0.2767	0.2264	0.1744	0.1432	0.1100	0.0733	0.0384	0.0135	0.0617	0.1008
SARS(X)	0.0543	0.1152	0.1820	0.2560	0.3318	0.4028	0.4643	0.5285	0.5878	0.6728
children(X)	0.3568	0.3600	0.3844	0.4099	0.4596	0.4889	0.5197	0.5521	0.5862	0.5641
mask(X)	0.0845	0.0673	0.0085	0.0501	0.0935	0.1343	0.1755	0.2198	0.2606	0.3270
infection(X)	0.3254	0.3445	0.3643	0.3850	0.4067	0.4293	0.4529	0.4777	0.5037	0.5310
coronavirus(X)	0.6851	<b>0.6999</b>	0.6662	0.6332	0.6067	0.5765	0.5447	0.5075	0.4625	0.4275
disinfection(X)	0.4760	0.4863	0.4970	0.5082	0.5199	0.5322	0.5450	0.5584	0.5724	0.5872
nucleic acid(X)	0.4747	<b>0.4788</b>	0.4439	0.4052	0.3879	0.3619	0.3378	0.3191	0.3082	0.2635
lockdown of the city(X)	0.5136	0.5239	0.5347	0.5460	0.5578	0.5702	0.5833	0.5970	0.6114	0.6070
Wuhan(X)	0.0314	0.0200	0.0302	0.0896	0.1467	0.2205	0.2970	0.3582	0.4081	0.4848
Zhong Nanshan(X)	0.1943	0.1313	0.0665	0.0003	0.0457	0.0896	0.1510	0.2272	0.2729	0.3443
epidemic(X)	<b>0.5147</b>	0.4737	0.4293	0.3832	0.3426	0.3000	0.2527	0.1985	0.1800	0.1275
flu(X)	0.4335	<b>0.4382</b>	0.3982	0.3557	0.3322	0.3141	0.3016	0.2872	0.2789	0.2618
diarrhea(X)	0.0170	0.0241	0.0315	0.0392	0.0472	0.0556	0.0644	0.0737	0.0833	0.0935
stuffy nose(X)	0.1693	0.1045	0.0692	0.0169	0.0017	0.0327	0.0296	0.0658	0.0925	0.1160
dry cough(X)	0.2348	0.1833	0.1334	0.0788	0.0181	0.0485	0.1188	0.1901	0.2333	0.3076
NCP(X)	<b>0.2957</b>	0.2629	0.2310	0.2299	0.2288	0.1962	0.1932	0.1594	0.1542	0.1097
close contact(X)	<b>0.8517</b>	0.8320	0.8133	0.7973	0.7918	0.7718	0.7544	0.7486	0.7339	0.7111
suspected case(X)	<b>0.5151</b>	0.4743	0.4298	0.3845	0.3537	0.3055	0.2687	0.2260	0.2043	0.1638
pneumonia(X)	<b>0.5872</b>	0.5513	0.5122	0.4713	0.4380	0.3993	0.3590	0.3125	0.3018	0.2592
cough(X)	0.2399	0.2551	0.3295	0.4071	0.4057	0.4007	0.4178	0.4423	0.4500	0.4686
dyspnea(X)	<b>0.2556</b>	0.2045	0.1579	0.1067	0.0708	0.0348	0.0004	0.0327	0.0246	0.0308
virus pneumonia(X)	0.5305	0.5528	0.5601	0.5646	0.5768	0.5915	0.5839	0.6021	0.6111	0.6604
novel coronavirus(X)	<b>0.7286</b>	0.6994	0.6665	0.6337	0.6073	0.5767	0.5448	0.5073	0.5034	0.4704

Optimal values are shown in bold in the table

**Table 23** Correlation analysis between keywords and the cumulative number of confirmed cases—Tianmen

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Center for Disease Control and Prevention(S)	<b>0.2928</b>	0.2216	0.1864	0.1525	0.1500	0.1066	0.0634	0.0522	0.0524	0.1141
Red Cross Society(S)	0.1973	0.1569	0.1089	0.0636	0.0139	0.0697	0.1003	0.1619	0.2320	0.2701
Spring Festival(S)	0.4613	0.4748	0.6133	0.6928	0.6996	0.6962	<b>0.7007</b>	0.6880	0.6929	0.6484
Health Commission(S)	0.1361	0.0699	0.0465	0.0229	0.0618	0.1125	0.1891	0.2197	0.2775	0.3638
vaccine(S)	0.1632	0.1196	0.0793	0.0782	0.0680	0.0121	0.0754	0.0403	0.1334	0.1069
SARS(S)	0.0881	0.1480	0.2128	0.2926	0.3649	0.4298	0.5038	0.6000	0.7422	0.7693
hand washing(S)	0.1451	0.1436	0.0074	0.1703	0.1899	0.1863	0.0626	0.2074	0.2291	0.2433
children(S)	0.2288	0.2790	0.2337	0.2942	0.2715	0.2538	0.3234	0.2301	0.2167	0.2084
mask(S)	0.2317	0.1382	0.1410	0.0495	0.0078	0.0583	0.0974	0.1626	0.2496	0.3294
coronavirus(S)	0.0438	0.1123	0.1656	0.2463	0.3259	0.4111	0.4721	0.5706	0.6488	0.7431
novel(S)	0.1238	0.0791	0.0611	0.0226	0.1555	0.1043	0.2058	0.1922	0.2825	0.2884
disinfection(S)	0.0474	0.0446	0.0185	0.0690	0.0069	0.1177	0.0932	0.1824	0.2029	0.2642
lockdown of the city(S)	0.0387	0.0152	0.0112	0.0700	0.1520	0.2197	0.2235	0.2713	0.2982	0.3434
Wuhan(S)	0.0104	0.0776	0.1595	0.2354	0.2978	0.3770	0.4667	0.5360	0.6387	0.7244
Zhong Nanshan(S)	0.0177	0.0603	0.1250	0.2011	0.2718	0.3546	0.4313	0.4931	0.5906	0.6543
epidemic(S)	0.8447	<b>0.8601</b>	0.8272	0.8323	0.7863	0.7798	0.7790	0.7810	0.7659	0.8212
flu(S)	0.0533	0.0370	0.0518	0.0753	0.1355	0.2419	0.2926	0.3413	0.4231	0.5413
diarrhea(S)	0.1399	0.0676	0.0039	0.0516	0.1053	0.1264	0.2307	0.3264	0.3709	0.3524
dry cough(S)	0.1126	0.2082	0.2316	0.2967	0.3825	0.4543	0.5418	0.5610	0.5970	0.6910
NCPS(S)	<b>0.4385</b>	0.4075	0.3689	0.3301	0.3354	0.4043	0.3758	0.3473	0.3447	0.3198
novel coronary pneumonia(S)	0.8600	0.8742	<b>0.8817</b>	0.8598	0.8295	0.8300	0.8062	0.8072	0.7794	0.7743
COVID-19(S)	0.7333	<b>0.7510</b>	0.7293	0.7155	0.6921	0.6678	0.6549	0.6429	0.6391	0.7079
novel coronavirus pneumonia(S)	<b>0.4534</b>	0.4199	0.3765	0.3285	0.2682	0.2578	0.2154	0.1687	0.1087	0.0790
asymptomatic infection(S)	0.5315	<b>0.5446</b>	0.4972	0.4781	0.4804	0.4414	0.4492	0.4597	0.4002	0.4009
aerosol transmission(S)	0.1551	0.1220	<b>0.1704</b>	0.1199	0.0294	0.1241	0.1189	0.1069	0.0930	0.0507
pneumonia(S)	0.0867	0.0283	0.0344	0.1107	0.1769	0.2673	0.3431	0.4196	0.5008	0.6010
cough(S)	0.0743	0.0178	0.0279	0.0840	0.1985	0.2349	0.2316	0.2640	0.3677	0.4427
fever(S)	0.1159	0.1317	0.1998	0.2708	0.3245	0.4555	0.4349	0.5531	0.6527	0.6673
dyspnea(S)	0.0275	0.0533	0.1730	0.1505	0.1727	0.2453	0.2762	0.3189	0.3236	0.3913
high temperature(S)	0.2033	0.2260	0.2428	0.3174	0.4294	0.4576	0.4067	0.4762	0.5886	0.5895
novel coronavirus(X)	0.1206	0.0518	0.0020	0.0544	0.1220	0.2106	0.2914	0.3828	0.4715	0.5727
SARS-CoV-2(S)	0.2915	0.2975	<b>0.3038</b>	0.2585	0.2109	0.1691	0.1680	0.1173	0.1030	0.0676
Spring Festival(X)	0.7088	<b>0.7924</b>	0.7021	0.6977	0.7308	0.7073	0.7195	0.7920	0.7628	0.7014
Red Cross Society(X)	0.2849	0.2339	0.1730	0.1090	0.0303	0.0522	0.1014	0.1642	0.2189	0.2624
government(X)	0.3842	0.2799	0.3117	0.2734	0.1926	0.2583	0.3184	0.3122	0.3706	0.4060

**Table 23** (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Health Commission(X)	0.0530	0.0162	0.0893	0.1426	0.1915	0.3061	0.3448	0.4417	0.4619	0.5320
medical(X)	0.2807	0.1074	0.1598	0.0878	0.0211	0.0817	0.1133	0.1808	0.1850	0.2751
vaccine(X)	0.2265	0.2371	0.1968	0.1277	0.0610	0.0024	0.0368	0.0896	0.1878	0.1448
SARS(X)	0.0235	0.1532	0.1161	0.1685	0.2639	0.3531	0.4174	0.5000	0.6445	0.6245
children(X)	0.2838	0.4058	0.3821	0.4054	0.4297	0.4552	0.5099	0.5405	0.5696	0.6032
mask(X)	0.1945	0.0245	0.0947	0.0538	0.0463	0.1414	0.1663	0.1723	0.2071	0.3494
infection(X)	0.3221	0.2493	0.3540	0.2963	0.3988	0.4260	0.4496	0.4743	0.5002	0.5273
coronavirus(X)	<b>0.7003</b>	0.6295	0.6707	0.6790	0.6237	0.6104	0.5846	0.5427	0.4008	0.4671
disinfection(X)	0.4166	0.4255	0.4348	0.4445	0.4446	0.4652	0.4762	0.4878	0.4999	0.5126
nucleic acid(X)	<b>0.5800</b>	0.4368	0.5041	0.5031	0.3893	0.4164	0.4186	0.4058	0.3809	0.2232
lockdown of the city(X)	0.4402	0.4549	0.5296	0.5355	0.4892	0.4967	0.5076	0.5192	0.5313	0.5441
Wuhan(X)	0.0965	0.0488	0.0052	0.0780	0.1742	0.2173	0.2840	0.4259	0.4910	0.5335
Zhong Nanshan(X)	0.2355	0.0615	0.1396	0.0886	0.0039	0.0543	0.0828	0.1290	0.2001	0.3511
epidemic(X)	<b>0.6251</b>	0.4791	0.5271	0.5139	0.4581	0.4089	0.3731	0.3326	0.2981	0.2848
flu(X)	0.4264	<b>0.4350</b>	0.3762	0.3751	0.3803	0.3566	0.3060	0.2430	0.1742	0.2651
diarrhea(X)	0.0968	0.0864	0.0992	0.1100	0.1276	0.1373	0.1293	0.0124	0.1453	0.1658
stuffy nose(X)	0.2649	0.1739	0.2276	0.1834	0.1012	0.0701	0.0392	0.0193	0.1635	0.0912
dry cough(X)	0.2411	0.1227	0.2115	0.1757	0.1181	0.0530	0.0001	0.0852	0.2637	0.3057
NCP(X)	0.2160	0.2167	0.2134	0.2069	0.2121	<b>0.3185</b>	0.3076	0.2651	0.1607	0.1523
close contact(X)	<b>0.7577</b>	0.7384	0.7552	0.7333	0.7061	0.6937	0.6765	0.7193	0.7312	0.6640
suspected case(X)	<b>0.5847</b>	0.4578	0.5337	0.5146	0.4063	0.3794	0.3237	0.3415	0.2966	0.1903
pneumonia(X)	0.5664	0.4719	<b>0.5779</b>	0.5631	0.5155	0.4857	0.4611	0.4144	0.2578	0.2520
cough(X)	0.0838	0.1426	0.2345	0.3086	0.3522	0.3432	0.3430	0.3900	0.4555	0.4511
dyspnea(X)	0.2718	0.2431	0.2257	0.2076	0.1426	0.0907	0.0627	0.0100	0.0606	0.0663
virus pneumonia(X)	0.4841	0.6011	0.4848	0.5193	0.5658	0.5742	0.5678	0.5310	0.5845	<b>0.6720</b>
novel coronavirus(X)	<b>0.6901</b>	0.6283	0.6523	0.6663	0.6233	0.5916	0.5628	0.5338	0.5097	0.5041

Optimal values are shown in bold in the table

**Table 24** Correlation analysis between keywords and the cumulative number of confirmed cases—Xiangyang

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Center for Disease Control and Prevention(S)	0.1518	0.1157	0.0678	0.0262	0.0403	0.1109	0.1951	0.2652	0.3318	0.3949
Red Cross Society(S)	0.2356	0.1783	0.1275	0.0781	0.0192	0.0472	0.0836	0.1588	0.2199	0.3001
Spring Festival(S)	0.7922	0.8042	0.8334	0.8526	0.8556	0.8592	<b>0.8683</b>	0.8492	0.8291	0.8109
Health Commission(S)	0.0588	0.0087	0.0702	0.1450	0.2297	0.3080	0.3947	0.4726	0.5529	0.6446
vaccine(S)	0.3382	0.2899	0.2443	0.1877	0.1303	0.0967	0.0599	0.0063	0.0523	0.1095
SARS(S)	0.1218	0.1918	0.2626	0.3402	0.4223	0.5079	0.5984	0.6970	0.7969	0.8988
hand washing(S)	<b>0.4566</b>	0.4147	0.3725	0.3515	0.3473	0.3244	0.2879	0.2437	0.2642	0.2378
children(S)	<b>0.4075</b>	0.3623	0.3111	0.2652	0.3472	0.3183	0.3616	0.3479	0.3188	0.2981
mask(S)	0.0042	0.0701	0.1382	0.2094	0.2857	0.3636	0.4427	0.5332	0.6155	0.7076
coronavirus(S)	0.0752	0.1423	0.2127	0.2850	0.3655	0.4499	0.5384	0.6315	0.7284	0.8339
novel(S)	0.0628	0.0019	0.0525	0.1027	0.1777	0.2545	0.3341	0.3974	0.4746	0.5698
disinfection(S)	0.1339	0.0946	0.0505	0.0131	0.0806	0.1365	0.2076	0.2715	0.3354	0.4147
lockdown of the city(S)	0.1600	0.1188	0.0608	0.0019	0.0693	0.1360	0.2102	0.2737	0.3340	0.4057
Wuhan(S)	0.0572	0.1229	0.1932	0.2649	0.3515	0.4391	0.5261	0.6178	0.7124	0.8078
Zhong Nanshan(S)	0.0213	0.0822	0.1440	0.2230	0.2999	0.3815	0.4700	0.5601	0.6475	0.7594
epidemic(S)	<b>0.8915</b>	0.8829	0.8798	0.8753	0.8479	0.8215	0.8179	0.8130	0.8098	0.8175
flu(S)	0.0793	0.1442	0.2122	0.2592	0.3238	0.4057	0.4929	0.5726	0.6578	0.7545
diarrhea(S)	0.0994	0.0334	0.0207	0.0899	0.1626	0.1657	0.2361	0.3052	0.3766	0.4415
dry cough(S)	0.0065	0.0654	0.1386	0.2045	0.2762	0.3631	0.4404	0.5195	0.6144	0.7185
NCPS(S)	<b>0.8058</b>	0.7822	0.7565	0.7297	0.7019	0.6729	0.6425	0.6145	0.5817	0.5450
novel coronary pneumonia(S)	0.9053	0.9100	0.9136	<b>0.9164</b>	0.9000	0.8942	0.8967	0.8963	0.8771	0.8728
COVID-19(S)	0.8169	0.8365	<b>0.8571</b>	0.8368	0.8182	0.7956	0.7720	0.7519	0.7352	0.7234
novel coronavirus pneumonia(S)	0.0793	0.0149	0.0517	0.1220	0.1987	0.2756	0.3394	0.4199	0.5104	0.6012
asymptomatic infection(S)	<b>0.7008</b>	0.66674	0.6476	0.6357	0.6090	0.5906	0.5719	0.5263	0.4840	0.4391
aerosol transmission(S)	<b>0.6684</b>	0.6342	0.5979	0.5602	0.5264	0.5028	0.4625	0.4249	0.3846	0.3400
pneumonia(S)	0.0000	0.0625	0.1299	0.1985	0.2835	0.3704	0.4551	0.5427	0.6356	0.7303
cough(S)	0.0207	0.0485	0.1225	0.2038	0.2793	0.3414	0.4103	0.4931	0.5635	0.6236
fever(S)	0.2455	0.2433	0.2848	0.2895	0.3367	0.3551	0.4317	0.4845	0.5593	0.6466
dyspnea(S)	0.0057	0.0521	0.1236	0.1941	0.2664	0.3431	0.4142	0.4913	0.5557	0.6330
high temperature(S)	0.0344	0.0207	0.0864	0.1621	0.2421	0.3134	0.3844	0.4747	0.5541	0.6359
novel coronavirus(S)	0.0508	0.0136	0.0780	0.1437	0.2228	0.3026	0.3859	0.4742	0.5658	0.6621
SARS-CoV-2(S)	0.4929	0.5035	0.5146	<b>0.5262</b>	0.5048	0.4722	0.4544	0.4325	0.4030	0.3811
Spring Festival(X)	0.7912	<b>0.8003</b>	0.7877	0.7704	0.7771	0.7781	0.7747	0.7656	0.7661	0.7524
Red Cross Society(X)	0.2085	0.1568	0.1037	0.0818	0.0197	0.0517	0.1217	0.1845	0.2499	0.2890
government(X)	0.1556	0.1799	0.1497	0.1370	0.1323	0.1177	0.2041	0.2927	0.3081	0.2932

**Table 24** (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Health Commission(X)	0.0325	0.0775	0.1110	0.1799	0.2656	0.3344	0.3422	0.4095	0.4944	0.5908
medical(X)	0.2129	0.1744	0.1050	0.0344	0.0408	0.1162	0.1808	0.2356	0.3021	0.3685
vaccine(X)	0.1675	0.1307	0.1181	0.1049	0.0501	0.0147	0.0184	0.0603	0.0987	0.1543
SARS(X)	0.0584	0.1181	0.1839	0.2566	0.3365	0.4167	0.4844	0.5511	0.6368	0.7250
children(X)	0.4808	0.4642	0.4410	0.4727	0.4899	0.5256	0.4981	0.4709	0.4629	0.4984
mask(X)	0.1026	0.0549	0.0049	0.0286	0.0937	0.1617	0.2066	0.2529	0.3015	0.3528
infection(X)	0.3829	0.4054	0.4289	0.4534	0.4790	0.5058	0.5338	0.5632	0.5940	0.6263
coronavirus(X)	<b>0.7232</b>	0.6935	0.6570	0.6196	0.5773	0.5356	0.5007	0.4610	0.4187	0.3704
disinfection(X)	0.4472	0.4565	0.4663	0.4764	0.4871	0.5508	0.5640	0.5779	0.5924	0.6077
nucleic acid(X)	<b>0.5088</b>	0.4652	0.4713	0.4351	0.3897	0.3467	0.3300	0.3025	0.2746	0.2567
lockdown of the city(X)	0.4611	0.4677	0.4765	0.4838	0.4922	0.5011	0.5105	0.5204	0.5309	0.5421
Wuhan(X)	0.0137	0.0172	0.0660	0.0957	0.1618	0.2333	0.3030	0.3767	0.4635	0.5358
Zhong Nanshan(X)	0.1415	0.0822	0.0133	0.0100	0.0821	0.1608	0.2138	0.2650	0.3363	0.4247
epidemic(X)	<b>0.4563</b>	0.4146	0.3660	0.3636	0.3107	0.2514	0.2025	0.1500	0.0904	0.0247
flu(X)	<b>0.4386</b>	0.4222	0.4261	0.3838	0.3346	0.2894	0.2581	0.2363	0.2195	0.1988
diarrhea(X)	<b>0.2397</b>	0.1957	0.1496	0.1483	0.1468	0.1032	0.0966	0.0936	0.0864	0.0801
stuffy nose(X)	0.1962	0.1517	0.1107	0.0801	0.0236	0.0347	0.0691	0.1347	0.1883	0.2335
dry cough(X)	0.2573	0.2052	0.1502	0.1352	0.0799	0.0188	0.0461	0.1190	0.1957	0.2709
NCP(X)	<b>0.3854</b>	0.3502	0.3135	0.2792	0.2458	0.2130	0.2091	0.1748	0.1394	0.1325
close contact(X)	0.8166	0.8095	<b>0.8278</b>	0.8067	0.7820	0.7656	0.7587	0.7356	0.7104	0.6935
suspected case(X)	<b>0.5657</b>	0.5312	0.4916	0.4455	0.3937	0.3409	0.3029	0.2474	0.2006	0.1509
pneumonia(X)	<b>0.5407</b>	0.5038	0.5126	0.4707	0.4198	0.3725	0.3338	0.2873	0.2373	0.1819
cough(X)	0.2886	0.3416	0.4142	0.4419	0.4504	0.4598	0.4508	0.4600	0.4977	0.5228
dyspnea(X)	<b>0.2395</b>	0.2049	0.1558	0.0987	0.0463	0.0111	0.0481	0.0879	0.0636	0.0441
virus pneumonia(X)	0.5558	0.5838	0.5490	0.5317	0.5673	0.5799	0.5762	0.5742	0.6000	0.6326
novel coronavirus(X)	<b>0.6971</b>	0.6670	0.6771	0.6891	0.6527	0.6152	0.5889	0.5557	0.5203	0.4802

Optimal values are shown in bold in the table

**Table 25** Correlation analysis between keywords and the cumulative number of confirmed cases—Xiamen

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Center for Disease Control and Prevention(S)	0.1306	0.0368	0.0047	0.0347	0.0408	0.0611	0.1314	0.2077	0.2949	0.3177 <b>0.4007</b>
Red Cross Society(S)	0.4881	0.4365	0.3791	0.3101	0.2717	0.2140	0.1405	0.0358	0.0587	0.1095 0.1936
Spring Festival(S)	0.5408	0.5754	0.6284	0.6840	0.7160	0.7341	0.7396	0.7162	0.7071	0.6904 0.6892 <b>0.7557</b>
Health Commission(S)	0.2971	0.1916	0.1133	0.0273	0.0038	0.0426	0.1264	0.2297	0.2779	0.3497 0.3818
vaccine(S)	0.3997	0.3951	0.4460	<b>0.4538</b>	0.3718	0.3359	0.1832	0.1068	0.0631	0.0800 0.1205
SARS(S)	0.0942	0.0176	0.0598	0.1484	0.2436	0.3216	0.4067	0.5013	0.5937	0.7091 0.7696
hand washing(S)	<b>0.3271</b>	0.2872	0.2866	0.1675	0.1653	0.1114	0.1068	0.1019	0.1172	0.0542 0.0338
children(S)	0.4579	0.3781	0.3332	0.3376	0.3820	<b>0.4592</b>	0.3823	0.2932	0.2975	0.2688 0.3054
mask(S)	0.2326	0.1771	0.0849	0.0504	0.0715	0.1560	0.2334	0.3347	0.4102	0.5113 0.5285
coronavirus(S)	0.1649	0.0762	0.0041	0.0818	0.1513	0.2194	0.3114	0.3934	0.5065	0.6270 0.7207
novel(S)	0.2660	0.2453	0.2289	0.1710	0.1195	0.0502	0.0794	0.1176	0.1396	0.1950 0.2303
disinfection(S)	0.1238	0.0651	0.0132	0.0187	0.0710	0.0497	0.0667	0.1494	0.2421	0.2636 0.3164
lockdown of the city(S)	0.2303	0.1874	0.1401	0.1384	0.1087	0.0800	0.0248	0.0795	0.2505	0.3789 0.4227
Wuhan(S)	0.1657	0.0965	0.0250	0.0424	0.1244	0.2264	0.3373	0.4464	0.5483	0.6398 0.6982
Zhong Nanshan(S)	0.1797	0.0947	0.0010	0.0960	0.1502	0.2331	0.3123	0.4151	0.5213	0.5921 0.6894
epidemic(S)	0.7211	0.6953	0.6831	0.6973	0.7503	0.7743 <b>0.7890</b>	0.7395	0.6902	0.6637	0.7175 0.7841
flu(S)	0.1930	0.1373	0.0474	0.0194	0.0824	0.1480	0.2215	0.3311	0.4525	0.5472 0.6490
diarrhea(S)	0.2217	0.1804	0.1547	0.1190	0.0307	0.0195	0.0989	0.1507	0.2167	0.2900 0.3375
dry cough(S)	0.0496	0.0001	0.0240	0.0558	0.1266	0.2268	0.2924	0.4005	0.4596	0.5097 0.5929
NCPS(S)	0.5631	0.5747	0.5353	0.4982	0.4256	0.4993	0.5684	0.6392 <b>0.7070</b>	0.6650	0.6348 0.5819
novel coronary pneumonia(S)	0.7155	0.6980	0.6993	0.6887	0.6995	0.7161	0.7506	0.7328	0.7462	0.7669 0.7742 <b>0.7944</b>
COVID-19(S)	0.5149	0.5269	0.5395	0.5526	0.5664	0.5809	0.5453	0.5411	0.5101	0.5924 0.6211
novel coronavirus pneumonia(S)	0.4487	0.3869	0.3349	0.2836	0.2183	0.1640	0.0923	0.0061	0.0947	0.1899 0.2889
asymptomatic infection(S)	0.59238	0.5746	<b>0.5890</b>	0.4824	0.4802	0.4526	0.4604	0.5174	0.4642	0.3666 0.2987
aerosol transmission(S)	0.35887	0.3991	0.4292	<b>0.4756</b>	0.4299	0.4256	0.4154	0.3877	0.3834	0.3259 0.2598
pneumonia(S)	0.2345	0.1506	0.0783	0.0051	0.0672	0.1429	0.2332	0.3385	0.4549	0.5617 0.6392
cough(S)	0.1451	0.0844	0.0540	0.0598	0.1175	0.2169	0.2777	0.3764	0.4055	0.4126 0.4652
fever(S)	0.1246	0.0443	0.0212	0.0666	0.1369	0.1717	0.2834	0.2319	0.2514	0.3765 0.4652 <b>0.5228</b>
dyspnea(S)	0.0370	0.0712	0.1087	0.1451	0.1903	0.1700	0.2337	0.3791	0.4462	0.5833 0.6387
high temperature(S)	0.0659	0.0824	0.0119	0.0121	0.0993	0.2111	0.2253	0.2979	0.3135	0.3365 0.3302
novel coronavirus(S)	0.2912	0.2300	0.1663	0.1100	0.0642	0.0170	0.0933	0.2052	0.3399	0.4120 0.4766
SARS-CoV-2(S)	0.1843	0.1880	0.1918	0.1958	0.1999	0.2042	0.2087	0.2134	0.1586	0.1641 0.1755
Spring Festival(X)	0.5344	0.6303	0.6696	0.6795	<b>0.7244</b>	0.6648	0.6330	0.6581	0.6492	0.7095 0.7036
Red Cross Society(X)	<b>0.5548</b>	0.4053	0.3594	0.3184	0.2533	0.2956	0.2555	0.1559	0.0509	0.1112 0.1411
government(X)	0.2896	0.1951	0.1725	0.1482	0.0492	0.1478	0.1995	0.2311	0.2212	0.2466 0.2344

**Table 25** (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Health Commission(X)	0.2383	0.0493	0.0108	0.0224	0.0613	0.0805	0.1250	0.2747	0.3639	0.3980
medical(X)	0.4103	0.2620	0.2471	0.1944	0.1002	0.1547	0.0523	0.1039	0.2459	0.2738
vaccine(X)	0.3158	0.1677	0.1963	0.2095	0.1467	0.1534	0.0143	0.1127	0.1904	0.1880
SARS(X)	0.1427	0.0287	0.0670	0.0850	0.1557	0.1400	0.2332	0.3525	0.4711	0.5593
children(X)	0.2428	0.3023	0.3840	0.4625	0.5935	0.6253	0.6084	0.5917	0.6579	0.6918
mask(X)	0.3208	0.1648	0.1143	0.0700	0.0352	0.0231	0.0974	0.2108	0.3631	0.4432
infection(X)	0.0142	0.0426	0.0626	0.1684	0.2786	0.3884	0.4100	0.4800	0.5033	0.5235
coronavirus(X)	0.5655	0.4988	0.5202	0.5886	0.6067	<b>0.7312</b>	0.7309	0.6734	0.5173	0.5088
disinfection(X)	0.2867	0.2927	0.2989	0.3054	0.3121	0.3192	0.3265	0.3342	0.3423	0.3507
nucleic acid(X)	<b>0.5996</b>	0.5274	0.5451	0.5150	0.3974	0.4374	0.4380	0.3967	0.4235	0.4125
lockdown of the city(X)	0.3143	0.3252	0.3992	0.4384	0.4443	0.4528	<b>0.4547</b>	0.4040	0.4103	0.4170
Wuhan(X)	0.2322	0.1044	0.0467	0.0452	0.1408	0.0780	0.1234	0.2340	0.3789	0.5385
Zhong Nanshan(X)	0.3753	0.2128	0.1652	0.1154	0.0138	0.0858	0.0576	0.0363	0.2108	0.3517
epidemic(X)	<b>0.6177</b>	0.5192	0.5382	0.5338	0.4925	0.5424	0.5126	0.4431	0.4026	0.3756
flu(X)	0.4444	0.3139	0.2567	0.2807	0.3588	0.4898	<b>0.4986</b>	0.4092	0.2286	0.1812
diarrhea(X)	0.2681	0.2694	<b>0.3842</b>	0.2474	0.0454	0.0224	0.1346	0.0121	0.1000	0.0855
stuffy nose(X)	0.2962	0.1611	0.0991	0.0673	0.0013	0.0774	0.1426	0.0969	0.0842	0.0807
dry cough(X)	<b>0.4662</b>	0.3283	0.3053	0.2759	0.2112	0.2586	0.1994	0.1241	0.0500	0.0307
NCP(X)	0.2455	0.2505	0.2022	0.1512	0.1066	0.2236	0.2979	<b>0.4068</b>	0.3658	0.3523
close contact(X)	0.5275	0.5398	0.5528	0.5663	0.5444	0.5593	0.5780	0.6154	0.5861	0.5777
suspected case(X)	<b>0.6602</b>	0.5668	0.5968	0.5900	0.4970	0.5018	0.4079	0.3418	0.3440	0.3585
pneumonia(X)	0.5309	0.4647	0.4966	0.5519	0.5188	0.5763	<b>0.6085</b>	0.5478	0.5015	0.3581
cough(X)	0.1222	0.2561	0.2910	0.3331	0.3502	0.1928	0.1424	0.2064	0.3576	0.4677
dyspnea(X)	<b>0.4577</b>	0.2856	0.2484	0.2027	0.1410	0.1226	0.0812	0.0335	0.0739	0.1068
virus pneumonia(X)	0.3238	0.2885	0.2755	0.3359	0.4292	0.5190	0.5639	0.5479	0.5394	0.5194
novel coronavirus(X)	0.6502	0.5922	0.6028	0.6645	0.7038	0.7555	<b>0.7620</b>	0.7090	0.6852	0.6783

Optimal values are shown in bold in the table

**Table 26** Correlation analysis between keywords and the cumulative number of confirmed cases—Xiantao

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Center for Disease Control and Prevention(S)	0.0048	0.0417	0.0577	0.0727	0.0909	0.1485	0.1553	0.2056	0.2642	0.2937
Red Cross Society(S)	0.2052	0.1609	0.1227	0.0704	0.0227	0.0007	0.0458	0.1070	0.1532	0.2110
Spring Festival(S)	0.6092	0.6173	0.6296	0.6645	0.6680	<b>0.6719</b>	0.6576	0.6479	0.6198	0.6106
Health Commission(S)	0.1098	0.0621	0.0075	0.0805	0.1616	0.2221	0.2595	0.3274	0.3738	0.4372
vaccine(S)	0.2401	0.2282	0.2068	0.1502	0.1069	0.0561	0.0055	0.0527	0.1113	0.1774
SARS(S)	0.1203	0.1922	0.2633	0.3398	0.4226	0.5075	0.5992	0.6949	0.7761	0.8457
hand washing(S)	0.1660	0.1744	0.1832	0.1923	0.2018	0.2117	0.2220	0.2328	0.2440	0.2558
children(S)	0.4079	<b>0.4491</b>	0.4183	0.4207	0.4050	0.3707	0.3903	0.3555	0.3331	0.3312
mask(S)	0.0696	0.0294	0.0192	0.0882	0.1415	0.2079	0.2845	0.3440	0.4045	0.4745
coronavirus(S)	0.0601	0.1318	0.2027	0.2748	0.3580	0.4397	0.5260	0.6195	0.7170	0.8124
novel(S)	0.0405	0.0204	0.0202	0.0584	0.1200	0.1777	0.2467	0.2848	0.3231	0.3671
disinfection(S)	0.0773	0.0701	0.0653	0.0257	0.0269	0.0381	0.0487	0.0632	0.0768	0.0912
lockdown of the city(S)	0.0891	0.0516	0.0337	0.0093	0.0535	0.1125	0.1549	0.1833	0.2537	0.2920
Wuhan(S)	0.0252	0.0868	0.1588	0.2307	0.3154	0.4006	0.4814	0.5551	0.6398	0.7282
Zhong Nanshan(S)	0.0501	0.1085	0.1735	0.2547	0.3231	0.4038	0.4929	0.5826	0.6741	0.7787
epidemic(S)	<b>0.8340</b>	0.8324	0.8247	0.8121	0.7809	0.7553	0.7572	0.7436	0.7398	0.7273
flu(S)	0.0171	0.0929	0.1702	0.2468	0.3198	0.3984	0.4729	0.5549	0.6016	0.6773
diarrhea(S)	0.0097	0.0416	0.0994	0.1537	0.2109	0.2910	0.3454	0.4078	0.5003	0.5875
dry cough(S)	0.1254	0.1771	0.2254	0.2939	0.3662	0.4402	0.5068	0.5687	0.6201	0.6886
NCPS(S)	<b>0.5207</b>	0.4986	0.4664	0.4327	0.4003	0.3643	0.3399	0.3062	0.2781	0.2502
novel coronary pneumonia(S)	0.9045	0.9102	0.9158	0.9174	0.9139	0.9123	0.9122	0.9171	0.9200	<b>0.9264</b>
COVID-19(S)	0.6834	0.6992	<b>0.7156</b>	0.6919	0.6687	0.6424	0.6272	0.6052	0.5873	0.5777
novel coronavirus pneumonia(S)	0.2980	0.2439	0.1868	0.1308	0.0670	0.0001	0.0581	0.1273	0.1859	0.2483
asymptomatic infection(S)	<b>0.5446</b>	0.5167	0.4866	0.4652	0.4432	0.4018	0.3700	0.3252	0.2776	0.2388
aerosol transmission(S)	<b>0.2759</b>	0.2359	0.1978	0.1592	0.1135	0.0768	0.0659	0.0544	0.0408	0.0028
pneumonia(S)	0.0214	0.0427	0.1063	0.1687	0.2507	0.3365	0.4154	0.5020	0.5972	0.6890
cough(S)	0.0251	0.0490	0.1004	0.1635	0.2265	0.3023	0.3314	0.4088	0.4879	0.5389
fever(S)	0.0903	0.1552	0.1980	0.2276	0.2877	0.3251	0.3824	0.4260	0.4821	0.5610
dyspnea(S)	0.1876	0.2137	0.2603	0.3356	0.4073	0.4201	0.4790	0.5421	0.5206	0.5858
high temperature(S)	0.3397	0.3898	0.4200	0.4729	0.5303	0.5995	0.6408	0.6357	0.6802	0.7262
novel coronavirus(S)	0.1362	0.0712	0.0147	0.0517	0.1332	0.2146	0.2925	0.3719	0.4674	0.5585
SARS-CoV-2(S)	0.2786	0.2843	0.2902	0.2963	<b>0.3027</b>	0.2544	0.2072	0.2104	0.2132	0.1653
Spring Festival(X)	0.7891	0.8005	0.8254	0.8195	0.8302	0.8342	0.8340	0.8280	<b>0.8346</b>	0.8275
Red Cross Society(X)	0.2069	0.1563	0.1017	0.0418	0.0228	0.0963	0.1741	0.2420	0.3096	0.3596
government(X)	0.3477	0.3641	0.3312	0.2924	0.2768	0.2438	0.3327	<b>0.4250</b>	0.4211	0.3992

**Table 26** (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Health Commission(X)	0.0002	0.0447	0.0801	0.1474	0.2309	0.3003	0.3155	0.3814	0.4665	0.5642
medical(X)	0.1814	0.1325	0.1183	0.0509	0.0222	0.0862	0.1467	0.1957	0.2512	0.3092
vaccine(X)	0.1864	0.1487	0.1132	0.0969	0.0404	0.0019	0.0321	0.0759	0.1156	0.1723
SARS(X)	0.0707	0.1289	0.1594	0.2292	0.3084	0.3863	0.4493	0.5124	0.5968	0.6812
children(X)	0.4480	0.4737	0.4519	0.4781	0.5055	0.5342	0.5949	0.6293	0.6117	0.6474
mask(X)	0.1371	0.0884	0.0707	0.0031	0.0652	0.1278	0.1783	0.2235	0.2710	0.3201
infection(X)	0.3200	0.3374	0.3555	0.3744	0.3941	0.4147	0.4362	0.4587	0.4822	0.5069
coronavirus(X)	<b>0.7256</b>	0.6993	0.6647	0.6291	0.5890	0.5495	0.5161	0.4778	0.4408	0.3965
disinfection(X)	0.4535	0.4632	0.4733	0.4838	0.4948	0.5063	0.5183	0.5309	0.5441	0.5579
nucleic acid(X)	<b>0.5297</b>	0.4848	0.4358	0.4371	0.3956	0.3481	0.3325	0.2996	0.2666	0.2443
lockdown of the city(X)	0.4720	0.4806	0.4897	0.4992	0.5091	0.5196	0.5305	0.5421	0.5543	0.5671
Wuhan(X)	0.0001	0.0358	0.0609	0.1147	0.1862	0.2588	0.3300	0.4214	0.5186	0.5997
Zhong Nanshan(X)	0.2193	0.1625	0.1467	0.0795	0.0063	0.0655	0.1181	0.1662	0.2342	0.3201
epidemic(X)	<b>0.5344</b>	0.4975	0.4537	0.4568	0.4096	0.3565	0.3134	0.2688	0.2188	0.1614
flu(X)	<b>0.4134</b>	0.3944	0.3976	0.3997	0.3525	0.3066	0.2782	0.2586	0.2452	0.2319
diarrhea(X)	0.0330	0.1009	0.1123	0.1187	0.1309	0.1418	0.1531	0.1650	0.1774	0.1904
stuffy nose(X)	0.0973	0.0591	0.0569	0.0763	0.0229	0.0334	0.1075	0.1321	0.1976	0.2251
dry cough(X)	0.2678	0.2216	0.1639	0.1498	0.0962	0.0375	0.0249	0.0993	0.1742	0.2474
NCP(X)	<b>0.3211</b>	0.2836	0.2422	0.2072	0.1674	0.1628	0.1597	0.1538	0.1489	0.1437
close contact(X)	0.7973	0.7880	0.8056	<b>0.8266</b>	0.8052	0.7856	0.7804	0.7587	0.7377	0.7323
suspected case(X)	<b>0.5967</b>	0.5633	0.5256	0.4818	0.4322	0.3819	0.3475	0.2941	0.2484	0.2010
pneumonia(X)	<b>0.5825</b>	0.5500	0.5606	0.5215	0.4737	0.4298	0.3920	0.3490	0.3067	0.2559
cough(X)	0.2030	0.2670	0.2970	0.3265	0.3463	0.3569	0.3514	0.3617	0.3970	0.4294
dyspnea(X)	0.1337	0.0889	0.0750	0.0603	0.0060	0.0506	0.0800	0.1255	0.1229	0.1168
virus pneumonia(X)	0.5053	0.5290	0.5319	0.5526	0.5932	0.6120	0.6057	0.6029	0.6263	<b>0.6655</b>
novel coronavirus(X)	<b>0.7274</b>	0.7006	0.7122	0.6797	0.6401	0.6045	0.5746	0.5401	0.5061	0.4647

Optimal values are shown in bold in the table

**Table 27** Correlation analysis between keywords and the cumulative number of confirmed cases—Xiaogan

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Center for Disease Control and Prevention(S)	0.1706	0.1457	0.1004	0.0354	0.0242	0.0570	0.1093	0.1472	0.2257	0.3002
Red Cross Society(S)	0.1681	0.1163	0.0627	0.0059	0.0586	0.1103	0.1737	0.2348	0.2977	0.3578
Spring Festival(S)	0.8011	0.8235	0.8467	<b>0.8484</b>	0.8410	0.8304	0.8374	0.8402	0.8234	0.8265
Health Commission(S)	0.0399	0.0157	0.0850	0.1377	0.1966	0.2599	0.3342	0.4098	0.4951	0.5750
vaccine(S)	0.2236	0.1937	0.1510	0.1077	0.0748	0.0442	0.0051	0.0474	0.1135	0.1913
SARS(S)	0.1686	0.2350	0.3070	0.3828	0.4611	0.5457	0.6328	0.7264	0.8191	0.9230
hand washing(S)	0.1803	0.1730	<b>0.2071</b>	0.1689	0.1617	0.1342	0.1248	0.1168	0.1083	0.0755
children(S)	0.3337	0.3857	0.3672	0.3737	0.3827	<b>0.3967</b>	0.3773	0.3459	0.3144	0.2732
mask(S)	0.0671	0.1331	0.1793	0.2310	0.2897	0.3524	0.4188	0.4843	0.5677	0.6553
coronavirus(S)	0.1115	0.1758	0.2438	0.3136	0.3902	0.4710	<b>0.5459</b>	0.6457	0.7411	0.8379
novel(S)	0.0276	0.0695	0.1335	0.1846	0.2498	0.3141	0.3923	0.4608	0.5395	0.6229
disinfection(S)	0.0088	0.0656	0.1006	0.1397	0.2084	0.2712	0.3206	0.3615	0.4243	0.4991
lockdown of the city(S)	0.1226	0.0768	0.0384	0.0074	0.0600	0.1065	0.1654	0.2358	0.3078	0.3807
Wuhan(S)	0.1064	0.1698	0.2371	0.3070	0.3718	0.4493	0.5333	0.6261	0.7191	0.8172
Zhong Nanshan(S)	0.0866	0.1438	0.2083	0.2797	0.3514	0.4348	0.5205	0.6044	0.6923	0.7906
epidemic(S)	0.8707	0.8624	0.8662	0.8725	0.8752	0.8857	<b>0.8900</b>	0.8812	0.8756	0.8769
flu(S)	0.1195	0.1824	0.2495	0.2961	0.3668	0.4409	0.5143	0.6012	0.6973	0.7944
diarrhea(S)	0.1294	0.0729	0.0246	0.0361	0.0859	0.1383	0.1987	0.2725	0.3327	0.4138
dry cough(S)	0.1157	0.1813	0.2357	0.2987	0.3383	0.4098	0.4929	0.5699	0.6548	0.7358
NCPS(S)	<b>0.7156</b>	0.6933	0.6695	0.6436	0.6206	0.5945	0.5635	0.5363	0.5048	0.4772
novel coronary pneumonia(S)	0.9257	0.9295	0.9319	0.9310	0.9326	0.9370	0.9422	<b>0.9466</b>	0.9465	0.9431
COVID-19(S)	<b>0.8302</b>	0.8121	0.7981	0.7873	0.7784	0.7787	0.7623	0.7552	0.7568	0.7327
novel coronavirus pneumonia(S)	0.2162	0.1714	0.1439	0.0950	0.0583	0.0130	0.0375	0.0947	0.1616	0.2251
asymptomatic infection(S)	<b>0.7057</b>	0.6781	0.6722	0.6385	0.6280	0.6165	0.5893	0.5580	0.5268	0.4897
aerosol transmission(S)	<b>0.4399</b>	0.4132	0.3876	0.3513	0.3213	0.2906	0.2542	0.2096	0.1737	0.1366
pneumonia(S)	0.0573	0.1212	0.1862	0.2548	0.3275	0.4027	0.4825	0.5695	0.6606	0.7547
cough(S)	0.1918	0.2373	0.2985	0.2998	0.2918	0.3493	0.4456	0.5451	0.6479	0.6461
fever(S)	0.2901	0.2834	0.3462	0.4065	0.4401	0.4887	0.4948	0.5387	0.6159	0.6809
dyspnea(S)	0.0782	0.0197	0.0241	0.0727	0.1380	0.1879	0.2415	0.3051	0.3888	0.4533
high temperature(S)	0.0221	0.0911	0.1394	0.2073	0.2454	0.3307	0.3890	0.4631	0.5346	0.6144
novel coronavirus(S)	0.0167	0.0444	0.1069	0.1733	0.2444	0.3184	0.3958	0.4801	0.5687	0.6599
SARS-CoV-2(S)	<b>0.5920</b>	0.5692	0.5799	0.5911	0.5787	0.5894	0.5688	0.5441	0.5261	0.5133
Spring Festival(X)	0.8039	0.8073	0.8169	0.8210	<b>0.8257</b>	0.8222	0.8086	0.8117	0.8115	0.7973
Red Cross Society(X)	0.2373	0.1850	0.1285	0.0805	0.0330	0.0224	0.0780	0.1418	0.2104	0.2934
government(X)	0.3292	0.3165	0.3091	0.2909	0.2538	0.2245	0.3077	0.3964	0.4051	0.3909

Table 27 (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Health Commission(X)	0.0088	0.0461	0.0492	0.0911	0.1516	0.2186	0.2670	0.3380	0.4195	0.4995
medical(X)	0.1735	0.1188	0.0712	0.0317	0.0038	0.0379	0.0649	0.1184	0.1740	0.2157
vaccine(X)	0.1955	0.1662	0.1347	0.1008	0.0677	0.0211	0.0069	0.0434	0.0977	0.1604
SARS(X)	0.1058	0.1721	0.2417	0.3126	0.3871	0.4566	0.4928	0.5531	0.6306	0.7141
children(X)	0.4338	0.4117	0.4406	0.4711	0.5210	0.5732	0.5750	0.5561	0.5350	0.5726
mask(X)	0.0584	0.0033	0.0353	0.0751	0.1170	0.1592	0.1982	0.2613	0.3216	0.3899
infection(X)	0.3539	0.3758	0.3963	0.4222	0.4470	0.4730	0.5003	0.5289	0.5589	0.5905
coronavirus(X)	<b>0.6950</b>	0.6603	0.6384	0.6082	0.5780	0.5422	0.4994	0.4696	0.4263	0.3852
disinfection(X)	0.4987	0.5097	0.5211	0.5330	0.5454	0.5585	0.5721	0.5864	0.6013	0.6170
nucleic acid(X)	<b>0.5321</b>	0.4988	0.4917	0.4689	0.4537	0.4424	0.4101	0.3646	0.3180	0.2628
lockdown of the city(X)	0.5254	0.5372	0.5496	0.5596	0.5716	0.5842	0.5782	0.5910	0.6046	0.6189
Wuhan(X)	0.0489	0.0052	0.0502	0.1226	0.1955	0.2548	0.3442	0.4100	0.4807	0.5541
Zhong Nanshan(X)	0.1225	0.0623	0.0232	0.0164	0.0712	0.1400	0.1754	0.2453	0.3055	0.3836
epidemic(X)	<b>0.4747</b>	0.4292	0.3938	0.3568	0.3132	0.2639	0.2466	0.2036	0.1452	0.0923
flu(X)	<b>0.4409</b>	0.4046	0.3841	0.3712	0.3622	0.3501	0.3424	0.3329	0.3075	0.2779
diarrhea(X)	0.0618	0.0562	0.0479	0.0466	0.0377	0.0280	0.0240	0.0196	0.0087	0.0005
stuffy nose(X)	0.1686	0.1221	0.0899	0.0767	0.0361	0.0107	0.0299	0.0784	0.1222	0.1751
dry cough(X)	0.2177	0.1759	0.1250	0.0672	0.0057	0.0542	0.0900	0.1572	0.2211	0.2945
NCP(X)	<b>0.2778</b>	0.2499	0.2188	0.1860	0.1547	0.1492	0.1460	0.0993	0.0548	0.0145
close contact(X)	<b>0.8143</b>	0.8024	0.7997	0.7830	0.7674	0.7600	0.7765	0.7570	0.7331	0.7130
suspected case(X)	<b>0.4854</b>	0.4454	0.4197	0.3755	0.3422	0.3026	0.2853	0.2509	0.2030	0.1556
pneumonia(X)	<b>0.6024</b>	0.5685	0.5455	0.5132	0.4807	0.4435	0.4007	0.3588	0.3149	0.2684
cough(X)	0.3078	0.3762	0.4317	0.4908	0.4930	0.4876	0.4609	0.4779	0.5027	0.5173
dyspnea(X)	<b>0.2301</b>	0.1881	0.1596	0.1310	0.0994	0.0715	0.0507	0.0038	0.0096	0.0146
virus pneumonia(X)	0.6014	0.6026	0.5858	0.5922	0.6070	0.6397	0.6244	0.6438	0.6775	<b>0.6971</b>
novel coronavirus(X)	<b>0.6839</b>	0.6538	0.6342	0.6069	0.5800	0.5467	0.5437	0.5164	0.4790	0.4424

Optimal values are shown in bold in the table

**Table 28** Correlation analysis between keywords and the cumulative number of confirmed cases—Yichang

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Center for Disease Control and Prevention(S)	<b>0.2973</b>	0.2656	0.2338	0.1586	0.1058	0.0726	0.0306	0.0233	0.0574	0.1112
Red Cross Society(S)	0.2334	0.1874	0.1335	0.0760	0.0412	0.0183	0.0726	0.1437	0.2001	0.2765
Spring Festival(S)	0.8581	0.8626	0.8867	<b>0.8891</b>	0.8816	0.8811	0.8757	0.8652	0.8621	0.8593
Health Commission(S)	0.1093	0.0363	0.0390	0.1066	0.1735	0.2507	0.3143	0.3830	0.4660	0.5456
vaccine(S)	0.2283	0.1916	0.1453	0.1051	0.0504	0.0074	0.0300	0.0894	0.1369	0.1822
SARS(S)	0.2104	0.2821	0.3589	0.4371	0.5213	0.6064	0.6974	0.7906	0.8597	0.9260
hand washing(S)	<b>0.3582</b>	0.3251	0.3131	0.2739	0.2183	0.2115	0.1608	0.1452	0.0978	0.0608
children(S)	0.5504	0.5269	0.5023	0.4896	0.4517	0.4889	<b>0.5820</b>	0.5613	0.5195	0.4906
mask(S)	0.0654	0.1330	0.1999	0.2733	0.3415	0.4032	0.4718	0.5476	0.6233	0.7081
coronavirus(S)	0.1046	0.1709	0.2392	0.3120	0.3895	0.4692	0.5501	0.6394	0.7388	0.8400
novel(S)	0.0064	0.0524	0.1123	0.1798	0.2386	0.3143	0.4062	0.4859	0.5788	0.6819
disinfection(S)	<b>0.2895</b>	0.2560	0.2028	0.1863	0.1297	0.0948	0.0768	0.0341	0.0551	0.0845
lockdown of the city(S)	0.1438	0.1149	0.0629	0.0027	0.0471	0.0968	0.1669	0.2381	0.2979	0.3787
Wuhan(S)	0.0907	0.1649	0.2370	0.3075	0.3810	0.4604	0.5361	0.6255	0.7256	0.8191
Zhong Nanshan(S)	0.0678	0.1369	0.2056	0.2728	0.3493	0.4281	0.5111	0.6018	0.6936	0.7868
epidemic(S)	<b>0.6942</b>	0.6585	0.6218	0.5965	0.5608	0.5411	0.5269	0.5096	0.4570	0.4126
flu(S)	0.0838	0.1225	0.1923	0.2631	0.3360	0.4060	0.4808	0.5577	0.6560	0.7517
diarrhea(S)	0.0434	0.0154	0.0805	0.1277	0.1849	0.2577	0.3182	0.3704	0.4404	0.5074
dry cough(S)	0.0940	0.1602	0.2271	0.2883	0.3508	0.4174	0.4984	0.5762	0.6715	0.7603
NCP(S)	<b>0.7519</b>	0.7287	0.6940	0.6567	0.6342	0.6127	0.5820	0.5585	0.5445	0.5100
novel coronary pneumonia(S)	<b>0.9276</b>	0.9181	0.9122	0.9103	0.9074	0.9099	0.9058	0.8920	0.8885	0.8901
COVID-19(S)	<b>0.8404</b>	0.8245	0.8043	0.7893	0.7756	0.7699	0.7401	0.7245	0.7150	0.7007
novel coronavirus pneumonia(S)	0.2147	0.1610	0.1065	0.0701	0.0260	0.0329	0.0920	0.1602	0.2138	0.2783
asymptomatic infection(S)	<b>0.6790</b>	0.6694	0.6503	0.6150	0.5787	0.5628	0.5304	0.4863	0.4438	0.4107
aerosol transmission(S)	<b>0.5176</b>	0.4851	0.4573	0.4336	0.4031	0.3702	0.3345	0.2890	0.2481	0.2059
pneumonia(S)	0.0663	0.1411	0.2129	0.2830	0.3592	0.4398	0.5230	0.6106	0.7126	0.8072
cough(S)	0.2371	0.1942	0.1436	0.0694	0.0293	0.0156	0.0873	0.1404	0.1776	0.2319
fever(S)	0.1348	0.2055	0.2808	0.2767	0.3188	0.3671	0.4319	0.4988	0.5626	0.5796
dyspnea(S)	0.0455	0.1005	0.1527	0.2119	0.2989	0.3265	0.3799	0.4638	0.5157	0.6099
high temperature(S)	0.1110	0.1708	0.1849	0.2514	0.3068	0.3938	0.4730	0.5361	0.6087	0.6968
novel coronavirus(S)	0.0170	0.0541	0.1180	0.1868	0.2627	0.3364	0.4088	0.4954	0.5925	0.6827
SARS-CoV-2(S)	<b>0.6330</b>	0.6117	0.5892	0.5653	0.5533	0.5297	0.5108	0.4707	0.4605	0.4492
Spring Festival(X)	0.8399	0.8492	0.8579	0.8648	0.8734	0.8725	0.8780	<b>0.8856</b>	0.8781	0.8604
Red Cross Society(X)	0.2758	0.2338	0.1888	0.1284	0.0784	0.0681	0.0262	0.0716	0.1123	0.1727
government(X)	0.2940	0.2999	0.2678	0.2205	0.2130	0.2059	0.2414	0.2940	0.3445	0.3287

Table 28 (continued)

Keywords	Delay days									
	14	13	12	11	10	9	8	7	6	5
Health Commission(X)	0.0630	0.1412	0.1564	0.1977	0.2644	0.3267	0.3460	0.3919	0.4968	0.5683
medical(X)	0.1467	0.0957	0.0513	0.0147	0.0660	0.1037	0.1437	0.1814	0.2602	0.3074
vaccine(X)	0.1968	0.1513	0.1313	0.0999	0.0552	0.0401	0.0167	0.0601	0.0892	0.1354
SARS(X)	0.1007	0.1639	0.2219	0.2936	0.3750	0.4103	0.4524	0.5043	0.5992	0.6687
children(X)	0.4189	0.4484	0.4778	0.4718	0.5169	<b>0.5513</b>	0.5276	0.5014	0.4943	0.4679
mask(X)	0.1186	0.0663	0.0233	0.0182	0.0774	0.1089	0.1370	0.1999	0.2889	0.3309
infection(X)	0.3301	0.3392	0.3616	0.3924	0.4207	0.4385	0.4855	0.5139	0.5269	0.5590
coronavirus(X)	<b>0.6514</b>	0.6147	0.5918	0.5683	0.5279	0.5216	0.4943	0.4691	0.4433	0.4133
disinfection(X)	0.5295	0.5412	0.5534	0.5423	0.5544	0.5670	0.5803	0.5941	0.6087	0.6160
nucleic acid(X)	<b>0.5181</b>	0.4873	0.4691	0.4404	0.4024	0.3938	0.3958	0.3529	0.2937	0.2551
lockdown of the city(X)	0.5070	0.5274	0.5433	0.5488	0.5534	0.5823	0.5898	0.5927	0.6094	0.6091
Wuhan(X)	0.0200	0.0103	0.0645	0.1375	0.2188	0.2766	0.3507	0.4376	0.5156	0.5755
Zhong Nanshan(X)	0.1379	0.0734	0.0131	0.0373	0.0865	0.1389	0.2118	0.3257	0.3949	0.4533
epidemic(X)	<b>0.4238</b>	0.3740	0.3402	0.2927	0.2347	0.1966	0.1594	0.1385	0.0657	0.0079
flu(X)	<b>0.4846</b>	0.4318	0.3863	0.3661	0.3540	0.3450	0.3272	0.3260	0.3107	0.2866
diarrhea(X)	0.0405	0.0394	0.0541	0.0432	0.0045	0.0028	0.0168	0.0075	0.0041	0.0008
stuffy nose(X)	0.0895	0.0185	0.0277	0.0765	0.1049	0.1404	0.1723	0.1928	0.2444	0.2931
dry cough(X)	0.2406	0.1964	0.1567	0.0975	0.0251	0.0232	0.0821	0.1209	0.1842	0.2510
NCP(X)	<b>0.4199</b>	0.3944	0.3566	0.3188	0.2944	0.2848	0.2694	0.2242	0.1803	0.1462
close contact(X)	<b>0.8282</b>	0.8102	0.7950	0.7899	0.7639	0.7464	0.7535	0.7519	0.7159	0.7091
suspected case(X)	<b>0.5090</b>	0.4664	0.4393	0.4039	0.3434	0.3054	0.2675	0.2426	0.2059	0.1583
pneumonia(X)	<b>0.5050</b>	0.4587	0.4229	0.3985	0.3530	0.3068	0.2762	0.2618	0.1942	0.1474
cough(X)	0.3112	0.3754	0.4428	0.5117	0.5158	0.4920	0.4970	0.4912	0.5242	0.5402
dyspnea(X)	0.2093	0.1655	0.1260	0.0846	0.0402	0.0199	0.0170	0.0463	0.0374	0.0155
virus pneumonia(X)	0.5725	0.5896	0.5660	0.5759	0.6024	0.5902	0.5988	0.6370	<b>0.6459</b>	0.6435
novel coronavirus(X)	<b>0.7394</b>	0.7076	0.6838	0.6633	0.6313	0.6264	0.6168	0.5572	0.4823	0.4454

Optimal values are shown in bold in the table

## Appendix B

Critical values for the two-tailed Bonferroni–Dunn test:

Methods	2	3	4	5	6	7
q0.05	1.960	2.241	2.394	2.498	2.576	2.638
q0.10	1.645	1.960	2.128	2.241	2.326	2.394
Methods	8	9	10	11	12	13
q0.05	2.690	2.724	2.774	3.219	3.268	3.313
q0.10	2.450	2.498	2.539	2.978	3.030	3.077

where *methods* is the number of methods used for comparison.

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**Data availability** The datasets generated during and/or analyzed during the current study are available from the corresponding author on reasonable request.

## Declarations

**Conflict of interest** The authors declare that they have no conflict of interest.

**Ethical approval** This article does not contain any studies with human participants performed by any of the authors.

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