Original Article

# Mixture Distribution for Modeling Turkey Covid-19 Case

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Abstract - The COVID-19 pandemic hit the world in early 2020 and caused many deaths in various countries, one of which was Turkey. In March 2020, the first case of COVID-19 was confirmed by the Turkish Ministry of Health. In May 2021, the rise of cases continued to occur until it reached 5.2 million cases with a death toll of 47,271 people. In this research, a model for Turkey's COVID-19 positive cases had built using a mixed distribution, namely log-normal, gamma and Weibull. Parameter estimation for each distribution was carried out using the maximum likelihood method. Then by using the Bayesian Information Criterion (BIC) and Akaike Information Criterion (AIC) tests, it was found that the 3-component gamma distribution is the best distribution to describe the pattern of spikes in positive cases of COVID-19 in Turkey.

Keywords - Covid-19, Gamma distribution, Log-normal distribution, Mixed distribution, Turkey, Weibull distribution.

#### **1. Introduction**

At the beginning of 2020, Sars-Cov-2 or known as covid-19 was shocking the world. On March 11, 2020, the World Health Organization (WHO) announced covid-19 virus as pandemic. A month later, in the middle of the month April 2020, WHO reported that 311,847 death from 4,619,477 cases in the world. Until the end of May 2021, there have been 169.5 million cases with 3,530,582 deathl [16].

This virus also spread in Turkey when Turkish Ministry of Health reported the first case on March 11, 2020. Until May 2021, Turkey's Covid-19 cases have reached 5.2 million with 47,271 death cases. This situation forced Turkey to lockdown or restrictions on people's movements in April 2021.

One of the important parameters to control covid-19 is the number of positive patients. Some researchers have been modelled of the number of covid-19 positive patients using simple distributions. Qi fang [12] used a negative binomial distribution to modelling the transmission of the covid-19 virus in Shenzhen, China. But, somehow it's difficult to use that simple technique since the data may also contain more than one distribution. Therefore, we need a mixture distribution to apply in covid-19 cases.

Kremer et al [10]., have modelled the spread of secondary cases of covid-19 using the Poisson mixture distribution in Hong Kong, India and Rwanda. The study stated that mixture of Poisson and log-normal distribution was the best distribution. Then in August 2020, Polymenis [11] used an exponential mixture distribution to measure the risk level of covid-19 in Italy. In this study, we will model the first three months of positive covid-19 cases in Turkey by using the weibull, gamma log-normal mixture distribution.

### 2. The Data Set

We collect Turkey daily positive cases of Covid-19 data for 3 months (March to May 2020) that present in Figure 1 and Figure 2. The first case occurred on March 12, 2020 and the death cases began a week after the first positive case. On March 19, 2020, the number of positive cases reached 93 cases and fluctuated until the end of the March 2020. In April, it reached 5138 cases and forced Turkish government to implemented the large-social restrictions or known as lockdown.



Fig. 2 The Histogram of daily new cases covid 19 of Turkey

#### **3. Model Selection**

Goodness of fit test is used to compare several different models distribution with the aim of determining the most representative model for the data. In this study, we use Akaike Information Criterion (AIC) and Bayesian test Information Criterion (BIC). The AIC defined as [1]

$$AIC = -2\ln(L) + 2a,$$

where L is likelihood function of model and a is the number of parameter.

The BIC test is derived from the Bayesian framework as an estimate of the Bayes factor for two models and defined as follow [1]

$$BIC = -2\ln(L) + s\ln(n),$$

where *n* is number of sample.

# 4. Materials and Methods

#### 4.1. Model Distribution

In this study, we have 9 distribution models to approximate Turkey daily covid-19 positive cases, namely: unmixed distributions (log-normal (L), gamma (G) and Weibull (W)); two-component mixture distribution models (2-component log-normal (ML2), 2-component gamma (MG2) and 2-component Weibull (MW2)) and three-components mixture distributions (3-component log-normal (ML3), 3- component gamma (MG3) and 3-component Weibull (MW3)).

Table 1. Probability density function of models							
Distributions	Density Function						
L	$f(x;\mu,\sigma) = \frac{1}{x\sqrt{2\pi(\sigma^2)}}e^{-\frac{(\ln x-\mu)^2}{2\sigma^2}}$						
G	$f(x;\alpha,\beta) = \frac{\beta}{\Gamma(\alpha)} x^{\alpha-1} e^{-\beta x}$						
W	$f(x;k,c) = \frac{k}{c} \left(\frac{x}{c}\right)^{k-1} e^{-\left(\frac{x}{c}\right)^{k}}$						
ML2	$w_1 f(x; \mu_1, \sigma_1) + w_2 f(x; \mu_2, \sigma_2),  w_1 + w_2 = 1$						
MG2	$w_1 f(x; \alpha_1, \beta_1) + w_2 f(x; \alpha_2, \beta_2),  w_1 + w_2 = 1$						
MW2	$w_1 f(x; k_1, c_1) + w_2 f(x; k_2, c_2),  w_1 + w_2 = 1$						
ML3	$w_1 f(x; \mu_1, \sigma_1) + w_2 f(x; \mu_2, \sigma_2) + w_3 f(x; \mu_3, \sigma_3),  w_1 + w_2 + w_3 = 1$						
MG3	$w_1 f(x; \alpha_1, \beta_1) + w_2 f(x; \alpha_2, \beta_2) + w_3 f(x; \alpha_3, \beta_3),  w_1 + w_2 + w_3 = 1$						
MW3	$w_1 f(x; k_1, c_1) + w_2 f(x; k_2, c_2) + w_3 f(x; k_3, c_3),  w_1 + w_2 + w_3 = 1$						

#### 4.2. Maximum Likelihood

The maximum likelihood method is a method to estimate parameter of a distributor by using the maximum probability. First, make likelihood function for joint probability of density function  $f(x_i|a_1, a_2, ..., a_k)$  that defined as [9]

$$L(a|x) = \prod_{i=1}^{n} f(x_i | a_1, a_2, \dots, a_k).$$

Then make log-likehood function that defined as [4]

$$l(a|x) = \ln L(a|x).$$

After that, using partial derivation to maximum log-likehood function

#### 5. Results and Discussion

By using R application, we get the value of parameters for each model that present in Table 2.

	Table 2. Computed parameters of each model										
	L	G	W	ML2	MG2	MW2	ML3	MG3	<i>MW</i> 3		
μ	7.101	—	—	—	—	—	-	—	—		
σ	1.851	—	—	—	—	—		—	—		
α	_	0.886	_	_	_	_		—	_		
β	_	0.00037	—	—	—	—		—	—		
k	—	—	1.069	—	—	—	-	—	—		
С	_	—	2391.22	—	—	—		—	—		
$\mu_1$	—	—	—	4.602	—	—	4.55	—	—		
$\sigma_1$	—	—	—	2.241	—	—	2.217	—	—		
$\mu_2$	_	_	_	7.92	_	_	7.73	_	_		
$\sigma_2$	_	_	_	0.379	_	_	0.281	_	—		

Table 2. Computed parameters of each model

$\mu_3$	_	—	—	—	—	—	8.382	—	—
$\sigma_3$	-	—	—	—	—	—	0.087	—	—
$\alpha_1$		—	—	—	0.568	—	—	0.568	—
$\beta_1$		—	—	—	0.0026	—	—	0.0026	—
$\alpha_2$		—	—	—	7.085	—	—	13.602	—
$\beta_2$		—	—	—	0.0023	—	—	0.0058	—
α3		—	—	—	—	—	—	120	—
$\beta_3$		—	—	—	—	—	—	0.027	—
$k_1$		—	—	—	—	0.635	—	—	0.654
$c_1$		—	—	—	—	191.402	—	—	195.09
$k_2$		—	—	—	—	2.872	—	—	4.402
<i>c</i> <sub>2</sub>		—	—	—	—	3318.16	—	—	2498.894
$k_3$	—	—	_	—	—	—	—	—	10.276
<i>C</i> <sub>3</sub>	_	—	—	—	—	—	—	—	4495.856
$w_1$	_	_	_	0.2487	0.2235	0.2234	0.246	0.225	0.229
$w_2$	_	_	_	0.7513	0.77654	0.7766	0.530	0.539	0.503
<i>W</i> <sub>3</sub>	_	_	_	_	_	_	0.224	0.236	0.268

The probability density function for each research model can be formed based on Table 1 and Table 2. Then, we use the graphical technique namely probability density function (pdf) plot to observe the model distributions.



Fig. 3 Pdf plot for comparisons Predicted and observed daily new cases covid 19 in Turkey for L, G and W distribution

In Figure 3, it can be seen that the Log-normal distribution is unmixed (model L, G and W) is not fit to approximating the Turkey daily covid-19 positive cases histogram. The unmixed model only has 1 wave, while there are 3 waves in observation data. Further on, we also graph the pdf plot for two-component mixture distribution that present in Figure 4.



Fig. 4 Pdf plot for comparisons Predicted and observed daily new cases covid 19 in Turkey for ML2, MG2 and MW2 distribution



Figure 4 describe that the two-component mixture distribution are better than unmixed distribution since has 2 waves in pdf plot. But these models are not fit to for the data. Finally, we have 3-component mixture distribution plot in Figure 5.

Fig. 5 Pdf plot for comparisons Predicted and observed daily new cases covid 19 in Turkey for ML3, MG3 and MW3 distribution

It is clearly seen that the 3-component mixture distributions (ML3, MG3 and MW3) approximate Turkey covid-19 cases very well. The goodness fit test was carried out by using AIC) and BIC to select the best model in Table 3.

Table 5. The Goodness and fit test result of the daily new cases covid 19 in Turkey										
	L	G	W	ML2	MG2	MW2	ML3	MG3	MW3	
AIC	1082.1	1037.3	1037.6	1063.8	1029.8	1003.4	999.7	996.3	998.8	
BIC	1086.2	1041.5	1041.7	1074.2	1040.2	1013.8	1016.3	1012.9	1015.4	
Log(L)	-539.03	-516.7	-516.8	-526.9	-509.9	-496.7	-491.8	-490.2	-491.4	

Table 3. The Goodness and fit test result of the daily new cases covid 19 in Turkey

## 6. Conclusion

Based on the discussion, 3-component log-normal mixture distribution, 3-component gamma mixture distribution and 3-component Weibull mixture distribution can be used for illustrates Turkey covid-19. From the goodness fit results, we obtain that the 3-component gamma distribution model is the best model.

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