Modelling the spread of HIV/AIDS in Morocco

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Abstract

In this article, we propose a non-linear mathematical model with a variable size population to study the spread of HIV/AIDS in Morocco. We estimate the parameters that include this model based on real data. An estimation of the basic reproductive number R_0 and some numerical simulations that provide insights into the future of HIV/AIDS in Morocco are also presented in this study.

Keywords: Mathematical modelling, HIV/AIDS, Parameter estimation, Basic reproductive number R_0 .

1. Introduction

HIV (human immunodeficiency virus) is the virus that causes AIDS (acquired immunodeficiency syndrome). By killing or damaging cells of the body's immune system, HIV progressively destroys the body's ability to fight infections and certain cancers. People diagnosed with AIDS may get life-threatening diseases called opportunistic infections, which are caused by microbes such as viruses, bacteria, fungi, or parasites. These infections do not usually make healthy people sick. Those with HIV/AIDS are also at an increased risk of developing certain cancers and a variety of other neurological disorders.

HIV/AIDS is one of the most destructive diseases humankind has ever faced, with profound social, economic and public health consequences. It has become, for three decades, a full-blown pandemic affecting all parts of the world. As reported by the latest UNAIDS report [1], the number of people living with HIV in the world in 2010 was estimated to be 34 million, including 2.7 million new infections occurred during the same year, that is 7400 cases per day. The number of AIDS deaths occurred in 2010 was about 1.8 million, according to the same source. In Morocco, the first case of AIDS was recorded in 1986. Since then, the number of new AIDS cases has continued to rise. HIV is gaining ground and according to the official data of the Ministry of Public Health, Morocco has up to 31st December 2011, 2284 and 4169 cumulative reported cases of HIV and AIDS.

To fight HIV/AIDS, mathematical models play an important role to better understand this disease's dynamics; they provide information on the essential parameters of this dynamic, which can not be observed and predict its evolution in a given population (see for example [2-3-4-5-6]).

Following our contacts with the direction of the epidemiology and fight against diseases, within the Ministry of Public Health, we noted a lack of some national data related to the spread of HIV/AIDS in Morocco, such as the time between HIV diagnosis and the onset of AIDS, and the mean time from AIDS diagnosis to death because most of patient die outside hospitals. To overcome this problem, we propose an estimation to these parameters by fitting a mathematical model, modelling the HIV/AIDS transmission, to the statistics available in Morocco.

This paper is organized as follows: Section 2, describes a mathematical model that fits the case of HIV/AIDS. The parameter estimation and the estimated basic reproductive number R_0 are presented in section 3. In section 4, we give some numerical simulations that predict the future of this epidemic in Morocco. Finally, the conclusions are summarized in section 5.

2. The model

In this section we consider a simple model, with a variable size population, where

- t = time in years
- S(t) = number of susceptible individuals;
- I(t) = number of declared HIV-positive cases without AIDS;
- A(t) = number of reported AIDS cases.



We suppose that the AIDS patients are removed from the population since, unlike other classes (S and I), they do not have sex anymore, and have no children. In fact, their relationships with the rest of the population are insignificant. The model is given by the following nonlinear system of differential equation:

$$\begin{cases} \frac{dS}{dt} = A - \mu S - \beta S \frac{I}{N} \\ \frac{dI}{dt} = \beta S \frac{I}{N} - (\mu + \gamma)I \\ \frac{dA}{dt} = \gamma I - (\mu + \delta)A \end{cases}$$
(1)

where $S(0) \ge 0$, $I(0) \ge 0$ and $A(0) \ge 0$ are given and N(t) = S(t) + I(t) + A(t) for all t. The definitions of the above model parameters are as follow

Table 1: Parameter definition				
Parameter	er Definition			
Λ	Λ Recruitment rate			
β	β Transmission coefficient			
μ	Natural mortality rate			
γ	Rate of developing AIDS of (I) individuals			
δ	AIDS induced mortality rate			

Note that the system (1) is dissipative, that is all solutions are uniformly bounded in a proper subset $\Omega \subset \mathbb{R}^3_+$ where

$$\Omega = \left\{ \left(S, I, A \right) \in \mathbb{R}^{3}_{+} : N(t) \leq \frac{\Lambda}{\mu} \right\}$$

Also, the continuity of the right side of the system (1) and its derivatives implies that unique solutions exist. Therefore, the model is mathematically and epidemiologically well posed. It should also be noted that the basic reproductive number R_0 of (1) is given by

$$R_0 = \frac{\beta}{\mu + \gamma} \tag{2}$$

3. Parameter estimation

An important step for validating a given mathematical model is to estimate its parameters. In this section we give estimation for the parameters used in the model (1); the aim is to reconstruct the HIV/AIDS in Morocco in the past and simulate its future development. An estimation of the basic reproductive number R_0 will also be provided in this section.

In our model we fix the natural mortality rate, μ , which is supposed to be the inverse of life expectancy at birth [7]. So, according to the database of the World Bank [8], Morocco has made, for the entire period 1987-2011 (Table 2), an average life expectancy at birth of 68 years. Hence,

$$\mu = \frac{1}{68} = 0.01470588$$
 per year

Table 2	Life	expectancy	at	birth	in	Morocco
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Year	Life expectancy at birth
1987	62,47
1988	63,04
1989	63,60
1990	64,15
1991	64,68
1992	65,20
1993	65,69
1994	66,16
1995	66,61
1996	67,05
1997	67,47
1998	67,88
1999	68,28
2000	68,68
2001	69,05
2002	69,42
2003	69,77
2004	70,10
2005	70,42
2006	70,72
2007	71,02
2008	71,31
2009	71,59
2010	71,86
2011	72,82*

For the remaining parameters $(\Lambda, \beta, \gamma, \delta)$, they are estimated by the least squares method using data from the High Commission to the Plane [9] and the Ministry of Public Health [10] (Table 3). For the fitting procedure by the least squares method, we used the Levenberg-Marquardt method implemented in MATLAB [11] in the

^{*} Estimation calculated using a simple linear regression model.

built-in routine "Isqcurvefit" which is part of the optimization toolbox. The resulting parameter estimates are listed in Table 4, and the best model fit to the data is shown in Fig. 1.

Year	Population	Susceptible	VIH	AIDS
1 cai	(N)	(S)	(I)	(A)
1987	22749000	22748989	2	9
1988	23227000	23226965	21	14
1989	23703000	23702973	7	20
1990	24177000	24176932	42	26
1991	24649000	24648938	34	28
1992	25117000	25116924	46	30
1993	25582000	25581933	23	44
1994	26073000	26072899	24	77
1995	26386000	26385933	10	57
1996	26848000	26847933	1	66
1997	27310000	27309907	1	92
1998	27775000	27774863	44	93
1999	28238000	28237774	61	165
2000	28705000	28704846	42	112
2001	29170000	29169832	39	129
2002	29631000	29630786	64	150
2003	29520000	29519696	99	205
2004	29840000	29839589	140	271
2005	30172000	30171612	100	288
2006	30509000	30508581	129	290
2007	30850000	30849386	247	367
2008	31195000	31194381	203	416
2009	31543000	31542398	190	412
2010	31894000	31893231	309	460
2011	32245000	32244247	406	347

Table 4: Parameter estimation		
Parameter	Estimations	
Λ	805479	
β	0,883856382	
γ	0,654004963	
δ	0,163045919	



Fig. 1 The best-fit solution obtained by fitting the solution of the model (1) (solid line) to the real data (dotted line) in Table 3

We find that in Morocco, the mean time between HIV diagnosis and the onset of AIDS is about one year and a half; this result can be interpreted by the fact that most diagnoses occur in more advanced stages, hence the necessity to accentuate efforts of HIV screening in Morocco. Also, the results obtained show that for the Moroccan case, the mean time from AIDS diagnosis to death is nearly six years.

Using the formula (2) and the values of the parameters β , μ and γ , previously calculated, we obtain an estimate of the basic reproductive number R_0 which is

$$R_0 \approx 1.32$$

This means that the number of secondary cases which an infected person would produce in a completely susceptible Moroccan population is about 1,32, that is to say, the epidemic will grow even in Morocco. However, the extent of the disease remains lower than that recorded in other

African countries that are highly affected by HIV/AIDS such as South Africa, where R_0 is 7 [12].

4. Numerical simulation

In this section, we give some numerical simulations which predict the evolution of the disease in Morocco. First of all, we note that the system (1) admits an endemic equilibrium

$$E^{*} = \left(S^{*}, I^{*}, A^{*}\right), \text{ where}$$

$$S^{*} = \frac{1}{R_{0}}N^{*}$$

$$I^{*} = \frac{(\mu + \delta)(R_{0} - 1)}{(\mu + \delta + \gamma)R_{0}}N^{*}$$

$$A^{*} = \frac{\gamma(R_{0} - 1)}{(\mu + \delta + \gamma)R_{0}}N^{*}$$

$$N^{*} = \frac{\Lambda(\mu + \delta + \gamma)R_{0}}{\mu(\mu + \delta + \gamma)R_{0} + \delta\gamma(R_{0} - 1)}$$

The Jacobian matrix of the linearized system in the neighbourhood of E^* is

$$J\left(E^{*}\right) = \begin{pmatrix} -\mu - \alpha_{1} & -\alpha_{2} & \alpha_{3} \\ \alpha_{1} & -\alpha_{2} & -(\mu + \gamma) & -\alpha_{3} \\ 0 & \gamma & -(\mu + \delta) \end{pmatrix}$$

where

$$\alpha_{I} = \beta \frac{I^{*}}{N^{*}} \left(I - \frac{S^{*}}{N^{*}} \right)$$
$$\alpha_{2} = \beta \frac{S^{*}}{N^{*}} \left(I - \frac{I^{*}}{N^{*}} \right)$$
$$\alpha_{3} = \beta \frac{S^{*}I^{*}}{N^{*2}}$$

Numerically, using Matlab, we find

$$S^{*} = 1.3273 \times 10^{7}$$
$$I^{*} = 9.1263 \times 10^{5}$$
$$A^{*} = 3.3578 \times 10^{6}$$
$$N^{*} = 1.7544 \times 10^{7}$$

and the eigenvalues corresponding to $J(E^*)$ are as follows:

$$\lambda_{1} = -0.0561$$

$$\lambda_{2} = -0.0912 + 0.1516 i$$

$$\lambda_{3} = -0.0912 - 0.1516 i$$

We conclude that E^* is locally-asymptotically stable as the eigenvalues obtained have strictly negative real part. This result is also confirmed by Figure 2, where we observe that the population of susceptible individuals, HIV individuals and AIDS individuals converge asymptotically to the endemic equilibrium state as time increases.



Fig. 2 Evolution of the HIV/AIDS population in Morocco

Regarding the disease's evolution in Morocco during the coming years, Figure 3 shows that the number of HIV individuals will continue to increase; it will reach its maximum in 2060, which is about 2.3×10^6 cases. Then it will decrease asymptotically to the endemic equilibrium state.





Fig. 3 Evolution of the number of HIV individuals without AIDS

In Figure 4, we observe that the number of people with AIDS will continue to rise; it will reach its maximum in 2065 which is about 7.375×10^6 cases. Then it will decrease asymptotically to the endemic equilibrium state.



Fig. 4 Evolution of the number of AIDS cases

5. Conclusion

In this paper we presented a simple model, with varying population size, describing the spread of HIV/AIDS. We then estimated the parameters of this model in order to reconstruct the epidemic in the past, to simulate its future development in Morocco, and to estimate the basic reproductive number for the Moroccan case. Although our estimates were based only on reported cases of HIV/AIDS, we note that the number of people suffering from HIV/AIDS continues to rise quickly and more intensely. Therefore, vigilance is still required and strengthening the national strategy against HIV/AIDS is needed more than before; especially that estimates of the number of people living with HIV, published since 2003 by the Ministry of Public Health in Morocco in collaboration with UNAIDS (Fig 5), show a large discrepancy with the number of reported cases.



Fig. 5 Estimated number of people living with HIV. Morocco, 2003-2009 according to [13]

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