

## Bifurcation Analysis for Some Fractional SIRS Models

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### Abstract

In this paper , some fractional-order epidemic models are investigated. In particular , a fractional SIRS epidemic model is considered to describe the transmission dynamics of infectious diseases incorporating memory effects through fractional derivatives . The existence and stability of equilibrium points of the proposed model are analyzed .Moreover, analytical solution are obtained for some special cases , highlighting the impact of fractional-order dynamics on the behavior of the epidemic system.

**Keywords:** *Dynamical Systems, Fractional Epidemic Models, Stability, Mittag-Leffler Function.*

### Introduction

Fractional-order differential equations have received considerable attention in recent years due to their ability to model memory and hereditary properties inherent in many real-world phenomena , including epidemiological processes In contrast to classical integer-order models , fractional epidemic models provide a more accurate description of disease transmission dynamics .

In the following sections we shall study some fractional epidemic models.

### The Fractional SIRS Epidemic Model

Let  $S(t)$  .  $I(t)$  . and  $R(t)$  denote the numbers of susceptible infected , and recovered individuals at time  $t$  , respectively . The fractional SIRS model is given by :

$$\begin{cases} \frac{d^\alpha S(t)}{dt^\alpha} = b - cS(t) - f(I)S^* + \delta R(t), \\ \frac{d^\alpha I(t)}{dt^\alpha} = f(I)S^* - (c + \mu)I(t), \\ \frac{d^\alpha R(t)}{dt^\alpha} = \mu I(t) - (c + \delta)R(t), \end{cases} \quad (2 \cdot 1)$$

$$S^*(t) = \lim_{t \rightarrow \infty} N(t) - I(t) - R(t),$$

, see [1-5]. Where  $0 < \alpha \leq 1$  .  $b > 0$

The total population size is defined as :

$$N(t) = S(t) + I(t) + R(t)$$

### Description of Parameters

The parameters appearing in system (1.1) are defined as follows :

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$b > 0$  : the recruitment rate of the population

$c > 0$  : the natural death rate

$\mu > 0$  : the recovery rate of infected individuals

$\delta > 0$  : the rate at which recovered individuals lose immunity and return to the susceptible class

$f(I)$  : the incidence rate function describing the interaction between susceptible and infected individuals .

The incidence rate is assumed to be of the nonlinear form :

$$f(I)S = \frac{BI}{I + VI^k} S$$

Where  $B > 0$  .  $V > 0$  and  $k > 0$  are constants

Let  $S(t)$  .  $I(t)$  . and  $R(t)$  denote the numbers of susceptible , infective, and removed individuals at time  $t$  , respectively .

The parameter  $\beta$  represents the average number of new infections generated per unit time in a fully susceptible population .

For convenience , we introduce the following notations :

$$\lambda = \frac{b}{a} \quad \Lambda_0 = \frac{\beta}{C + \delta} \Lambda \quad \cdot \quad R_0 = \frac{\Lambda_0}{\delta}$$

Where all parameters are assumed to be positive constants.

This work is organized as follows . In Section 1 , we investigate the fractional-order system given by equation (1.1) . In Section 2 , we analyze several epidemic models , highlighting their qualitative behavior and mathematical properties .

#### 4. Population Dynamics

By summing the equation of system (1.1) , we obtain the fractional differential equation governing the total population :

$$\frac{d^\alpha N(t)}{dt^\alpha} = b - c N(t) \quad (4 \cdot 1)$$

This equation implies that the total population approaches the steady-state value  $N^* = \frac{b}{c}$  as  $t \rightarrow \infty$  . ensuring the biological feasibility of the model .

Where  $N(t) = S(t) + I(t) + R(t)$

Using the result in [6] , we can obtain the solution of (4.1) in the form:

$$N(t) = \int_0^\infty \xi_\alpha(\theta) e^{-c(t-\theta)^\alpha} d\theta + N(0) + \alpha \int_0^t \int_0^\infty \theta b \xi_\alpha(\theta) (t-y)^{\alpha-1} e^{-c(t-y)^\alpha} d\theta dy$$

Consequently

$$N(t) = \frac{b}{c} + \left( N(0) - \frac{b}{c} \right) E_{\alpha}(-ct^{\alpha}) . t \geq 0 \quad (4.2)$$

Type equation here.

Where

$$E_{\alpha}(t) = \sum_{j=0}^{\infty} \frac{t^j}{\Gamma(\alpha j + 1)} . \quad \alpha > 0$$

Is the Mittag-Leffler function

So

$$\lim_{t \rightarrow \infty} N(t) = \frac{b}{c}$$

Now we can write

$$\begin{cases} \frac{d^{\alpha}I(t)}{dt^{\alpha}} = f(t)[\beta N(t) - R(t) - I(t)] - (c + \mu)I(t). \\ \frac{d^{\alpha}R(t)}{dt^{\alpha}} = \mu I(t) - (c + \delta)R(t) \end{cases} \quad (4.3)$$

System (1.5) can be written in the form

$$\begin{cases} \frac{d^{\alpha}x(t)}{dt^{\alpha}} = \alpha(1 + \rho x^{k-1})(N^*(t) - x - y)x(t) - \gamma x \\ \frac{d^{\alpha}y(t)}{dt^{\alpha}} = \gamma x - y \end{cases} \quad (4.4)$$

Where the dimensionless variables and parameters are defined by

$$I = \frac{c + \delta}{\beta} x . \quad R = \frac{C + \delta}{\beta} y . \quad t = \frac{1}{c + \delta} \tau .$$

$$P = \left( \frac{C + \delta}{\beta} \right)^{k-1} . \quad N^* = \frac{\beta}{c + \delta} N .$$

$$\gamma = \frac{C + M}{C + \delta} . \quad \eta = \frac{\mu}{C + \delta}$$

According to the results in [1 – 3]

We can deduce that (0.0) is a unique disease free equilibrium for system (4.4)

The disease free equilibrium (0.0) is a stable node if

$$R_0 < 1 . R_0 > 1$$

If  $R_0 = 1$  . then (0.0) is a saddle node .

2. Some fractional modified SIR models

Consider the following epidemic fractional epidemic model

$$\frac{d^\alpha S}{dt^\alpha} = -bSI - aS$$

$$\frac{d^\alpha I}{dt^\alpha} = bSI - aI$$

$$\frac{d^\alpha R}{dt^\alpha} = a(I + S)$$

Let us suppose that the initial conditions

$$S(0) < 1 \text{ . } I(0) > 0 \text{ .}$$

$$R(0) = 0 \text{ and}$$

$$S(0) + I(0) = 1$$

We have

$$\frac{d^\alpha N(t)}{dt^\alpha} = 0$$

Thus

$$N(t) = S(0) + I(0) + R(0) = 1$$

$$\frac{d^\alpha R}{dt^\alpha} = a[1 - R]$$

$$= a - aR$$

Using the results in [6] , we can get

$$R(t) = a \int_0^t \int_0^\infty \alpha \theta (t - \eta)^{\alpha-1} e^{-a(t-\eta)^\alpha \theta} \xi_\alpha d\theta dy$$

$$= 1 - E_\alpha(-at^\alpha) .$$

Where

$$E_\alpha(-at^\alpha) = \sum_{j=0}^{\infty} \frac{(-at^\alpha)^j}{\Gamma(\alpha j + 1)} \text{ . } \alpha > 0$$

Is the Mittag-Leffler function .

Thus

It is clear that

$$\lim_{t \rightarrow \infty} R(t) = 1$$

$$S(t) + I(t) = E_\alpha(-at^\alpha) .$$

It follows that

$$\lim_{t \rightarrow \infty} S(t) = \lim_{t \rightarrow \infty} I(t) = 0$$

3. The fractional SIR model

Consider

$$\begin{cases} \frac{d^\alpha S}{dt^\alpha} = -bSI \\ \frac{d^\alpha I}{dt^\alpha} = bSI - aI \\ \frac{d^\alpha R}{dt^\alpha} = aI \end{cases} \quad (3 \cdot 1)$$

$S > 0$  .  $I > 0$  and  $R \geq 0$  .

for all  $t \geq 0$  .  $0 < b < a$

$$S(0) + I(0) = 1$$

$$R(0) = 0$$

$$S(t) + I(t) + R(t) = 1$$

Thus

$$\frac{d^\alpha R(t)}{dt^\alpha} \leq a - aR(t).$$

So

$$R(t) \leq a \int_0^t \int_0^\infty \alpha \theta (t - \eta)^{\alpha-1} e^{-a(t-\eta)\alpha\theta} \xi_\alpha d\theta dy$$

$$R(t) \leq 1 - E_\alpha(at^\alpha)$$

Thus the proportion of removal  $\mathcal{R}$  starts at zero and steadily increases , eventually approaching .

As a limit

$$\lim_{t \rightarrow \infty} R(t) \leq 1$$

Notice that

$$\begin{aligned} \frac{d^\alpha I}{dt^\alpha} &= b [1 - R - I] I - aI \\ &\leq -(a - b)I \end{aligned}$$

Thus

$$I(t) \leq I(0) \int_0^\infty \xi_\alpha(\theta) e^{-(a-b)(t-\theta)\alpha} d\theta$$

Consequently

$$\lim_{t \rightarrow \infty} I(t) = 0$$

This means that the proportion  $I(t)$  of infected approaches zero as  $t$  tends to  $\infty$

We notice that

$$\frac{d^\alpha S(t)}{dt^\alpha} > -bS(t)$$

Thus

$$S(t) > S(0) \int_0^\infty \xi_\alpha(\theta) e^{-bt^\alpha} d\theta$$

This means that the proportion of susceptible  $S(t)$  starts at  $S(0)$  and steadily decreases , eventually approaching a limit

$$\lim_{t \rightarrow \infty} S(t) > 0 , \text{ see [7-12].}$$

## Conclusion

Some fractional epidemic models are studied. Analytical solutions are obtained for some special cases. The stability of solutions are obtained.

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