

THOMAS DIERKES

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by a highly flexible modelling approach**

Herausgegeben vom
Konrad-Zuse-Zentrum für Informationstechnik Berlin
Takustraße 7
D-14195 Berlin-Dahlem

Telefon: 030-84185-0
Telefax: 030-84185-125

e-mail: bibliothek@zib.de
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Construction of ODE systems from time series data by a highly flexible modelling approach

T Dierkes

Konrad-Zuse-Zentrum für Informationstechnik Berlin (ZIB)
Takustraße 7, 14195 Berlin-Dahlem, Germany

E-mail: dierkes@zib.de

Abstract. In this paper, a down-to-earth approach to purely data-based modelling of unknown dynamical systems is presented. Starting from a classical, explicit ODE formulation $y' = f(t, y)$ of a dynamical system, a method determining the unknown right-hand side $f(t, y)$ from some trajectory data $y_k(t_j)$, possibly very sparse, is given. As illustrative examples, a semi-standard predator-prey model is reconstructed from a data set describing the population numbers of hares and lynxes over a period of twenty years [1], and a simple damped pendulum system with a highly non-linear right-hand side is recovered from some artificial but very sparse data [2].

AMS classification scheme numbers: 65L09, 37M10, 92C42, 92D25

1. Introduction

Modelling of dynamical systems is one of the key topics not only in systems biology, but also in other disciplines [3, 4]. Most often, a desired behaviour of a dynamical system or, at least, parts of it, has already been anticipated. In this paper we will present a down-to-earth approach to determine a completely unknown dynamical system from given, possibly very sparse, data sets; yet open enough for easy incorporation of additional knowledge about the dynamical system to be considered. Previous work related to ours include [2, 5] and [6], describing a software package called **Data2Dynamics**. In the former two papers, a trajectory method is presented that first replaces the right-hand side of the ODE by a linear combination of a family of known functions, and tries then to find the coefficients of the linear combination by fitting the trajectory solutions to the given data. The title of the third paper, “Data2Dynamics : a modeling environment ...”, is a classical misnomer: The complete dynamical model has already to be known in terms of an ODE system, only a possible control function for the ODE appears in their method as variable and unknown and could, in principle, be estimated, simultaneously with the classical fit approach, from the given data. In contrast to all three cited papers, our approach is slightly different, yet quite robust and fast and, therefore, extremely versatile, as will be explained in this paper.

The dynamical system is supposed to be given as an explicit ODE,

$$y' = f(t, y)$$

where the dash ' represents the derivative w.r.t. the time variable $t \in \mathbb{R}$ of the solution trajectory, $y = y(t)$, $y \in \mathbb{R}^n$. The right-hand side, $f : \mathbb{R} \times \mathbb{R}^n \rightarrow \mathbb{R}^n$, presumably smooth enough, has to be determined from data sampled of trajectory solutions, $y_k(t_{k,j})$, to this ODE, for $k \in \{1, \dots, n\}$ and for some discrete time points $t_{k,1} < t_{k,2} < t_{k,3} < \dots < t_{k,m}$. Note that the sample time points can totally be asynchronous.

Since any non-autonomous system is readily transformed to an equivalent autonomous system, w.l.o.g. we can assume f to be independent of the time variable t ,

$$f(t, y) = f(y).$$

If a trajectory component, y_k , has no sample points, we have to stipulate some values for that component describing a desired behaviour of the dynamical system in this component. The general idea we have in mind here, is that only a subsystem of a *known*, but apart from that, parametrised dynamical system,

$$y' = f(y; p)$$

with a fixed parameter vector $p \in \mathbb{R}^q$ is unknown, i.e. only some of the solution trajectory components are given by sampled measurement points, or else by values describing a new, desired evolution of the dynamical system under investigation.

2. Modelling Approach

Having $m \in \mathbb{N}$ sample points of each solution trajectory component $y_{k,j} := y_k(t_{k,j})$, i.e. nm data values, of the unknown ODE system

$$y' = f(y) \tag{2.1}$$

the basic idea of our approach to find f from the given data is to first construct a suitable approximation $y^*(t)$ to the data points $y_{k,j}$ for that the derivative can readily be computed as well. Then, we can construct a (discrete) mapping

$$F : y^* \mapsto (y^*)'$$

with arbitrary many samples over the interval $[t_{min}, t_{max}]$, determined by the range of the sample time points of the given data.

From this, by making use of a set of ansatz functions $\{\varphi_\ell, \ell = 0, 1, \dots\}$ over \mathbb{R}^n (or $[a, b]^n$, if bounds of the components y_k are available), it is possible to reconstruct the unknown function $f : \mathbb{R}^n \rightarrow \mathbb{R}^n$ by linear, or even non-linear, combinations of the φ_ℓ : For as many $t_j \in [t_{min}, t_{max}]$ as necessary, in a linear case, we eventually obtain

$$(y_k^*)'(t_j) = \sum_{\ell=0}^L c_\ell^k \varphi_\ell(y^*(t_j)), \quad k = 1, \dots, n \tag{2.2}$$

where, for each $k \in \{1, \dots, n\}$, the c_ℓ^k , $\ell = 0, \dots, L$, are coefficients to be determined in the least-squares sense. Note that the right-hand side depends on all n trajectory components whereas the left-hand side includes only the k -th component.

The solution of (2.2) finally yields the description of the unknown f in terms of the (deliberately chosen) ansatz functions $\varphi_\ell(\cdot)$.

2.1. Best Polynomial Approximation

Considering the set

$$\mathcal{P}_N = \{ \text{polynomials on } [-1, 1] \text{ of degree at most } N \}$$

it is well-known that for a function $g(t)$, Lipschitz continuous on $[-1, 1]$, it is very difficult to find its corresponding *minimax polynomial*, i.e. the polynomial $p^*[g](t)$ that has the smallest maximum deviation from $g(t)$, see [7],

$$\|g - p^*[g]\|_\infty = \min_{p \in \mathcal{P}_N} \|g - p\|_\infty.$$

As it turns out, most remarkably, the corresponding Chebyshev series to g , which is absolutely and uniformly convergent,

$$g(t) = \sum_{k=0}^{\infty} a_k T_k(t)$$

yields very nearly the same polynomial of degree N as the *minimax polynomial* $p^*[g]$ when truncated at the N -th term,

$$p^*[g](t) \approx \sum_{k=0}^N a_k T_k(t).$$

Here, $T_k(t)$ denotes the k -th Chebyshev polynomial, defined as real part of the complex function z^k on the unit circle (see, e.g., [8]),

$$T_k(t) = \operatorname{Re} z^k = \frac{1}{2}(z^k + \bar{z}^k) = \cos(k\theta), \quad t = \cos \theta, \quad \text{and} \quad |z| = 1,$$

and the coefficients a_k are given for $k \geq 1$ by the formula

$$a_k = \frac{2}{\pi} \int_{-1}^1 \frac{g(t) T_k(t)}{\sqrt{1-t^2}} dt \tag{2.3}$$

and for $k = 0$ by the very same formula with $2/\pi$ changed to $1/\pi$.

The discrete analogue to (2.3), when $g(t)$ is only given at the $M + 1$ Chebyshev nodes

$$t_j = \cos \theta_j, \quad \theta_j = \frac{\pi(j + 1/2)}{M + 1}, \quad j = 0, \dots, M,$$

reads [9, 7]

$$\tilde{a}_k = \frac{2}{M+1} \sum_{j=0}^M g(t_j) T_k(t_j) \quad (2.4)$$

$$= \frac{2}{M+1} \sum_{j=0}^M g \left(\cos \frac{\pi(j+1/2)}{M+1} \right) \cos \frac{\pi k(j+1/2)}{M+1}. \quad (2.5)$$

In the very likely case that the sample time points t_j of the given data do *not* coincide with the Chebyshev nodes $\cos \theta_j$, we can use a simple linear interpolation scheme to transform the given data values to the required Chebyshev nodes. An even more simple table lookup procedure would in some cases do equally well. In the case of very rough data, alternatively, we could think of an additional smoothing step by making use of a suitable spline interpolation where the data points are the control points of the spline interpolation scheme, for example [10]. Having such a smooth interpolation of the given data at hand, values at the desired Chebyshev nodes are then readily available.

Thus, as building block for the first step, we arrive at the discrete Chebyshev approximation to g of order M ,

$$g(t) \approx \frac{1}{2} \tilde{a}_0 + \sum_{k=1}^M \tilde{a}_k T_k(t), \quad (2.6)$$

evidently a polynomial of degree M . Moreover, the derivative of $g(t)$ can readily be approximated in turn,

$$g'(t) \approx \frac{1}{2} \tilde{a}'_0 + \sum_{k=1}^M \tilde{a}'_k T_k(t), \quad (2.7)$$

where the new coefficients \tilde{a}'_k are given by the simple backwards recurrence formula

$$\begin{aligned} \tilde{a}'_{M+1} &= \tilde{a}'_M = 0, \\ \tilde{a}'_{k-1} &= \tilde{a}'_{k+1} + 2k \tilde{a}_k, \quad k = M, M-1, \dots, 1. \end{aligned}$$

Now, making use of (2.6) and (2.7), we can readily set up the discrete mapping $F : y^* \mapsto (y^*)'$ from the data $y_{k,j}$ by generating one separate Chebyshev approximation for each component y_k . Depending on the smoothness of the given data, the maximal degree M of these approximations can be adjusted accordingly that, as an intriguing side effect, comes in handy as a cheap filtering device in this step.

2.2. Least-squares Solution: Linear and Non-linear Case

For each $k \in \{1, \dots, n\}$, equation (2.2) can be rewritten as

$$b^{(k)} = A c^{(k)} \quad (2.8)$$

where the m -vector $b^{(k)}$ and the $(m \times (L+1))$ -matrix A (sometimes known as *Gram matrix*) are given respectively by

$$\begin{aligned} b_j^{(k)} &:= (y_k^*)'(t_j), & j &= 1, \dots, m \\ A_{j,\ell} &:= \varphi_\ell \left(y_1^*(t_j), \dots, y_n^*(t_j) \right), & \ell &= 0, \dots, L. \end{aligned}$$

Here, as explained in the previous section, the approximations $y_k^*(\cdot)$ and its derivatives are evaluated in terms of their truncated Chebyshev series of order M ,

$$y_k^*(t) = \left[\sum_{j=0}^M \tilde{y}_{k,j} T_j(t) \right] - \frac{1}{2} \tilde{y}_{k,0}$$

with the discrete Chebyshev coefficients $\tilde{y}_{k,j}$ computed by (2.4) with the given $(n \times m)$ -data matrix $(y_{k,j})$.

The solution $(c^{(k)})^*$, in a least-squares sense, to the linear system (2.8) is readily obtained by making use of the QR-decomposition $A = QR P^T$ with pivoting, P being a permutation matrix,

$$(c^{(k)})^* = P \tilde{R}^{-1} \tilde{Q}^T b^{(k)}, \quad R = \begin{bmatrix} \tilde{R} \\ 0 \end{bmatrix}, \quad Q = \begin{bmatrix} \tilde{Q} & * \end{bmatrix},$$

$$\tilde{R} \in \mathbb{R}^{(L+1) \times (L+1)}, \quad \tilde{Q} \in \mathbb{R}^{m \times (L+1)},$$

provided the *Gram matrix* A has full rank, and $m \geq L + 1$. If one or both of these conditions are violated, there are adequate formulae available as well, see [11].

In the not so unlikely case, some (or all) of the unknown coefficients c^k enter *non-linearly* the ansatz function family $\{\varphi_\ell\}$, we have to resort to an iterative scheme, such as Gauss-Newton, in order to get a decent estimate of $(c^k)^*$. For the details we refer to [12].

2.3. Verification

Up to now, we have not touched any of the dynamical properties of the underlying, but unknown ODE system (2.1), with the only exception of (2.7) in determining the left-hand side $b^{(k)}$ in (2.8). Consequently, we can take advantage of this fact in the following manner if we want to verify the resulting solution of the previous section.

With the least-squares solutions $(c^{(k)})^*$, $k = 1, \dots, n$, at hand, it is possible to compute numerically the solution $y^b(t)$ to the initial value problem of the approximated ODE,

$$y' = \sum_{\ell=0}^L c_\ell \varphi_\ell(y), \quad y(t_0) = (y_{1,1}, \dots, y_{n,1})^T, \quad (2.9)$$

$$c_\ell = \left(c_\ell^{(1)}, \dots, c_\ell^{(n)} \right)^T \quad (2.10)$$

provided all sample time points $t_{k,j}$ start exact synchronously. For this numerical task any suitable integration scheme can be applied, i.e., in most cases, a decent one-step Runge-Kutta scheme or, alternatively, a linearly implicit scheme (e.g. LIMEX [13, 14]) will do.

If the discrepancy between the approximated solution $y^b(\cdot)$ and the given data $y_k(t_{k,j})$ is satisfactory we could say the unknown model has successfully been verified.

Or else, if the discrepancy is too high, we could invoke yet another parameter identification run: This time with the recovered ODE system (2.9), and the unknown coefficients c_ℓ as parameters with an obvious starting guess.

If all else fails, another ansatz function family has to be chosen, or the filtering parameters have to be adjusted differently, or the problem at hand needs just more (measurement) data.

3. Numerical Results and Discussion

In this section, we will study a predator-prey model and a damped pendulum model. For the ansatz function family we choose in both cases multi-variate polynomials,

$$\varphi_\ell(x) := \varphi_{(\ell_1, \dots, \ell_n)}(x_1, \dots, x_n) := x_1^{\ell_1} \cdot x_2^{\ell_2} \cdot \dots \cdot x_n^{\ell_n} \quad (3.1)$$

where $\ell = (\ell_1, \dots, \ell_n)$ denotes a multi-index with the usual meaning, as indicated. Consequently, for a polynomial of maximal degree d , i.e. $|\ell| = \ell_1 + \dots + \ell_n \leq d$, there are

$$\binom{n+d}{d} = \binom{n+d}{n}$$

different monomial terms $\varphi_\ell(\cdot)$ with the determining coefficients $c_\ell \in \mathbb{R}$, specifying uniquely the Gram matrix A in (2.8).

The runtime protocols for each component of the example computations, that will be given below, have the following structure. In the first and second column, two different representations of the multi-index are listed (of which the one in the second column is the more familiar notation). In the third column the percentage taken with respect to the absolute maximum of the coefficients c_ℓ found is given, and finally, in the last column, the actual value of each coefficient is printed. Additionally, the resulting equation for the investigated component is recorded in which only monomial terms are included according to the indicated percentage threshold. Last but not least, the weighted ℓ_2 -norm of the residual of the least-squares equation (2.8) is displayed.

3.1. Hare and Lynx

Commonly, the dynamics of a predator-prey system is modelled by Lotka-Volterra-type equations [1],

$$y'_{\text{prey}} = y_{\text{prey}} (\alpha - \beta y_{\text{pred}}), \quad (3.2)$$

$$y'_{\text{pred}} = -y_{\text{pred}} (\gamma - \delta y_{\text{prey}}) \quad (3.3)$$

where y_{prey} and y_{pred} denote the population number of the prey and the predator, respectively. Here, the coupling constants $\alpha, \beta, \gamma, \delta \in \mathbb{R}_{\geq 0}$ describe the influence of the species on each other. We want to recover such a system from some historical data of numbers of lynxes (predator) and hares (prey), collected by the Hudson Bay

Company in Canada during the years 1900–1920. This data set can be seen, more or less, as representative for the total population number of both species [1].

For basic biological reasons a constant term is missing in both components of the anticipated predator-prey model (3.2)–(3.3). Hence, we exclude the constant monomial term in both reconstruction runs as well. The maximal degree of the multi-variate polynomials is set to 3.

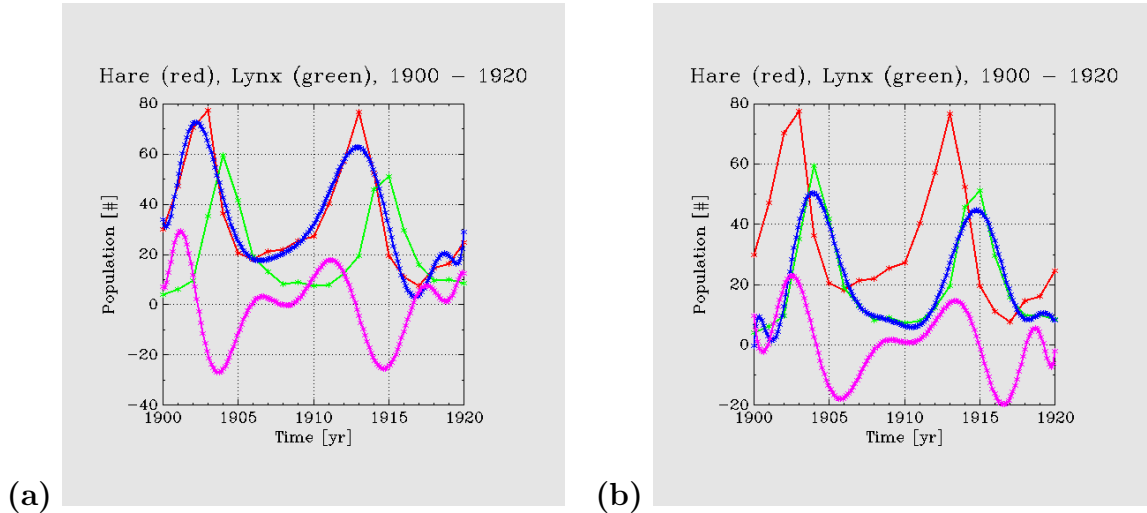


Figure 1. $T_n(x)$ approximations (blue) and its derivative (magenta) to (a) hare data, and (b) lynx data ($n=80$, evaluation of the approximations with $n=11$ each).

Runtime protocol for $y_0 = y_{\text{prey}}$ component.

```
#total = 16 (max. deg. 3)
[ 0 2 ] --> [0, 1] 100.00 % c_k = 8.999962e-01
[ 0 3 ] --> [0, 2] 3.79 % c_k = -3.408105e-02
[ 0 4 ] --> [0, 3] 0.00 % c_k = 1.703492e-04
[ 1 2 ] --> [1, 0] 19.66 % c_k = -1.769268e-01
[ 1 3 ] --> [1, 1] 3.44 % c_k = -3.092972e-02
[ 1 4 ] --> [1, 2] 0.00 % c_k = 3.976574e-04
[ 2 3 ] --> [2, 0] 2.65 % c_k = 2.384418e-02
[ 2 4 ] --> [2, 1] 0.00 % c_k = -4.353977e-05
[ 3 4 ] --> [3, 0] 0.00 % c_k = -2.065367e-04
m = 9 monomial(s) 0.10 % threshold
f(y0,y1) = + ( 9.0e-01) y1^1 + (-3.4e-02) y1^2 + (-1.8e-01) y0^1
           + (-3.1e-02) y0^1 y1^1 + ( 2.4e-02) y0^2
LSQ: ||residual/sqrt(n)||_2 = 3.095545072754599
```

The y_{prey} is found to be described by the equation

$$y_0' = +y_0((ay_0 - b) - cy_1) + dy_1 - ey_1^2 \quad (3.4)$$

with $a = 0.024$, $b = 0.18$, $c = 0.031$, $d = 0.9$, and $e = 0.034$.

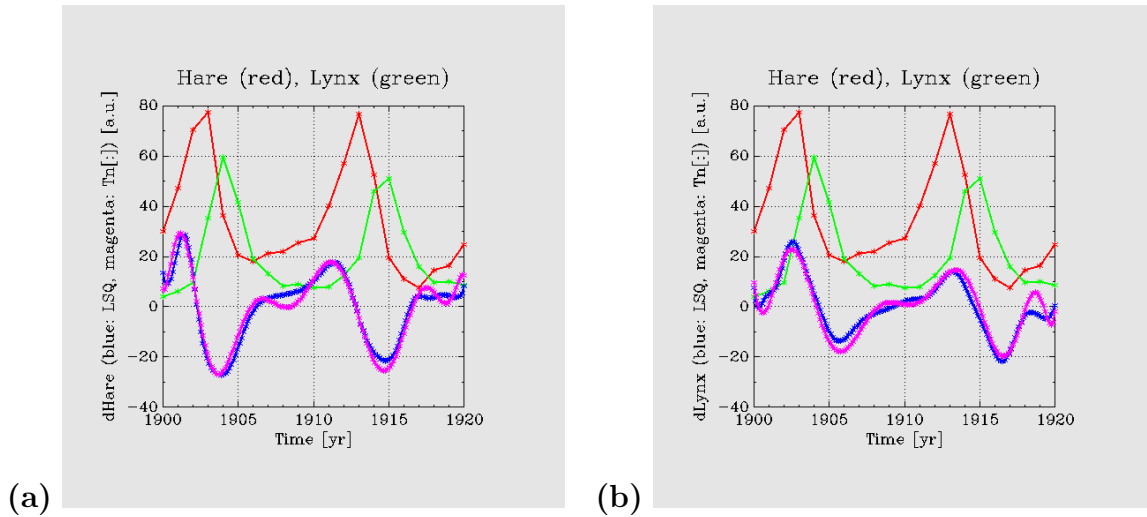


Figure 2. Least-squares solution (blue) with $(c^{(k)})^*$, $k = 1, 2$, and degree = 3 to the unknown $F : y^* \mapsto (y^*)'$ for (a) hare data, and (b) lynx data (magenta each, and evaluation with 150 equidistant time point each).

Runtime protocol for $y_1 = y_{\text{pred}}$ component.

```
#total = 16 (max. deg. 3)
[ 0 2 ] --> [0, 1] 100.00 % c_k = -1.590145e+00
[ 0 3 ] --> [0, 2] 1.35 % c_k = 2.146557e-02
[ 0 4 ] --> [0, 3] 0.00 % c_k = 1.283903e-04
[ 1 2 ] --> [1, 0] 45.70 % c_k = 7.266958e-01
[ 1 3 ] --> [1, 1] 0.87 % c_k = 1.379691e-02
[ 1 4 ] --> [1, 2] 0.00 % c_k = -4.190196e-04
[ 2 3 ] --> [2, 0] 1.35 % c_k = -2.140643e-02
[ 2 4 ] --> [2, 1] 0.00 % c_k = 2.619544e-04
[ 3 4 ] --> [3, 0] 0.00 % c_k = 1.834174e-04
m = 9 monomial(s) 0.10 % threshold
f(y0,y1) = + (-1.6e+00) y1^1 + ( 2.1e-02) y1^2 + ( 7.3e-01) y0^1
           + ( 1.4e-02) y0^1 y1^1 + (-2.1e-02) y0^2
LSQ: ||residual/sqrt(n)||_2 = 3.1046555639780498
```

Here, the y_{pred} component is found to be given by the equation

$$y_1' = -y_1 (a - b y_0) + c y_1^2 + d y_0 - e y_0^2 \quad (3.5)$$

with $a = 1.6$, $b = 0.014$, $c = 0.021$, $d = 0.73$, and $e = 0.021$.

It would be interesting to see if an additional, iterative Gauss-Newton fit would improve the identified parameters further and, possibly, establish some link to the standard predator-prey model (3.2)–(3.3). Unfortunately, for the case with multi-variate polynomials of maximal degree 3, as presented here, such an additional verification has not been successful at all. Instead, the Gauss-Newton iteration stops after a few steps with the ODE system being not integrable any more, i.e. the Newton path of the coefficient sets during the iteration inexplicably leads to an ODE system that can not be solved numerically. A systematic and comprehensive investigation of this unexpected behaviour would certainly go beyond the scope of this article and hence, must be left open for now.

However, a repeated computation, completely analogous to the one presented here, only with the maximal polynomial degree restricted to 2, nicely demonstrates a relationship to the standard predator-prey model, as we have previously gathered, see Table 1. In particular, there is an intriguing match between the coefficients of the second component, i.e. the ODE equation of the lynx. The incompatibility factor in the last row of Table 1 is determined by the ratio of the norms of successive updates during the Gauss-Newton iteration and thus, can be seen as an estimate of the convergence speed, especially at the end of the iteration. This factor, if strictly below 1.0, can be thought to measure how compatible a model to given data is (see [12, sec. 4.3.2] for the mathematical background). The values in the last column of Table 1 can be found in [1]. All the computational details are given in the Appendix.

Table 1. Identified coefficients of the standard model [1] and our data-based approach.

Component	Multi-index	Data-based	Standard
$y'_0 = f_{\text{prey}}(y_0, y_1)$	[0, 1]	1.848730e-01	
	[0, 2]	3.675366e-03	
	[1, 0]	2.531473e-01	5.475337e-01 (= α)
	[1, 1]	-3.789005e-02	-2.811932e-02 (= $-\beta$)
	[2, 0]	7.856857e-03	
$y'_1 = f_{\text{pred}}(y_0, y_1)$	[0, 1]	-9.587181e-01	-8.431750e-01 (= $-\gamma$)
	[0, 2]	6.845210e-04	
	[1, 0]	5.296776e-02	
	[1, 1]	2.408610e-02	2.655759e-02 (= δ)
	[2, 0]	-1.113486e-04	
Scaled residual (Normf)		3.0712e-00	4.2362e-00
κ (incomp. factor, see [12])		1.0274e-01	2.9798e-02

To sum up the hare and lynx case, the reconstructed components y_0 and y_1 seem to follow only partly the standard predator-prey model. Instead, our findings here show that the data might be more involved than previously assumed. Our conclusion is also supported by the verification result, as can be seen in Figure 3.

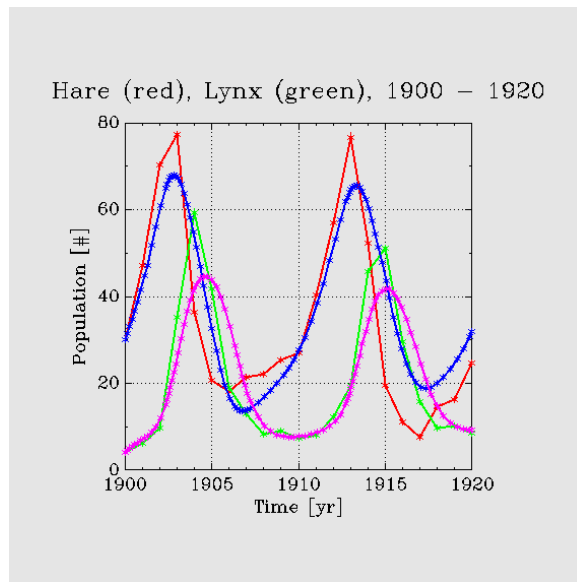


Figure 3. Comparison between given data (red, green) and solution (blue, magenta) to the approximated ODE with multi-variate polynomials of maximal degree 3, as shown in Fig. 2 (by using the adaptive time stepping integrator LIMEX).

3.2. Damped Pendulum

Here, we consider the initial value problem

$$\theta_1' = \theta_2, \quad (3.6)$$

$$\theta_2' = -u\theta_2 - \frac{g}{l} \sin \theta_1 \quad (3.7)$$

with the initial condition $(\theta_1, \theta_2)(0) = (1, 0)$, and the constants $u = 0.25$, $l = 2.0$, and $g = 9.81$ (the gravitational acceleration on earth) are used [2]. Very sparse data has been generated by solving numerically (3.6)–(3.7) in the time interval $[0, 10]$, and subsequently taking 49 equidistant samples of the solution trajectories in this interval.

In order to be able to catch a glimpse of the non-linearity on the right-hand side in (3.7), for the reconstruction trial the maximal degree of the multi-variate polynomials is set to 4.

As seen in the following runtime protocol for the θ_1 component, the equation

$$y_0' = a y_1 \quad (3.8)$$

with $a = 1.0$ is found.

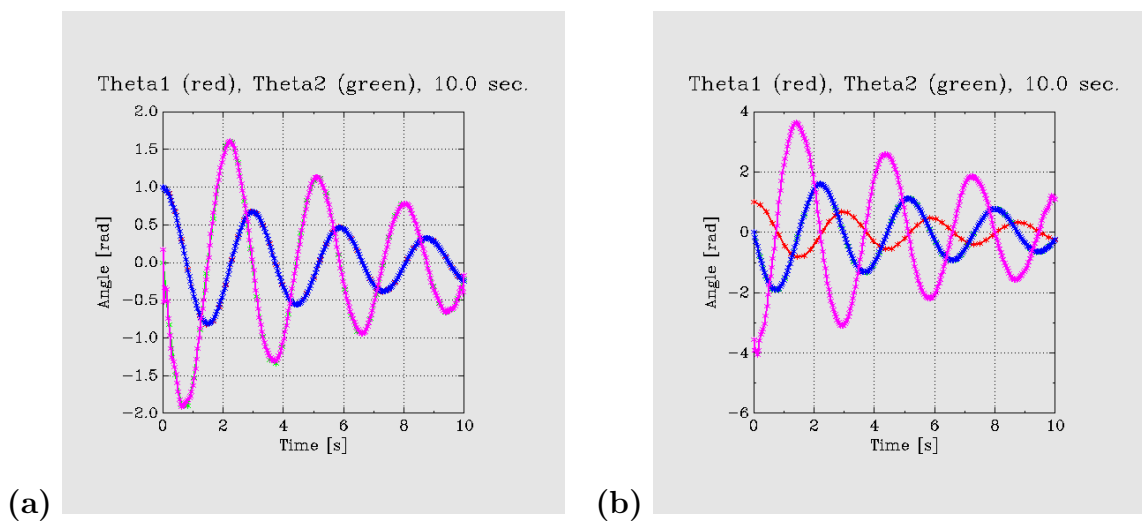


Figure 4. $T_n(x)$ approximations (blue) and its derivative (magenta) to (a) θ_1 data, and (b) θ_2 data ($n=80$, evaluation of the approximations with $n=62$ each).

Runtime protocol for $y_0 = \theta_1$ component.

```
#total = 25 (max. deg. 4)
[ 0 1 ] --> [0, 0] 0.00 % c_k = 2.001792e-03
[ 0 2 ] --> [0, 1] 100.00 % c_k = 9.982190e-01
[ 0 3 ] --> [0, 2] 0.00 % c_k = -6.094577e-03
[ 0 4 ] --> [0, 3] 0.00 % c_k = -2.859434e-04
[ 0 5 ] --> [0, 4] 0.00 % c_k = 1.027003e-03
[ 1 2 ] --> [1, 0] 0.00 % c_k = 1.384047e-03
[ 1 3 ] --> [1, 1] 0.00 % c_k = 2.514082e-02
[ 1 4 ] --> [1, 2] 0.00 % c_k = 4.947748e-03
[ 1 5 ] --> [1, 3] 0.00 % c_k = -5.810996e-03
[ 2 3 ] --> [2, 0] 0.00 % c_k = -2.636188e-02
[ 2 4 ] --> [2, 1] 0.00 % c_k = 3.051582e-02
[ 2 5 ] --> [2, 2] 0.00 % c_k = 2.399808e-02
[ 3 4 ] --> [3, 0] 0.00 % c_k = -2.561132e-03
[ 3 5 ] --> [3, 1] 0.00 % c_k = -1.691789e-02
[ 4 5 ] --> [4, 0] 0.00 % c_k = 3.776657e-02
m = 15 monomial(s) 5.00 % threshold
f(y0,y1) = + ( 1.0e+00) y1^1
LSQ: ||residual/sqrt(n)||_2 = 0.035592552825152
```

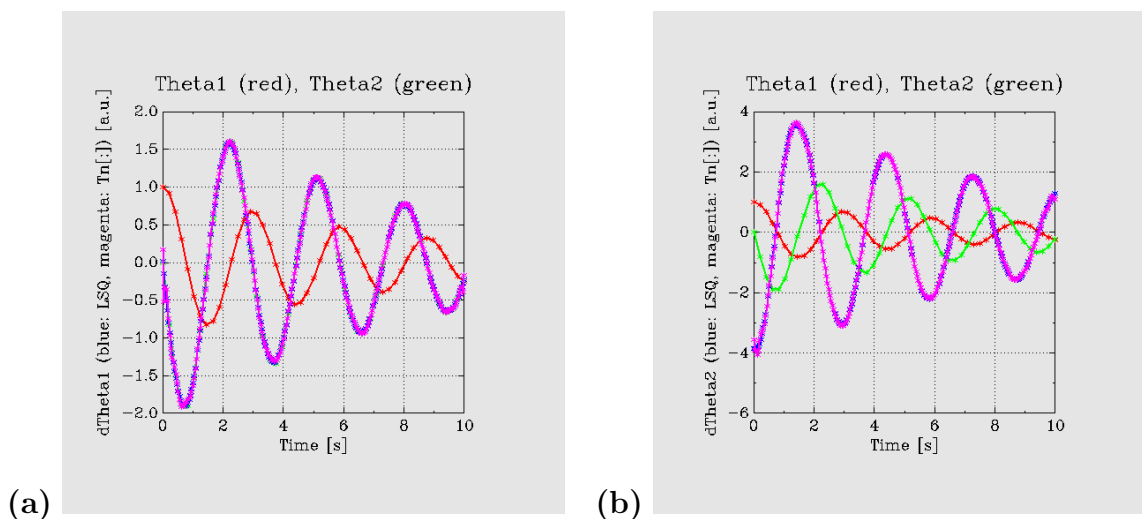


Figure 5. Least-squares solution (blue) with $(c^{(k)})^*$, $k = 1, 2$, and degree = 4 to the unknown $F : y^* \mapsto (y^*)'$ for (a) θ_1 data, and (b) θ_2 data (magenta each, and evaluation with 250 equidistant time point each).

Runtime protocol for $y_1 = \theta_2$ component.

```
#total = 25 (max. deg. 4)
[ 0 1 ] --> [0, 0] 0.00 % c_k = 5.555385e-03
[ 0 2 ] --> [0, 1] 5.13 % c_k = -2.535032e-01
[ 0 3 ] --> [0, 2] 0.00 % c_k = -1.796957e-02
[ 0 4 ] --> [0, 3] 0.00 % c_k = -6.444822e-03
[ 0 5 ] --> [0, 4] 0.00 % c_k = -3.850649e-04
[ 1 2 ] --> [1, 0] 100.00 % c_k = -4.942789e+00
[ 1 3 ] --> [1, 1] 0.00 % c_k = -1.564249e-01
[ 1 4 ] --> [1, 2] 0.00 % c_k = 2.928157e-02
[ 1 5 ] --> [1, 3] 0.00 % c_k = 1.852611e-02
[ 2 3 ] --> [2, 0] 0.00 % c_k = -9.883681e-02
[ 2 4 ] --> [2, 1] 0.00 % c_k = 1.441293e-01
[ 2 5 ] --> [2, 2] 0.00 % c_k = 1.787311e-01
[ 3 4 ] --> [3, 0] 19.50 % c_k = 9.639487e-01
[ 3 5 ] --> [3, 1] 9.90 % c_k = 4.893068e-01
[ 4 5 ] --> [4, 0] 0.00 % c_k = 2.128485e-01
m = 15 monomial(s) 5.00 % threshold
f(y0,y1) = + (-2.5e-01) y1^1 + (-4.9e+00) y0^1 + ( 9.6e-01) y0^3
          + ( 4.9e-01) y0^3 y1^1
LSQ: ||residual/sqrt(n)||_2 = 0.04531770760243261
```

In this case, for the θ_2 component, we obtain

$$y_1' = -a y_1 - b y_0 + c y_0^3 + d y_0^3 y_1 \quad (3.9)$$

with $a = 0.25$, $b = 4.9$, $c = 0.96$, and $d = 0.49$.

Summarising these findings, the recovered equations in this non-linear case agree convincingly well with the pendulum model that has been used to generate the given sparse data set. Especially for the second component $y_1 = \theta_2$, the reconstructed coefficients $b = 4.943 \sim g/l = 4.905$ and $c = 0.9639 \sim -g/6l = 0.8175$ match the Taylor coefficients of the sine in the right-hand side of the original damped pendulum model. The discrepancy in c is compensated by the additional coefficient d , indicating that the approximated model is still not complete. Nevertheless, the verification result, as can be seen clearly in Figure 6, is already almost perfect.

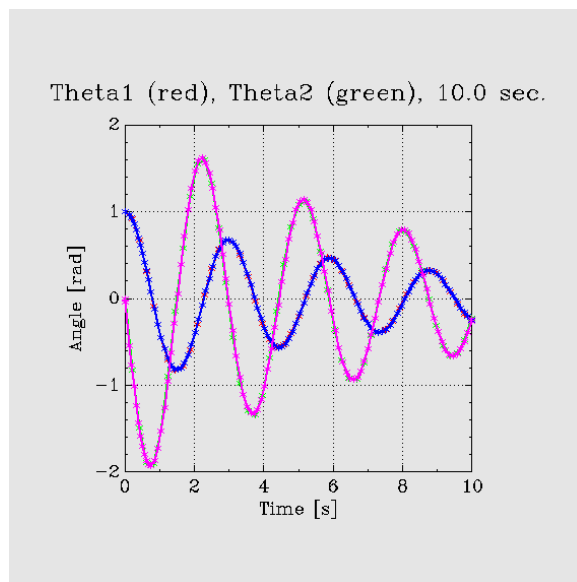


Figure 6. Comparison between given data (red, green) and solution (blue, magenta) to the approximated ODE with multi-variate polynomials of maximal degree 4, as shown in Fig. 5 (by using the adaptive time stepping integrator LIMEX).

4. Conclusion

In the present paper, a highly flexible method for the reconstruction of an unknown dynamical system, to be described in terms of an explicit ODE system, is presented. The reconstruction method is based on sampled, possibly sparse, trajectory solution points as given data.

In principle, any kind of relationship between the available information is tried to be recovered automatically, if different components of the unknown system are given. Yet, because of the extreme simple structure of the presented method, and the fact that

the ODE itself is not being solved during the reconstruction, it is possible to readily adapt the approach to any anticipated behaviour of the unknown dynamical system. Moreover, the approach is fast and, most importantly, highly reliable.

Additionally, the results of our fast reconstruction method could be used as a very good starting guess, for example, of more sophisticated iterative identification procedures such as well-established Gauss-Newton codes [12].

Acknowledgments

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Appendix

Here, we repeat the computations of section 3.1 for the predator-prey case, only the maximal degree of the multi-variate polynomials is set to 2. This results in the following two runtime protocols.

Runtime protocol for $y_0 = y_{\text{prey}}$ component.

```
#total = 9 (max. deg. 2)
[ 0 2 ] --> [0, 1]    7.72 %   c_k = -3.591668e-02
[ 0 3 ] --> [0, 2]    0.93 %   c_k = -4.319190e-03
[ 1 2 ] --> [1, 0]   100.00 %  c_k =  4.653563e-01
[ 1 3 ] --> [1, 1]    3.58 %   c_k = -1.665442e-02
[ 2 3 ] --> [2, 0]    0.00 %   c_k = -9.424325e-05
m = 5 monomial(s)      0.10 %   threshold
f(y0,y1) = + (-3.6e-02) y1^1 + (-4.3e-03) y1^2 + ( 4.7e-01) y0^1
           + (-1.7e-02) y0^1 y1^1
LSQ: ||residual/sqrt(n)||_2 = 3.7658055884421944
```

Runtime protocol for $y_1 = y_{\text{pred}}$ component.

```
#total = 9 (max. deg. 2)
[ 0 2 ] --> [0, 1]   100.00 %  c_k = -9.800874e-01
[ 0 3 ] --> [0, 2]    0.68 %   c_k =  6.673957e-03
[ 1 2 ] --> [1, 0]    7.26 %   c_k =  7.117392e-02
[ 1 3 ] --> [1, 1]    1.52 %   c_k =  1.487619e-02
[ 2 3 ] --> [2, 0]    0.16 %   c_k =  1.545017e-03
m = 5 monomial(s)      0.10 %   threshold
f(y0,y1) = + (-9.8e-01) y1^1 + ( 6.7e-03) y1^2 + ( 7.1e-02) y0^1
           + ( 1.5e-02) y0^1 y1^1 + ( 1.5e-03) y0^2
LSQ: ||residual/sqrt(n)||_2 = 3.666500740586521
```

Taking these coefficients c_ℓ^{prey} and c_ℓ^{pred} , $|\ell| \leq 2$, as starting values for a non-linear least squares method, in order to identify parameters of the ODE model build by these two polynomials

$$y_0' = \sum_{|\ell| \leq 2} c_\ell^{\text{prey}} y^\ell, \quad (\text{A.1})$$

$$y_1' = \sum_{|\ell| \leq 2} c_\ell^{\text{pred}} y^\ell \quad (\text{A.2})$$

with initial conditions $y_0(1900) = 30$ and $y_1(1900) = 4$ kept fixed, will result in a runtime protocol as follows, trying to match the given measurement data best in the

least squares sense by successively varying these 10 coefficients. Here, we choose an error-oriented Gauss-Newton scheme with adaptive trust region and rank strategy, NLSCON, as fitting routine. Solutions to the IVP of (A.1)–(A.2) are computed by the linearly-implicit extrapolation integrator LIMEX [13, 14] during the Gauss-Newton iteration, with $\text{RTOL} = 1.0\text{E} - 9$ and $\text{ATOL} = 1.0\text{E} - 9$. All other relevant settings are included in the resulting runtime protocol of the parameter identification task for the data-based model (A.1)–(A.2).

Runtime protocol for an error-oriented Gauss-Newton scheme (NLSCON).

```

      N L S C O N  *****  V e r s i o n  2 . 3 . 3  ***

Gauss-Newton-Method for the solution of nonlinear least squares
problems

Real    Workspace declared as 1742 is used up to 1578 ( 90.6 percent)
Integer Workspace declared as   62 is used up to   50 ( 80.6 percent)

Number of parameters to be estimated (N) :    10
Number of data to fitted, e.g. observations (MFIT) :    42
Number of equality constraints (MCON) :        0

Prescribed relative precision (PTOL) :    0.10D-02

The Jacobian is supplied by numerical differentiation
                                     (feedback strategy included)
Automatic row scaling of the Jacobian is allowed

Rank-1 updates are inhibited
Problem is specified as being highly nonlinear
Bounded damping strategy is off
Maximum permitted number of iteration steps :    40

Internal parameters:

Starting value for damping factor FCSTART =  0.10D-01
Minimum allowed damping factor FCMIN =  0.10D-01
Rank-1 updates decision parameter SIGMA =  0.10D+04
Initial Jacobian pseudo-rank IRANK =    10

```

Maximum permitted subcondition COND = 0.45D+16

It	Normf		Normx	Damp.Fct.	New	Rank
0	0.8345461D+01		0.251D+00		0	10
1	0.8277889D+01	*	0.248D+00	0.010		
1						
1	0.8277889D+01		0.241D+00		0	10
2	0.5289067D+01	*	0.916D-01	0.375		
2						
2	0.5289067D+01		0.991D-01		0	10
2	0.9455332D+01	*	0.133D+00	0.819		
3	0.4570706D+01	*	0.755D-01	0.232		
3						
3	0.4570706D+01		0.827D-01		0	10
4	0.6342464D+01	*	0.732D-01	1.000		
4						
4	0.6342464D+01		0.229D-01		0	10
5	0.3444394D+01	*	0.723D-02	1.000		
5						
5	0.3444394D+01		0.171D-01		0	10
6	0.3083738D+01	*	0.315D-02	0.955		
6						
6	0.3083738D+01		0.652D-02		0	10
7	0.3071453D+01	*	0.120D-03	1.000		
7						
7	0.3071453D+01		0.176D-02		0	10
8	0.3071223D+01	*	0.252D-05	1.000		
8						
8	0.3071223D+01		0.180D-03		0	10
9	0.3071220D+01	*	0.957D-07	1.000		

Solution of nonlinear least squares problem obtained within 9 iteration steps

Incompatibility factor kappa 0.103D+00

Achieved relative accuracy 0.207D-04

Additional statistical analysis of the result of the identification task.

Standard deviation of parameters

```
-----
No.  Estimate      +/-  sigma(X)      =  %
  1   0.185D+00    +/-  0.174D+00    =  94.25 %
  2   0.368D-02    +/-  0.463D-02    = 125.87 %
  3   0.253D+00    +/-  0.107D+00    =  42.40 %
  4  -0.379D-01    +/-  0.646D-02    =  17.04 %
  5   0.786D-02    +/-  0.271D-02    =  34.46 %
  6  -0.959D+00    +/-  0.198D+00    =  20.67 %
  7   0.685D-03    +/-  0.491D-02    = 716.79 %
  8   0.530D-01    +/-  0.725D-01    = 136.89 %
  9   0.241D-01    +/-  0.465D-02    =  19.29 %
 10  -0.111D-03    +/-  0.124D-02    = 1109.90 %
```

Independent confidence intervals

(on 95%-probability level using F-distribution $F(\alpha, 1, m-n) = 4.15$)

```
1 ( -0.170D+00 ,  0.540D+00 )
2 ( -0.575D-02 ,  0.131D-01 )
3 (  0.345D-01 ,  0.472D+00 )
4 ( -0.510D-01 , -0.247D-01 )
5 (  0.234D-02 ,  0.134D-01 )
6 ( -0.136D+01 , -0.555D+00 )
7 ( -0.931D-02 ,  0.107D-01 )
8 ( -0.947D-01 ,  0.201D+00 )
9 (  0.146D-01 ,  0.336D-01 )
10 ( -0.263D-02 ,  0.241D-02 )
```

```
***** Statistics * NLSCON *****
*** Gauss-Newton iter.:      9 ***
*** Corrector steps   :      1 ***
*** Rejected rk-1 st. :      0 ***
*** Jacobian eval.    :     10 ***
*** Function eval.    :     12 ***
*** ... for Jacobian :    100 ***
*****
```

In order to have a visual inspection of these findings, the resulting population curves are plotted in Figure A1. In particular it seems that the curves of our approximated model match more the given data in the time period 1905 – 1915.

Note that, although the Gauss-Newton iteration converges nicely, and with full rank of the Jacobians, the overall estimated standard deviation of the identified coefficients is relatively high. This means that the linearisation at the minimum point found is nearly flat and hence, the corresponding confidence intervals are comparatively wide.

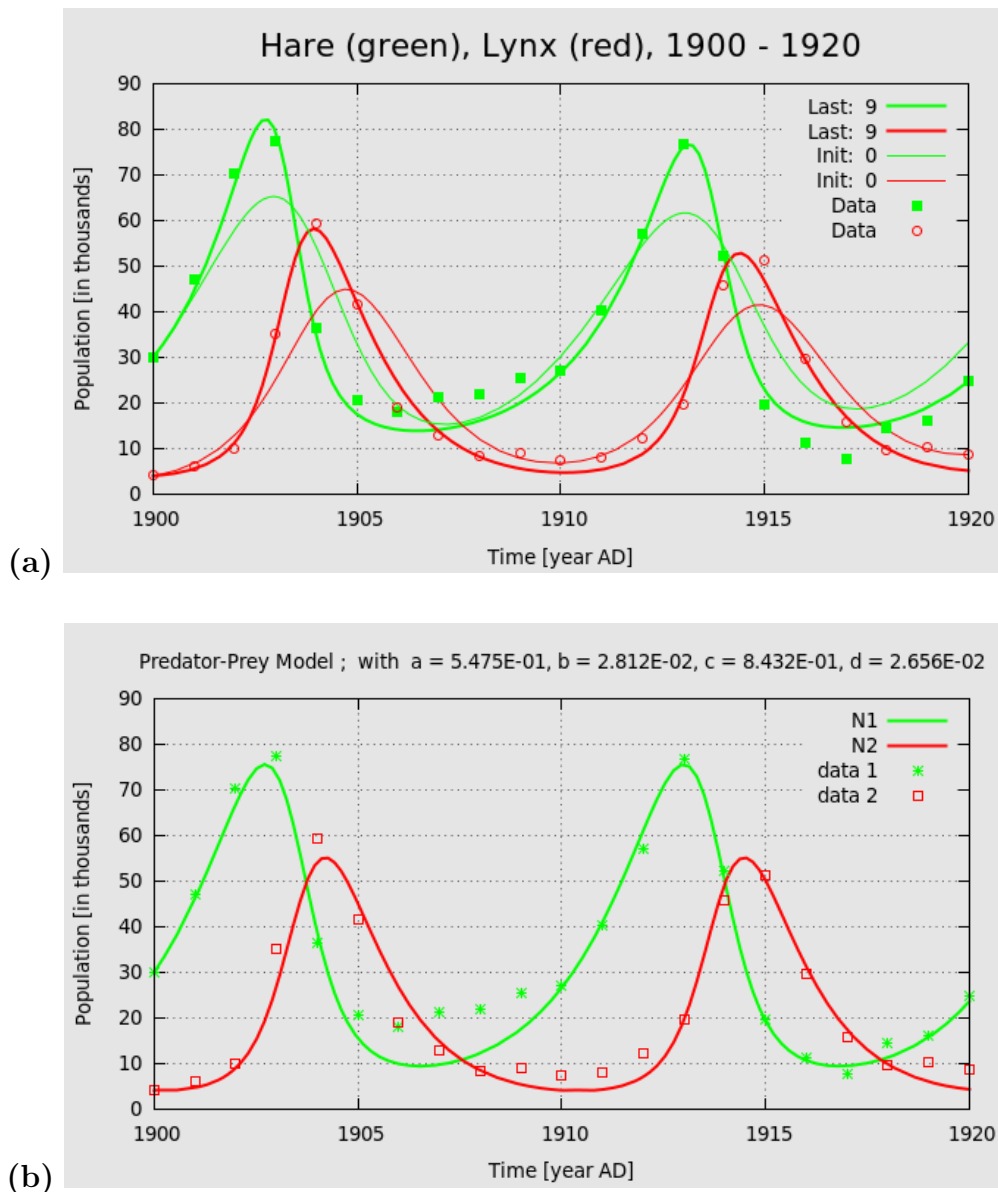


Figure A1. Fitted population curves (a) of the data-based model (with maximal polynomial degree = 2) and (b) of the standard model [1].

The implementation of the presented approach that, written in Ruby, has been applied to perform all example computations in this paper is available upon request.

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