

Team Control Number

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Problem Chosen

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Summary

With the rapid development of China's economy and society, and the aging trend, the demand for hospital health service continues to increase. The core objective of this paper is to establish mathematical models to describe the relationship between the common disease, the medical development and the hospitals. Thus we can allocate social resources more appropriately.

In the first task, on the basis of the traditional grey prediction model, we use the Markov chain to correct the result residual. We first established the traditional GM (1,1) model to predict and analyze the proportion of elderly people and medical demand in the period of 2009-2021. Then we calculate the expected value of predicted residual through Markov model. Finally, residual expectation is used to correct the inherent deviation of traditional grey prediction. The prediction residual of the model is less than 7%, and the NSE value approaches to 1, which indicates that the model has high reliability.

In the second task, according to characteristics of each disease, we select the most common disease by TOPSIS. Firstly, via the grey Markov model of question one, we predict number of disease and death in Hubei Province. The morbidity and mortality, the growth rate of morbidity and mortality and the morbidity and mortality in 2019-2021 are obtained. The above five factors are brought into the TOPSIS model, with weights of 0.15, 0.3, 0.15, 0.3 and 0.1 respectively. The comprehensive evaluation scores of each disease and each region are calculated. The ranking results show that heart disease is the key monitored disease in Hubei Province.

In the third task, according to the data of real world hospital, we first establish the $M/M/c/\infty$ queuing model and then abstract the medical statistic into model parameters. Then based on this queuing model, we adopt the optimized systems design by figuring out the number of doctors in each outpatient department where the social cost gets to the minimum.

In the fourth task, we use latent class analysis (LCA) to identify each patient segments based on detailed information on their use of hospital services. Then we described the resultant patient types in terms of their characteristics. Finally, we exploit the availability of hospital data in the post-survey period to test the relevance of the patient types in order to optimize the future hospital utilisation.

keyword: GM (1,1)-Markov; TOPSIS; LCA; queuing model;

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1 Introduction

1.1 Background

With the rapid development of China's economy and the deepening trend of aging, people's demand for hospital health services is also getting higher and higher. Therefore, it is of great significance to establish an appropriate model to study the trend of aging in the future and the trend of people's demand for medical care. At the same time, the establishment of a reasonable competition and cooperation mechanism between private and public hospitals can also maximize the utilization of resources.

1.2 Work

The problem requires us to make rational use of network resources?answer the following questions

1. The aging trend of China and the people's medical needs to make a reasonable forecast.
2. Take a province as an example to analyze the most common disease in the future.
3. According to the medical needs of patients, an optimal queuing method is proposed to equalize the number of queues between different hospitals and different departments of the same hospital.
4. Put forward the best cooperation and competition strategy between private hospital and public hospital.
5. Write recommendations for relevant medical departments and prepare "14 five-year plans" for their reference.

2 Problem Analysis

Analysis of task one According to the relevant data from 2009 to 2018 in the National Bureau of statistics, the group first selects appropriate indicators and then establishes a grey prediction model to predict and analyze the population aging trend and residents' medical needs of the epidemic in 2009-2018. Then, through the Markov model, the data from 2009-2018 simulates the distribution of residual in each interval and calculates the expectation of the predicted residual in 2019-2021 Value. Finally, the prediction results and residual expectation are made to be different, and the inherent deviation of traditional grey prediction is corrected.

Through the combination of the two models, the goal of scientific prediction of the future development of population aging and the trend of residents' medical needs is achieved.

Analysis of task two According to the second question, we are asked to analyze the most common diseases in a specific province in the future . we took the Hubei province as an example to collect data on common diseases such as heart disease ,and established a model with Markov model.And calculated The number of cases, growth rate of cases, growth rate of deaths and death rate of each disease ,Then substituted the five index data into the principal component analysis model. The disease with the highest score was the most common disease in the province in the future.

Analysis of task three The task three ask us to propose a common queuing theory method and figure its related optimal queuing for this kind of queuing problem.Via referring to the operational mode of the real world hospital, we choose to adopt the $M/M/c/\infty$ model as the required theory. Then we acquire the hospital statistics and abstract the data into the parameter in model. Finally, to get the social cost to be lowest, we manage to find the optimized systems design for the queuing model.

Analysis of task four The task four ask us to analyse the complex cooperation and competition between private hospitals and public hospitals then propose the optimal cooperation and competition strategies among multiple hospitals. After referring to a deal of data and document, we are aware of that there exist complicated partnerships. Based on latent class analysis (LCA),we would provide the best strategies we come up with.

3 Symbol and Assumptions

3.1 Symbol Description

symbols	definitions
$X^{(i)}$	People time series
a	Developing grayscale
u	Endogenous control gray
$\hat{\alpha}$	Parameter vector to be estimated
ε	Residual sequence
P	State transition matrix
E_k	State interval
η	Error expectation
$\bar{x}(k)$	GM-Markov combination forecast value
$\hat{x}(k)$	GM prediction value
t'_0	Initial state probability vector
A	Multiple attribute decision matrix
B	Normalized decision matrix
W	Weight vector
C	Weighted gauge matrix
C^*	Positive ideal solution
C^0	Negative ideal solution
d_i	Attribute decision vector
f_i^*	Comprehensive evaluation index
δ_i	Incidence growth rate
η_i	Morbidity and mortality
C	amount of doctors
λ	average arrival rate
μ	average inspection rate
P_n	probability of number of people reach n
W_q	average waiting time
W_s	average time of stay
L_s	average length of queuing
Z	social cost
c_s	cost of each doctor
c_w	cost of waiting

3.2 Fundamental assumptions

1. In order to ensure the accuracy of the prediction results, it is assumed that the data given in the database is authentic.

2. It is assumed that the growth rate of morbidity and mortality is more important than its base in the key prevention and control of diseases.
3. Ignore the influence of different diagnosis methods on the total number of patients and deaths.
4. The hourly wage of patient could be considered as the cost dissipated in queuing. And the hourly wage of doctor could be seen as the cost of cost of service.

4 Establishment and solution of the model

4.1 The model of Problem 1

4.1.1 Establishment of model

GM(1,1) The total number of cases in 2009-2018 is time series:

$$X^{(0)} = [x^{(0)}(1), x^{(0)}(2), \dots, x^{(0)}(10)]$$

Generate a 1-AGO sequence by one accumulation:

$$X^{(1)} = [x^{(1)}(1), x^{(1)}(2), \dots, x^{(1)}(10)]$$

In the formula: $x^{(1)}(k) = \sum_{i=1}^k x^{(0)}(i)$, $k = 1, 2, \dots, 10$.

Establish a differential equation based on the 1-AGO sequence as:

$$\frac{dX^{(1)}}{dt} + aX^{(1)} = u \quad (1)$$

In the formula: a is Develop grayscale, u is Endogenous control grayscale. Let $\hat{\alpha}$ be the parameter vector to be estimated and $\hat{\alpha} = [a, u]^T$, be found by least squares method:

$$\hat{\alpha} = (B^T B)^{-1} B^T Y_n \quad (2)$$

Solving equation 1, The preliminary prediction model for the $k + 1$ aging is available:

$$\hat{X}(k + 1) = [X^{(0)}(1) - \frac{u}{a}]e^{-ak} + \frac{u}{a}, k = 1, 2, \dots, 10 \quad (3)$$

Similarly, the death number is taken as the vector $X^{(0)} = [x^{(0)}(1), x^{(0)}(2), \dots, x^{(0)}(10)]$ Bring in the model to obtain the 2019-2021 death number grayscale prediction value.

Markov correction The Markov model is used to estimate the state and state probability of the GM(1,1) prediction error term, and the predicted value of the predicted state is used to correct the GM(1,1) prediction value. The state is divided by the 2009-2018 forecast data and the real data residual, and the residual sequence is:

$$\varepsilon = [\varepsilon(1), \varepsilon(2), \dots, \varepsilon(10)]$$

Absolute maximum residual value $\delta_{max} = \max_{1 \leq i \leq 10} |\varepsilon(i)|$. The prediction error is divided into three states. Let $\lambda = \frac{\delta_{max}}{6}$. The status is $E_1 : (-3\lambda, -\lambda)$, $E_2 : (-\lambda, \lambda)$ and $E_3 : (\lambda, 3\lambda)$. The formula for calculating the initial state probability vector is:

$$\begin{cases} p_{Ek} = \frac{n_{Ek}}{13} \\ t_0 = [p_{E1}, p_{E2}, p_{E3}] \end{cases} \quad (4)$$

In the formula: n_{Ek} is number of E_k occurrences in 2008-2019. Replace the probability E_k of its occurrence with the frequency at which the state p_{Ek} appears. And construct the state transition matrix as:

$$P = \begin{pmatrix} P_{11} & P_{12} & P_{13} \\ P_{21} & P_{22} & P_{23} \\ P_{31} & P_{32} & P_{33} \end{pmatrix}$$

In the formula: P_{ij} is E_i transition probability transferred to E_j after a period.

That is, the Markov model can be expressed as:

$$t_{k+1} = t_k \cdot P \quad (5)$$

Let the middle value of the status interval be \bar{E}_1, \bar{E}_2 and \bar{E}_3 , so the error expectation of GM(1,1) in the kth year is: .

$$\eta = [p_{E1} \quad p_{E2} \quad p_{E3}] \cdot \begin{bmatrix} \bar{E}_1 \\ \bar{E}_2 \\ \bar{E}_3 \end{bmatrix} \quad (6)$$

When the predicted value of GM(1,1) for the number of patients in the k year is $\hat{x}(k)$, Modified grey Markov combination forecast model $\bar{x}(k)$ Can be recorded as?

$$\bar{x}(k) = \hat{x}(k) - \eta \quad (7)$$

Forecast result evaluation index Root mean square error (RMSE), average phase error absolute value (MAPE), and Nash efficiency coefficient (NSE) are commonly used to measure prediction results. The RMSE can evaluate the high-value predictions of the number of patients and the number of deaths. The calculation formula is:

$$\text{RMSE} = \sqrt{\frac{1}{n} \sum_{i=1}^n (y_i - y_i^*)^2}$$

The smaller the root mean square error, the higher the reliability of the model and the more accurate the result.

MAPE is used to evaluate the prediction results of the stationary part of the prediction data. The calculation formula is:

$$\text{MAPE} = \frac{1}{n} \sum_{i=1}^n \left| \frac{y_i - y_i^*}{y_i} \right| \times 100\%$$

The value obtained by MAPE is an absolute value, which is a relative index. When two MAPE values are compared, the smaller the value, the higher the reliability of the model.

The NSE can be used to evaluate the predictive power of the model. The formula is as follows:

$$\text{NSE} = 1 - \frac{\sum_{i=1}^n (y_i - y_i^*)^2}{\sum_{i=1}^n (y_i - \bar{y})^2}$$

The closer the NSE value is to 1, the better the model quality and the higher the model's credibility. Close to 0, indicating that the simulation result is close to the average level of observations, that is, the overall result is credible, but the simulation error is large. Far less than 0, the model is not credible.

4.1.2 Solution of Grey Markov Model

The predicted values of the Chinese elderly population forecast for 2009-2018 by GM(1,1) are as follows:

$$\hat{X}(k+1) = 266856.0905e^{0.042782k} - 255354.5841, k = 1, 2, \dots, 10 \quad (8)$$

The error status range is shown in the table 1 .

Table 1: The age range division of the elderly

Status	E_1	E_2	E_3
Residual interval	$[-469, -221]$	$(-221, 221]$	$(221, 469]$

According to the error interval range, the predicted number of elderly people in 2009-2018 is classified into the error interval as shown in the table 2 .

Table 2: The age range division of the elderly

year	2009	2010	2011	2012	2013	2014	2015	2016	2017	2018
Residual interval	E_2	E_2	E_1	E_2	E_3	E_2	E_1	E_1	E_2	E_2

From this, the initial state probability vector t_0 is obtained, and the transfer matrix P is:

$$t'_0 = [3/10, 3/5, 1/10]$$

$$P' = \begin{pmatrix} 1/3 & 2/3 & 0 \\ 1/4 & 5/8 & 1/8 \\ 0 & 1 & 0 \end{pmatrix} \quad (9)$$

The predicted solution obtained by gray prediction and Markov correction is shown in the figure 1 .

Similarly, the calculation of the medical demand forecast value for 2009-2018 is obtained as follows:

$$\hat{X}(k+1) = 258.7846e^{-0.043809k} - 246.9258, k = 1, 2, \dots, 10 \quad (10)$$

The error status range is shown in the table 3 .

Table 3: Medical demand status interval division

Status	E_1	E_2	E_3
Residual interval	$[-164.45, -58.74]$	$(-58.74, 58.74]$	$(58.74, 164.45]$

The medical demand forecast is classified into the error interval as shown in the table ?? .

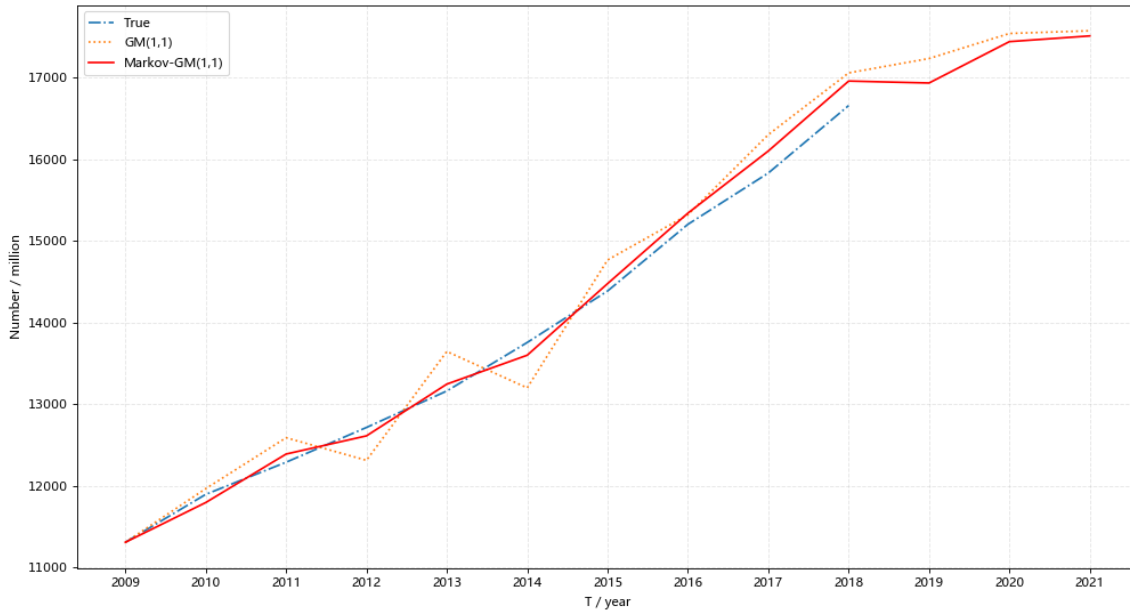


Figure 1: Comparison curve of prediction value of the number of the elderly

Table 4: Medical demand error status interval

year	2009	2010	2011	2012	2013	2014	2015	2016	2017	2018
Residual interval	E_2	E_3	E_1	E_2	E_3	E_1	E_1	E_3	E_2	E_2

From this, the initial state probability vector t_0 is obtained, and the transfer matrix P is:

$$t'_0 = [3/10, 2/5, 3/10]$$

$$P' = \begin{pmatrix} 0 & 1/2 & 1/2 \\ 1/8 & 3/4 & 1/8 \\ 1/2 & 1/2 & 0 \end{pmatrix} \quad (11)$$

The predicted solution obtained by gray prediction and Markov correction is shown in the figure 2 .

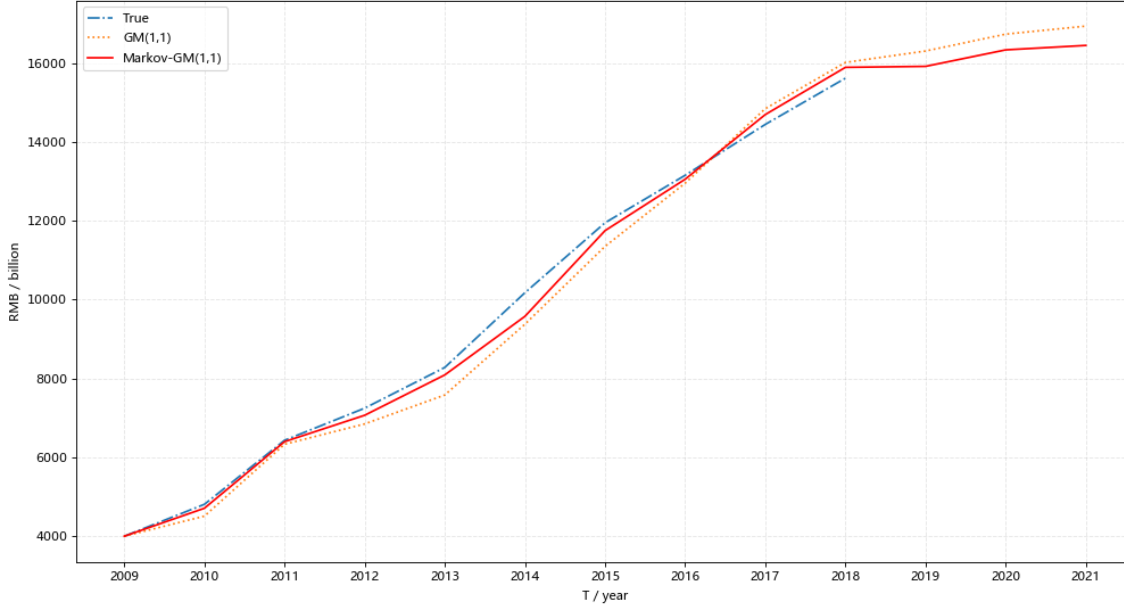


Figure 2: Comparison curve of medical demand forecast

4.1.3 Conclusion

According to the direct comparison of the prediction solution curves in Figure 1 and figure 2, the predicted value corrected by Markov model has higher fitting degree and consistent volatility compared with the traditional gray prediction value, and can reflect the actual value fluctuation better than the traditional gray model prediction value. The prediction indexes of the two models are shown in table 5:

Table 5: Test of prediction results

Test parameters	RMSE	MAPE	NSE
GM(1,1) for elderly people	3040.04	0.0213	0.9455
Markov-GM(1,1) for elderly people	1238.64	0.0095	0.9900
GM(1,1) for medical demand	265.88	0.0628	0.8178
Markov-GM(1,1) for medical demand	101.13	0.0273	0.9736

From the above prediction results, it can be concluded that the RMSE of the number of patients and the number of deaths calculated by using the modified grey Markov model is smaller than that of the traditional grey model, which shows that the modified results are more reliable. The MPAE value of the modified model is closer to 0 and the NSE value is closer to 1 compared with the traditional model, which shows that the improved gray Markov model has a higher fitting degree and

better prediction effect, which is suitable for the short-term prediction of the number of infectious diseases and deaths.

4.2 The model of Problem 2

4.2.1 Establishment of model

According to the different data of each year, we modified the grey markov model of the first question.

$$\begin{cases} \bar{x}(k) = \hat{x}(k) - \eta \\ \hat{X}(k+1) = [X^{(0)}(1) - \frac{u}{a}]e^{-ak} + \frac{u}{a}, k = 1, 2, \dots, 13 \\ \eta = [p_{E1} \ p_{E2} \ p_{E3}] \cdot [\bar{E}_1 \ \bar{E}_2 \ \bar{E}_3]^T \end{cases} \quad (12)$$

The revised predicted number of patients in year k was $\bar{x}(k)$, $\hat{x}(k)$ is the traditional predicted value of $GM(1, 1)$, η is the expected error of GM in year k .

4.2.2 Calculation of predictors

We set X_i as the incidence data of 13 diseases from 2009 to 2021 in hubei province, $X_i = [x_{i1}, x_{i2}, x_{i3}, x_{i4}, x_{i5}] (i = 1, 2, \dots, 13)$, The prediction results are obtained by using markov model. Similarly, Put the death toll data $X'_i = [x'_{i1}, x'_{i2}, x'_{i3}, x'_{i4}, x'_{i5}] (i = 1, 2, \dots, 13)$ into the forecasting results $X'_i = [x'_{i1}, x'_{i2}, x'_{i3}, x'_{i4}, x'_{i5}, x'_{i6}] (i = 1, 2, \dots, 13)$.

According to the prediction, the growth rate of the number of cases, the growth rate of the number of deaths and the mortality rate of the sick population in 2019 are respectively:

$$\begin{cases} \delta_i = \frac{x_{i6} - x_{i5}}{x_{i5}} \times 100\% \\ \delta'_i = \frac{x'_{i6} - x'_{i5}}{x'_{i5}} \times 100\% \\ \eta_i = \frac{x_{i6}}{x_{i5}} \times 100\% \end{cases} \quad (13)$$

Get the decision attribute vector of each group:

$$d_i = [x_{i6}, x'_{i6}, \delta_i, \delta'_i, \eta_i]$$

Similarly, the number of morbidity and mortality of various diseases in Hubei province were brought into the model to obtain the decision attribute vectors of various diseases $d'_i (i = 1, 2, \dots, 13)$.

4.2.3 TOPSIS

Let the disease multi-attribute decision matrix $A = (a_{ij})_{13 \times 5}$ be expressed as:

$$A = [d_1^T, d_2^T, \dots, d_{13}^T]^T$$

Normalize A to obtain normalized decision matrix $B = (b_{ij})_{13 \times 5}$, where:

$$b_{ij} = a_{ij} / \sqrt{\sum_{i=1}^{13} a_{ij}^2}, i = 1, 2, \dots, 13; j = 1, 2, \dots, 5$$

It is assumed that regions with high growth rates of the number of patients and deaths need to focus on prevention and control, construct weight vectors:

$$W = [0.15, 0.3, 0.15, 0.3, 0.1] \quad (14)$$

The weighted canonical matrix, positive ideal solution, negative ideal solution can be obtained as follows: $C = (c_{ij})_{13 \times 5}$, $C^* = [c_1^*, c_2^*, c_3^*, c_4^*, c_5^*]$, $C^0 = [c_1^0, c_2^0, c_3^0, c_4^0, c_5^0]$, where in:

$$\begin{cases} c_{ij} = w_j \cdot b_{ij}, i = 1, 2, \dots, 13; j = 1, 2, \dots, 5 \\ c_j^* = \max_i(c_{ij}), j = 1, 2, \dots, 5 \\ c_j^0 = \min_i(c_{ij}), j = 1, 2, \dots, 5 \end{cases}$$

Calculating the distance from each disease attribute decision vector to the positive and negative ideal solution. The distance from d_i to the positive ideal solution and from d_i to the negative ideal solution is:

$$\begin{cases} s_j^* = \sqrt{\sum_{j=1}^n (c_{ij} - c_j^*)^2}, i = 1, 2, \dots, 13 \\ s_j^0 = \sqrt{\sum_{j=1}^n (c_{ij} - c_j^0)^2}, i = 1, 2, \dots, 13 \end{cases}$$

Calculating the comprehensive evaluation index of each program:

$$f_i^* = s_j^0 / (s_j^0 + s_j^*), i = 1, 2, \dots, 13 \quad (15)$$

Obtain the order of key prevention and control diseases according to f_i^* (range from large to small). Similarly, the decision vector $d'_i (i = 1, 2, \dots, 13)$ of each disease can be brought into TOPSIS model to obtain the priority of disease prevention and control

4.2.4 Results and Conclusion

The number of cases and deaths of each occupation were put into the grayscale markov model to obtain the predicted values, as shown in table 6 .

Table 6: Predicted results of patients with various diseases

disease/10*thousand	2009	2010	...	2019	2020	2021
Immunodeficiency	1.5686	1.5021	...	1.3377	1.2777	1.2673
Endocrine disease	20.3322	18.129	...	20.3322	19.2462	18.6414
Nervous system	6.8927	5.8433	...	8.6325	8.9582	7.8506
Urogenital diseases	7.3363	7.2032	...	6.892	6.943	7.042
...
Heart disease	128.8231	129.1872	...	149.45	153.454	155.6778
infectious disease	7.9431	7.421	...	5.437	4.824	4.534
Perinatal diseases	1.5351	2.0284	...	1.124	0.943	0.7423
Digestive system diseases	16.578	16.9643	...	14.45	14.43	14.34
Other	10.7305	9.5811	...	6.745	7.345	7.123

Calculate the positive ideal solution and the negative ideal solution as:

$$C^0 = [0.0066, 0.0048, 0.0653, 0.0261, 0.0331]$$

$$C^* = [-0.1856, -0.1847, -0.0809, -0.1163, -0.0929]$$

That is, the comprehensive evaluation index of the disease of 13 is shown in the figure 3 :

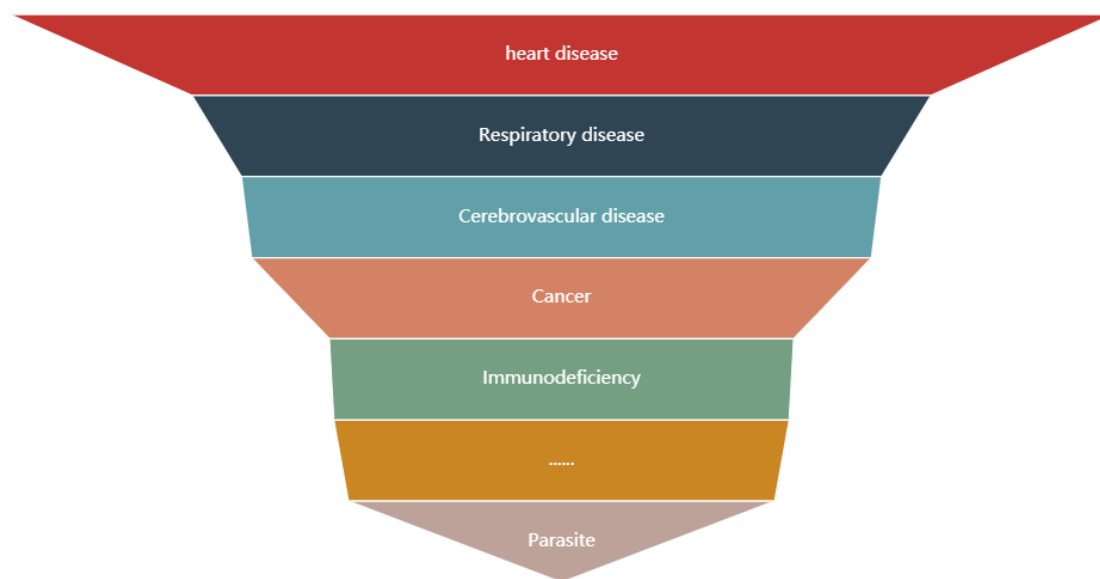


Figure 3: TOPSIS scores for different diseases

According to TOPSIS ranking, the number of heart attack and death in Hubei Province is on the rise, and the rising rate is high, which belongs to the key population of prevention and control.

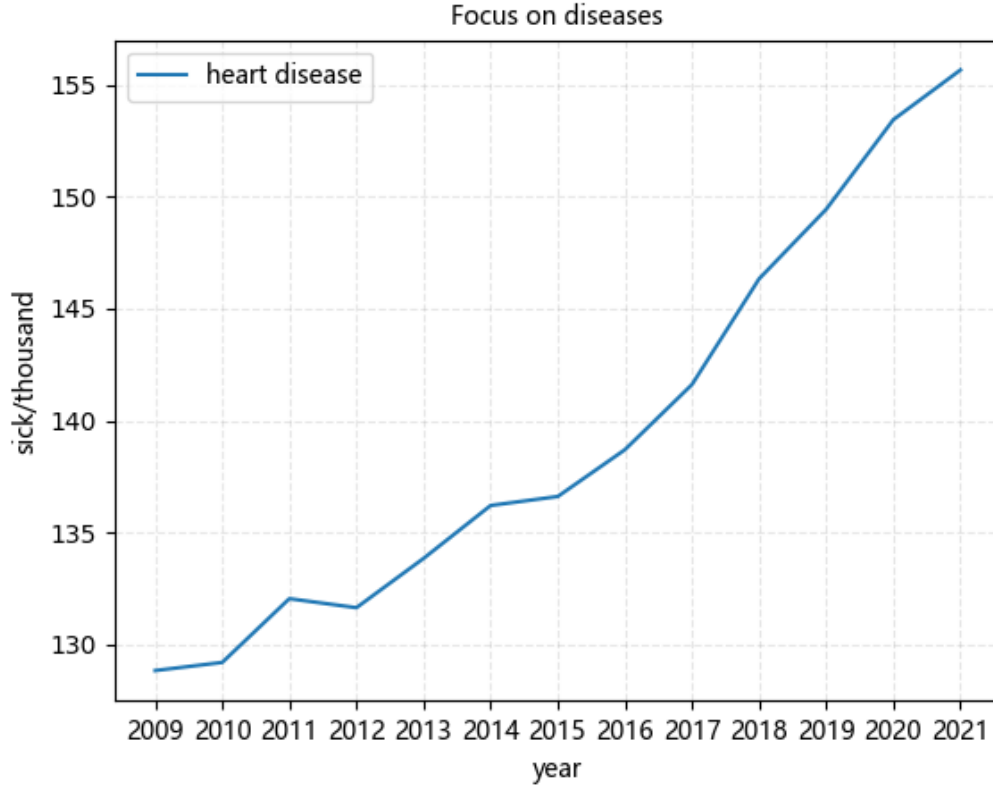


Figure 4: Focus on diseases

4.3 The model of Problem 3

4.3.1 Establishment of model

$M/M/c/\infty$ model As the theory of $M/M/c/\infty$ illustrate:

$$\begin{cases} P_0 = \left[\sum_{k=0}^{c-1} \frac{1}{k!} \left(\frac{\lambda}{\mu} \right)^k + \frac{1}{c!} \cdot \frac{1}{1-\rho} \cdot \left(\frac{\lambda}{\mu} \right)^c \right]^{-1} \\ P_n = \begin{cases} \frac{1}{n!} \left(\frac{\lambda}{\mu} \right)^n P_0 & (n \leq c) \\ \frac{1}{c!c^{n-c}} \left(\frac{\lambda}{\mu} \right)^n P_0 & (n > c) \end{cases} \end{cases} \quad (16)$$

In the above expression, n is the amount of patient. p_n is the probability of number of patient reach n in the moment t . C represents the amount of doctors in an outpatient department. λ means the average arrival rate of patients. And the μ means the average inspection rate of the doctor.

Based on this theory, the following relation is set up:

$$\begin{cases} L_s = L_q + \frac{\lambda}{\mu} \\ L_q = \sum_{n=c+1}^{\infty} (n-c)P_n = \frac{(c\rho)^c \rho}{c!(1-\rho)^2} P_0 \end{cases} \quad (17)$$

In the above expression, L_s is the average length of the queuing which means the number of patient waiting in the system. Depending on it, we could derive the expression of average waiting time W_q and average time of stay W_s :

$$\begin{cases} W_q = \frac{L_q}{\lambda} \\ W_s = \frac{L_s}{\lambda} \end{cases} \quad (18)$$

4.3.2 Solution of model

Optimized systems design For $M/M/c/\infty$ system, the expectation of the social cost per hour in the steady state is set up as Z . And Z could be calculated as the following expression:

$$z = c_s \cdot c + c_w \cdot L_s \quad (19)$$

In this expression, C is the number of doctors in the outpatient department. $C(s)$ represents the average hourly wage of the doctors while $C(w)$ represents the patient's L_s is the amount of patient waiting in the system witch is function of C . Thus, Z is the function of C . To get the optimal solution of C^* which make Z the minimum, we adopt the marginal analysis to solve it. And accroding to the feature that Z is the minimum, the following relation exists:

$$\begin{cases} Z(c^*) \leq Z(c^* - 1) \\ Z(c^*) \leq Z(c^* + 1) \end{cases} \quad (20)$$

Take expression (15) into expression (16) to gain:

$$L(C^*) - L(C^* + 1) \leq \frac{C(s)}{C(w)} \leq L(C^* - 1) - L(C^*) \quad (21)$$

Thus, we utilize the ergodic algorithm to obtain the value of C^* .

4.4 The model of Problem 4

4.4.1 Establishment of model

Latent class analysis (LCA) we use latent class analysis (LCA) to identify distinct consumer segments in the hospital care markets based on detailed information on their use of hospital services. By examining the hospital utilisation pattern of

each type, we assign putative names for all types. For indicators of hospital use, we consider not only the location and frequency of admissions, but also resource use during admissions, which reflects patient health. The indicators of resource use that we considered are the number of secondary procedures, which reflect the complexity of procedures for example due to comorbidities, the length of stay and any record of emergency department presentation. The resultant patient types are described in terms of their demographics and background characteristics, as well as variables that can be manipulated such as income and lifestyle-related factors. And in the end, we exploit the availability of hospital data in the post-survey period to test the relevance of the patient types in predicting future hospital utilisation.

Let $Y_i = (Y_{i1}, \dots, Y_{iM})$ denote patient's response to M hospital indicator variables, where the possible values of Y_{iM} are $1, \dots, r_m$. Let $L_i = 1, 2, \dots, n_c$ be the latent type of patient. Where n_c is the possible number of latent types. And let X_i denote the covariates of patient i that affects the market segmentation. The contribution by patient i to the likelihood is:

$$(1) P(Y_i = y | X_i = x) = \sum_{l=1}^{n_c} \gamma_l(x) \prod_{m=1}^M \prod_{k=1}^{r_m} \rho_{mk|l}^{I(y_m=k)} \quad (22)$$

where $I(y_m = k)$ is an indicator function that is equal to 1 if y_m is equal to k and 0 otherwise. The parameters to estimate are the probability of membership in each latent type, which is the gamma parameters, γ , and the item-response probabilities conditional on type membership, which is the rho parameter, β . The gamma parameter can be given by a multinomial logit (MNL) model such as:

$$\gamma_l(x) = P(L_i = l | X_i = x) = \frac{\exp(x\beta_l)}{\sum_{j=1}^{n_c} \exp(x\beta_j)} \quad (23)$$

where β is the MNL coefficient.

4.4.2 Solution of model

From 22, we can see that the model will increase in size very quickly with M and r_m . For example, a model with eight binary indicators would result in 28 or 256 cells. In our case, we have six hospital indicator variables which are a mix of continuous and count variables at many levels. Therefore, some grouping of the levels is needed. We base ours on the inspection of the distribution of each indicator with the intention that each level represents a distinct case of hospital use.

The next part of our analysis concerns the extent to which the identified patient types can predict future demand for hospital services. The patient types are included as covariates, and the identifying restriction is satisfied using lifestyle variables; lifestyles affect the probability of an admission but do not directly affect the

choice between public or private hospital. We use hospital data from a year after the survey date. Because only a few patients in our data were hospitalised for the same health problems again in the following year, for prospective use, we consider both repeat procedures and the overall demand for any procedure. In this case, we find that private health insurance can lower the utilisation of public hospitals, thereby reducing the public health care burden. A reduction in health costs may also be achieved through public policies that promote a general improvement in health to reduce use of resources per admission.

As a consequence, the strategies is to moderate the private health insurance and encourage the private hospital to establish the outpatient department the public hospital have little of while improving the quality of service of private hospital.

4.5 A letter to the medical department

Dear Medical leaders:

According to our modeling research, the aging trend in China is increasingly strengthened, and the majority of people's medical needs are increasingly high. It is estimated that the aging population over 65 will reach 170 million in 2021, which means that China will enter the aging society. According to the data, China's aging is on the rise, and meeting the medical and old-age needs of the elderly will be a huge social problem. Therefore, we should take measures as soon as possible in the field of gerontology scientific research, clinical medicine, rehabilitation nursing and public health policy, health management to actively carry out the diagnosis and treatment of the elderly related diseases difficult critical illness, demonstration and promotion of appropriate and effective high-level diagnosis and treatment technology; We will carry out teaching and training of high-level geriatric medical personnel, and train backbone clinical technicians and academic leaders. Undertake national geriatric medicine clinical translational research, in view of the elderly health has a major impact on disease organizations to carry out the relevant scientific research, in a timely manner to translate into clinical application and scientific research achievements at home and abroad for the next "five-year plan", and establish a medical, nursing, scientific research, teaching, prevention, management and policy making of the "seven" function of the major disease prevention and health management at the core of the organization.

Strengthening cooperation between public and private hospitals is the focus of improving medical efficiency. In order to alleviate the "three difficulties" of patients in public hospitals, such as difficulty in seeing a doctor, difficulty in hospitalization and difficulty in operation, public hospitals and private hospitals can cooperate and learn from each other. The cooperation between public hospitals and private hospitals includes public hospitals exporting management and technology to private

hospitals, public hospitals borrowing beds from private hospitals, public hospitals hosting private hospitals, private hospitals or private capital participating in public hospitals. The cooperative hospital carries out standardized management and operation according to the standards of public hospitals to provide medical services for ordinary patients. At the same time, public hospitals' technical advantages, talent advantages and management advantages are utilized to enhance the popularity and credibility of private hospitals, improve the management and technical level of private hospitals, and create a win-win situation for public hospitals, private hospitals, patients and the society. It is suggested to set up a hospital management group with large public hospitals as the leading role to promote the development of private hospitals or small hospitals and at the same time make rational use of resources.

Finally, according to the model in question 2, we can predict the key disease prevention of each province in the next few years, and the corresponding province can take emergency measures on the medical problems that should be solved most, strengthen the prevention and treatment of related diseases, establish an appropriate medical insurance system, and the corresponding hospital should also make preparations in advance. I believe that with the hard work of all medical staff, China's medical environment will be better and better.

This is for your reference only.

5 Strengths and Weaknesses

5.1 Strengths

1. The gray prediction value improved by Markov model has higher fitting degree with the actual value, and the volatility is consistent, so the prediction effect is better
2. the queue model of more service system based on the analysis of queue Doctor-patient system with the method of queueing theory.

5.2 Weaknesses

1. The GM(1,1) prediction model can only make short-term prediction, and is not suitable for long-term prediction.
2. Effective coefficient of restitution can not be calculated accurately. This affects the accuracy of the result of the model.

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```

1  x = [970279 1259308 1127571 1163959 1169540 1076938 991350 953275 951508
2
3  %?????GM(1,1)?
4  sizexd2 = size(x,2);
5  %????
6
7  k=0;
8  for y1=x
9  k=k+1;
10 if k>1
11 x1(k)=x1(k-1)+x(k);
12 %???
13 z1(k-1)=-0.5*(x1(k)+x1(k-1));
14 %z1???1????B
15 yn1(k-1)=x(k);
16 else
17 x1(k)=x(k);
18 end
19 end
20 %x1, z1, k, yn1
21
22 sizez1=size(z1,2);
23 %size(yn1);
24 z2 = z1';
25 z3 = ones(1,sizez1)';
26
27 YN = yn1'; %
28 %YN
29
30 B=[z2 z3];
31 au0=inv(B'*B)*B'*YN;
32 au = au0';
33 %B, au0, au
34
35 afor = au(1);
36 ufor = au(2);
37 ua = au(2)./au(1);
38 %afor, ufor, ua
39 %????? a u ? u/a??
40
41 constant1 = x(1)-ua;

```

```

42  afor1 = -afor;
43  x1t1 = 'x1(t+1)';
44  estr = 'exp';
45  tstr = 't';
46  leftbra = '(';
47  rightbra = ')';
48  %constant1, afor1, x1t1, estr, tstr, leftbra, rightbra
49
50  strcat(x1t1, '=', num2str(constant1), estr, leftbra, num2str(afor1), tstr, rightbra);
51  %????????
52
53  %*****
54  %????
55
56  k2 = 0;
57  for y2 = x1
58  k2 = k2 + 1;
59  if k2 > k
60  else
61  ze1(k2) = exp(-(k2-1)*afor);
62  end
63  end
64  %ze1
65
66  sizeze1 = size(ze1, 2);
67  z4 = ones(1, sizeze1)';
68  G=[ze1' z4];
69  X1 = x1';
70  au20=inv(G'*G)*G'*X1;
71  au2 = au20';
72  %z4, X1, G, au20
73
74  Aval = au2(1);
75  Bval = au2(2);
76  %Aval, Bval
77  %????? A,B??
78
79  strcat(x1t1, '=', num2str(Aval), estr, leftbra, num2str(afor1), tstr, rightbra, ', ');
80  %????????
81
82  nfinal = sized2-1 + 3;

```

```

83 %????????5    ?????????????
84
85 %nfinal = size xd2 - 1 + 1;
86 %?????? 1
87
88 for k3=1:nfinal
89 x3fcast(k3) = constant1*exp(afor1*k3)+ua;
90 end
91 %x3fcast
92 %????????
93
94 for k31=nfinal:-1:0
95 if k31>1
96 x31fcast(k31+1) = x3fcast(k31)-x3fcast(k31-1);
97 else
98 if k31>0
99 x31fcast(k31+1) = x3fcast(k31)-x(1);
100 else
101 x31fcast(k31+1) = x(1);
102 end
103 end
104
105 end
106 x31fcast
107 %????????
108
109
110 for k4=1:nfinal
111 x4fcast(k4) = Aval*exp(afor1*k4)+Bval;
112 end
113 %x4fcast
114
115 for k41=nfinal:-1:0
116 if k41>1
117 x41fcast(k41+1) = x4fcast(k41)-x4fcast(k41-1);
118 else
119 if k41>0
120 x41fcast(k41+1) = x4fcast(k41)-x(1);
121 else
122 x41fcast(k41+1) = x(1);
123 end

```



```

124 end
125
126 end
127 x41fcast ,x
128 %????????
129
130 %***???p C*****////////////////////////////////////
131 k5 = 0;
132 for y5 = x
133 k5 = k5 + 1;
134 if k5 > sized2
135 else
136 err1(k5) = x(k5) - x41fcast(k5);
137 end
138 end
139 %err1
140 %????
141
142
143 xavg = mean(x);
144 %xavg
145 %x???
146
147 err1avg = mean(err1);
148 %err1avg
149 %err1???
150
151 k5 = 0;
152 sltotal = 0 ;
153 for y5 = x
154 k5 = k5 + 1;
155 if k5 > sized2
156 else
157 sltotal = sltotal + (x(k5) - xavg)^2;
158 end
159 end
160 slsuquare = sltotal ./ sized2;
161 slsqr = sqrt(slsuquare);
162 %slsuquare ,slsqr
163 %slsuquare ???x??? slsqr ?x?????S1
164

```

```

165 k5 = 0;
166 s2total = 0 ;
167 for y5 = x
168 k5 = k5 + 1;
169 if k5 > sized2
170 else
171 s2total = s2total + (err1(k5) - err1avg)^2;
172 end
173 end
174 s2suquare = s2total ./ sized2;
175 %s2suquare      ???err1???S2
176
177 Cval = sqrt(s2suquare ./ s1suquare);
178 %nnn = 0.6745 * s1sqrt
179 %Cval C???
180
181 k5 = 0;
182 pnum = 0 ;
183 for y5 = x
184 k5 = k5 + 1;
185 if abs( err1(k5) - err1avg ) < 0.6745 * s1sqrt
186 pnum = pnum + 1;
187 %ppp = abs( err1(k5) - err1avg )
188 else
189 end
190 end
191 pval = pnum ./ sized2;
192
193 %p???
194
195 %arr1 = x41fcast(1:6)

```

```

1 from example.common import Faker
2 from pyecharts import options as opts
3 from pyecharts.charts import Bar, Funnel
4 from pyecharts.globals import ThemeType
5 import pandas as pd
6 import numpy as np
7 import random
8 import matplotlib
9 from matplotlib import pyplot as plt

```

```

10 from matplotlib import font_manager
11 my_font = font_manager.FontProperties(fname="C:\Windows\Fonts\msyh.ttc")#
12 from sklearn.metrics import explained_variance_score, mean_squared_error
13
14
15 def funnel_label_inside() -> Funnel:
16     c = (
17     Funnel()
18     .add(
19     "",
20     [list(z) for z in zip(["heart disease", "Respiratory disease", "Cerebrovasc
21     label_opts=opts.LabelOpts(position="inside"),
22     )
23     .set_global_opts(title_opts=opts.TitleOpts(title=""))
24     )
25     return c
26
27 # funnel_label_inside().render()
28 # plt.figure(figsize=(15, 8), dpi=80)
29 x = [2009, 2010, 2011, 2012, 2013, 2014, 2015, 2016, 2017, 2018, 2019, 2020, 2021]
30 # ????????,
31 plt.plot(x, [128.8231, 129.1872, 132.0385, 131.6355,
32 149.45, 153.454, 155.6778
33 ], label='heart disease') # ??
34 # plt.plot(x, [54916, 67490.08141, 78063.99174, 81181.69584, 1
35 # ], label='????/?') # ??
36 # plt.plot(x, [47991, 48952.06066, 43076.23039, 40811.81632,
37 # ], label='????/?', color="red") # ???
38
39
40 # ???
41 _xtick_labels = ["{}".format(int(i)) for i in x]
42 plt.xticks(x, _xtick_labels, fontproperties=my_font)
43 # plt.yticks(range(0, 9))
44
45 # ???
46 plt.grid(alpha=0.3, linestyle="—") # alpha???? 0-1
47 plt.title("Focus on diseases", fontproperties=my_font)
48 plt.xlabel("year", fontproperties=my_font)
49 plt.ylabel("sick/thousand", fontproperties=my_font)
50 # ???

```

```
51 plt.legend(prop=my_font, loc=0)
52 plt.show()

1 # import matplotlib.pyplot as plt
2 import pandas as pd
3 import numpy as np
4
5 if __name__ == "__main__":
6     filename = r'data.xlsx'
7     data = pd.read_excel(filename)
8     y = data['???'].values[0:13]
9     predict = data["?????"].values[0:13]
10
11     exc = y-predict
12     # print(exc)
13     lamada = max(exc)/3
14     print(-3*lamada,-lamada,lamada,3*lamada)
15
16     lable = [0]*len(exc)
17     for i in range(len(exc)):
18         if exc[i]>=-3*lamada and exc[i]<-lamada:
19             lable[i]=1
20         elif exc[i]<=lamada and exc[i]>=-lamada:
21             lable[i]=2
22         else:
23             lable[i]=3
24
25     print("???????" + str(lable))
26     e1=0
27     e2=0
28     e3=0
29     for i in lable:
30         if i==1:
31             e1+=1
32         elif i==2:
33             e2 += 1
34         else:
35             e3+=1
36     P = np.array([e1/len(lable),e2/len(lable),e3/len(lable)])
37     print("???????P?" + str(P))
38     E = np.array([-2*lamada,0,2*lamada])
```

```
39
40 print("??????E?" + str(E))
41
42 diyihang_fenmu = 0
43 diyihang_fenzi = np.array([0]*3)
44
45 dierhang_fenmu = 0
46 dierhang_fenzi = np.array([0]*3)
47
48 disanhang_fenmu = 0
49 disanhang_fenzi = np.array([0]*3)
50 for i in range(len(lable)-1):
51     #?????
52     if lable[i] == 1:
53         diyihang_fenmu += 1
54     elif lable[i] == 2:
55         dierhang_fenmu += 1
56     else:
57         disanhang_fenmu += 1
58
59     #?????
60     if lable[i] == 1 and lable[i+1] == 1:
61         diyihang_fenzi[0] += 1
62     if lable[i] == 1 and lable[i+1] == 2:
63         diyihang_fenzi[1] += 1
64     if lable[i] == 1 and lable[i+1] == 3:
65         diyihang_fenzi[2] += 1
66
67     if lable[i] == 2 and lable[i+1] == 1:
68         dierhang_fenzi[0] += 1
69     if lable[i] == 2 and lable[i+1] == 2:
70         dierhang_fenzi[1] += 1
71     if lable[i] == 2 and lable[i+1] == 3:
72         dierhang_fenzi[2] += 1
73
74     if lable[i] == 3 and lable[i+1] == 1:
75         disanhang_fenzi[0] += 1
76     if lable[i] == 3 and lable[i+1] == 2:
77         disanhang_fenzi[1] += 1
78     if lable[i] == 3 and lable[i+1] == 3:
79         disanhang_fenzi[2] += 1
```

```

80
81 P0 = np.array([diyihang_fenzi/diyihang_fenmu, dierhang_fenzi/dierhang_fenmu])
82 print("???P0:" + str(P0))
83
84 pre2017 = np.dot(P, E)
85 # print(pre2017)
86 print(2239.50918495696 - pre2017)
87 #
88 print(2150.39689699288 - np.dot(np.dot(P, P0), E))
89
90 print(2064.83047520321 - np.dot(np.dot(np.dot(P, P0), P0), E))

1 def Identification_Algorithm(x):
2 B = np.array([[1]*2]*10)
3 tmp = np.cumsum(x)
4
5 print("x len:", len(x))
6 print("tmp len:", len(tmp))
7 for i in range(len(x)-1):
8 B[i][0] = ( tmp[i] + tmp[i+1] ) * (-1.0) / 2
9 print(i)
10 Y = np.transpose(x[1:])
11 BT = np.transpose(B)
12 a = np.linalg.inv(np.dot(BT, B))
13 a = np.dot(a, BT)
14 a = np.dot(a, Y)
15 a = np.transpose(a)
16 return a;
17
18
19 def GM_Model(X0, a, tmp):          #GM(1,1)
20 A = np.ones(len(X0))
21 for i in range(len(A)):
22 A[i] = a[1]/a[0] + (X0[0] - a[1]/a[0]) * np.exp(a[0]*(tmp[i]-1)*(-1))
23 print('GM(1,1)???:\nX(k) = ', X0[0] - a[1]/a[0], 'exp( ', -a[0], ' (k-1)) ', a[1]/a[0])
24 XK = Series(A, index=pd.period_range('2003', '2013', freq = 'A-DEC'))
25 print('GM(1,1)?????: ')
26 print(XK)
27 return XK;
28
29

```

```

30 def Return(XK):
31     tmp = np.ones(len(XK))
32     for i in range(len(XK)):
33         if i == 0:
34             tmp[i] = XK[i]
35         else:
36             tmp[i] = XK[i] - XK[i-1]
37     X_Return = Series(tmp, index=pd.period_range('2003', '2013', freq = 'A-DEC'))
38     print('?????:\n')
39     print(X_Return)
40     return X_Return
41
42
43 if __name__ == '__main__':
44     # ???????
45     filename = r'data.xlsx'
46     data = pd.read_excel(filename, index_col=u'??')
47     date = pd.period_range('2003', '2013', freq = 'A-DEC')
48     tmp = np.array([1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11])
49     data = np.array([129227, 129988, 130756, 131448, 132129, 132802, 133450,
50     data = data ** (1 / 10)
51     X0 = Series(data, index = date)
52     X0_copy = Series(data, index=tmp)
53     print('?????:\n')
54     print(X0)
55
56     # ?????????????
57     X1 = np.cumsum(X0)
58     print('???????:')
59     print(X1)
60
61     # ???
62     a = Identification_Algorithm(data)
63     print('a???:')
64     print(a)
65
66     # GM(1,1)??
67     XK = GM_Model(X0, a, tmp)
68
69     # ???
70     X_Return = Return(XK)

```

```

71
72 # ??????????
73 X_Compare1 = np.ones(len(X0))
74 X_Compare2 = np.ones(len(X0))
75 for i in range(len(data)):
76     X_Compare1[i] = data[i]-X_Return[i]
77     X_Compare2[i] = X_Compare1[i]/data[i]*100
78     Compare = { 'GM':XK, '1?AGO':np.cumsum(data), 'Returnvalue':X_Return, 'Realit',
79 X_Compare = DataFrame(Compare,index=date)
80 print ( '?????????' )
81 print (X_Compare)
82
83 # ???
84 error_square = np.dot(X_Compare,np.transpose(X_Compare))
85 # ?????
86 error_avg = np.mean(error_square)
87 # ?????
88 S = 0
89 # X0????
90 for i in range(1,len(X0)-1,1):
91     S += X0[i]-X0[0]+(XK[-1]-XK[0])/2
92     S = np.abs(S)
93
94 SK = 0
95 # XK????
96 for i in range(1,len(XK)-1,1):
97     SK += XK[i]-XK[0]+(XK[-1]-XK[0])/2
98     SK = np.abs(SK)
99
100 S_Sub = 0
101 # /S-SK/b
102 for i in range(1, len(XK)-1, 1):
103     S_Sub += X0[i]-X0[0]-(XK[i]-XK[0])+((X0[-1]-X0[0])-(XK[i]-XK[0]))/2
104 S_Sub = np.abs(S_Sub)
105
106 T = (1+S+SK)/(1+S+SK+S_Sub)
107
108 if T >= 0.9:
109     print ( '?????' )
110     print ( '???GM(1,1)??\nX(k) = ',X0[0]-a[1]/a[0], 'exp( ', -a[0], ' (k-1)) ',

```



```

107 elif T >= 0.8:
108     print ( '?????' )
109     print ( '???GM(1,1)??\nX(k) = ',X0[0]-a[1]/a[0] , 'exp( ', -a[0] , ' (k-1)) ',
110     elif T >= 0.7:
111     print ( '?????' )
112     print ( '???GM(1,1)??\nX(k) = ',X0[0]-a[1]/a[0] , 'exp( ', -a[0] , ' (k-1)) ',
113     elif T >= 0.6:
114     print ( '?????' )
115     print ( '?????GM(1,1)??\nX(k) = ',X0[0]-a[1]/a[0] , 'exp( ', -a[0] , ' (k-1)) ',
116
117     print( '??T?', T)
118
119 X2014 = Series(np.array([136782]) ** (1 / 10), index=pd.period_range('201
120 X_Return = X_Return.append(X2014)
121 print (X_Return)
122
123 B = pd.DataFrame([X0, X_Return], index=['X0', 'X_Return'])
124 B = np.transpose(B)
125 B.plot()
126 plt.plot(B)
127 plt.show()

1 import pandas as pd
2 import numpy as np
3 import random
4 import matplotlib
5 from matplotlib import pyplot as plt
6 from matplotlib import font_manager
7 my_font = font_manager.FontProperties(fname="C:\Windows\Fonts\msyh.ttc")#
8 from sklearn.metrics import explained_variance_score,mean_squared_error
9 def evalmape(y_true, y_pred):
10     n = len(y_true)
11     mape = sum(np.abs((y_true - y_pred)/y_true))/n
12     return mape
13
14 def nse(y_true, y_pred):
15     return 1 - (sum((y_true-y_pred)**2)/sum((y_true-np.mean(y_true))**2))
16
17 if __name__ == "__main__":
18     filename = r'data.xlsx'
19     data = pd.read_excel(filename)

```

```

20 x = [2009,2010,2011,2012,2013,2014,2015,2016,2017,2018,2019,2020,2021]
21 y1 = [11307,11894,12288 ,12714 ,13161, 13755 ,14386 ,15203, 15831 ,
22 y2 = [11307,11964,12588 ,12311 ,13645, 13199 ,14766 ,15320, 16301 ,
23 y3 =[11307,11794,12388 ,12611 ,13245, 13599 ,14476 ,15340, 16101 ,
24 # ???????
25 plt.figure(figsize=(15, 8), dpi=80)
26
27 # ?????????
28 plt.plot(x[0:10],y1 , label='True ',linestyle="--") # ??
29 plt.plot(x, y2, label='GM(1,1) ', linestyle=":") # ???
30 plt.plot(x, y3, label='Markov-GM(1,1) ',color="red") # ???
31
32 # ???
33 _xtick_labels = [ "{}".format(int(i)) for i in x]
34 plt.xticks(x, _xtick_labels , fontproperties=my_font)
35 # plt.yticks(range(0, 9))
36
37 # ???
38 plt.grid(alpha=0.3, linestyle="--") # alpha???? 0-1
39 plt.title("Comparison curve of prediction value of the number of the elde
40 plt.xlabel("T / year", fontproperties=my_font)
41 plt.ylabel("Number / million", fontproperties=my_font)
42 # ???
43 plt.legend(prop=my_font, loc=0)
44
45 # ??
46 plt.show()
47
48 mse_predict = mean_squared_error(y1, y2[0:10])
49 print(np.sqrt(mse_predict))
50 MAPE = evalmape(pd.DataFrame(y1), pd.DataFrame(y2)[0:10])
51 print(MAPE)
52 # NSE = nse(pd.DataFrame(y1), pd.DataFrame(y2)[0:10])
53 # print(NSE)

```