# SEMI-ANALYTIC METHODS FOR THE SOLUTION OF TWO EPIDEMIOLOGICAL MODELS 

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#### Abstract

In this paper, we apply three semi-analytical methods, viz: the Differential Transform Method (DTM), Homotopy Perturbation Method (HPM) and the Variational Iteration Method (VIM) to compute approximate solutions of a continuous mathematical model of Shigella diarrhea comprising of a non-constant population and a deterministic model on the impact of stress on the dynamics and treatment of Tuberculosis.


## 1. Introduction

Most epidemiological models are formulated using systems of linear and non-linear differential equations. After the stability analyses of the models are qualitatively obtained, the numerical solution of such models are in most cases conducted using the fourth-order Runge-Kutta method embedded in either MAPLE or MATLAB mathematical softwares. In a view to diversifying the approach, we employed the Differential Transform Method (DTM), the Homotopy Perturbation Method (HPM) and the Variational Iteration Method (VIM) to solve the models by by Ojaswita et al. [1] and Namawejje et al. [2].
The concept of DTM was first proposed by Zhou [3] in 1986. This method constructs a semi-analytical numerical technique that uses Taylor series for the solution of differential equations in the form a polynomial. It is different from the high-order Taylor series method which requires symbolic computation of the necessary derivatives of the data functions. The differential transform is an iterative procedure for obtaining analytic Taylor series solutions of differential equations. The method is very effective and powerful for solving various kinds of differential equations [4]. For examples, it was used to find the analytic approximate solution of the Volterra's model population growth of a species in a closed system [5], to solve a system of nonlinear differential equations, namely, the Coullet system [6], to find the solution of nonlinear PDEs [7], to solve Volterra integral and integro-differential equations with proportional delays [8] and so on.
The Homotopy Perturbation Method (HPM) was first proposed by He [9] and was developed and improved by He [10, 11, 12]. HPM is a coupling of the traditional perturbation method in topology, He [13]. The method provides an approximate analytical solution in a series form. HPM has been widely used by numerous researchers successfully for physical systems such as bifurcation, asymptotology, nonlinear wave equations, oscillations with discontinuities by He [14, 15, 16, 17]. It has also been applied to the reaction-diffusion equation [18] and the SIR infectious disease model [19].
The Variational Iteration Method (VIM), first envisioned by He [20] modifying the approach by Inokuti et al. [21], has been applied to many linear and nonlinear ODEs and PDEs. For example, He [22] employed VIM to give approximate solutions for some well - known nonlinear problems and in [23], he successfully applied VIM to autonomous systems of ODEs. Also, He [24] gave a solution for a seepage flow problem with fractional derivatives in porous media using VIM. Other researchers demonstrated further applications of VIM. For instance, Soliman [25] applied VIM to solve KdV Burger's and Lax's seventh - order KdV equations. Momani and Abuasad [26] also used VIM to solve the Helmholtz equation and Odibat and Momani [27] solved nonlinear fractional differential equations via VIM; while Bildik [28] used VIM, DTM AND ADM to solve different types of nonlinear partial differential equations, and Abbasbandy [29] solved the quadratic Riccati differential equations by He's VIM with considering Adomian polynomials. Jufeng [30] applied VIM to solve two - point boundary value problems.

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The paper is organized as follows: In section 2, we give a brief description of VIM and in section 3, two mathematical models are solved using these methods. Some concluding remarks are given in section 4.

### 2.1 The Differential Transformation Method (DTM)

An arbitrary function $f(x)$ can be expanded in Taylor series about a point $x=0$ as
$f(x)=\sum_{k=0}^{\infty} \frac{x^{k}}{k!}\left[\frac{d^{k} f}{d x^{k}}\right]_{x=0}$
The differential transformation of $f(x)$ is defined as
$F(x)=\frac{1}{k!}\left[\frac{d^{k} f}{d x^{k}}\right]_{x=0}$
Then the inverse differential transform is
$f(x)=\sum_{k=0}^{\infty} x^{k} F(k)$
Table 1 - The Fundamental Mathematical Operations Performed by DTM [31]

Original Function
$y(x)=g(x) \pm h(x)$
$y(x)=\alpha g(x)$
$y(x)=\frac{d g(x)}{d x}$
$y(x)=\frac{d^{2} g(x)}{d x^{2}}$
$y(x)=\frac{d^{m} g(x)}{d x^{m}}$
$y(x)=1$
$y(x)=x$
$y(x)=x^{m}$
$y(x)=g(x) h(x)$
$y(x)=e^{(\lambda x)}$
$y(x)=(1+x)^{m}$

Transformed Function
$Y(k)=G(k) \pm H(k)$
$Y(k)=\alpha G(k)$
$Y(k)=(k+1) G(k+1)$
$Y(k)=(k+1)(k+2) G(k+2)$
$Y(k)=(k+1)(k+2) \ldots(k+m) G(k+2)$
$Y(k)=\delta(k)$
$Y(k)=\delta(k-1)$
$Y(k)=\delta(k-m)=\left\{\begin{array}{l}1, k=m \\ 0, k \neq m\end{array}\right.$
$Y(k)=\sum_{m=0}^{k} H(m) G(k-m)$
$Y(k)=\frac{\lambda^{k}}{k!}$
$Y(k)=\frac{m(m-1) \ldots(m-k+1)}{k!}$

### 2.2 Homotopy Perturbation Method (HPM)

To illustrate the ideas of this method, the following nonlinear equation was considered, He [32].
$A(u)-f(r)=0, r \in \Omega$
Subject to the boundary condition of:
$B\left(u, \frac{\partial u}{\partial n}\right)=0, \quad r \in \Gamma$
Where $A$ is a general differential operator, $B$ is a boundary operator, $f(r)$ a known analytical function and $\Gamma$ is the boundary of the domain $\Omega$. The operator $A$ can be divided into two parts of $L$ and $N$, where $L$ is the linear part, while $N$ is a nonlinear one. Equation (4) can therefore be rewritten as follows:

$$
\begin{equation*}
L(u)+N(r)-f(r)=0, \quad r \in \Omega \tag{6}
\end{equation*}
$$

Using the homotopy technique, we construct a homotpy $v(r, p): \Omega \times[0,1] \rightarrow R$, which satisfies
$H(v, p)=(1-p)\left[L(v)-L\left(u_{0}\right)\right]+p[L(v)+N(v)-f(r)]=0$
$p \in[0,1], \quad r \in \Omega$
Or
$H(v, p)=L(v)-L\left(u_{0}\right)+p L\left(u_{0}\right)+p[N(u)-f(r)]=0$
Where
$L(u)$ is the part
$L(u)=L(v)-L\left(u_{0}\right)+p L\left(u_{0}\right)$
And $N(u)$ is the nonlinear part
$N(u)=p N(v)$
In eqn (7), $p \in[0.1]$ is an embedding parameter and $u_{0}$ is an initial approximation of (4) and (9), we will have

```
\(H(v, 0)=L(v)-L\left(u_{0}\right)=0\)
\(H(v, 1)=L(v)+N(v)-f(r)=0\)
\(H(v, 1)=L(v)+N(v)-f(r)=0\)
```

The changing process of p from zero to unity is just of $H(v, r)$ from $u_{0}(r)$ to $u(r)$. In topology, this is called deformation and $L(v)-L\left(u_{0}\right)$ and $A(v)-f(r)$ are called homotopy. According to HPM, we can first use the embedding parameter p as a small parameter and assume that the solution of eqn (7) and (9) can be written as power series in p :
$v=v_{0}+p v_{1}+p^{2} v_{2}+\cdots$
Setting $p=1$ results in the approximate solution (4) as
$u=\lim _{p \rightarrow 1} v=v_{0}+v_{1}+v_{2}+\cdots$
The series (13) is convergent for most cases. However, the convergence rate depends on the nonlinear operator $A(v)$. The following observations are made by He [11]:

- The derivatives of $N(v)$ with respect to $v$ must be small because the parameter p may be large, i.e. $p \rightarrow 1$.
- The norm of $L^{-1} \frac{\partial N}{\partial V}$ must be smaller than one so that the series converges.


### 2.3 The Variational Iteration Method (VIM)

To illustrate the basic concept of the Variational Iteration Method, we consider
the following general differential equation:
$L u+N u=g(x, t)$
where $L$ and $N$ are linear and non-linear operators respectively, and $g(x, t)$ is the source inhomogeneous term. He proposed the variational iteration method where a correction functional for (14) can be written as
$u_{i, n+1}(x, t)=u_{i, n}(x, t)+\int_{t_{0}}^{t} \lambda\left\{L u_{i, n}(s)+N \tilde{u}_{i, n}(s)-g(s)\right\} \mathrm{d} s$,
where $i=1,2, \ldots, m, \lambda$ is a general Lagrange multiplier, which can be identified optimally via the variational theory, n denotes the nth approximation and $\tilde{u}_{i, n}$ is considered as a restricted variation which means $\delta \tilde{u}_{i, n}=0$. It is required first to determine the Lagrange multiplier $\lambda$ [32] that will be identified optimally via integration by parts. The successive approximations $u_{i, n+1}(x, t), n \geq 0$, of the solution $u(x, t)$ will be readily obtained upon using the Lagrange multiplier obtained and by using any selective function $u_{0}$. Having $\lambda$ determined, then several approximations $u_{j}(x, t), j \geq 0$, can be determined. Consequently, the solution is given by
$u=\lim _{n \rightarrow \infty} u_{n}$

## 3. Application

3.1. Considering the Shigella diarrhea model by Ojaswita et al. [1],
$\frac{d S}{d t}=\pi-\mu S-\beta S I+\alpha R$
$\frac{d I}{d t}=\beta S I-\mu I-\gamma I$
$\frac{d R}{d t}=\gamma I-\mu R-\alpha$
Subject to the initial conditions $S(0)=100, I(0)=40, R(0)=10$
where the parameters $\pi$ is the population renewal rate, $\beta$ is the infectivity rate, $\mu$ is the natural death rate, $r$ is the recovery rate and $\alpha$ is immunity loss rate with their values given as $\pi=10, \mu=0.012$ [33], $\alpha=0.25, r=0.14$.
By applying the DTM to the model, we obtained the following recurrence relations:
$S(k+1)=\frac{1}{k+1}\left[10-0.012 S(k)-0.0001\left(\sum_{m=o}^{k} S(m) I(k-m)\right)+0.25 R(k)\right]$
$I(k+1)=\frac{1}{k+1}\left[0.0001\left(\sum_{m=0}^{k} S(m) I(k-m)\right)-0.152 I(k)\right]$
$R(k+1)=\frac{1}{k+1}[0.14 I(k)-0.262 R(k)]$
Then the closed form of the solutions of order 8 are
$s(t)=\sum_{k=0}^{8} S(k) t^{k}$
Transactions of the Nigerian Association of Mathematical Physics Volume 17, (October - December, 2021), 51-60

$$
\begin{aligned}
& s(t)=100+10.9000 t+5.3137 t^{2}+3.2399754 t^{3}+2.493257432 t^{4}+1.992024714 t^{5}+1.661583302 t^{6} \\
& +1.424919360 t^{7}+1.247257316 t^{8} \\
& i(t)=\sum_{k=0}^{8} I(k) t^{k} \\
& i(t)=40-5.6800 t+0.42508 t^{2}-0.01509925333 t^{3}+0.003137287795 t^{4}+0.001579329081 t^{5} \\
& +0.001076797522 t^{6}+0.0007809196739 t^{7}+0.0005911186595 t^{8} \\
& r(t)=\sum_{k=0}^{8} R(k) t^{k} \\
& r(t)=10+2.980 t-0.78798 t^{2}+0.08865398667 t^{3}-0.006335309995 t^{4}+0.0004198143020 t^{5} \\
& +0.00001851912070 t^{6}+0.00002084280621 t^{7}+0.00001298349239 t^{8}
\end{aligned}
$$

Applying HPM to the model, we obtain the following:
$(1-p) \frac{d S}{d t}+p\left[\frac{d S}{d t}+\mu S+\beta S I-\pi-\alpha R\right]=0$
$(1-p) \frac{d I}{d t}+p\left[\frac{d I}{d t}+(\mu+\gamma) I-\beta S I\right]=0$
$(1-p) \frac{d R}{d t}+p\left[\frac{d R}{d t}-\gamma I+(\mu+\alpha) R\right]=0$
Let $S=x_{0}+p x_{1}+p^{2} x_{2}+\cdots$
$I=y_{0}+p y_{1}+p^{2} y_{2}+\cdots$
$R=z_{0}+p z_{1}+p^{2} z_{2}+\cdots$
Solving and substituting $x_{0}, x_{1}, x_{2}, y_{0}, y_{1}, y_{2}, z_{0}, z_{1}, z_{2}$ into $\mathrm{S}, \mathrm{I}$ and R above gives
$S(t)=S_{0}+p\left[\pi+\alpha R_{0}-\mu S_{0}-\beta S_{0} I_{0}\right] t-p^{2}-\left\{\beta S_{0} I_{0}\left[\beta S_{0}-(\mu+\gamma)\right]+\right.$
$\left.\left(\pi+\alpha R_{0}-\mu S_{0}-\beta S_{0} I_{0}\right)\left(\mu+\beta I_{0}\right)\right\} \frac{t^{2}}{2}$
$I(t)=I_{0}+p\left[\left(\beta S_{0} I_{0}-(\mu+\gamma) I_{0}\right) t\right]+p^{2}\left\{\beta S_{0} I_{0}\left[\beta S_{0}-(\mu+\gamma)\right]+\beta I_{0}\left[\pi+\alpha R_{0}-\mu S_{0}-\beta S_{0} I_{0}\right]-(\mu+\gamma)\left[\beta S_{0}-\right.\right.$
$\left.(\mu+\gamma)] I_{0}\right\} \frac{t^{2}}{2}$
$R(t)=R_{0}+p\left[\left(\gamma I_{0}-(\mu+\alpha) R_{0}\right) t\right]+p^{2}\left\{\gamma I_{0}\left[\beta S_{0}-(\mu+\alpha)\right]-(\mu+\alpha)\left[\alpha I_{0}-(\mu\right.\right.$
$\left.\left.+\alpha) R_{0}\right]\right\} \frac{t^{2}}{2}$
Setting $p=1$, the results follow.
Applying VIM to the same model, we construct the correction functional as follows
$S^{n+1}(t)=S^{n}(t)+\int_{\substack{0 \\ t}}^{t} \lambda_{1}(\xi)\left\lfloor\frac{d}{d \xi} S^{n}(\xi)-\mu S^{n}(\xi)-\pi+\beta S^{\mathrm{n}}(\xi) I^{\mathrm{n}}(\xi)-\alpha \mathrm{R}^{\mathrm{n}}(\xi)\right] d \xi$
$I^{n+1}(t)=I^{n}(t)+\int_{0}^{t} \lambda_{2}(\xi)\left\lfloor\frac{d}{d \xi} I^{n}(\xi)-\beta \mathrm{S}^{\mathrm{n}}(\xi) I^{\mathrm{n}}(\xi)+(\gamma+\mu) I^{\mathrm{n}}(\xi)\right\rfloor d \xi$
$R^{n+1}(t)=R^{n}(t) \int_{0}^{t} \lambda_{3}(\xi)\left[\frac{d}{d \xi} R^{n}(\xi)+(\alpha+\mu) \mathrm{R}^{\mathrm{n}}(\xi)-\gamma \mathrm{I}^{\mathrm{n}}(\xi)\right] d \xi$
Making the functional stationary, we obtain the following Lagrange multipliers
$\lambda_{1}(\xi)=-e^{\mu(\xi-t)}, \lambda_{2}(\xi)=-e^{(\mu+\gamma)(\xi-t)}$ and $\lambda_{3}(\xi)=-e^{(\mu+\alpha)(\xi-t)}$
Substituting these Lagrange multipliers into the correction functional to obtain
$S^{n+1}(t)=S^{n}(t)+\int_{0}^{t}-e^{\mu(\xi-t)}\left\lfloor\frac{d}{d \xi} S^{n}(\xi)-\mu S^{n}(\xi)-\pi+\beta \mathrm{S}^{\mathrm{n}}(\xi) \mathrm{I}^{\mathrm{n}}(\xi)-\alpha \mathrm{R}^{\mathrm{n}}(\xi)\right\rfloor d \xi$

Transactions of the Nigerian Association of Mathematical Physics Volume 17, (October - December, 2021), 51 -60
$I^{n+1}(t)=I^{n}(t)+\int_{0}^{t}-e^{(\mu+\gamma)(\xi-t)}\left\lfloor\frac{d}{d \xi} I^{n}(\xi)-\beta S^{\mathrm{n}}(\xi) \mathrm{I}^{\mathrm{n}}(\xi)+(\gamma+\mu) \mathrm{I}^{\mathrm{n}}(\xi)\right\rfloor d \xi$
$R^{n+1}(t)=R^{n}(t) \int_{0}^{t}-e^{(\mu+\alpha)(\xi-t)}\left\lfloor\frac{d}{d \xi} R^{n}(\xi)+(\alpha+\mu) \mathrm{R}^{\mathrm{n}}(\xi)-\gamma \mathrm{I}^{\mathrm{n}}(\xi)\right\rfloor d \xi$
The VIM, DTM and HPM algorithms are coded in the MAPLE software and the solutions are shown in Tables 1a and 1 b . The graphs are also displayed in Fig. 1, 2, and 3.
3.2 Considering the model by Namawejje et al. [2];
$\frac{d S}{d t}=\pi-(1-k) \alpha_{1} \lambda S-k \alpha_{2} \lambda S-\mu S$
$\frac{d L}{d t}=(1-k) \alpha_{1} \lambda S-\left(\varphi \alpha_{3} \lambda+\gamma\right) L-\mu L$
$\frac{d I}{d t}=k \alpha_{2} \lambda S+\left(\varphi \alpha_{3} \lambda+\gamma\right) L-(v+\mu+d) I+\alpha_{4} \lambda \omega R$
$\frac{d R}{d t}=\nu I-\alpha_{4} \lambda \omega R-\mu R$
where $\lambda=\beta c \frac{I}{N}$
subject to the initial conditions $S(0)=95000, I(0)=25000, I(0)=19500, R(0)=13500$
where the parameters $\pi$ is the recruitment rate of population, $\beta$ is the probability of getting tuberculosis infection, $\mu$ is the per capita death rate due to the disease, $\gamma$ is the progression rate of disease to active TB, $\varphi$ is the level of re-infection, $\alpha_{1}$ is the enhancement stress factor for slow progressors, $\alpha_{2}$ is the enhancement stress factor for fast progressors, $\alpha_{3}$ is the enhancement stress factor for re-infection (exogenous reactivation), $\alpha_{4}$ is the enhancement stress factor for re-infection (endogenous reactivation), c is per capita contact rate, $v$ is the recovery rate, k is the proportion from S to $\mathrm{I}, \omega$ is the rate of endogenous reactivation (from R to I ) with their values given as $\pi=2041, \alpha_{1}=1.6, \alpha_{2}=2, \alpha_{3}=1.3, \alpha_{4}=1, \beta=$ $0.35, c=2, \mu=0.02041, k=0.05, \varphi=0.06, \gamma=0.05, \omega=0.21, v=0.2, d=0.3$.
Applying the VIM to this model, we construct the correction functional as follows

$$
\begin{aligned}
& S^{n+1}(t)=S^{n}(t)+\int_{0}^{t} \lambda_{1}(\xi)\left\lfloor\frac{d}{d \xi} S^{n}(\xi)-\pi+(1-k) \alpha_{1} \beta c \frac{S I}{N}+k \alpha_{2} \beta c \frac{S I}{N}+\mu S\right\rfloor d \xi \\
& L^{n+1}(t)=L^{n}(t)+\int_{0}^{t} \lambda_{2}(\xi)\left\lfloor\frac{d}{d \xi} L^{n}(\xi)-(1-k) \alpha_{1} \beta c \frac{S I}{N}+\varphi \alpha_{3} \beta c \frac{L I}{N}+(\gamma+\mu) L\right\rfloor d \xi \\
& I^{n+1}(t)=I^{n}(t)+\int_{0}^{t} \lambda_{3}(\xi)\left\lfloor\frac{d}{d \xi} I^{n}(\xi)-k \alpha_{2} \beta c \frac{S I}{N}-\varphi \alpha_{3} \beta c \frac{L I}{N}-\gamma L+(v+\mu+d) I-\alpha_{4} \omega \beta c \frac{I R}{N}\right\rfloor d \xi \\
& R^{n+1}(t)=R^{n}(t)+\int_{0}^{t} \lambda_{4}(\xi)\left\lfloor\frac{d}{d \xi} R^{n}(\xi)-v I+\alpha_{4} \omega \beta c \frac{I R}{N}+\mu R\right\rfloor d \xi
\end{aligned}
$$

Applying DTM to the model, we get
$S(k+1)=\frac{1}{k+1}\left[\pi-(1-k) \alpha_{1} \frac{\beta c}{N}\left(\sum_{m=o}^{k} S(m) I(k-m)\right)-k \alpha_{2} \frac{\beta c}{N}\left(\sum_{m=o}^{k} S(m) I(k-m)\right)-\mu S(k)\right]$
$L(k+1)=\frac{1}{k+1}\left[(1-k) \alpha_{1} \frac{\beta c}{N}\left(\sum_{m=o}^{k} S(m) I(k-m)\right)-\varphi \alpha_{3} \frac{\beta c}{N}\left(\sum_{m=o}^{k} L(m) I(k-m)\right)-\gamma L(k)-\mu L(k)\right]$
$I(k+1)=\frac{1}{k+1}\left[k \alpha_{2} \frac{\beta c}{N}\left(\sum_{m=o}^{k} S(m) I(k-m)\right)+\varphi \alpha_{3} \frac{\beta c}{n}\left(\sum_{m=o}^{k} L(m) I(k-m)\right)+\gamma L(k)-(v+\mu+d) I(k)+\alpha_{4} \omega \frac{\beta c}{N}\left(\sum_{m=o}^{k} R(m) I(k-m)\right)\right]$
$R(k+1)=\frac{1}{k+1}\left[v I(k)-\alpha_{4} \omega \frac{\beta c}{N}\left(\sum_{m=o}^{k} R(m) I(k-m)\right)-\mu R(k)\right]$

The VIM, DTM and the HPM algorithms are codded in MAPLE software and the solutions are shown in Tables 2a, 2b and 2c. The graphs are also displayed in Fig. 4, 5, 6 and 7.

## Conclusion

Three semi-analytic methods, VIM, DTM and HPM have been used to solve two mathematical models and the results have shown that these methods are powerful mathematics tools for solving systems of Ordinary Differential Equations which appear in mathematical modelling of differential phenomena.


Fig. 1: Graph of the Solution Obtained using VIM, DTM and HPM for the Susceptible Population


Fig. 3: Graph of the Solution Obtained using VIM, DTM and HPM for the Recovered Population


Fig. 5: Graph of the Solution Obtained using VIM, DTM and HPM for the Susce Population


Fig. 2: Graph of the Solution Obtained using VIM, DTM and HPM for the Infected Population


Fig. 4: Graph of the Solution Obtained using VIM, DTM and HPM for the Susce Population


Fig. 6: Graph of the Solution Obtained using VIM, DTM and HPM for the Susce Population

Transactions of the Nigerian Association of Mathematical Physics Volume 17, (October - December, 2021), 51-60


Fig. 7: Graph of the Solution Obtained using VIM, DTM and HPM for the Susce Population
Table 2a: Solution of the Shigella diarrhea model Using VIM and DTM

| t | VIM |  |  | DTM |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | S(t) | I(t) | R(t) | S(t) | $\mathrm{I}(\mathrm{t})$ | $\mathrm{R}(\mathrm{t})$ |
| 0 | 99.9999999 | 39.99999677 | 9.999999492 | 100 | 40 | 10 |
| 0.01 | 100.1090278 | 39.94324289 | 10.02972116 | 100.1095346 | 39.94324249 | 10.02972129 |
| 0.02 | 100.2181235 | 39.88656695 | 10.05928483 | 100.2201518 | 39.88656991 | 10.05928552 |
| 0.03 | 100.3272825 | 39.82998138 | 10.08869207 | 100.3318718 | 39.82998216 | 10.08869320 |
| 0.04 | 100.4365000 | 39.77347752 | 10.11794549 | 100.4447159 | 39.77347917 | 10.11794488 |
| 0.05 | 100.5457728 | 39.71705893 | 10.14704066 | 100.5587054 | 39.71706083 | 10.14704109 |
| 0.06 | 100.6551114 | 39.66072304 | 10.17598418 | 100.6738630 | 39.66072707 | 10.17598234 |
| 0.07 | 100.7645151 | 39.60447376 | 10.20476965 | 100.7902118 | 39.60447779 | 10.20476916 |
| 0.08 | 100.8739780 | 39.54830790 | 10.23340166 | 100.9077756 | 39.54831292 | 10.23340206 |
| 0.09 | 100.983491 | 39.49222588 | 10.26187979 | 101.0265793 | 39.49223236 | 10.26188157 |
| 0.1 | 101.0930687 | 39.43623225 | 10.29021066 | 101.1466480 | 39.43623603 | 10.29020822 |

Table 2b: Solution of the Shigella diarrhea model Using DTM

| HPM | $\mathrm{I}(\mathrm{t})$ | $\mathrm{R}(\mathrm{t})$ |
| :--- | :--- | :--- |
| $\mathrm{S}(\mathrm{t})$ | 40 | 10 |
| 100 | 39.94324251 | 10.02966356 |
| 100.1089941 | 39.88657003 | 10.05905425 |
| 100.2179765 | 39.82998257 | 10.08817206 |
| 100.3269471 | 39.77348013 | 10.11701699 |
| 100.4359059 | 39.71706270 | 10.14558905 |
| 100.5448530 | 39.66073029 | 10.17388823 |
| 100.6537883 | 39.60448289 | 10.20191454 |
| 100.7627119 | 39.54832051 | 10.22966797 |
| 100.8716237 | 39.49224315 | 10.25714852 |
| 100.9805237 | 39.43625080 | 10.28435620 |
| 101.0894120 |  |  |

Table 3a: Solution of the Tuberculosis Model Using DTM

| t | DTM |  |  |  |
| :--- | :--- | :--- | :--- | :--- |
|  | $\mathrm{S}(\mathrm{t})$ | $\mathrm{I}(\mathrm{t})$ | $\mathrm{R}(\mathrm{t})$ | $\mathrm{L}(\mathrm{t})$ |
| 0 | 95000 | 19500 | 13500 | 25000 |
| 0.01 | 94791.65963 | 19433.50262 | 13530.64988 | 25175.08021 |
| 0.02 | 94584.74029 | 19367.37239 | 13561.16726 | 25348.92599 |
| 0.03 | 94379.23582 | 19301.60709 | 13591.55291 | 25521.54805 |
| 0.04 | 94175.14025 | 19236.20451 | 13621.80764 | 25692.95697 |
| 0.05 | 93972.44791 | 19171.16247 | 13651.93223 | 25863.16321 |
| 0.06 | 93771.15328 | 19106.47879 | 13681.92745 | 26032.17721 |
| 0.07 | 93571.25117 | 19042.15132 | 13711.79408 | 26200.00921 |
| 0.08 | 93372.73658 | 18978.17794 | 13741.53286 | 26366.66943 |
| 0.09 | 93175.60478 | 18914.55650 | 13771.14459 | 26532.16795 |
| 0.1 | 92979.85129 | 18851.28490 | 13800.63001 | 26696.51482 |

Table 3b: Solution of the Tuberculosis Model Using HPM

| HPM |  |  |  |
| :--- | :--- | :--- | :--- |
| $\mathrm{S}(\mathrm{t})$ | $\mathrm{I}(\mathrm{t})$ | $\mathrm{R}(\mathrm{t})$ | $\mathrm{L}(\mathrm{t})$ |
| 95000 | 19500 | 13500 | 25000 |
| 94791.54439 | 19433.51028 | 13530.64975 | 25175.07840 |
| 94584.28355 | 19367.40452 | 13561.16619 | 25348.91161 |
| 94378.21748 | 19301.68272 | 13591.54932 | 25521.49963 |
| 94173.34618 | 19236.34489 | 13621.79915 | 25692.84245 |
| 93969.66966 | 19171.39101 | 13651.91567 | 25862.94007 |
| 93767.18791 | 19106.82109 | 13681.89888 | 26031.79251 |
| 93565.90093 | 19042.63514 | 13711.74879 | 26199.39975 |
| 93365.80873 | 18978.83314 | 13741.46538 | 26365.76179 |
| 93166.91130 | 18915.41511 | 13771.04868 | 26530.87864 |
| 92969.20864 | 18852.38104 | 13800.49866 | 26694.75030 |

Table 3c: Solution of the Tuberculosis Model Using VIM

| t | VIM |  |  |  |
| :--- | :--- | :--- | :--- | :--- |
|  | $\mathrm{S}(\mathrm{t})$ | $\mathrm{I}(\mathrm{t})$ | $\mathrm{R}(\mathrm{t})$ | $\mathrm{L}(\mathrm{t})$ |
| 0 | 95000 | 19500 | 13500 | 25000 |
| 0.01 | 94791.55697 | 19433.50261 | 13530.64988 | 25175.08014 |
| 0.02 | 94584.32723 | 19367.37235 | 13561.16726 | 25348.92543 |
| 0.03 | 94378.30081 | 19301.60696 | 13591.55291 | 25521.54614 |
| 0.04 | 94173.46787 | 19236.20420 | 13621.80764 | 25692.95242 |
| 0.05 | 93969.81868 | 19171.16188 | 13651.93223 | 25863.15431 |
| 0.06 | 93767.34359 | 19106.47778 | 13681.92744 | 26032.16178 |
| 0.07 | 93566.03307 | 19042.14971 | 13711.79406 | 26199.98464 |
| 0.08 | 93365.87765 | 18978.17551 | 13741.53285 | 26366.63266 |
| 0.09 | 93166.86801 | 18914.55305 | 13771.14457 | 26532.11546 |
| 0.1 | 92968.99484 | 18851.28015 | 13800.62997 | 26696.44263 |

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