

IDENTIFICATION OF GUMEL-MICKENS HIV MODEL WITH INCOMPLETE DATA ON A POPULATION

M. Ngungu¹, MY. Shatalov², CR. Kikawa³

Tshwane University of Technology

¹Merciend@yahoo.com

²Shatalovm@tut.ac.za

³Richard.kikawa@gmail.com

Department of Mathematics and Statistics

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OUTLINE

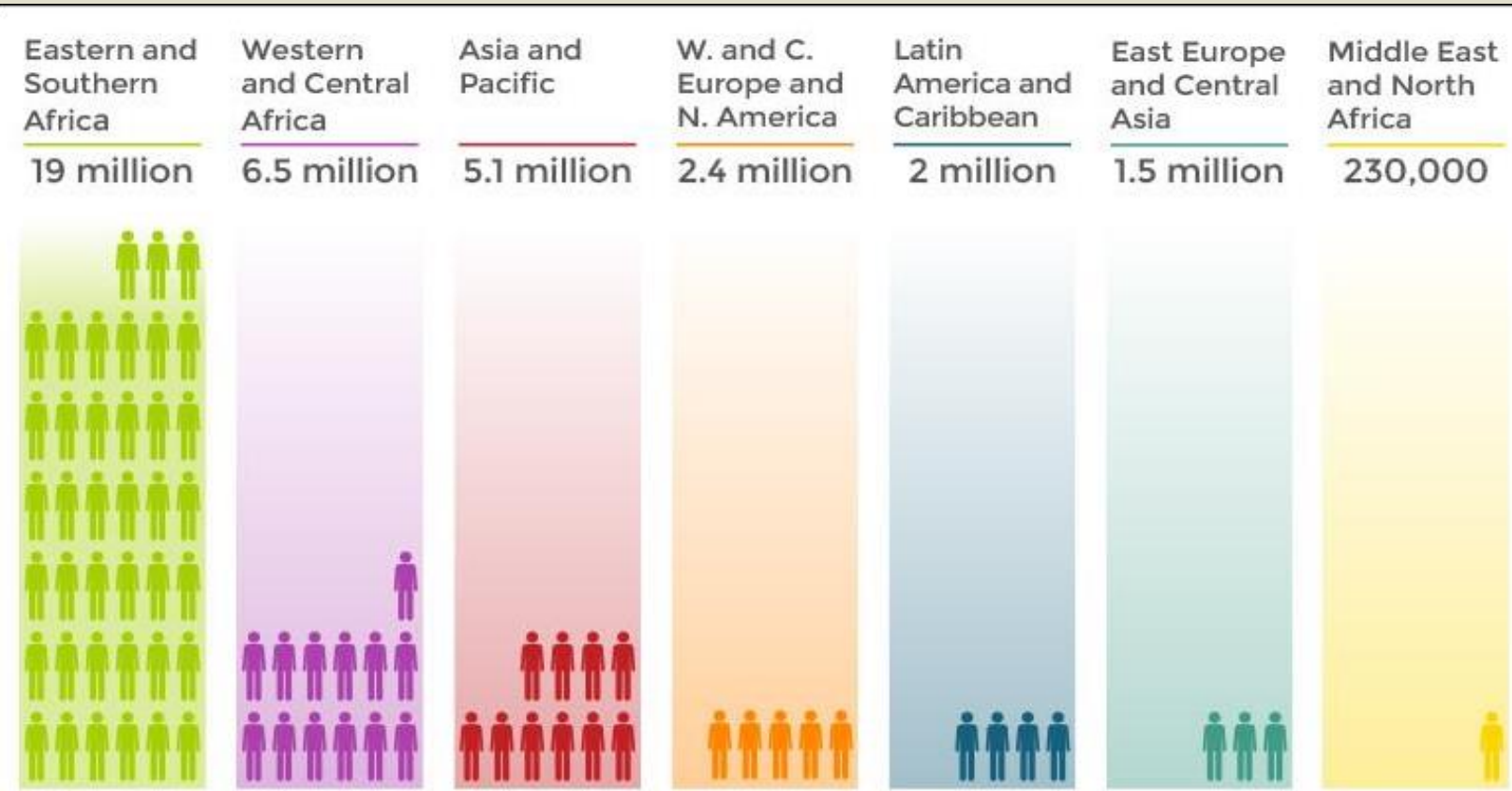
- Introduction
- Objectives
- The Gumel-Mickens Model
- Solution of the problem & numerical results
- Conclusions and applications
- References

INTRODUCTION

- Challenges/problems of **incomplete information or unavailable data** in modelling of disease dynamics
- Previously, (Fedotov & Shatalov, 2007) problems on estimation of dynamic system parameters from experimental data in linear systems with respect to their unknown coefficients have been discussed
- This talk presents a **method of identifying mathematical models** that can be used to model HIV/AIDs transmission dynamics in the presence of incomplete data.

INTRODUCTION-CONT...

HIV/AIDS population statistics by end 2015



Geographical region	World	Sub-Saharan Africa	South Africa
Living with HIV/AIDS	36.7 million	25.6 million	6.9 million
New Infections	2.1 million	0.97 million	0.38 million
Deaths	1.1 million	0.47 million	0.18 million

HIV/AIDS related challenges

- Affects health of a population,
- consequently, impacts development and social economic growth of households, communities and nations at large.
- Owing to these statistics there is a need to study, design, analyse and solve mathematical models for determining indicators that could give insight into eradicating HIV/AIDSs.

OBJECTIVES

- In “*On identification of dynamical system parameters from experimental data*” the authors, (Mathye, Fedotov and Shatalov, 2004) the Gumel-Mickens HIV transmission model was considered with the aim of estimating the unknown parameters and restoring the information
- Analogous, with different parameters and coefficients unknown;
 - Fully **identify** the Gumel-Mickens HIV transmission model,
 - find all **coefficients** of this mathematical model and
 - **restore** information on
 - HIV- susceptible population
 - and uninfected-vaccinated population
- Ultimately, with the key objective of giving insights in eradicating HIV from the community.

THE GUMEL-MICKENS MODEL

➤ HIV transmission model that monitors the dynamics of HIV in four sexually active populations in the presence of vaccines as described by Gumel Moghadas and Mickens

$$D(T) = \left\{ \begin{array}{l} \dot{X} = P_1 - \mu_1 X - \alpha_v \frac{XY_V}{N} - \alpha_w \frac{X(Y_W + Y_{WV})}{N} \\ \dot{Y}_V = P_2 - \mu_2 Y_V + \alpha_v \frac{XY_V}{N} - \gamma \frac{Y_V(Y_W + Y_{WV})}{N} \\ \dot{Y}_W = -\mu_3 Y_W + \alpha_w \frac{X(Y_W + Y_{WV})}{N} \\ \dot{Y}_{WV} = -\mu_4 Y_{WV} + \gamma \frac{Y_V(Y_W + Y_{WV})}{N} \end{array} \right.$$

With $t > t_0$ and initial conditions:

$$X(t_0) = X_0,$$

$$Y_W(t_0) = Y_{W0},$$

$$Y_V(t_0) = Y_{V0},$$

$$Y_{WV}(t_0) = Y_{WV0},$$

Parameter	Interpretation
$X(t)$	Population susceptible to HIV
$Y_V(t)$	Individuals uninfected by wild type and vaccinated population
$Y_W(t)$	Individuals infected by wild type and unvaccinated population
$Y_{WV}(t)$	Individuals infected by wild type and vaccinated population
$Z = Y_W(t) + Y_{WV}(t)$	Infected by wild type and vaccinated population
$ZZ = X(t) + Y_V(t)$	Healthy population
$N(t) = X(t) + Y_V(t) + Y_W(t) + Y_{WV}(t)$	Total (sexually active) population
P	Recruitment rate of susceptible population
$\mu_1, \mu_2, \mu_3, \mu_4$	Natural cessation of sexual activity
α_w and α_v	Rate of transmission of virus and vaccine, respectively
γ	Degree of protection against virus

Above parameters are taken from "Effect of a preventive vaccine on dynamics of HIV transmission" by A.B. Gumel, S.M. Moghadas and R.E. Mickens // *Communications in nonlinear Science and Numerical Simulations*, 9 2004, pp.649-659.

THE GUMEL-MICKENS MODEL-CONT...

$$N = X + Y_V + Y_W + Y_{WV}$$

P. Mathye : N, X, Y_V - are known

$$Z = Y_W + Y_{WV} = N - X - Y_V - \text{also known}$$

But: $Y_W = Y_W(t)$ and $Y_{WV} = Y_{WV}(t)$ - are not known

In this case: N, Y_W, Y_{WV} - are known

$$ZZ = N - (Y_W + Y_{WV}) = X + Y_V - \text{also known}$$

But: $X = X(t)$ and $Y_V = Y_V(t)$ - are not known

and coefficients are unknown

THE GUMEL-MICKENS MODEL-CONT...

Assumptions:

- all dependent variables of G-M model and parameters are non-negative,
- all new individuals recruited into the society (either via birth or immigration) at a rate P per year are considered to be susceptible analogous to Gumel *et al.*,
- population is reduced by the natural cessation of sexual activity at a constant rate l , and by infection with the virus which may be acquired from each new sexual partner at a rate α_w .

SOLUTION OF PROBLEM

➤ Stage 1:

➤ Use direct method of initial problem solving

➤ assume that all parameters and initial conditions of the four populations are known as below and solve the system numerically using Computer Algebra System (CAS)

```
pi := 2000      p := 0.8      c := 5.0      beta_v := 0.5      beta_w := 0.45      psi := 0.6
mu := 0.031     d1 := 0.3      d2 := 0.25     d3 := 0.2
```

➤ Recalculation of the parameters into new parameters

```
p1 := (1 - p) * pi      mu1 := mu      mu2 := mu + d1      mu3 := mu + d2      mu4 := mu + d3
p2 := p * pi            alpha_v := c * beta_v      alpha_w := c * beta_w      g := (1 - psi) * alpha_w      ( g for gamma )
```

SOLUTION OF PROBLEM-CONT...

- System of ODE with known parameters

Right hand side of the Cauchy system ($u_0 = X, u_1 = Y_v, u_2 = Y_w, u_3 = Y_{vw}$):

$$D(t, u) := \begin{bmatrix} p1 - \left[\alpha v \cdot \frac{u_0 \cdot u_1}{(u_0) + (u_1) + (u_2) + (u_3)} \right] - \left[\alpha w \cdot \frac{u_0 \cdot (u_2 + u_3)}{(u_0) + (u_1) + (u_2) + (u_3)} \right] - \mu 1 \cdot u_0 \\ p2 + \left[\alpha v \cdot \frac{u_0 \cdot u_1}{(u_0) + (u_1) + (u_2) + (u_3)} \right] - \left[g \cdot \frac{u_1 \cdot (u_2 + u_3)}{(u_0) + (u_1) + (u_2) + (u_3)} \right] - \mu 2 \cdot u_1 \\ \left[\alpha w \cdot \frac{u_0 \cdot (u_2 + u_3)}{(u_0) + (u_1) + (u_2) + (u_3)} \right] - \mu 3 \cdot u_2 \\ \left[g \cdot \frac{u_1 \cdot (u_2 + u_3)}{(u_0) + (u_1) + (u_2) + (u_3)} \right] - \mu 4 \cdot u_3 \end{bmatrix}$$

Initial conditions are also taken from the abovementioned paper:

$$u := (80 \cdot 10^3 \quad 2 \cdot 10^3 \quad 8 \cdot 10^3 \quad 8 \cdot 10^3)^T$$

SOLUTION OF PROBLEM-CONT...

Time interval (**T**) is different (shorter than in the previously cited paper). **NN** is number of interval onto given time interval.

T := 5 **NN** := 25 **i** := 0..**NN**

Solution of system of equation by the specific method:

U := Adams(**u**, 0, **T**, **NN**, **D**) **TOL** = 1×10^{-3}

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SOLUTION OF PROBLEM-CONT...

➤ Stage 2:

- Forget the values of the parameter and assume that we know only N - total population and Y_W - population infected, $Y_{WV}(t)$ -population infected-vaccinated
- It is also known that the sum $N - Y_W - Y_{WV} = N - Z = X + Y_V$, that is the sum of healthy population

$$\begin{array}{llll}
 N_i := (U^{(1)})_i + (U^{(2)})_i + (U^{(3)})_i + (U^{(4)})_i & Z_i := (U^{(3)})_i + (U^{(4)})_i & U^{(1)} = X & \\
 ZZ_i := N_i - Z_i & YW := U^{(3)} & U^{(2)} = Y_V & \\
 & YVW := U^{(4)} & U^{(3)} = Y_W & \\
 & & U^{(4)} = Y_{VW} &
 \end{array}$$

- Interpolation and subsequent differentiation using CAS

$$\begin{array}{ll}
 VYw := \text{cspline}[(U^{(0)}), U^{(3)}] & VYvw := \text{cspline}(U^{(0)}, U^{(4)}) \\
 Yw(t) := \text{interp}(VYw, U^{(0)}, U^{(3)}, t) & Yvw(t) := \text{interp}(VYvw, U^{(0)}, U^{(4)}, t)
 \end{array}$$

NUMERICAL RESULTS

➤ Numerical solutions

$N =$

	0
0	$9.8 \cdot 10^4$
1	$9.679 \cdot 10^4$
2	$9.52 \cdot 10^4$
3	$9.32 \cdot 10^4$
4	$9.078 \cdot 10^4$
5	$8.798 \cdot 10^4$
6	$8.489 \cdot 10^4$
7	$8.159 \cdot 10^4$
8	$7.82 \cdot 10^4$
9	$7.479 \cdot 10^4$
10	$7.143 \cdot 10^4$
11	$6.816 \cdot 10^4$
12	$6.501 \cdot 10^4$
13	$6.2 \cdot 10^4$
14	$5.913 \cdot 10^4$
15	...

$YW =$

	0
0	$8 \cdot 10^3$
1	$1.397 \cdot 10^4$
2	$2.086 \cdot 10^4$
3	$2.823 \cdot 10^4$
4	$3.541 \cdot 10^4$
5	$4.17 \cdot 10^4$
6	$4.655 \cdot 10^4$
7	$4.977 \cdot 10^4$
8	$5.143 \cdot 10^4$
9	$5.181 \cdot 10^4$
10	$5.121 \cdot 10^4$
11	$4.995 \cdot 10^4$
12	$4.826 \cdot 10^4$
13	$4.633 \cdot 10^4$
14	$4.428 \cdot 10^4$
15	...

$YVW =$

	0
0	$8 \cdot 10^3$
1	$7.725 \cdot 10^3$
2	$7.544 \cdot 10^3$
3	$7.493 \cdot 10^3$
4	$7.601 \cdot 10^3$
5	$7.869 \cdot 10^3$
6	$8.269 \cdot 10^3$
7	$8.745 \cdot 10^3$
8	$9.236 \cdot 10^3$
9	$9.689 \cdot 10^3$
10	$1.007 \cdot 10^4$
11	$1.037 \cdot 10^4$
12	$1.058 \cdot 10^4$
13	$1.07 \cdot 10^4$
14	$1.076 \cdot 10^4$
15	...

SOLUTION OF PROBLEM-CONT...

➤ Goal function

$$G = G(a_1, a_2, a_3, a_4)$$

$$G = \frac{1}{2} \sum_{k=0}^n \left\{ a_1 \dot{Y}_{w,k} + a_2 Y_{w,k} + a_3 \dot{Y}_{wv,k} + a_4 Y_{wv,k} - \frac{ZZ_k - Z_k}{N_k} \right\}^2$$

Where;

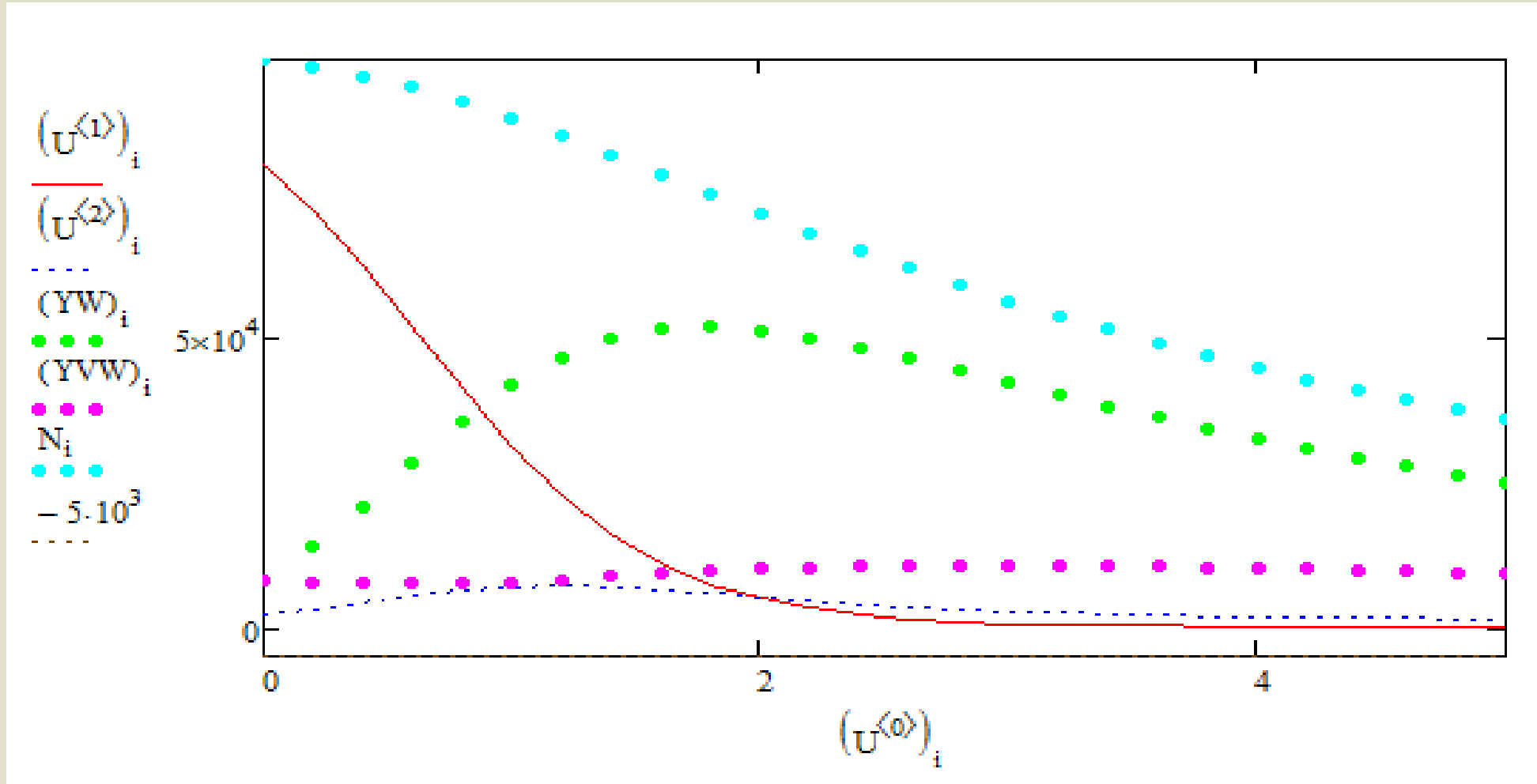
$$a_1 = \alpha_w^{-1}, \quad a_2 = \mu_3 \alpha_w^{-1}, \quad a_3 = \gamma^{-1}, \quad a_4 = \mu_4 \gamma^{-1}$$

➤ We calculate,

$$X_k = \frac{\dot{Y}_{w,k}(t) + \mu_3 Y_{w,k}(t)}{\alpha_w} \frac{N_k}{Z_k}$$

$$Y_{v,k} = \frac{\dot{Y}_{wv,k}(t) + \mu_4 Y_{wv,k}(t)}{\gamma} \frac{N_k}{Z_k}$$

NUMERICAL RESULTS-CONT...



Key	
—	Susceptible pop.
- - -	Uninfected-vaccinated pop.
•••	Infected-unvaccinated pop.
•••	Infected-vaccinated pop.
•••	Total pop.

Figure 1: Population graph against time with solution of known information

NUMERICAL RESULTS-CONT...

➤ Solution for the known on interpolation and differentiation

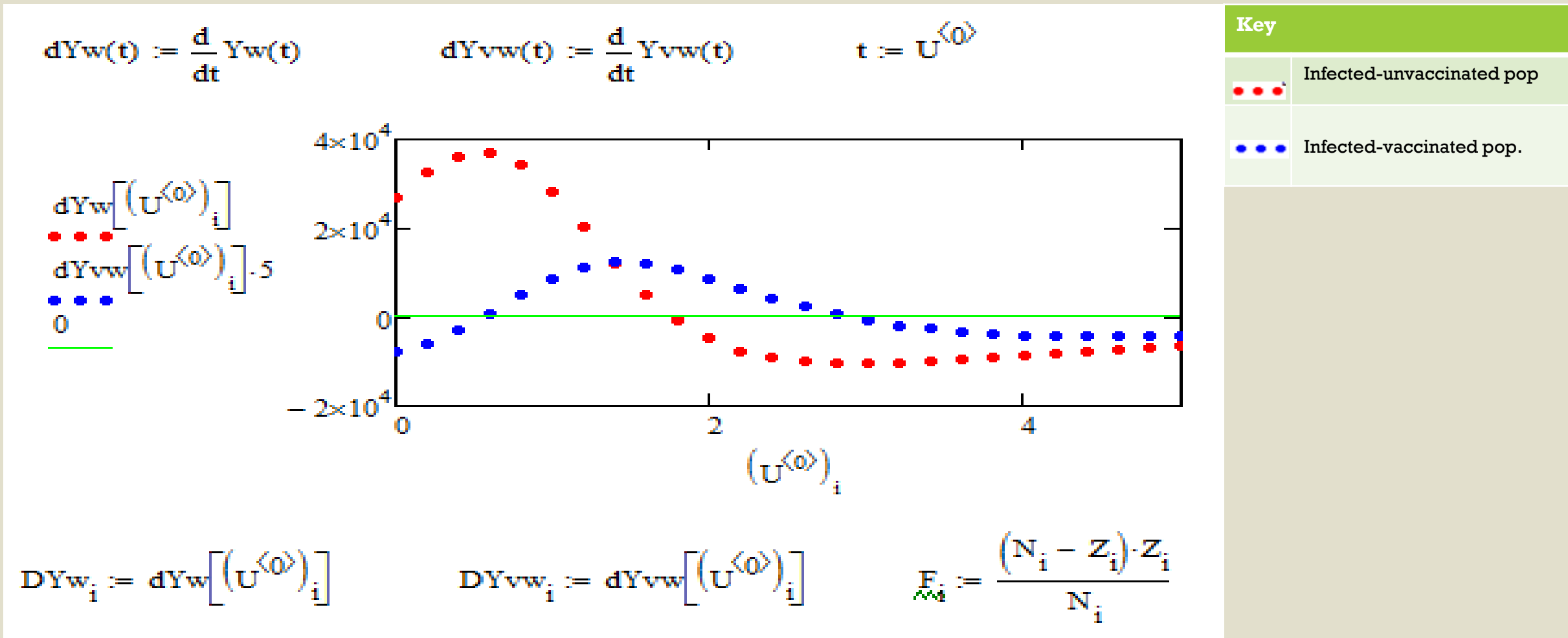


Figure 2: Population graph against time with known information ($Z = Y_W(t) + Y_{WV}(t)$)

SOLUTION OF PROBLEM-CONT...

➤ By means of **Least square method** we produce matrix of coefficients and RHS matrices for obtaining of unknown parameters:

$$\begin{aligned}
 a_{0,0} &:= \sum_i [(DYw_i)^2] & a_{0,1} &:= \sum_i [(DYw_i) \cdot (Yw_i)] & a_{0,2} &:= \sum_i [(DYw_i) \cdot (DYvw_i)] & a_{0,3} &:= \sum_i [(DYw_i) \cdot (Yvw_i)] & b_0 &:= \sum_i [(DYw_i) \cdot (F_i)] \\
 & & a_{1,1} &:= \sum_i [(Yw_i)^2] & a_{1,2} &:= \sum_i [(Yw_i) \cdot (DYvw_i)] & a_{1,3} &:= \sum_i [(Yw_i) \cdot (Yvw_i)] & b_1 &:= \sum_i [(Yw_i) \cdot (F_i)] \\
 & & & & a_{2,2} &:= \sum_i [(DYvw_i)^2] & a_{2,3} &:= \sum_i [(DYvw_i) \cdot (Yvw_i)] & b_2 &:= \sum_i [(DYvw_i) \cdot (F_i)] \\
 & & & & & & a_{3,3} &:= \sum_i [(Yvw_i)^2] & b_3 &:= \sum_i [(Yvw_i) \cdot (F_i)]
 \end{aligned}$$

$$\underline{\underline{A}} := \begin{pmatrix} a_{0,0} & a_{0,1} & a_{0,2} & a_{0,3} \\ a_{0,1} & a_{1,1} & a_{1,2} & a_{1,3} \\ a_{0,2} & a_{1,2} & a_{2,2} & a_{2,3} \\ a_{0,3} & a_{1,3} & a_{2,3} & a_{3,3} \end{pmatrix}$$

$$|A| = 9.155 \times 10^{34}$$

$$B := b$$

-1

NUMERICAL RESULTS-CONT...

➤ Solution of parameters and coefficients

$$a1234 := A^{-1} \cdot B$$

$$a1234 = \begin{pmatrix} 0.444 \\ 0.12 \\ 1.144 \\ 0.274 \end{pmatrix}$$

$$\begin{aligned} \alpha w^{-1} &= 0.444 \\ \mu_3 \cdot \alpha w^{-1} &= 0.125 \\ g^{-1} &= 1.111 \\ \mu_4 \cdot g^{-1} &= 0.257 \end{aligned}$$

$$X_i := \left(a1234_0 \cdot DYw_i + a1234_1 \cdot Yvw_i \right) \cdot \frac{N_i}{Z_i}$$

$$Yv_i := \left(a1234_2 \cdot DYvw_i + a1234_3 \cdot Yvw_i \right) \cdot \frac{N_i}{Z_i}$$

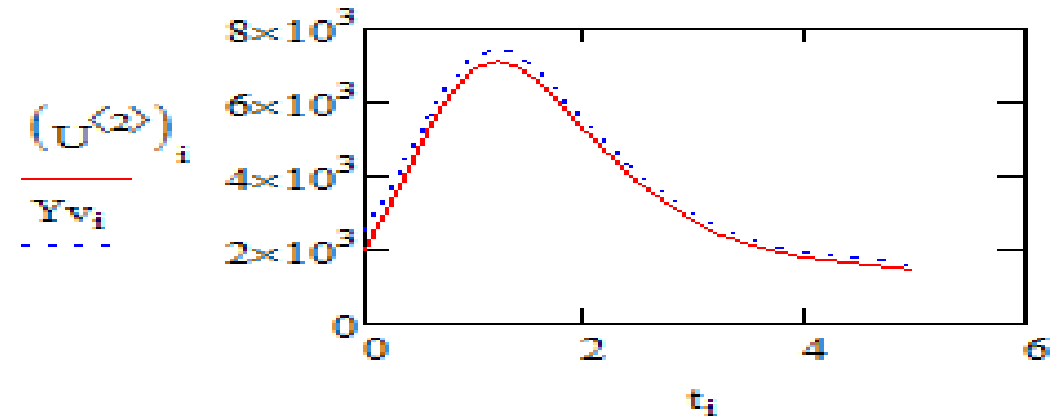
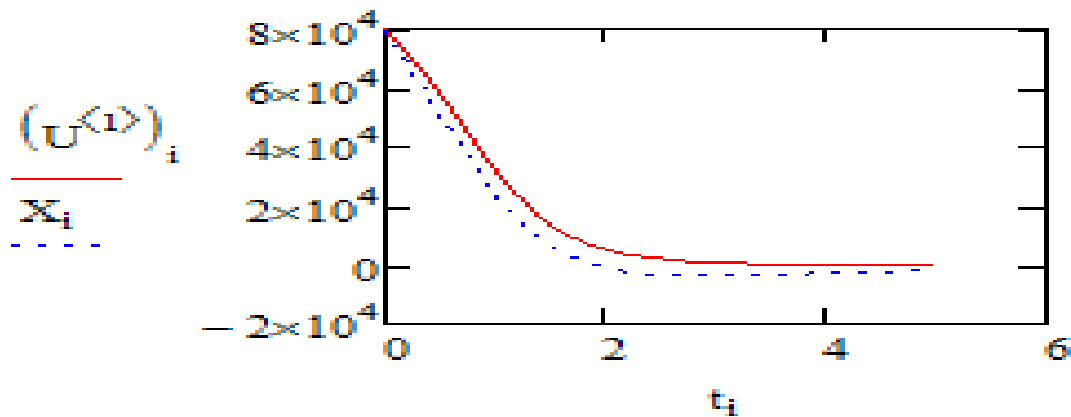


Figure 3: Population graph against time comparing the estimated and actual results

Results

➤ Relative error of estimation in %

$$\left| \frac{\left[\alpha w - \left(a_{1234_0} \right)^{-1} \right]}{\alpha w} \right| \cdot 100 = 0.036 \quad (\%)$$
$$\left| \frac{\left(\mu_3 - \frac{a_{1234_1}}{a_{1234_0}} \right)}{\mu_3} \right| \cdot 100 = 3.551 \quad (\%)$$

$$\left| \frac{\left[g - \left(a_{1234_2} \right)^{-1} \right]}{g} \right| \cdot 100 = 2.868 \quad (\%)$$
$$\left| \frac{\left(\mu_4 - \frac{a_{1234_3}}{a_{1234_2}} \right)}{\mu_4} \right| \cdot 100 = 3.57 \quad (\%)$$

Note:

- relative error can be further minimised by increasing the NN points
- All other parameters can be also estimated

DISCUSSIONS AND CONCLUSIONS

- The estimated parameters are calculated accurately
- The obtained $X(t)$ and $Y_v(t)$ values are in good qualitative and quantitative correspondence and possible improvement of accuracy could be achieved by proposing other estimation methods
- Applications,
 - by researchers in the medical or epidemiology filed for estimation
 - big data solutions aimed at restoring incomplete data or information
 - predictive analytics

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ASANTE

THANK YOU

MERCI

OBRIGADO

ARIGATO

KIITOS

ASANTE

KIITOS

KIA ORA

SALAMAT

MATONDO

VINAKA JUSPAXAR

