



Modified Patankar Linear Multistep Methods for Production-Destruction Systems

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Abstract

Modified Patankar schemes are linearly implicit time integration methods designed to be unconditionally positive and conservative. In the present work we extend the Patankar-type approach to linear multistep methods and prove that the resulting discretizations retain, with no restrictions on the step size, the positivity of the solution and the linear invariant of the continuous-time system. Moreover, we provide results on arbitrarily high order of convergence and we introduce an embedding technique for the Patankar weights denominators to achieve it.

Keywords Patankar-type schemes · Positivity-preserving · High order · Conservativity · Linear multistep methods

1 Introduction

We address Production-Destruction Systems (PDS) of ordinary differential equations of the form

$$y'_i(t) = P_i(y(t)) - D_i(y(t)), \quad y_i(0) = y_i^0, \quad i = 1, \dots, N, \quad (1.1)$$

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where $\mathbf{y}(t) = (y_1(t), \dots, y_N(t))^T \in \mathbb{R}^N$ is the vector of constituents at time $t \geq 0$. The right-hand side of (1.1) incorporates the non-negative terms $P_i(\mathbf{y}(t))$ and $D_i(\mathbf{y}(t))$ defined as follows

$$P_i(\mathbf{y}) = \sum_{j=1}^N p_{ij}(\mathbf{y}) \quad \text{and} \quad D_i(\mathbf{y}) = \sum_{j=1}^N d_{ij}(\mathbf{y}), \quad i = 1, \dots, N, \quad (1.2)$$

where the functions $p_{ij}(\mathbf{y}) \geq 0$ and $d_{ij}(\mathbf{y}) \geq 0$ represent, respectively, the rates of production and destruction processes for each constituent and component. Here, we introduce the matrices $P(\mathbf{y}) = \{p_{ij}(\mathbf{y})\} \in \mathbb{R}^{N \times N}$, $D(\mathbf{y}) = \{d_{ij}(\mathbf{y})\} \in \mathbb{R}^{N \times N}$ and equivalently reformulate the PDS (1.1) as

$$\mathbf{y}'(t) = (P(\mathbf{y}(t)) - D(\mathbf{y}(t)))\mathbf{e}, \quad \mathbf{y}(0) = \mathbf{y}^0, \quad (1.3)$$

where here, and in the following sections, $\mathbf{e} = (1, \dots, 1)^T$ represents a vector of all ones with the appropriate number of components.

The mathematical modeling of various real life phenomena leads to differential systems of the form (1.1). PDS find relevant applications into chemical [7, 16, 29] and biogeochemical [3, 39] processes, as well as into epidemic [6, 38], ecosystem [11, 15, 44] and astrophysical [25] models (see also [14, Table 1] for a comprehensive list of applications).

Our investigation is restricted to positive and fully conservative production-destruction systems, for which we assume that

$$y_i^0 > 0 \quad \text{implies} \quad y_i(t) > 0, \quad \forall t > 0, \quad i = 1, \dots, N \quad (1.4)$$

and that for each component-wise positive vector $\mathbf{z} \in \mathbb{R}^N$,

$$p_{ij}(\mathbf{z}) = d_{ji}(\mathbf{z}) \quad \text{and} \quad p_{ii}(\mathbf{z}) = d_{ii}(\mathbf{z}) = 0, \quad 1 \leq i, j \leq N. \quad (1.5)$$

The positivity condition (1.4) is guaranteed as long as the initial value problem (1.3) with positive initial value has a unique solution and the matrix $D(\mathbf{y})$ vanishes as $\mathbf{y} \rightarrow \mathbf{0} = (0, \dots, 0)^T \in \mathbb{R}^N$ (further discussions on this topic can be found in [4, 12, 42]). General theoretical results about the existence, uniqueness and positivity of the solution of a production-destruction system have been outlined in [12, Theorem 3.3] and [42, Theorem 1.2].

The assumption (1.5) leads to the linear invariant $\mathbf{e}^T \mathbf{y}(t)$ and to the following conservation law

$$\mathbf{e}^T(\mathbf{y}(t) - \mathbf{y}^0) = 0, \quad \forall t \geq 0. \quad (1.6)$$

As a matter of fact, for a fully conservative production-destruction system, the equality

$$\mathbf{e}^T(P(\mathbf{y}(t)) - D(\mathbf{y}(t)))\mathbf{e} = 0, \quad \forall t \geq 0,$$

holds true and from (1.3), $(\mathbf{e}^T \mathbf{y}(t))' = 0$.

When applied to a positive and fully conservative PDS, a numerical method ought to satisfy a discrete counterpart of (1.4) and (1.6). However, the requirement for such properties in standard schemes usually yields a severe restriction on the stepsize. This motivates the interest in devising unconditionally positive methods which, provided the initial values are positive, produce positive numerical solutions independently of the steplength. Moreover, we will refer to a numerical method as unconditionally conservative if it retains the linear invariant of the system (1.3) whatever the stepsize.

The development of unconditionally positive and conservative numerical methods for production–destruction differential systems has been addressed in several scientific contributions (see, for instance, [9, 46]). Particularly effective in this context is the class of Modified Patankar methods, arising from the manipulations of explicit schemes via the *Patankar-trick* [37], with the aim of ensuring the desired preservation properties for the numerical solution. The origin of this class traces back to [4] where a modification of forward Euler and Heun’s methods resulted in a first and second order positivity-preserving and conservative scheme, respectively. More recently, the same approach has been extended to Runge–Kutta methods and a general definition of Modified Patankar Runge–Kutta (MPRK) schemes has been introduced. In [26–28], MPRK schemes of second and third order were presented and analyzed. Strong-Stability-Preserving MPRK (SSPMPRK) methods were designed in [17, 18] to solve convection equations with stiff source terms, starting from the Shu–Osher form [41] of Runge–Kutta discretizations. A comprehensive analysis through center manifold theory of the stability of the MPRK and SSPMPRK schemes was carried out in [19, 21–23], respectively.

The modified Patankar approach was not solely limited to Runge–Kutta methods. A predictor-corrector modified Patankar scheme was presented in [12]. The proposed discretization relied on a MPRK method as a predictor and on a modified Patankar three-steps backward differentiation formula for the correction step. High order Modified Patankar Deferred Correction (MPDeC) schemes were outlined in [36, 42] as effective integrators to mitigate common issues of MPRK discretizations, including oscillations around steady-state solutions and accuracy loss in cases of vanishing initial states. The geometrically conservative non-standard integrators proposed in [20, 34], specifically designed to preserve linear invariants and to ensure the positivity of more general biochemical systems, are effectively employed for addressing production–destruction systems. Two modified Patankar methods based on specific second and third order multistep schemes were presented in [47].

The leading purpose of this paper is to broaden the benefits of Patankar-type modifications to linear multistep methods, providing a general numerical framework for achieving high order of convergence. The manuscript is organized as follows: in Sect. 2, the class of unconditionally positive and conservative Modified Patankar Linear Multistep (MPLM) methods is formulated. With Sect. 3 a theoretical investigation of the approximation error is performed and general results on arbitrarily high order of convergence are provided. Furthermore, a recursive embedding technique for the efficient computation of the Patankar weight denominators is presented. Numerical experiments are reported in Sect. 4, while some remarks and future perspectives, in Sect. 5, conclude the paper.

2 The Modified Patankar Linear Multistep Scheme

Let $\alpha = (\alpha_1, \dots, \alpha_k)^\top \in \mathbb{R}^k$ and $\beta = (\beta_1, \dots, \beta_k)^\top \in \mathbb{R}^k$ be the coefficients of an explicit k -steps Linear Multistep (LM) method, with $k > 1$ positive integer. Assume that such a method is convergent of order $p \geq 1$, which implies that

$$\sum_{r=1}^k \alpha_r = 1 \quad \text{and} \quad \sum_{r=1}^k (r^q \alpha_r - q r^{q-1} \beta_r) = 0, \quad 1 \leq q \leq p. \quad (2.7)$$

Consider $h > 0$, $t_n = nh$ for $n \geq 0$ and $\mathbf{y}^n = (y_1^n, \dots, y_N^n)^\top \approx \mathbf{y}(t_n)$, for $n \geq k$. From now on, all the inequalities involving vectors are considered component-wise. In analogy with the

scientific literature on Patankar methods (see, for instance, [27]), we provide the following definition.

Definition 1 Given the LM coefficients $\alpha \geq 0$ and $\beta \geq 0$, the k -steps scheme

$$y_i^n = \sum_{r=1}^k \alpha_r y_i^{n-r} + h \sum_{r=1}^k \beta_r \sum_{j=1}^N \left(p_{ij}(y^{n-r}) \frac{y_j^n}{\sigma_j^n} - d_{ij}(y^{n-r}) \frac{y_i^n}{\sigma_i^n} \right), \quad 1 \leq i \leq N, \quad n \geq k, \tag{2.8}$$

where $y^0, \dots, y^{k-1} \in \mathbb{R}^N$ are given and $\sigma_i^n \in \mathbb{R}, i = 1, \dots, N$, is referred to as a Modified Patankar Linear Multistep (MPLM- k) method if

- σ_i^n are unconditionally positive for each $i = 1, \dots, N$ and $n \geq k$;
- σ_i^n are independent of y_i^n for each $i = 1, \dots, N$ and $n \geq k$.

The linearly implicit numerical method (2.8) is devised by applying to a LM discretization of (1.1) a modified version of the source term linearization technique [5] originally proposed by Patankar in [37, Section 7.2-2], with the aim of designing unconditionally positive and conservative schemes. More specifically, the terms

$$\sigma_i^n = \sigma_i^n(y^{n-1}, \dots, y^{n-k}), \quad n \geq k, \quad i = 1, \dots, N,$$

are referred to as Patankar-Weight Denominators (PWDs). Notice that the discrete equation (2.8) is general enough to encompass the linear methods presented in [47, Section 2.2]. As a matter of fact, in case of

$$\begin{aligned} k = 3, \quad \alpha &= \left(\frac{3}{4}, 0, \frac{1}{4} \right)^T, \quad \beta = \left(0, \frac{3}{2}, 0 \right)^T, \quad \text{and} \\ k = 4, \quad \alpha &= \left(\frac{16}{27}, 0, 0, \frac{11}{27} \right)^T, \quad \beta = \left(\frac{16}{9}, 0, 0, \frac{4}{9} \right)^T, \end{aligned}$$

(2.8) coincides with the second and third order schemes of [47], respectively.

Throughout this paper we refer, when needed, to the following equivalent compact notation of (2.8)

$$y^n = \sum_{r=1}^k \alpha_r y^{n-r} + h \sum_{r=1}^k \beta_r M^{n-r} y^n, \quad n \geq k, \tag{2.9}$$

where

$$M^{n-r} = (P(y^{n-r}) - \text{diag}(D(y^{n-r})e)) \text{diag}(S^n) \in \mathbb{R}^{N \times N}, \quad r = 1, \dots, k, \tag{2.10}$$

with $S^n = (1/\sigma_1^n, \dots, 1/\sigma_N^n)^T \in \mathbb{R}^N, n \geq k$.

As already pointed out, the Patankar-type transformation is adopted to ensure positivity for the approximation of the solution to (1.1). The following result provides some conditions on the coefficients of (2.8), or equivalently of (2.9), which lead to a positive numerical solution regardless of the discretization step-size.

Lemma 1 Let $\{y^n\}_{n \geq k}$ be the approximation of the solution to (1.3) computed by (2.9). Suppose that (1.4) and (1.5) hold true and assume that

$h_1)$ the given starting values y^0, \dots, y^{k-1} are positive.

Then, for all $h > 0$ and $n \geq 0, y^n > 0$.

Proof From (2.9), the numerical method reads $M \mathbf{y}^n = \sum_{r=1}^k \alpha_r \mathbf{y}^{n-r}$, $n \geq k$, with $M = I - h \sum_{r=1}^k \beta_r M^{n-r} \in \mathbb{R}^{N \times N}$. From (2.10), the entries of M^{n-r} are

$$M_{ii}^{n-r} = - \sum_{j=1}^N \frac{d_{ij}(\mathbf{y}^{n-r})}{\sigma_i^n}, \quad M_{ij}^{n-r} = \frac{p_{ij}(\mathbf{y}^{n-r})}{\sigma_j^n}, \quad 1 \leq i, j \leq N, \quad i \neq j,$$

for $n \geq k$ and $r = 1, \dots, k$, which implies that M^T is a Z-matrix, since

$$M_{ii} = 1 + h \sum_{r=1}^k \beta_r \sum_{j=1}^N \frac{d_{ij}(\mathbf{y}^{n-r})}{\sigma_i} > 0 \quad \text{and} \quad M_{ji} = -h \sum_{r=1}^k \beta_r \frac{p_{ji}(\mathbf{y}^{n-r})}{\sigma_i} < 0.$$

Furthermore, from (1.5), M^T is strictly diagonally dominant since

$$\sum_{j=1, j \neq i}^N |M_{ji}| = h \sum_{r=1}^k \beta_r \sum_{j=1}^N \frac{d_{ij}(\mathbf{y}^{n-r})}{\sigma_i} < M_{ii}, \quad 1 \leq i, j \leq N, \quad i \neq j.$$

Therefore M^T is a M-matrix (see, for instance, [1, Lemma 6.2]). It follows that M is a M-matrix as well, it is invertible and M^{-1} has all non-negative entries, independently of $n \geq k$. Consequently $\mathbf{y}^n = M^{-1} \sum_{r=1}^k \alpha_r \mathbf{y}^{n-r}$ and an inductive procedure, starting from h_1), yields the result. \square

An investigation of the discrete-time level preservation of the invariance property (1.5) of a positive and fully conservative PDS is carried out with the following lemma.

Lemma 2 *Let $\{\mathbf{y}^n\}_{n \geq k}$ be the approximation of the solution to (1.3) computed by (2.9). Suppose that (1.4) and (1.5) hold true and assume that*

$h_1)$ *for the given starting values, $\mathbf{e}^T \mathbf{y}^0 = \dots = \mathbf{e}^T \mathbf{y}^{k-1} = \eta$.*

Then, for each $h > 0$ and $n \geq 0$, $\mathbf{e}^T (\mathbf{y}^n - \mathbf{y}^0) = 0$.

Proof It suffices to show that $\mathbf{e}^T \mathbf{y}^n = \eta$ for each $n \geq k$. The conservativity property (1.5) yields, independently of $1 \leq r \leq k$ and $n \geq k$,

$$\sum_{i,j=1}^N \left(p_{ij}(\mathbf{y}^{n-r}) \frac{y_j^n}{\sigma_j} - d_{ij}(\mathbf{y}^{n-r}) \frac{y_i^n}{\sigma_i} \right) = \sum_{i,j=1}^N \left(p_{ij}(\mathbf{y}^{n-r}) \frac{y_j^n}{\sigma_j} - p_{ji}(\mathbf{y}^{n-r}) \frac{y_i^n}{\sigma_i} \right) = 0,$$

then, from (2.9), $\mathbf{e}^T \mathbf{y}^n = \sum_{r=1}^k \alpha_r (\mathbf{e}^T \mathbf{y}^{n-r})$. The result is therefore derived from this last relation and the first of (2.7) by inductive arguments. \square

Lemmas 1 and 2 establish conditions for the method (2.9) to be unconditionally positive and conservative. As a matter of fact, if their hypotheses are fulfilled, the numerical solution computed by the MPLM- k scheme retains, with no restrictions on the step-length h , the properties (1.4) and (1.6) of the continuous-time PDS. Therefore to attain these properties we assume, from now on, that the starting values satisfy both $h_1)$ and $h_1)$.

3 Error Analysis and Convergence

In this section we analyze the error arising from the approximation of the continuous-time solution to (1.1) by the numerical methods of the class (2.9) and investigate the conditions

that the Patankar weight denominators have to satisfy for attaining high order of convergence. A recursive practical technique, based on the embedding of different MPLM schemes, is then presented to efficiently compute the PWDs.

In order to investigate the consistency of the discretization (2.9), we address the corresponding local truncation error, here denoted by

$$\delta_i(h; t_n) = y_i(t_n) - \sum_{r=1}^k \alpha_r y_i(t_{n-r}) - h \sum_{r=1}^k \beta_r \sum_{j=1}^N \left(p_{ij}(\mathbf{y}(t_{n-r})) \frac{y_j(t_n)}{\sigma_j(\mathbf{y}(t_{n-1}), \dots, \mathbf{y}(t_{n-k}))} - d_{ij}(\mathbf{y}(t_{n-r})) \frac{y_i(t_n)}{\sigma_i(\mathbf{y}(t_{n-1}), \dots, \mathbf{y}(t_{n-k}))} \right), \tag{3.11}$$

for $i = 1, \dots, N$ and $n \geq k$.

The following result provides a condition on the PWDs which ensures the order p consistency of the MPLM methods.

Theorem 2 (Sufficient condition) *Assume that the given functions describing problem (1.1) belong to $C^p(\Omega_0)$, with $p \geq 1$ and $\Omega_0 = \{z \in \mathbb{R}^N : 0 \leq z_i \leq e^T \mathbf{y}^0, i = 1, \dots, N\}$. If the Patankar weight denominators satisfy*

$$\sigma_i(\mathbf{y}(t_{n-1}), \dots, \mathbf{y}(t_{n-k})) = y_i(t_n) + \mathcal{O}(h^p), \quad i = 1, \dots, N, \quad n \geq k, \tag{3.12}$$

then the MPLM- k method (2.9) is consistent with (1.1), of order p .

Proof It suffices to prove that $\delta_i(h; t_n) = \mathcal{O}(h^{p+1})$ for each $i = 1, \dots, N$ and $n \geq k$. From (3.12) it follows that

$$\frac{y_i(t_n)}{\sigma_i(\mathbf{y}(t_{n-1}), \dots, \mathbf{y}(t_{n-k}))} = \frac{y_i(t_n)}{y_i(t_n) + \mathcal{O}(h^p)} = 1 + \mathcal{O}(h^p), \quad i = 1, \dots, N$$

and the local truncation error defined in (3.11) reads, for $n \geq k$,

$$\delta_i(h; t_n) = \delta_i^{LM}(h; t_n) + \mathcal{O}(h^{p+1}), \quad i = 1, \dots, N, \quad \text{where}$$

$$\delta_i^{LM}(h; t_n) = y_i(t_n) - \sum_{r=1}^k \alpha_r y_i(t_{n-r}) - h \sum_{r=1}^k \beta_r \sum_{j=1}^N (p_{ij}(\mathbf{y}(t_{n-r})) - d_{ij}(\mathbf{y}(t_{n-r}))), \tag{3.13}$$

is the local truncation error of the underlying LM method with coefficients α and β . Finally, since from (2.7) $\delta_i^{LM}(h; t_n) = \mathcal{O}(h^{p+1})$, we get the result. \square

Theorem 2 guarantees the consistency of the MPLM methods when the PWDs are selected to fulfil (3.12). The following result, whose proof is reported in Appendix A, states that this condition represents the minimum and the least stringent requirement for attaining the order p consistency, as it constitutes a necessary prerequisite.

Theorem 3 (Necessary condition) *Assume that the MPLM- k method (2.9) is consistent with (1.1) of order $p \geq 1$. Then the Patankar weight denominators satisfy (3.12).*

To investigate the convergence properties of the scheme (2.9), we first establish some preparatory results. In what follows, given $\Omega \subset \mathbb{R}^N$, we denote by $\Omega^l = \Omega \times \dots \times \Omega$, l times.

Lemma 3 Let $l \geq 1$ be a positive integer and Ω be a compact subset of \mathbb{R}^N . Consider, for $1 \leq i, j \leq N$ and $\mathbf{x} = (\mathbf{x}^1, \dots, \mathbf{x}^l) \in \Omega^l$, the functions

$$\sigma_i(\mathbf{x}) \in C^1(\Omega^l), \text{ such that } \sigma_i(\mathbf{x}) > 0,$$

$$A(\mathbf{x}) = \{a_{ij}(\mathbf{x})\} \text{ with } a_{ij}(\mathbf{x}) \in C^1(\Omega^l), \text{ such that the matrix } I - A(\mathbf{x}) \text{ is invertible,}$$

independently of \mathbf{x} . Then, the functions

$$Z : \mathbf{x} \in \Omega^l \rightarrow \text{diag}(\mathbf{S}(\mathbf{x})) \in \mathbb{R}^{N \times N}, \text{ with } \mathbf{S}(\mathbf{x}) = \left(\frac{1}{\sigma_1(\mathbf{x})}, \dots, \frac{1}{\sigma_N(\mathbf{x})}\right)^T \in \mathbb{R}^N,$$

$$F : \mathbf{x} \in \Omega^l \rightarrow (I - A(\mathbf{x}))^{-1} \in \mathbb{R}^{N \times N},$$

$$\mathbf{g} : \mathbf{x} \in \Omega^l \rightarrow F(\mathbf{x}) \sum_{r=1}^l \alpha_r \mathbf{x}^r \in \mathbb{R}^N,$$

are continuously differentiable on Ω^l .

Proof The first statement comes from

$$\frac{\partial Z}{\partial x_j^v}(\mathbf{x}^1, \dots, \mathbf{x}^l) = -\text{diag}(\bar{\mathbf{S}}(\mathbf{x})) \text{diag}(\hat{\mathbf{S}}^{vj}(\mathbf{x})), \quad \begin{matrix} j = 1, \dots, N, \\ v = 1, \dots, l, \end{matrix} \quad \text{with}$$

$$\bar{\mathbf{S}}(\mathbf{x}) = \left(\frac{1}{\sigma_1^2(\mathbf{x})}, \dots, \frac{1}{\sigma_N^2(\mathbf{x})}\right)^T \in \mathbb{R}^N, \quad \hat{\mathbf{S}}^{vj}(\mathbf{x}) = \left(\frac{\partial \sigma_1(\mathbf{x})}{\partial x_j^v}, \dots, \frac{\partial \sigma_N(\mathbf{x})}{\partial x_j^v}\right)^T \in \mathbb{R}^N.$$

Denote with $\bar{A}(\mathbf{x})$ the adjoint matrix of $I - A(\mathbf{x})$, whose entries are continuous functions of the coefficients $a_{ij}(\mathbf{x})$. Because of the assumptions, $\bar{A}(\mathbf{x})$ and $D_A : \mathbf{x} \in \Omega^l \rightarrow 1/\det A(\mathbf{x}) \in \mathbb{R}$ are well posed and continuously differentiable functions. Therefore, from $F(\mathbf{x}) = D_A(\mathbf{x})\bar{A}(\mathbf{x})$, it follows $F \in C^1(\Omega^l)$. Finally,

$$\frac{\partial \mathbf{g}}{\partial x_j^v}(\mathbf{x}^1, \dots, \mathbf{x}^l) = \frac{\partial F}{\partial x_j^v}(\mathbf{x}) \sum_{r=1}^l \alpha_r \mathbf{x}^r + \alpha_v (F_{1j}(\mathbf{x}), \dots, F_{Nj}(\mathbf{x}))^T, \quad \begin{matrix} j = 1, \dots, N, \\ v = 1, \dots, l, \end{matrix}$$

yields the result. □

The following result, whose proof is deduced from the arguments in [30, sec. 3.4], facilitates the analysis of the approximation error of (2.9).

Lemma 4 Let $l \geq 1$ be a positive integer. Consider a sequence of non-negative numbers $\{a_n\}_{n \in \mathbb{N}_0}$ and assume that there exist $b \geq 0$ and $c_r \geq 0, r = 1, \dots, l$, such that $c = \sum_{r=1}^l c_r > 1$ and $a_n \leq b + \sum_{r=1}^l c_r a_{n-r}$ for $n \geq l$. Then

$$a_n \leq \left(a^* + \frac{b}{c-1}\right) \exp(n(c-1)), \quad n \geq 0, \tag{3.14}$$

where $a^* = \max_{0 \leq j \leq l-1} a_j$.

Proof For $n = 0, \dots, l-1$, the bound (3.14) directly follows from the definition of a^* and the non-negativity of the termes involved. To prove it for $n \geq l$, we firstly show by induction that $a_n \leq c^n a^* + b \sum_{j=0}^{n-1} c^j$. When $n = l$, from $a_l \leq c a^* + b$ and $l \geq 1$, we get the result. Consider now $n > l$ and assume that the statement holds for $j = l, \dots, n-1$. It follows that

$$a_n \leq b + \sum_{r=1}^l c_r \left(c^{n-r} a^* + b \sum_{j=0}^{n-r-1} c^j\right) \leq b + \sum_{r=1}^l c_r \left(c^{n-1} a^* + b \sum_{j=0}^{n-2} c^j\right) \leq c^n a^* + b \sum_{j=0}^{n-1} c^j.$$

Therefore, it turns out that $a_n \leq c^n a^* + b(c^n - 1)/(c - 1)$ for $n \geq l$. Finally $c^n \leq \exp(n(c - 1))$, which completes the proof. □

A thorough analysis of the global discretization error of (2.9) leads to the following convergence result.

Theorem 4 *Let $\mathbf{y}(t)$ be the continuous-time solution to (1.3) for $t \in [0, T]$, with $T > 0$ and let $\{\mathbf{y}^n\}_{n \geq 0}$ be its approximation computed by the k -steps MPLM scheme (2.9) with $h = T/\bar{n}$. Define*

$$\Omega = \left\{ \mathbf{x} \in \mathbb{R}^N : \mu \leq x_i \leq e^T \mathbf{y}^0, i = 1, \dots, N \right\}, \tag{3.15}$$

with μ positive constant. Assume that the given functions describing problem (1.1) belong to $C^p(\Omega)$, with $p \geq 1$ and that

- the starting values satisfy $\|\mathbf{y}(t_m) - \mathbf{y}^m\| = \mathcal{O}(h^p)$, $m = 0, \dots, k - 1$;
- the PWDs are continuously differentiable functions on Ω^k and satisfy (3.12).

Then, the method (2.9) is convergent of order p .

Proof Because of the properties of the continuous and the numerical solution to (1.3) outlined in (1.4) and Lemma 1, there exists $\mu > 0$ such that $\mathbf{y}(t)$ and \mathbf{y}^n belong to Ω , for all $t \geq 0$ and $n = 0, \dots, \bar{n}$. Define, for $\mathbf{x} = (x^1, \dots, x^k) \in \Omega^k$ and $\mathbf{S}^n(\mathbf{x}) = \left(\frac{1}{\sigma_1^n(\mathbf{x})}, \dots, \frac{1}{\sigma_N^n(\mathbf{x})} \right)^T \in \mathbb{R}^N$, the functions

$$\begin{aligned} \Phi^{nr} : \mathbf{x} \in \Omega^k &\rightarrow (P(\mathbf{x}^r) - \text{diag}(D(\mathbf{x}^r)\mathbf{e})) \text{diag}(\mathbf{S}^n(\mathbf{x})) \in \mathbb{R}^{N \times N} \\ \mathbf{G}^{nr} : (\mathbf{x}, \mathbf{x}^{k+1}) \in \Omega^{k+1} &\rightarrow \Phi^{nr}(\mathbf{x}) \mathbf{x}^{k+1} \in \mathbb{R}^N, \end{aligned} \quad \begin{aligned} r = 1, \dots, k, \\ n = k, \dots, \bar{n}. \end{aligned}$$

The global discretization error $\mathbf{e}(h; t_n) = \mathbf{y}(t_n) - \mathbf{y}^n$ then satisfies

$$\begin{aligned} \mathbf{e}(h; t_n) = \sum_{r=1}^k \alpha_r \mathbf{e}(h; t_{n-r}) + h \sum_{r=1}^k \beta_r \left(\mathbf{G}^{nr}(\mathbf{y}(t_{n-k}), \dots, \mathbf{y}(t_n)) - \mathbf{G}^{nr}(\mathbf{y}^{n-k}, \dots, \mathbf{y}^n) \right) \\ + \delta(h; t_n), \end{aligned} \quad n = k, \dots, \bar{n}, \tag{3.16}$$

where the components of $\delta(h; t_n) = (\delta_1(h; t_n), \dots, \delta_N(h; t_n))^T \in \mathbb{R}^N$ are defined in (3.11). Because of the regularity assumptions on the known functions and the first result of Lemma 3, $\Phi^{nr} \in C^1(\Omega^k)$ and $\mathbf{G}^{nr} \in C^1(\Omega^{k+1})$, for each $r = 1, \dots, k$ and $n = k, \dots, \bar{n}$. Therefore, from the mean value theorem

$$\mathbf{G}^{nr}(\mathbf{y}(t_{n-k}), \dots, \mathbf{y}(t_n)) - \mathbf{G}^{nr}(\mathbf{y}^{n-k}, \dots, \mathbf{y}^n) = J_{\mathbf{G}^{nr}}(\xi_r^{n-k}, \dots, \xi_r^n) \begin{pmatrix} \mathbf{e}(h; t_{n-k}) \\ \vdots \\ \mathbf{e}(h; t_n) \end{pmatrix}, \tag{3.17}$$

where $J_{\mathbf{G}^{nr}}(\xi_r^{n-k}, \dots, \xi_r^n) \in \mathbb{R}^{N \times (Nk+N)}$ is the Jacobian of \mathbf{G}^{nr} . Let

$$J_{\mathbf{G}^{nr}}(\xi_r^{n-k}, \dots, \xi_r^n) = \left(\underbrace{J_{\mathbf{G}^{nr}}^{(0)}(\xi_r^{n-k}, \dots, \xi_r^n)}_{N \times N} \mid \dots \mid \underbrace{J_{\mathbf{G}^{nr}}^{(k)}(\xi_r^{n-k}, \dots, \xi_r^n)}_{N \times N} \right).$$

Then, for each $n = k, \dots, \bar{n}$ and $r = 1, \dots, k$,

$$\|J_{\mathbf{G}^{nr}}^{(j)}(\xi_r^{n-k}, \dots, \xi_r^n)\| \leq \|J_{\mathbf{G}^{nr}}(\xi_r^{n-k}, \dots, \xi_r^n)\| \leq J, \quad j = 0, \dots, k, \tag{3.18}$$

with J positive constant depending on the bounds of the known functions and their derivatives on Ω^{k+1} . Substituting (3.17) into (3.16) leads to the discrete equation

$$e(h; t_n) = \sum_{r=1}^k \alpha_r e(h; t_{n-r}) + h \sum_{r=1}^k \beta_r \sum_{j=0}^k J_{G^{nr}}^{(j)}(\xi_r^{n-k}, \dots, \xi_r^n) e(h; t_{n-j}) + \delta(h; t_n),$$

for $n = k, \dots, \bar{n}$. Denoted $B = J \sum_{r=1}^k \beta_r$, for a sufficiently small h , from (3.18)

$$\|e(h; t_n)\| \leq \sum_{r=1}^k \frac{\alpha_r + hB}{1 - hB} \|e(h; t_{n-r})\| + \max_{0 \leq n \leq \bar{n}} \frac{\|\delta(h; t_n)\|}{1 - hB}, \quad n = k, \dots, \bar{n}.$$

Because of the first relation in (2.7), $\sum_{r=1}^k \frac{\alpha_r + hB}{1 - hB} = 1 + h \frac{(k+1)B}{1 - hB} > 1$, so that from Lemma 4 it follows

$$\|e(h; t_n)\| \leq \left(\max_{0 \leq m \leq k-1} \|y(t_m) - y^m\| + \frac{\max_{0 \leq n \leq \bar{n}} \|\delta(h; t_n)\|}{h(k+1)B} \right) \exp\left(\frac{(k+1)TB}{1 - hB}\right),$$

for $0 \leq n \leq \bar{n}$ and $T = \bar{n}h$. Because of the assumptions on the initial values and the order p consistency of (2.9) (Theorem 2), we have

$$\max_{0 \leq n \leq \bar{n}} \|e(h; t_n)\| \leq Ch^p,$$

with C positive constant not depending on h , which yields the result. □

3.1 The σ -Embedding Technique

Theorem 4 provides sufficient conditions for the order p convergence of the numerical method (2.9), but gives no clues on how to compute unconditionally positive and conservative Patankar weight denominators satisfying (3.12). To achieve this goal, here we introduce an embedding technique based on a recursive use of MPLM methods. More specifically, for $p = 1$, we just consider the Modified Patankar Euler (MPE) method (see [4] for further details), corresponding to (2.9) with

$$k = 1, \quad \alpha_1 = \beta_1 = 1, \quad \sigma_i^n = y_i^{n-1}, \quad \text{for } n \geq 1, \quad i = 1, \dots, N, \quad (3.19)$$

and then investigate MPLM- k schemes for $p \geq 2$ and $k > 1$. Let $\sigma^n = \sigma^{n(p-1)}$ be PWDs satisfying the condition (3.12). Our approach consists in recursively computing $\sigma^{n(p-1)}$ by a $\tilde{k} \leq k$ steps, order $p - 1$ convergent MPLM- \tilde{k} method, whose coefficients are denoted by $\alpha^{(p-1)} \in \mathbb{R}^{\tilde{k}}$ and $\beta^{(p-1)} \in \mathbb{R}^{\tilde{k}}$, as follows

$$\sigma^{n(p-1)}(y^{n-1(p)}, \dots, y^{n-\tilde{k}(p)}) = \left(I - h \sum_{r=1}^{\tilde{k}} \beta_r^{(p-1)} M^{n-r(p)} \right)^{-1} \sum_{r=1}^{\tilde{k}} \alpha_r^{(p-1)} y^{n-r(p)}, \quad (3.20)$$

for $n \geq k$, where

$$M^{n-r(p)} = \left(P(y^{n-r(p)}) - \text{diag}(D(y^{n-r(p)})e) \right) \text{diag}(S^{n(p-2)}) \in \mathbb{R}^{N \times N}, \quad r = 1, \dots, k$$

and $S^{n(p-2)} = (1/\sigma_1^{n(p-2)}, \dots, 1/\sigma_N^{n(p-2)})^\top \in \mathbb{R}^N$. Here $y^{n(p)}$, $n \geq k$, represents the numerical solution computed by the MPLM- k method.

With the following result we prove that the PWDs computed adopting the σ -embedding technique (3.20) meet the condition (3.12), which is sufficient for to the convergence of the MPLM- k scheme.

Theorem 5 *Let $\mathbf{y}(t)$ be the continuous-time solution to (1.3) for $t \in [0, T]$, with $T > 0$ and let $\{\mathbf{y}^{n(p)}\}_{n \geq 0}$ be its approximation computed by the k -steps MPLM scheme (2.9) with $h = T/\bar{n}$. Assume that the given functions describing problem (1.1) belong to $C^p(\Omega)$, with $p \geq 1$ and Ω in (3.15).*

Assume that

- *the starting values satisfy $\|\mathbf{y}(t_m) - \mathbf{y}^{m(p)}\| = \mathcal{O}(h^p)$, $m = 0, \dots, k - 1$;*
- *the σ -embedding strategy is implemented and the PWDs are computed by a \tilde{k} -steps, order $p - 1$ convergent MPLM- \tilde{k} scheme as detailed in (3.20).*

Then, the PWDs functions $\sigma^{n(p-1)}$, $n = k, \dots, \bar{n}$, are continuously differentiable on $\Omega^{\tilde{k}}$ and satisfy (3.12). Therefore, the numerical method (2.9) is convergent of order p .

Proof The case $p = 1$ corresponds to the first order convergent MPE method (3.19). For $p \geq 2$, we prove the result by induction. Firstly, we show that the MPLM- k scheme (2.9) is quadratically convergent ($p = 2$) if the PWDs are computed by the one step Modified Patankar Euler discretization (3.19). As a matter of fact, the consistency condition (3.12) comes from [4, Theorem 3.11], since

$$\sigma^{n(1)}(\mathbf{y}(t_{n-1})) = (I - h\Phi(\mathbf{y}(t_{n-1})))^{-1} \mathbf{y}(t_{n-1}) = \mathbf{y}(t_n) + \mathcal{O}(h^2), \quad n \geq 1,$$

where $\Phi(\mathbf{x}) = (P(\mathbf{x}) - \text{diag}(D(\mathbf{x})\boldsymbol{e})) \text{diag}\left(\frac{1}{x_1}, \dots, \frac{1}{x_N}\right) \in \mathbb{R}^{N \times N}$. Furthermore, from Lemma 3, $\sigma^{n(1)} \in C^1(\Omega)$. Therefore, all the hypotheses of Theorem 4 are fulfilled and the second order convergence is established.

Consider $p > 2$ and assume the statement to be true for each $s = 1, \dots, p - 1$. In this case, the PWDs $\sigma^{n(p-1)}$, $n = k, \dots, \bar{n}$, are computed by an order $p - 1$ convergent, then consistent [8, Theorem 3.5], MPLM- \tilde{k} method, accordingly to (3.20). Therefore, from Theorem 3, the condition (3.12) directly follows. Furthermore, from the inductive hypotheses, the functions $\sigma^{n(p-2)} \in C^1(\Omega^{\tilde{k}})$, $n = k, \dots, \bar{n}$ and hence, from Lemma 3,

$$\sigma^{n(p-1)}(\mathbf{x}) = \left(I - h \sum_{r=1}^{\tilde{k}} \beta_r^{(p-1)} \Phi^{nr(p-2)}(\mathbf{x}) \right)^{-1} \sum_{r=1}^{\tilde{k}} \alpha_r^{(p-1)} \mathbf{x}^r \in \mathbb{R}^N,$$

with $\Phi^{nr(p-2)}(\mathbf{x}) = (P(\mathbf{x}^r) - \text{diag}(D(\mathbf{x}^r)\boldsymbol{e})) \text{diag}\left(S^{n(p-2)}(\mathbf{x})\right) \in \mathbb{R}^{N \times N}$,

and $S^{n(p-2)}(\mathbf{x}) = \left(\frac{1}{\sigma_1^{n(p-2)}(\mathbf{x})}, \dots, \frac{1}{\sigma_N^{n(p-2)}(\mathbf{x})} \right)^\top \in \mathbb{R}^N, \quad \mathbf{x} = (\mathbf{x}^1, \dots, \mathbf{x}^{\tilde{k}})^\top,$

is continuously differentiable on $\Omega^{\tilde{k}}$ as well. Finally, an application of Theorem 4 yields the result. □

Theorem 5 establishes a practical and general framework to compute the PWDs by embedding different MPLM methods, starting from the modified Patankar Euler discretization. This machinery allows the construction of arbitrarily high order unconditionally positive and conservative numerical methods. However, the σ -embedding technique in (3.20) is specifically

designed for positive PDS and cannot be implemented if any of the components of the initial value is zero. This issue has been addressed and overcome in [26] for modified Patankar Runge–Kutta schemes by replacing zero components of the initial value by small quantities like $realmin \approx 2.26 \cdot 10^{-308}$. Our numerical experiments of Sect. 4 confirm the effectiveness of this expedient also for the modified Patankar linear multistep methods.

4 Numerical Experiments

In this section we report some experiments and compare the performances of the method (2.9) with that of some well-established modified Patankar schemes. Our investigation is conducted employing the modified Patankar linear multistep methods of order from 2 to 6, whose coefficients are reported in Table 1. Here, we adopt the notation MPLM- $k(p)$ to indicate a k -steps, order p scheme. The numerical simulations of some test problems performed by the methods (2.9)–Table 1 with the σ -embedding technique provide experimental evidence of convergence up to the sixth order. A self-starting embedding procedure is here implemented for computing accurate starting values satisfying h_1) and h_1). To assess the order of the MPLM- $k(p)$ schemes, we consider the maximum absolute error $E(h)$ and the experimental rate of convergence \hat{p} defined as follows

$$E(h) = \max_{0 \leq n \leq T/h} \left\| \mathbf{y}^{n(ref)} - \mathbf{y}^n \right\|_{\infty}, \quad \hat{p} = \log_2 \left(\frac{E(h)}{E(\frac{1}{2}h)} \right). \quad (4.21)$$

For each test problem, the reference solution $\mathbf{y}^{n(ref)}$ in (4.21) is obtained by the `ode15s` built-in Matlab function with absolute and relative tolerances of `AbsTol` = $\varepsilon_{mach} \approx 2.22 \cdot 10^{-16}$ and `RelTol` = $\varepsilon_{mach} \cdot 10^2$, respectively.

A direct comparison with a third order Modified Patankar Runge–Kutta (MPRK3) method (see [26, Lemma 6, Case II] with $\gamma = 0.5$) and with high order Modified Patankar Deferred Correction (MPDeC) schemes (see [36]), highlights the competitive performances of the MPLM- k discretizations. Incidentally, in the case of the MPDeC integrators, the computational efforts required to obtain the coefficients are disregarded.

All the numerical experiments are conducted on a single machine equipped with an Intel Core i7-7700HQ Octa-Core processor operating at 2.80GHz and supported by 8.00 GB of RAM. Both MPRK3 and MPLM- k algorithms are implemented and executed using MATLAB (version R2020b). Additionally, the MPLM- k schemes are implemented and executed with Julia (version 1.8.5) and compared against the MPDeC codes available in the repository [35].

4.1 Test 1: Linear Test

Our first example consists in the following linear test

$$\begin{aligned} y_1'(t) &= -ay_1(t) + y_2(t), \\ y_2'(t) &= ay_1(t) - y_2(t), \end{aligned} \quad \text{with } a = 5, \quad t \in [0, 2] \quad \text{and} \quad \mathbf{y}^0 = \begin{pmatrix} 0.9 \\ 0.1 \end{pmatrix}, \quad (4.22)$$

proposed in [17, 18, 26, 27]. The system (4.22) describes the exchange of mass between two constituents and fits the form of a fully conservative PDS, where the production and destruction terms in (1.2) are given by

$$p_{12}(\mathbf{y}) = y_2, \quad p_{21}(\mathbf{y}) = ay_1, \quad d_{12}(\mathbf{y}) = ay_1, \quad d_{21}(\mathbf{y}) = y_2.$$

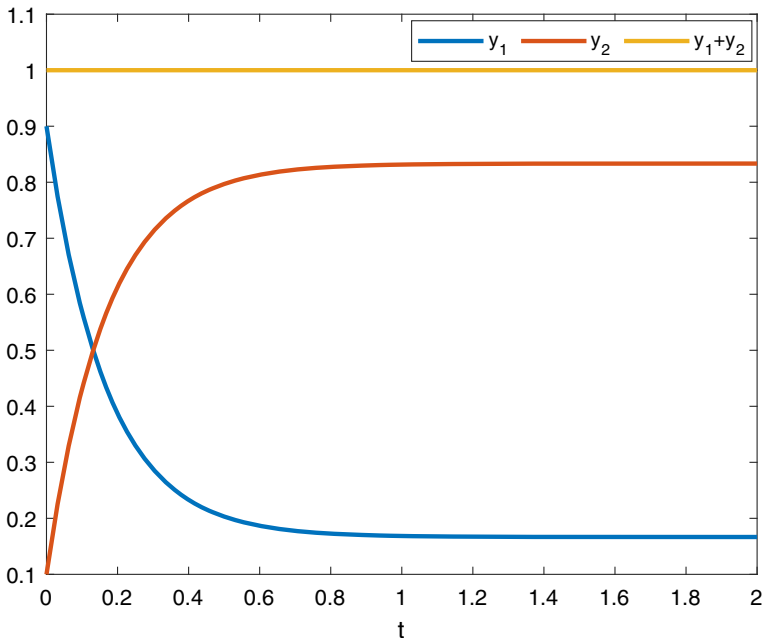


Fig. 1 Numerical solution of Test 1 computed by MPLM-10(6) with $h = 2^{-5}$

Since $p_{ij}(\mathbf{y})$ is continuously differentiable with bounded derivatives and $\lim_{\mathbf{y} \rightarrow \mathbf{0}} p_{ij}(\mathbf{y}) = 0$, for $i, j \in \{1, 2\}$, the existence of a unique and positive solution to (4.22) is guaranteed by [42, Theorem 1.2].

The approximation of the continuous-time solution to (4.22), computed by the MPLM-10(6) method with $h = 2^{-5}$, is reported in Fig. 1. In compliance with Lemmas 1 and 2, the numerical solution is positive and the linear invariant of the PDS (4.1) is retained. In Table 2 we list the maximum errors $E(h)$ on the integration interval $[0, 2]$ and the experimental rate of convergence \hat{p} for all the methods listed in Table 1. From Table 2, as well as from Figs. 2 and 3, it is clear that the experimental order agrees with the theoretical one established in Theorem 5. Furthermore, the work precision diagram of Fig. 4 shows the mean execution time over 10 runs against the approximation error for the MPLM- $k(p)$ methods with $2 \leq p \leq 6$. It is evident that, for $p \geq 4$, the methods in Table 1 are computationally more efficient than the benchmark scheme MPRK3, in terms of the accuracy-computational cost trade off.

The MPLM methods in Table 1 are then also compared to the modified Patankar deferred correction integrators outlined in [36], using the Julia implementation of the MPDeC schemes provided in [35]. For the sake of comparison, we consider the mean relative error taken over all time steps and all constituents,

$$\varepsilon(h) = \frac{1}{N} \sum_{i=1}^N \left(\frac{\sqrt{\bar{n} \sum_{n=1}^{\bar{n}} (y_i^{n(J)} - y_i^n)^2}}{\bar{n} \sum_{n=1}^{\bar{n}} y_i^{n(J)}} \right), \quad \bar{n} = T/h, \quad (4.23)$$

as defined in [26]. Here, the reference solution $\mathbf{y}^{n(J)}$ in (4.21) is obtained by the `radau` built-in Julia solver (implicit Runge–Kutta of variable order between 5 and 13) with absolute and relative tolerances of 10^{-14} . It turns out that for the Test 1, the MPDeC methods exhibit

Table 2 Experimental convergence of the numerical solutions to Test 1

h	$E(h)$	\hat{p}	h	$E(h)$	\hat{p}	h	$E(h)$	\hat{p}
MPE			MPLM-2(2)			MPLM-4(3)		
2^{-5}	$2.34 \cdot 10^{-2}$	–	2^{-5}	$4.92 \cdot 10^{-3}$	–	2^{-5}	$6.71 \cdot 10^{-4}$	–
2^{-6}	$1.21 \cdot 10^{-2}$	0.94	2^{-6}	$1.52 \cdot 10^{-3}$	1.70	2^{-6}	$1.41 \cdot 10^{-4}$	2.25
2^{-7}	$6.20 \cdot 10^{-3}$	0.97	2^{-7}	$4.24 \cdot 10^{-4}$	1.84	2^{-7}	$2.37 \cdot 10^{-5}$	2.57
2^{-8}	$3.13 \cdot 10^{-3}$	0.98	2^{-8}	$1.12 \cdot 10^{-4}$	1.92	2^{-8}	$3.48 \cdot 10^{-6}$	2.77
2^{-9}	$1.57 \cdot 10^{-3}$	0.99	2^{-9}	$2.89 \cdot 10^{-5}$	1.96	2^{-9}	$4.72 \cdot 10^{-7}$	2.88
2^{-10}	$7.88 \cdot 10^{-4}$	1.00	2^{-10}	$7.34 \cdot 10^{-6}$	1.98	2^{-10}	$6.16 \cdot 10^{-8}$	2.94
2^{-11}	$3.95 \cdot 10^{-4}$	1.00	2^{-11}	$1.85 \cdot 10^{-6}$	1.99	2^{-11}	$7.87 \cdot 10^{-9}$	2.97
MPLM-5(4)			MPLM-7(5)			MPLM-10(6)		
2^{-5}	$2.70 \cdot 10^{-4}$	–	2^{-5}	$1.12 \cdot 10^{-4}$	–	2^{-5}	$4.52 \cdot 10^{-5}$	–
2^{-6}	$3.02 \cdot 10^{-5}$	3.16	2^{-6}	$8.53 \cdot 10^{-6}$	3.71	2^{-6}	$3.51 \cdot 10^{-6}$	3.69
2^{-7}	$2.57 \cdot 10^{-6}$	3.56	2^{-7}	$4.64 \cdot 10^{-7}$	4.20	$3IMPE2^{-7}$	$1.15 \cdot 10^{-7}$	4.93
2^{-8}	$1.91 \cdot 10^{-7}$	3.75	2^{-8}	$1.93 \cdot 10^{-8}$	4.59	2^{-8}	$2.71 \cdot 10^{-9}$	5.40
2^{-9}	$1.36 \cdot 10^{-8}$	3.81	2^{-9}	$7.09 \cdot 10^{-10}$	4.77	2^{-9}	$5.30 \cdot 10^{-11}$	5.68
2^{-10}	$9.63 \cdot 10^{-10}$	3.82	2^{-10}	$2.49 \cdot 10^{-11}$	4.83	2^{-10}	$6.95 \cdot 10^{-13}$	6.26
2^{-11}	$6.88 \cdot 10^{-11}$	3.81	2^{-11}	$7.98 \cdot 10^{-13}$	4.97	2^{-11}	$3.34 \cdot 10^{-13}$	1.06

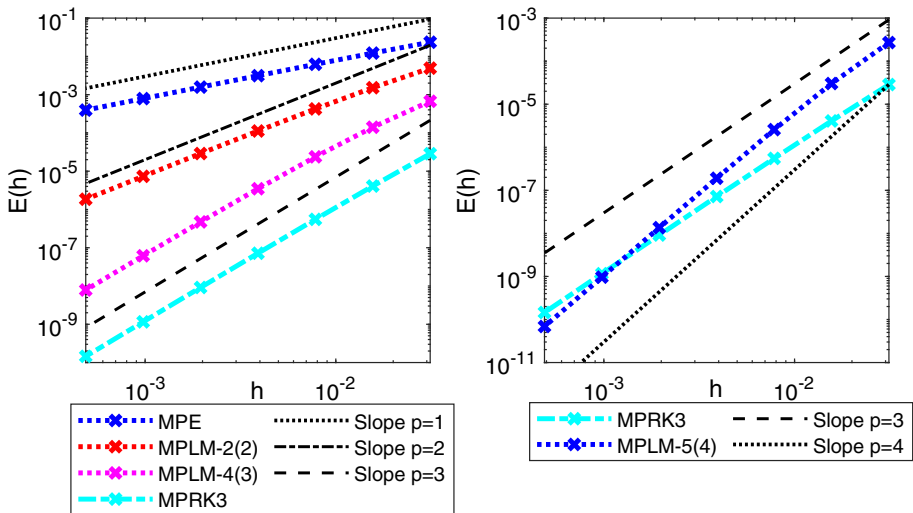


Fig. 2 Experimental order for MPE, MPLM-2(2), MPLM-4(3), MPLM-5(4) and MPRK3 applied to Test 1

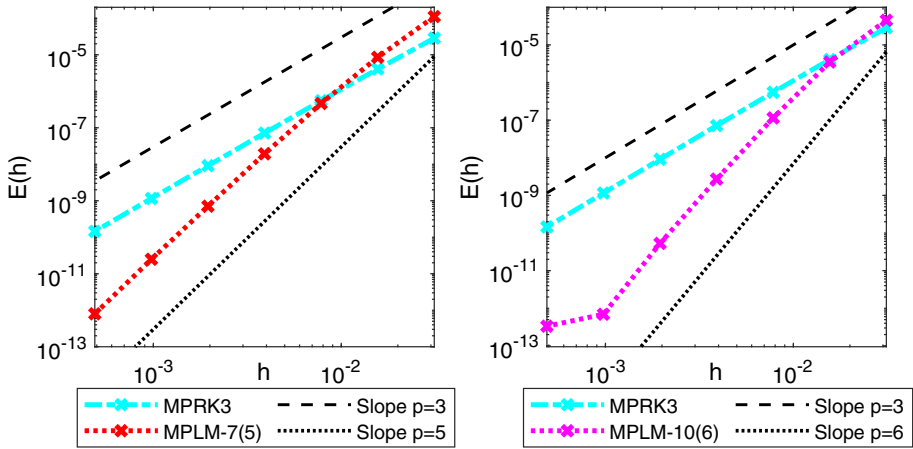


Fig. 3 Experimental order for MPLM-7(5), MPLM-10(6) and MPRK3 applied to Test 1

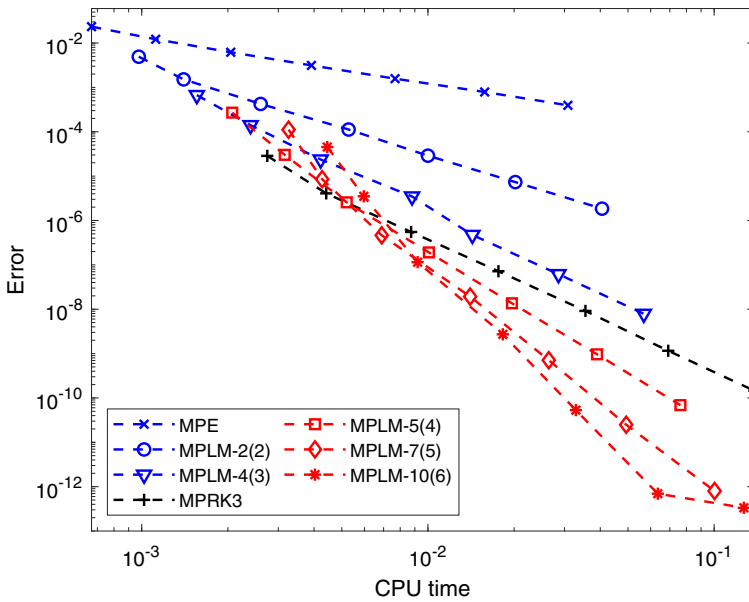


Fig. 4 Work precision diagram: error versus CPU time for the different methods applied to Test 1. $h = T/2^{5+m}$, $m = 1, \dots, 7$.

superior accuracy and performance with respect to the MPLM schemes, as shown in Figs. 5 and 6.

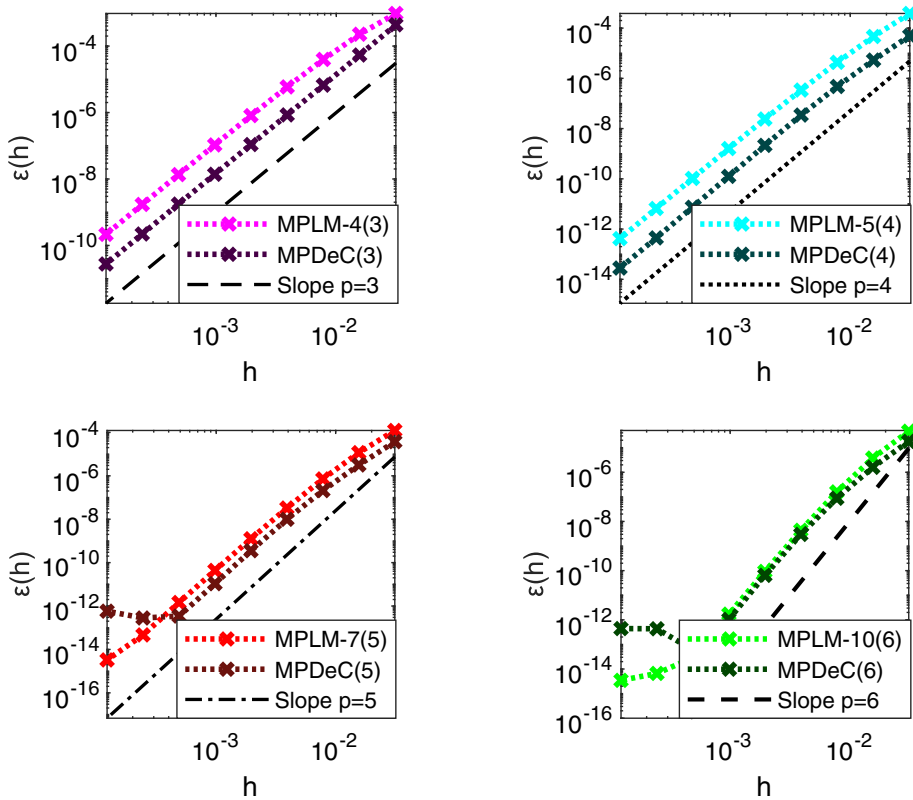


Fig. 5 Accuracies, in terms of (4.23), of MPLM and MPDeC schemes applied to Test 1. $h = T/2^{5+m}$, $m = 1, \dots, 7$.

4.2 Test 2: Nonlinear Test

For our second test problem, we consider the non-stiff nonlinear system [4, 26, 27]

$$\begin{aligned}
 y_1'(t) &= -\frac{y_1(t)y_2(t)}{y_1(t) + 1}, \\
 y_2'(t) &= \frac{y_1(t)y_2(t)}{y_1(t) + 1} - ay_2(t), \\
 y_3'(t) &= ay_2(t),
 \end{aligned}
 \quad \text{with } a = 0.3, \quad \text{and } y^0 = \begin{pmatrix} 9.98 \\ 0.01 \\ 0.01 \end{pmatrix}, \quad (4.24)$$

obtained from (1.1) by taking $p_{ij}(y) = 0 = d_{ji}(y)$ for all combination of $i, j = 1, \dots, 3$ other than

$$p_{21}(y) = d_{12}(y) = \frac{y_1 y_2}{y_1 + 1}, \quad p_{32}(y) = d_{23}(y) = ay_2.$$

The existence and the uniqueness of a positive solution to (4.24) comes from [42, Theorem 1.2]. As a matter of fact, given $0 < y = (y_1, y_2, y_3)^T \leq e^T y^0$,

$$\frac{\partial p_{ij}}{\partial y_l}(y) \leq \max\{a, e^T y^0\}, \quad \text{and} \quad p_{ij}(y) \xrightarrow{y \rightarrow 0} 0, \quad i, j, l \in \{1, 2, 3\}.$$

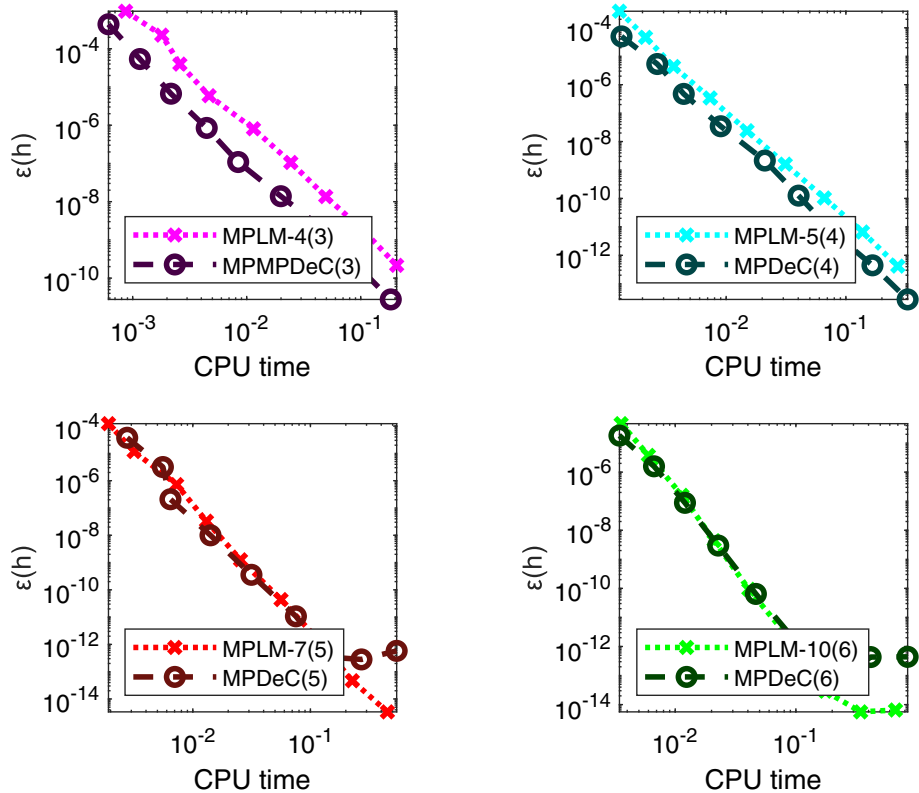


Fig. 6 Work precision diagram: mean error versus CPU time for the different methods applied to Test 1. $h = T/2^{5+m}$, $m = 1, \dots, 7$.

The PDS (4.24) models an algal bloom and might be interpreted as a geobiochemical model for the upper oceanic layer in spring, when nutrient rich surface water is captured in the euphotic zone. During the process, nutrients at time t , $y_1(t)$, are taken up by the phytoplankton $y_2(t)$ according to a Michaelis-Menten formulation. Concurrently, the phytoplankton biomass is converted to detritus $y_3(t)$, with a loss fixed rate $a > 0$, due the effects of mortality and zooplankton grazing. Thus the system is considered closed and the total biomass remains constant accordingly to the conservation law (1.6).

The outcomes of the numerical integration of (4.24) by MPLM-10(6) with $h = 1.88 \cdot 2^{-5}$ are shown in Fig. 7. The experimental results of Table 3 comply with the theoretical findings for this test as well and, from Figs. 8, 9, 10, it is clear that for $p \geq 3$ the MPLM- $k(p)$ methods outperform the MPRK3 discretization. The comparison of the MPLM methods with the MPDeC schemes, as illustrated in Fig. 11, highlights the higher accuracy of the latter. Also in this case, the mean error is computed as detailed in (4.23). For the sake of brevity, we omit the work precision diagrams comparing our methods with the MPDeC schemes, which exhibit superior performances also in this test.

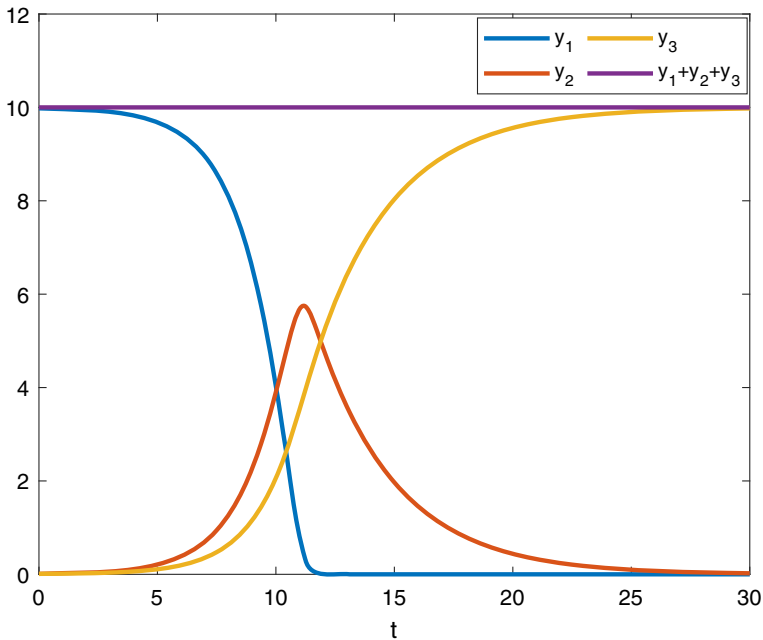


Fig. 7 Numerical solution of Test 2 by MPLM-10(6) with $h = 1.88 \cdot 2^{-5}$.

Table 3 Experimental convergence of the numerical solutions to Test 2

$1.07 \cdot h$	$E(h)$	\hat{p}	$1.07 \cdot h$	$E(h)$	\hat{p}	$1.07 \cdot h$	$E(h)$	\hat{p}
MPE			MPLM-2(2)			MPLM-4(3)		
2^{-3}	$2.57 \cdot 10^0$	–	2^{-3}	$1.76 \cdot 10^{-1}$	–	2^{-3}	$3.17 \cdot 10^{-2}$	–
2^{-4}	$1.40 \cdot 10^0$	0.88	2^{-4}	$4.83 \cdot 10^{-2}$	1.87	2^{-4}	$5.88 \cdot 10^{-3}$	2.43
2^{-5}	$7.28 \cdot 10^{-1}$	0.94	2^{-5}	$1.26 \cdot 10^{-2}$	1.93	2^{-5}	$9.29 \cdot 10^{-4}$	2.66
2^{-6}	$3.71 \cdot 10^{-1}$	0.97	2^{-6}	$3.23 \cdot 10^{-3}$	1.97	2^{-6}	$1.32 \cdot 10^{-4}$	2.82
2^{-7}	$1.88 \cdot 10^{-1}$	0.99	2^{-7}	$8.16 \cdot 10^{-4}$	1.98	2^{-7}	$1.76 \cdot 10^{-5}$	2.90
2^{-8}	$9.43 \cdot 10^{-2}$	0.99	2^{-8}	$2.05 \cdot 10^{-4}$	1.99	2^{-8}	$2.28 \cdot 10^{-6}$	2.95
2^{-9}	$4.73 \cdot 10^{-2}$	1.00	2^{-9}	$5.14 \cdot 10^{-5}$	2.00	2^{-9}	$2.90 \cdot 10^{-7}$	2.97
MPLM-5(4)			MPLM-7(5)			MPLM-10(6)		
2^{-3}	$1.64 \cdot 10^{-2}$	–	2^{-3}	$1.24 \cdot 10^{-2}$	–	2^{-3}	$1.06 \cdot 10^{-2}$	–
2^{-4}	$2.14 \cdot 10^{-3}$	2.94	2^{-4}	$1.23 \cdot 10^{-3}$	3.34	2^{-4}	$8.81 \cdot 10^{-4}$	3.59
2^{-5}	$2.02 \cdot 10^{-4}$	3.40	2^{-5}	$7.60 \cdot 10^{-5}$	4.02	2^{-5}	$3.92 \cdot 10^{-5}$	4.49
2^{-6}	$1.57 \cdot 10^{-5}$	3.69	2^{-6}	$3.57 \cdot 10^{-6}$	4.41	2^{-6}	$1.17 \cdot 10^{-6}$	5.07
2^{-7}	$1.10 \cdot 10^{-6}$	3.84	2^{-7}	$1.39 \cdot 10^{-7}$	4.68	2^{-7}	$2.63 \cdot 10^{-8}$	5.48
2^{-8}	$7.23 \cdot 10^{-8}$	3.92	2^{-8}	$4.91 \cdot 10^{-9}$	4.83	2^{-8}	$4.97 \cdot 10^{-10}$	5.72
2^{-9}	$4.64 \cdot 10^{-9}$	3.96	2^{-9}	$1.62 \cdot 10^{-10}$	4.92	2^{-9}	$9.40 \cdot 10^{-12}$	5.73

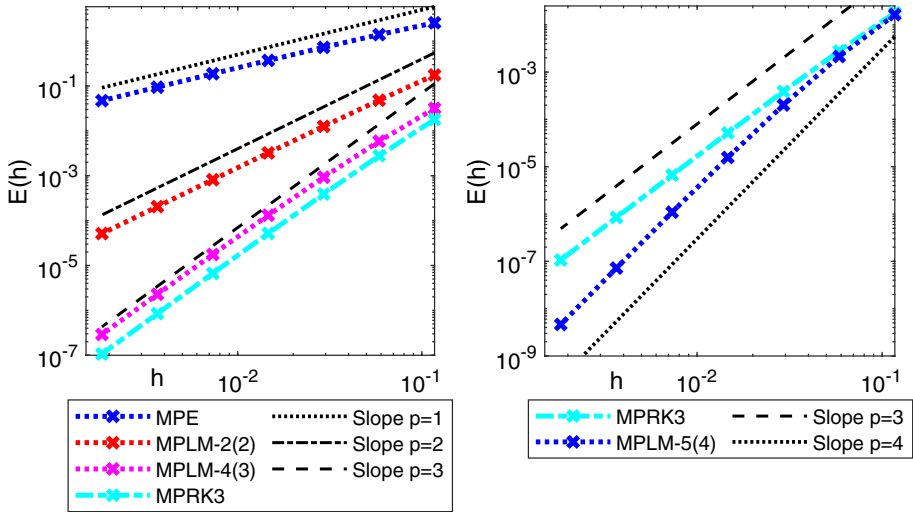


Fig. 8 Experimental order for MPE, MPLM-2(2), MPLM-4(3),MPLM-5(4) and MPRK3 applied to Test 2

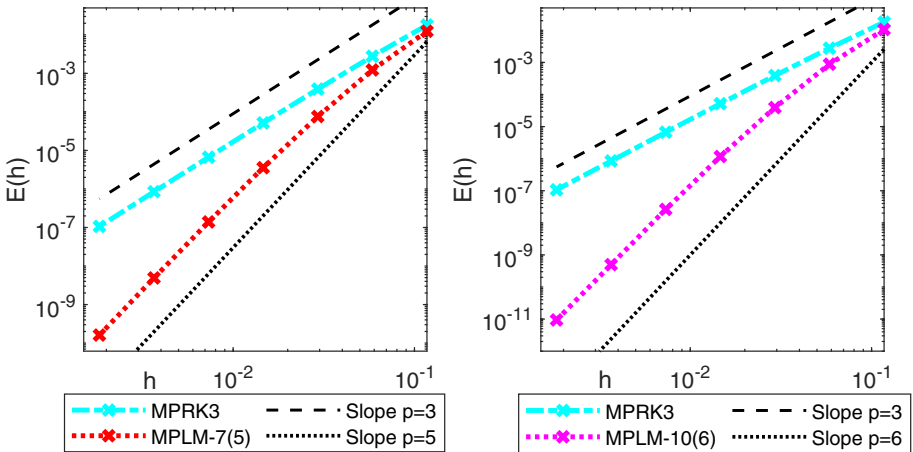


Fig. 9 Experimental order for MPLM-7(5), MPLM-10(6) and MPRK3 applied to Test 2

4.3 Test 3: Brusselator Test

Our next experiment addresses a typical nonlinear chemical kinetics problem modeled by the original Brusselator system [2, 13, 31]

$$\begin{aligned}
 y_1'(t) &= -k_1 y_1(t), \\
 y_2'(t) &= -k_2 y_2(t) y_5(t), \\
 y_3'(t) &= k_2 y_2(t) y_5(t), \\
 y_4'(t) &= k_4 y_5(t), \\
 y_5'(t) &= k_1 y_1(t) - k_2 y_2(t) y_5(t) + k_3 y_5^2(t) y_6(t) - k_4 y_5(t), \\
 y_6'(t) &= k_2 y_2(t) y_5(t) - k_3 y_5^2(t) y_6(t),
 \end{aligned}$$

with $y^0 = \begin{pmatrix} 10 \\ 10 \\ 0 \\ 0 \\ 0.1 \\ 0.1 \end{pmatrix}$.

(4.25)

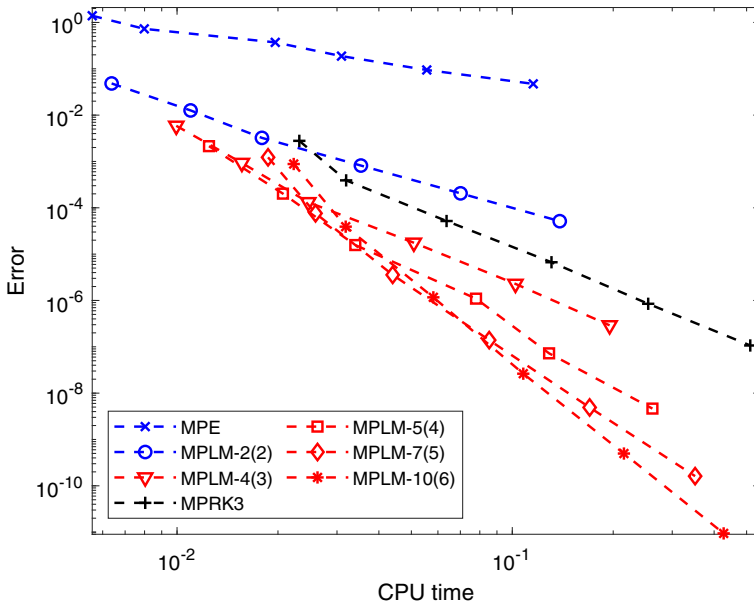


Fig. 10 Work precision diagram: error versus CPU time for the different methods applied to Test 2. $h = T/2^{8+m}$, $m = 1, \dots, 6$.

The differential system (4.25) falls in the form (1.1), setting

$$\begin{aligned}
 p_{32}(\mathbf{y}) = d_{23}(\mathbf{y}) &= k_2 y_2 y_5, & p_{45}(\mathbf{y}) = d_{54}(\mathbf{y}) &= k_4 y_5, & p_{51}(\mathbf{y}) = d_{15}(\mathbf{y}) &= k_1 y_1, \\
 p_{56}(\mathbf{y}) = d_{65}(\mathbf{y}) &= k_3 y_5^2 y_6, & p_{65}(\mathbf{y}) = d_{56}(\mathbf{y}) &= k_2 y_2 y_5,
 \end{aligned}$$

and $p_{ij}(\mathbf{y}) = d_{ji}(\mathbf{y}) = 0$ for all other combinations of i and j in $[1, 6] \cap \mathbb{N}$. Resorting to [42, Theorem 1.2] the existence of a unique non-negative solution to (4.25) can be proved.

For this test problem the σ -embedding technique (3.20) is not suitable for use due the presence of zero components in the initial value \mathbf{y}^0 . As already pointed out, we set $y_3^0 = y_4^0 = \text{realmin} \approx 2.26 \cdot 10^{-308}$. The simulation of the system (4.25) by the MPLM-10(6) method with $h = 1.25 \cdot 2^{-5}$, assuming $t \in [0, 10]$ and $k_l = 1$, $l = 1, \dots, 4$, results in the numerical solution of Fig. 12. Accordingly to the theoretical findings of Lemmas 1 and 2, both the positivity and the conservativity are guaranteed. Furthermore, the results of Table 4 and Figs. 13 and 14 underline the effectiveness of the choice to slightly variate the initial state. The work precision diagram of Fig. 15 confirms that, assuming the same mean execution time, the MPLM- $k(p)$ methods for $p \geq 3$ attain higher accuracy than the MPRK3 scheme.

Figure 16 illustrates the mean error, as defined in (4.23), plotted against the discretization stepsize for the MPLM- $k(p)$ and MPDeC(p) methods. The results reveal that for $p \in \{3, 4\}$ the latter exhibit greater accuracy, whereas for $p \in \{5, 6\}$ the MPLM integrators yield better results. Figure 17 shows, for $p \geq 3$, a comparison in terms of the accuracy-cost trade off, highlighting the advantages of the MPLM- $k(p)$ methods in terms of computational efficiency as well. The outcomes of the comparison, which differ from those obtained in Test 1 and Test 2, align with the inherent characteristics of each scheme. Specifically, MPDeC methods address multiple linear systems at each step, whereas MPLM methods necessitate an initialization procedure and the calculation of the PWDs. Consequently, for systems with smaller dimensions, the former demonstrate better efficiency. Conversely, as the number

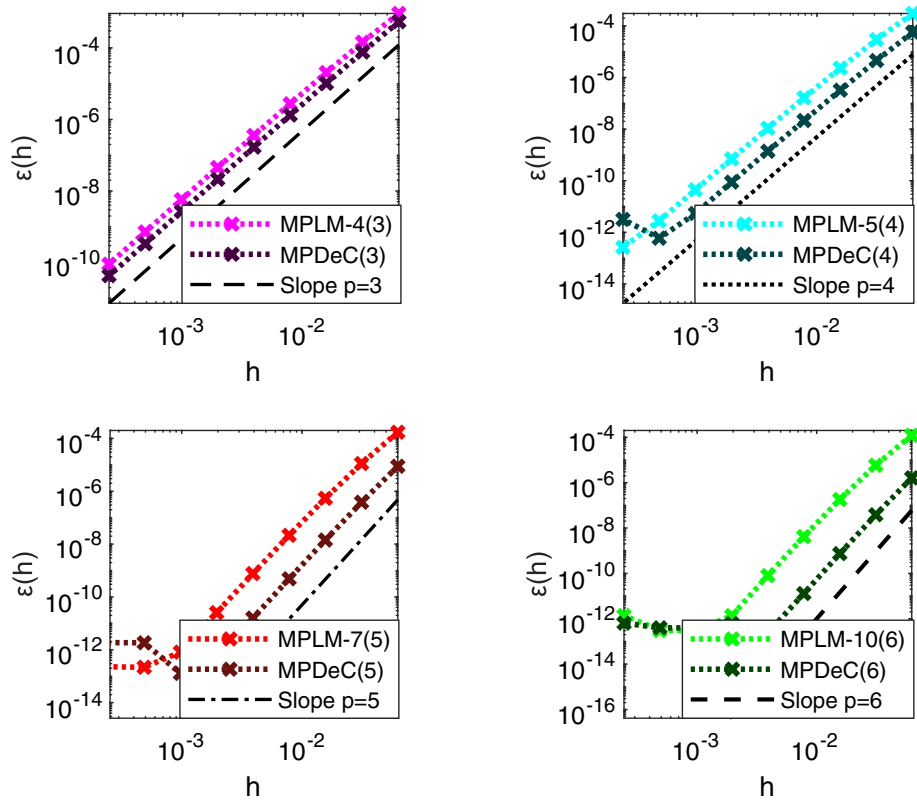


Fig. 11 Accuracies, in terms of (4.23), of MPLM and MPDeC schemes applied to Test 2. $h = T/2^{8+m}$, $m = 1, \dots, 6$.

of components increases, MPLM methods exhibit reduced computational complexity and become more advantageous.

4.4 Test 4: SACEIRQD COVID-19 Model

Our fourth test problem deals with the modified Susceptible-Infected-Recovered-Dead epidemic model

$$\begin{aligned}
 S'(t) &= - \left(\alpha + \frac{\beta I(t) + \sigma A(t)}{N_P} + \eta \right) S(t), \\
 A'(t) &= -\tau A(t) + \xi E(t), \\
 C'(t) &= \alpha S(t) - \mu C(t), \\
 E'(t) &= \left(\frac{\beta I(t) + \sigma A(t)}{N_P} + \eta \right) S(t) + \mu C(t) - (\gamma + \xi) E(t), \\
 I'(t) &= \tau A(t) + \gamma E(t) - \delta I(t), \\
 R'(t) &= \lambda Q(t), \\
 Q'(t) &= \delta I(t) - \lambda Q(t) - k_d Q(t), \\
 D'(t) &= k_d Q(t).
 \end{aligned}
 \tag{4.26}$$

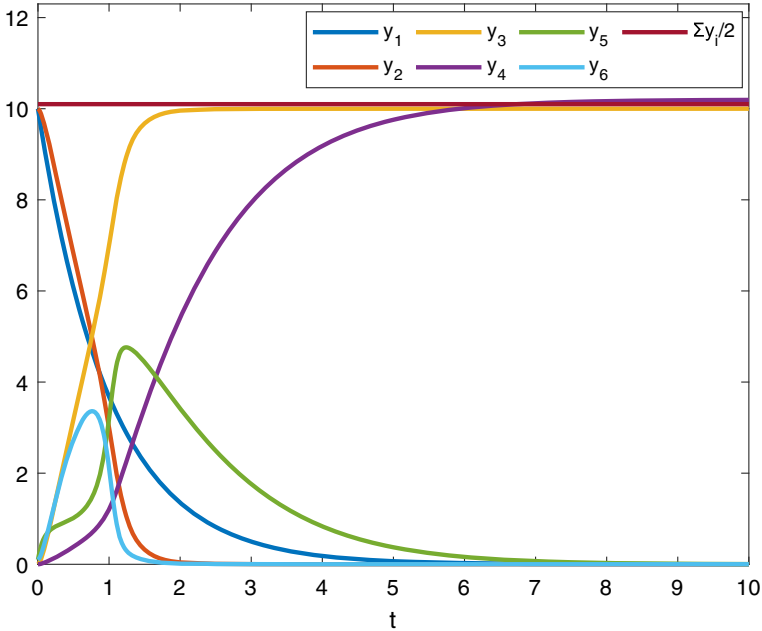


Fig. 12 Numerical solution of Test 3 by MPLM-10(6) with $h = 1.25 \cdot 2^{-5}$.

Table 4 Experimental convergence of the numerical solutions to Test 3

$0.80 \cdot h$	$E(h)$	\hat{p}	$0.80 \cdot h$	$E(h)$	\hat{p}	$0.80 \cdot h$	$E(h)$	\hat{p}
MPE			MPLM-2(2)			MPLM-4(3)		
2^{-5}	$2.30 \cdot 10^0$	–	2^{-5}	$5.44 \cdot 10^{-1}$	–	2^{-5}	$1.87 \cdot 10^{-1}$	–
2^{-6}	$1.31 \cdot 10^0$	0.82	2^{-6}	$1.77 \cdot 10^{-1}$	1.62	2^{-6}	$4.29 \cdot 10^{-2}$	2.13
2^{-7}	$6.86 \cdot 10^{-1}$	0.93	2^{-7}	$5.21 \cdot 10^{-2}$	1.77	2^{-7}	$8.04 \cdot 10^{-3}$	2.42
2^{-8}	$3.49 \cdot 10^{-1}$	0.97	2^{-8}	$1.43 \cdot 10^{-2}$	1.87	2^{-8}	$1.28 \cdot 10^{-3}$	2.65
2^{-9}	$1.76 \cdot 10^{-1}$	0.99	2^{-9}	$3.75 \cdot 10^{-3}$	1.93	2^{-9}	$1.83 \cdot 10^{-4}$	2.81
2^{-10}	$8.82 \cdot 10^{-2}$	1.00	2^{-10}	$9.62 \cdot 10^{-4}$	1.96	2^{-10}	$2.46 \cdot 10^{-5}$	2.90
2^{-11}	$4.42 \cdot 10^{-2}$	1.00	2^{-11}	$2.44 \cdot 10^{-4}$	1.98	2^{-11}	$3.19 \cdot 10^{-6}$	2.95
2^{-12}	$2.21 \cdot 10^{-4}$	1.00	2^{-12}	$6.13 \cdot 10^{-5}$	1.99	2^{-12}	$4.07 \cdot 10^{-7}$	2.97
MPLM-5(4)			MPLM-7(5)			MPLM-10(6)		
2^{-5}	$8.40 \cdot 10^{-2}$	–	2^{-5}	$5.80 \cdot 10^{-2}$	–	2^{-5}	$4.68 \cdot 10^{-2}$	–
2^{-6}	$1.54 \cdot 10^{-2}$	2.44	2^{-6}	$8.60 \cdot 10^{-3}$	2.75	2^{-6}	$5.66 \cdot 10^{-3}$	3.05
2^{-7}	$1.90 \cdot 10^{-3}$	3.03	2^{-7}	$7.48 \cdot 10^{-4}$	3.52	2^{-7}	$3.89 \cdot 10^{-4}$	3.86
2^{-8}	$1.78 \cdot 10^{-4}$	3.41	2^{-8}	$4.70 \cdot 10^{-5}$	3.99	2^{-8}	$1.75 \cdot 10^{-5}$	4.47
2^{-9}	$1.40 \cdot 10^{-5}$	3.67	2^{-9}	$2.27 \cdot 10^{-6}$	4.37	2^{-9}	$5.41 \cdot 10^{-7}$	5.02
2^{-10}	$9.93 \cdot 10^{-7}$	3.82	2^{-10}	$9.09 \cdot 10^{-8}$	4.64	2^{-10}	$1.27 \cdot 10^{-8}$	5.42
2^{-11}	$6.63 \cdot 10^{-8}$	3.90	2^{-11}	$3.26 \cdot 10^{-9}$	4.80	2^{-11}	$2.48 \cdot 10^{-10}$	5.68
2^{-12}	$4.30 \cdot 10^{-9}$	3.95	2^{-12}	$1.09 \cdot 10^{-10}$	4.90	2^{-12}	$1.63 \cdot 10^{-11}$	3.92

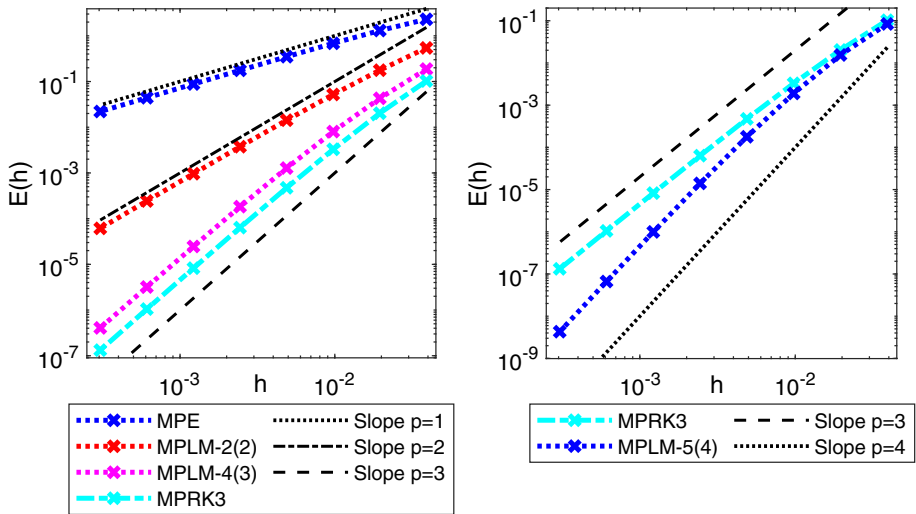


Fig. 13 Experimental order for MPE, MPLM-2(2), MPLM-4(3), MPLM-5(4) and MPRK3 applied to Test 3

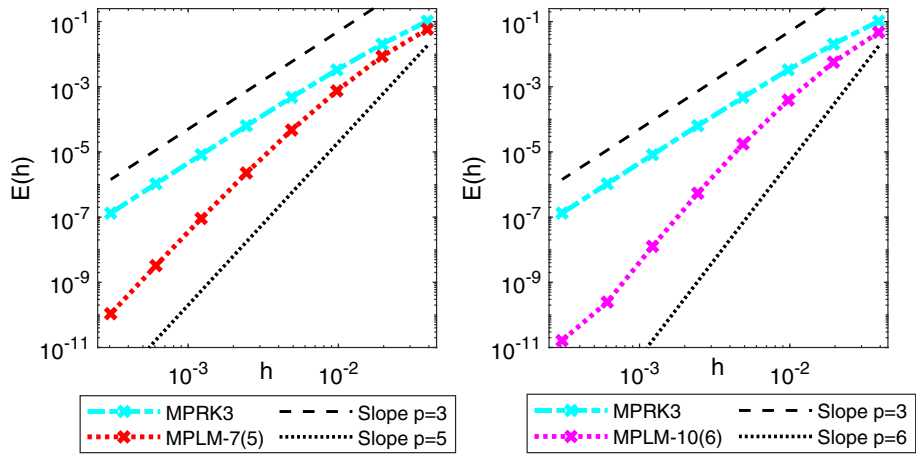


Fig. 14 Experimental order for MPLM-7(5), MPLM-10(6) and MPRK3 applied to Test 3

Originally introduced in [40] to analyze COVID-19 data, it incorporates the effect of asymptomatic infections and the influence of containment, isolation and quarantine measures on the spread of the disease. The non-negative states variables of the model (4.26) represent the sizes at time t (days) of the eight disjoint compartments in which a closed population of N_P individuals is partitioned: susceptible (S), asymptomatic (A), confined (C), exposed (E), infected (I), recovered (R), quarantined (Q), dead (D). The absence of migration turnover in the model leads to the conservation law

$$S(t) + A(t) + C(t) + E(t) + I(t) + R(t) + Q(t) + D(t) = N_P, \quad \forall t \geq 0.$$

We refer to [40] for the details on the physical interpretation of the positive constants $\alpha, \beta, \gamma, \delta, \sigma, \eta, \tau, \xi, \mu, \lambda$ and k_d .

In order to reformulate the system (4.26) as a PDS, we introduce the function

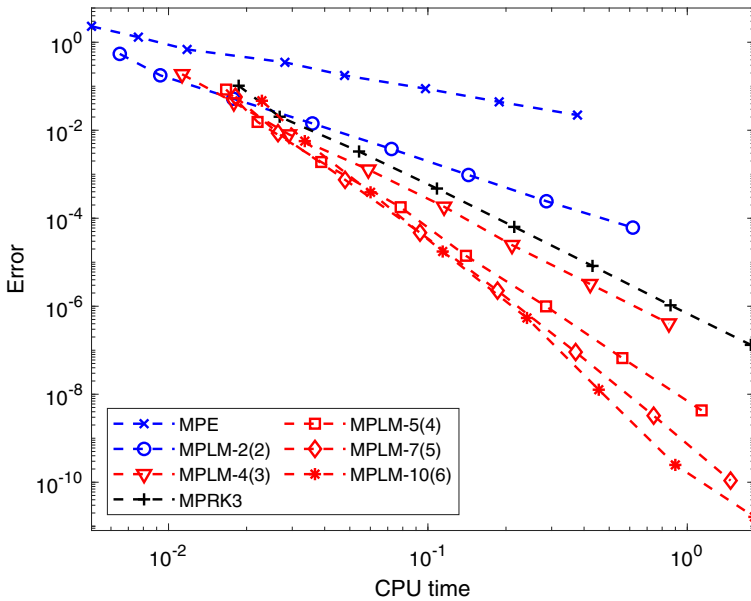


Fig. 15 Work precision diagram: error versus CPU time for the different methods applied to Test 3. $h = T/2^{8+m}$, $m = 0, \dots, 7$.

$y(t) = (S(t), A(t), C(t), E(t), I(t), R(t), Q(t), D(t))^T$ and the production-destruction terms as follows

$$\begin{aligned}
 p_{24}(y) &= d_{42}(y) = \xi y_4, & p_{31}(y) &= d_{13}(y) = \alpha y_1, \\
 p_{41}(y) &= d_{14}(y) = y_1 \left(\eta + \frac{\beta y_5 + \sigma y_2}{N_P} \right), & p_{43}(y) &= d_{34}(y) = \mu y_3, \\
 p_{52}(y) &= d_{25}(y) = \tau y_2, & p_{54}(y) &= d_{45}(y) = \gamma y_4, \\
 p_{67}(y) &= d_{76}(y) = \lambda y_7, & p_{75}(y) &= d_{57}(y) = \delta y_5, \\
 p_{87}(y) &= d_{78}(y) = K_d y_7
 \end{aligned}$$

and $p_{ij}(y) = d_{ji}(y) = 0$ for all other combinations of $i, j \in [1, 8] \cap \mathbb{N}$.

In [40, Table 1] the parameters of the model were fitted to the COVID-19 time series datasets of infected, recovered and death cases for different countries. Here, we adopt for the parameters of the model the values provided by the authors for Italy, i.e.

$$\begin{aligned}
 N_P &= 6.046 \cdot 10^7, & \alpha &= 0.0194, & \beta &= 7.567, & \mu &= 2.278 \cdot 10^{-6}, \\
 \eta &= 9.180 \cdot 10^{-7}, & \sigma &= 1.4633 \cdot 10^{-3}, & \tau &= 1.109 \cdot 10^{-4}, & \xi &= 0.263, \\
 \gamma &= 0.021, & \delta &= 0.077, & \lambda_0 &= 0.157, & \lambda_1 &= 0.025, \\
 k_{d0} &= 0.779, & k_{d1} &= 0.061, & y^0 &= (60459997, 0, 0, 1, 1, 0, 1, 0)^T.
 \end{aligned} \tag{4.27}$$

Furthermore, we set $\lambda = 10^{-4} \lambda_0 \int_0^{10^4} e^{-\lambda_1 t} dt$ and $k_d = 10^{-4} k_{d0} \int_0^{10^4} e^{-k_{d1} t} dt$.

The mathematical statements concerning the positivity (Lemma 1) and conservativity (Lemma 2) of the MPLM methods are confirmed in Fig. 18, where the numerical simulation of (4.26) by the MPLM-10(6) for $t \in [0, 180]$ is reported. For this test, to avoid the issues of null components in the initial value, we set $y_i^0 = \text{realmin} \approx 2.26 \cdot 10^{-308}$ for $i = 2, 3, 6, 8$.

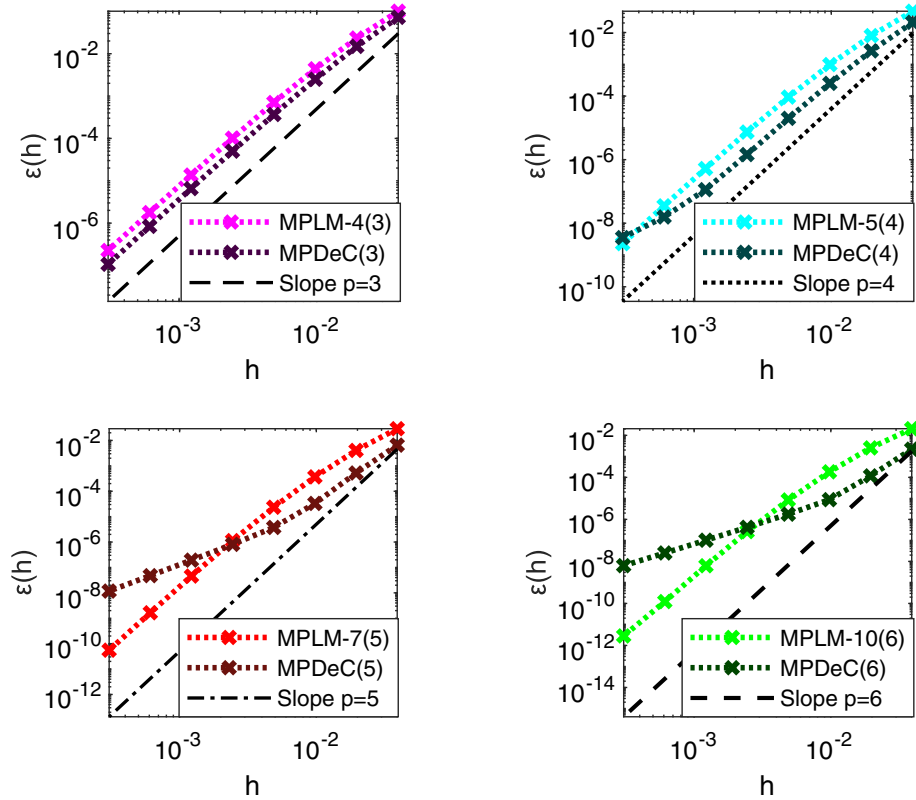


Fig. 16 Accuracies, in terms of (4.23), of MPLM and MPDeC schemes applied to Test 3. $h = 2^{-m}$, $m = 5, \dots, 12$.

Furthermore, due to the presence of large values in the components of the solution, we consider the relative maximum error

$$e(h) = \frac{E(h)}{\max_{0 \leq n \leq T/h} \|\mathbf{y}^{n(ref)}\|_\infty},$$

with $E(h)$ defined in (4.21). From Table 5 and Figs. 19 and 20 it is clear that the numerical solution behaves in accordance with the theoretical results and the experimental order of convergence coincides with the expected one. Moreover, the work precision diagram of Fig. 21 confirms, also for this test, the trends observed in Test 3.

The efficient integration of (4.26)–(4.27) using the MPDeC codes in [35] is not feasible and results in a numerical solution exhibiting several NAN. To overcome this issue, we consider the system (4.26) with the same parameters in (4.27) but with modified initial values

$$\bar{\mathbf{y}}^0 = (60459997, 10^{-10}, 10^{-10}, 1, 1, 10^{-10}, 1, 10^{-10})^\top, \quad \text{and} \quad \bar{\bar{\mathbf{y}}}^0 = 10^4 \mathbf{e}. \quad (4.28)$$

The comparisons of the MPLM- $k(p)$ and MPDeC(p) methods applied to Test 4, initialized with $\mathbf{y}(0) = \bar{\mathbf{y}}^0$ and $\mathbf{y}(0) = \bar{\bar{\mathbf{y}}}^0$, are presented in Figs. 22 and 24, respectively. Here, the error is computed as detailed in (4.23). The outcomes of the simulations show that MPLM methods are more accurate in the realistic scenario where $\mathbf{y}(0) = \bar{\mathbf{y}}^0$ (see Fig. 22), whereas

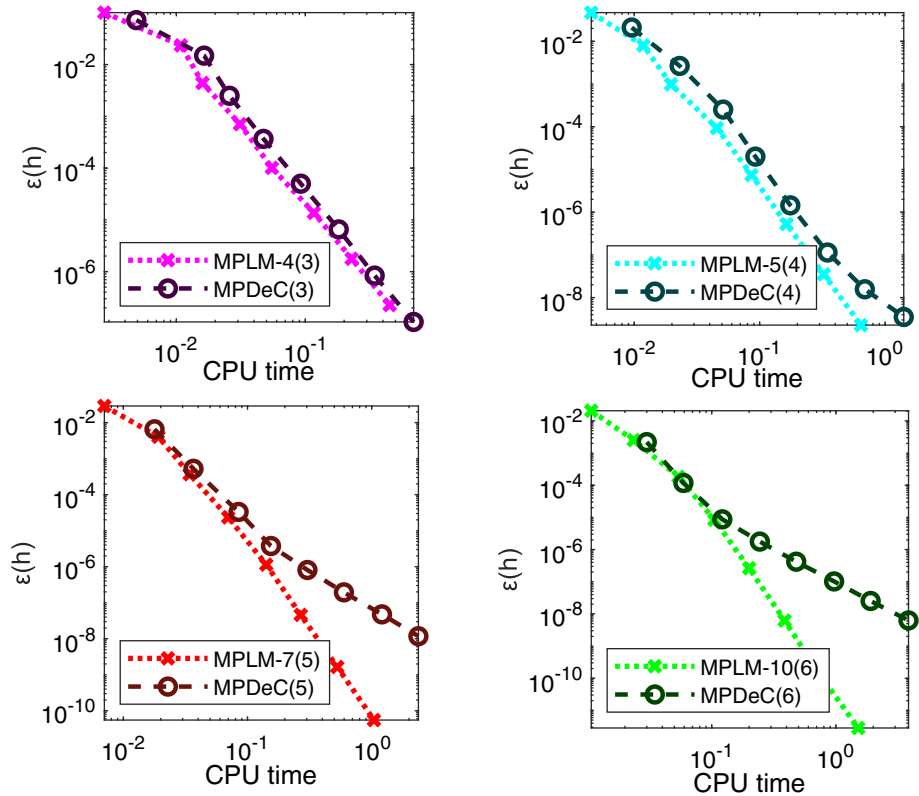


Fig. 17 Work precision diagram: mean error versus CPU time for the different methods applied to Test 3. $h = 2^{-m}$, $m = 5, \dots, 12$.

the MPDeC integrators are preferable in the case $y(0) = \bar{y}^0$ (see Fig. 24). The work precision diagrams associated with the MPLM and MPDeC simulations of Test 4, initialized with $y(0) = \bar{y}^0$ and $y(0) = \bar{\bar{y}}^0$, are depicted in Figs. 23 and 25 where it is clear that, at least for $p > 3$, MPLM methods require fewer computational resources than MPDeC schemes to achieve a predefined level of accuracy. On the basis of the comparative analysis we performed, the MPLM integrators emerge as more suitable for accurately simulating real-world scenarios of the infectious disease outbreak model (4.26). Furthermore, even in the unrealistic case of $y(0) = \bar{\bar{y}}^0$, they demonstrate superior efficiency compared to MPDeC methods.

4.5 Test 5: Spatially Heterogeneous Diffusion Equation

With the aim of assessing the performance of MPLM schemes on larger-scale problems, we turn our attention to the following Partial Differential Equation (PDE)

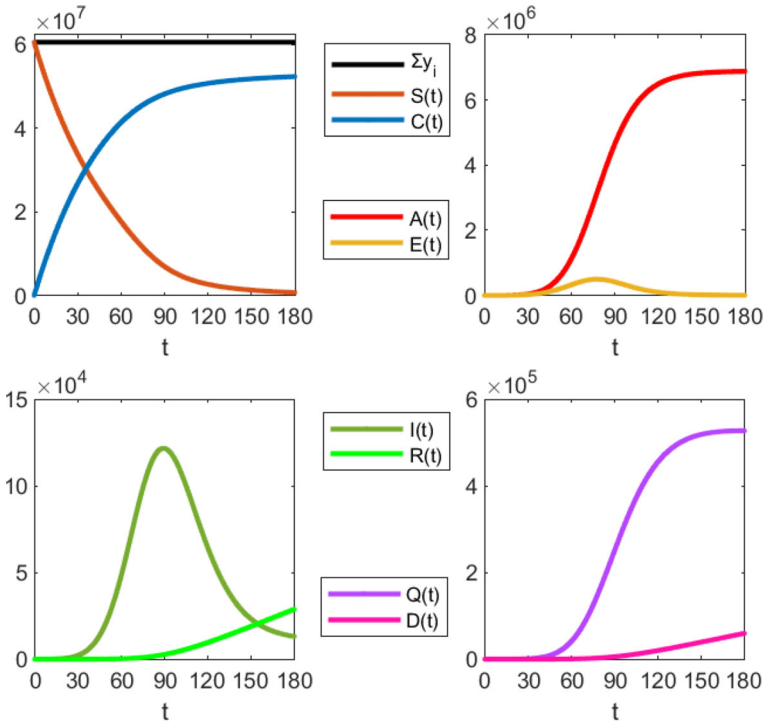


Fig. 18 Numerical solution of Test 4 by MPLM-10(6) with $h = 1.4 \cdot 2^{-1}$.

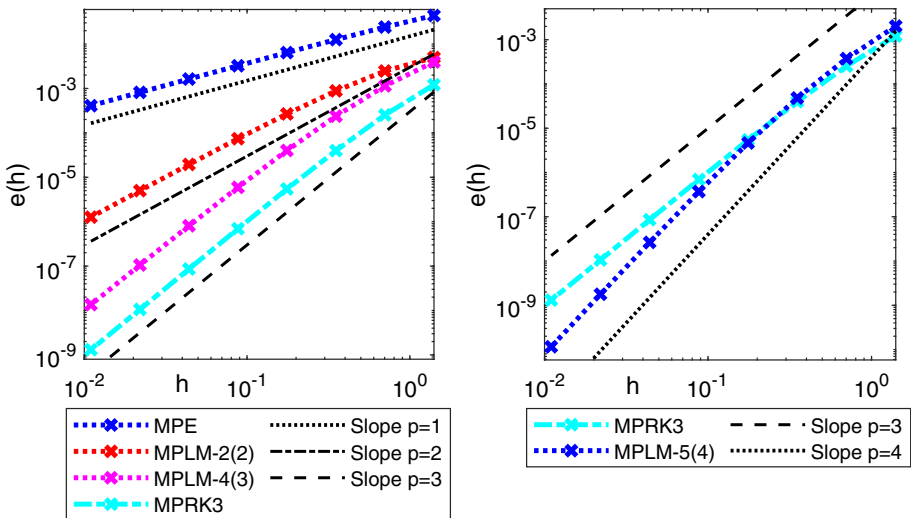


Fig. 19 Experimental order for MPE, MPLM-2(2), MPLM-4(3), MPLM-5(4) and MPRK3 applied to Test 4

Table 5 Experimental convergence of the numerical solutions to Test 4

$0.35 \cdot h$	$e(h)$	\hat{p}	$0.35 \cdot h$	$e(h)$	\hat{p}	$0.35 \cdot h$	$e(h)$	\hat{p}
MPE			MPLM-2(2)			MPLM-4(3)		
2^{-1}	$4.39 \cdot 10^{-2}$	–	2^{-1}	$4.98 \cdot 10^{-3}$	–	2^{-1}	$3.96 \cdot 10^{-3}$	–
2^{-2}	$2.41 \cdot 10^{-2}$	0.87	2^{-2}	$2.47 \cdot 10^{-3}$	1.01	2^{-2}	$1.14 \cdot 10^{-3}$	1.79
2^{-3}	$1.26 \cdot 10^{-2}$	0.94	2^{-3}	$8.82 \cdot 10^{-4}$	1.49	2^{-3}	$2.38 \cdot 10^{-4}$	2.26
2^{-4}	$6.42 \cdot 10^{-3}$	0.97	2^{-4}	$2.67 \cdot 10^{-4}$	1.72	2^{-4}	$4.02 \cdot 10^{-5}$	2.56
2^{-5}	$3.24 \cdot 10^{-3}$	0.99	2^{-5}	$7.38 \cdot 10^{-5}$	1.85	2^{-5}	$5.92 \cdot 10^{-6}$	2.76
2^{-6}	$1.63 \cdot 10^{-3}$	0.99	2^{-6}	$1.94 \cdot 10^{-5}$	1.92	2^{-6}	$8.09 \cdot 10^{-7}$	2.87
2^{-7}	$8.17 \cdot 10^{-4}$	1.00	2^{-7}	$4.99 \cdot 10^{-6}$	1.96	2^{-7}	$1.06 \cdot 10^{-7}$	2.93
2^{-8}	$4.09 \cdot 10^{-4}$	1.00	2^{-8}	$1.27 \cdot 10^{-6}$	1.98	2^{-8}	$1.36 \cdot 10^{-8}$	2.97
MPLM-5(4)			MPLM-7(5)			MPLM-10(6)		
2^{-1}	$2.03 \cdot 10^{-3}$	–	2^{-1}	$1.55 \cdot 10^{-3}$	–	2^{-1}	$8.56 \cdot 10^{-4}$	–
2^{-2}	$3.76 \cdot 10^{-4}$	2.43	2^{-2}	$1.50 \cdot 10^{-4}$	3.38	2^{-2}	$6.25 \cdot 10^{-5}$	3.78
2^{-3}	$4.83 \cdot 10^{-5}$	2.96	2^{-3}	$1.36 \cdot 10^{-5}$	3.46	2^{-3}	$4.43 \cdot 10^{-6}$	3.82
2^{-4}	$4.65 \cdot 10^{-6}$	3.38	2^{-4}	$8.41 \cdot 10^{-7}$	4.02	2^{-4}	$1.86 \cdot 10^{-7}$	4.57
2^{-5}	$3.71 \cdot 10^{-7}$	3.65	2^{-5}	$3.94 \cdot 10^{-8}$	4.42	2^{-5}	$5.32 \cdot 10^{-9}$	5.13
2^{-6}	$2.64 \cdot 10^{-8}$	3.81	2^{-6}	$1.54 \cdot 10^{-9}$	4.67	2^{-6}	$1.17 \cdot 10^{-10}$	5.50
2^{-7}	$1.78 \cdot 10^{-9}$	3.90	2^{-7}	$5.46 \cdot 10^{-11}$	4.82	2^{-7}	$2.39 \cdot 10^{-12}$	5.62
2^{-8}	$1.16 \cdot 10^{-10}$	3.94	2^{-8}	$1.89 \cdot 10^{-12}$	4.85	2^{-8}	$6.43 \cdot 10^{-13}$	1.90

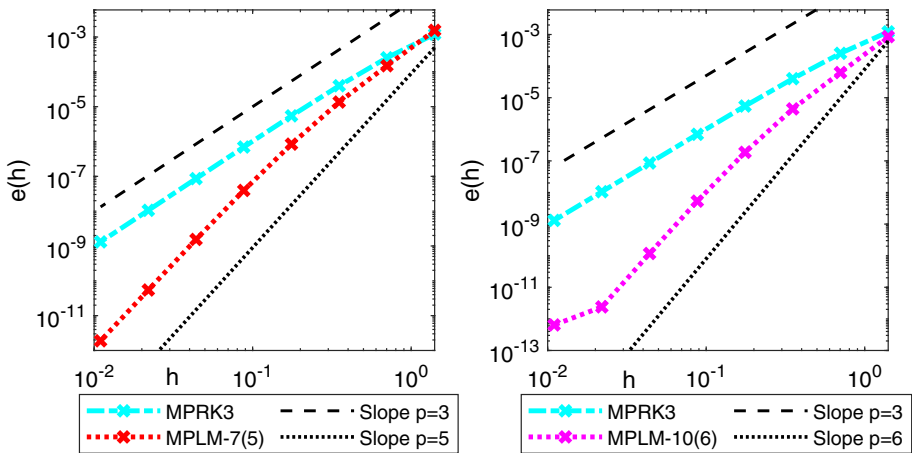


Fig. 20 Experimental order for MPLM-7(5), MPLM-10(6) and MPRK3 applied to Test 4

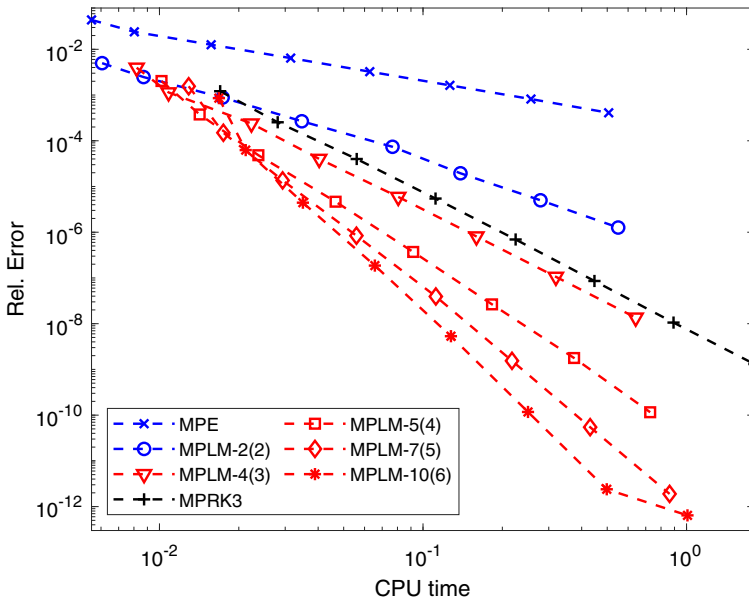


Fig. 21 Work precision diagram: error versus CPU time for the different methods applied to Test 4. $h = T/2^{7+m}$, $m = 0, \dots, 7$.

$$\frac{\partial u(x, t)}{\partial t} - \frac{\partial}{\partial x} \left(\mathfrak{D}(x) \frac{\partial u(x, t)}{\partial x} \right) = 0, \quad \begin{matrix} 0 \leq x \leq L, \\ 0 \leq t \leq T. \end{matrix} \tag{4.29}$$

The PDE (4.29) serves as a versatile mathematical framework for modeling a broad spectrum of phenomena extending beyond chemical diffusion and heat conduction [43]. For instance, in [24], it is derived from the Black-Scholes model, thereby broadening its applications to qualitative finance. Similarly, in [32], the same equation is utilized to describe nutrient uptake by plant root hairs. Furthermore, as argued in [10], it may be applicable in simulating the onset of corrosion in concrete bridge beams due to chloride ion diffusion.

Our experiment is conducted within a conservative setting, wherein we define initial conditions and Neumann zero-flux boundary conditions as follows

$$u(x, 0) = f(x), \quad \frac{\partial u}{\partial x}(0, t) = \frac{\partial u}{\partial x}(L, t) = 0. \tag{4.30}$$

In this scenario, the conservation law

$$\int_0^L u(x, t) dx = \int_0^L f(x) dx, \quad \forall t \geq 0, \tag{4.31}$$

holds true since integration with respect to x of both sides of (4.29) leads to

$$\frac{d}{dt} \int_0^L u(x, t) dx = \mathfrak{D}(L) \frac{\partial u(L, t)}{\partial x} - \mathfrak{D}(0) \frac{\partial u(0, t)}{\partial x} = 0.$$

In order to obtain a numerical solution to equation (4.29)–(4.30) that preserves positivity and satisfies a discrete equivalent of (4.31), we introduce a finite volume semi-discretization for the spatial variable (see, for instance, [33, Chapter 4]) and subsequently employ modified Patankar schemes to integrate the resulting system of ordinary differential equations.

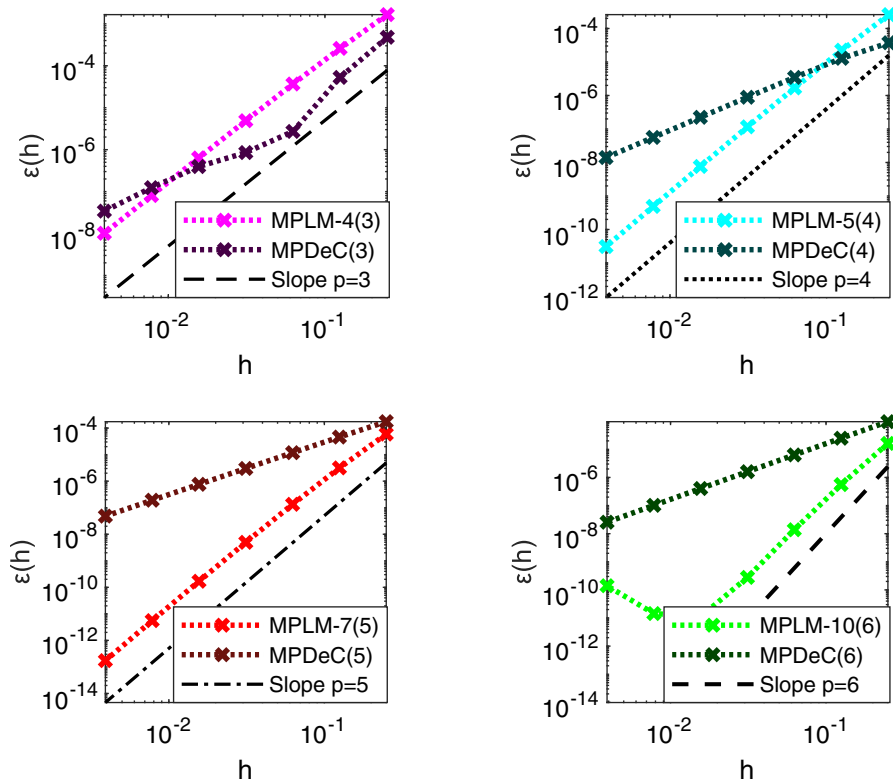


Fig. 22 Accuracies, in terms of (4.23), of the different methods applied to Test 4 with $y(0) = \bar{y}^0$ in (4.28). $h = 2^{-m}$, $m = 2, \dots, 8$.

4.5.1 Spatial Semi-discretization and Conservative PDS

Let $\Delta x > 0$ and $\{x_j\}_{j \geq 0}$ be a uniform mesh such that $x_j = (j + \frac{1}{2}) \Delta x$, represents the center of a one-dimensional cell with edges $x_{j-\frac{1}{2}}$ and $x_{j+\frac{1}{2}}$. Consider the approximations $v_j(t)$ of the average solution value over each grid cell, defined as

$$v_j(t) \approx u_j(t) = \frac{1}{\Delta x} \int_{x_{j-\frac{1}{2}}}^{x_{j+\frac{1}{2}}} u(x, t) dx, \quad j = 0, \dots, N_x, \tag{4.32}$$

with $\Delta x N_x = L$. Integrating the PDE (4.29) over the j -th cell and dividing by Δx yields the exact differential rule

$$u'_j(t) = -\frac{1}{\Delta x} \left(F_{j+\frac{1}{2}}(t) - F_{j-\frac{1}{2}}(t) \right), \quad j = 0, \dots, N_x, \tag{4.33}$$

where $F_{j \pm \frac{1}{2}}(t) = -\mathfrak{D}\left(x_{j \pm \frac{1}{2}}\right) \partial_x u\left(x_{j \pm \frac{1}{2}}, t\right)$ denotes the flux through the edges of the cell. A semi-discrete scheme is then derived by a midpoint approximation of the fluxes in (4.33). Specifically, for the interior cells ($j = 1, \dots, N_x - 1$), we set

$$v'_j(t) = \frac{\mathfrak{D}\left(x_{j+\frac{1}{2}}\right) v_{j+1}(t) - \left(\mathfrak{D}\left(x_{j+\frac{1}{2}}\right) - \mathfrak{D}\left(x_{j-\frac{1}{2}}\right)\right) v_j(t) + \mathfrak{D}\left(x_{j-\frac{1}{2}}\right) v_{j-1}(t)}{\Delta x^2}, \tag{4.34}$$

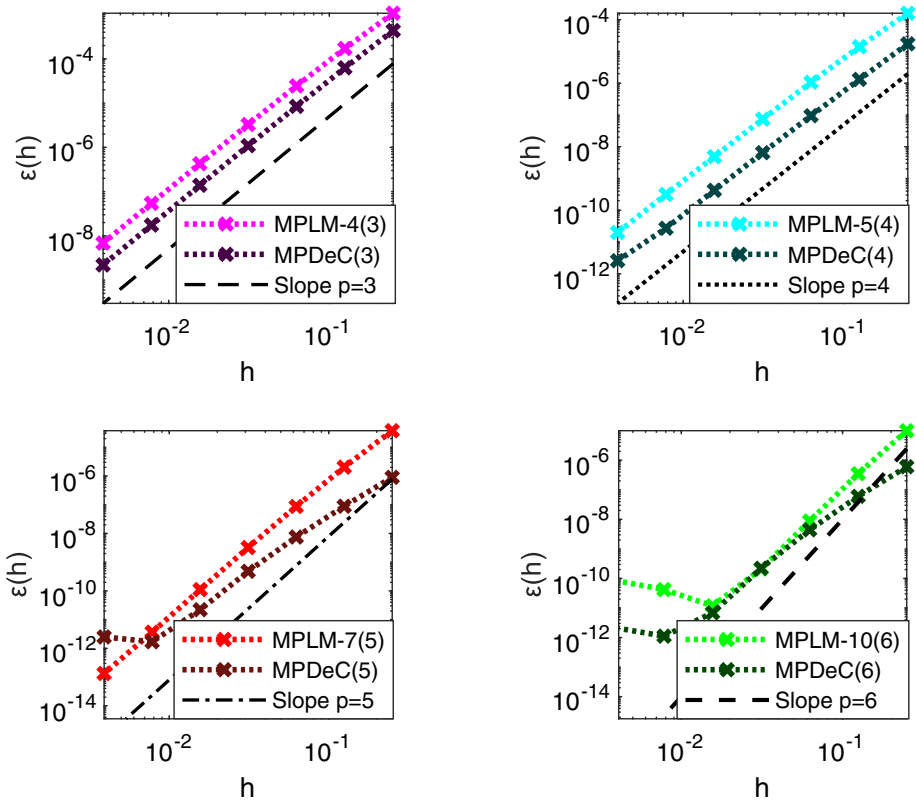


Fig. 24 Accuracies, in terms of (4.23), of the different methods applied to Test 4 with $y(0) = \bar{y}^0$ in (4.28). $h = 2^{-m}$, $m = 2, \dots, 8$.

with $\mathbf{v}(t) = (v_0(t), \dots, v_{N_x}(t))^T \in \mathbb{R}^{N_x+1}$ and $\mathfrak{D}_j = \mathfrak{D}(x_j)$, for each j . The following result proves, for the solution to (4.36), the semi-discrete counterpart of the conservation law (4.31).

Theorem 6 *Let $u(x, t)$ be the continuous solution to (4.29)–(4.30) for $(x, t) \in [0, L] \times [0, T]$, with positive L and T . Let $\{v_j(t)\}_{j \geq 0}$ be its approximation (in the sense of (4.32)) computed by (4.36) with $\Delta x = L/N_x$. Then, independently of $\Delta x > 0$,*

$$\Delta x \sum_{j=0}^{N_x} v_j(t) = \Delta x \sum_{j=0}^{N_x} f(x_j), \quad \forall t \geq 0. \tag{4.37}$$

Proof Given $\mathbf{f} = (f(x_0), \dots, f(x_{N_x}))^T$, the equality $\Delta x \mathbf{e}^T \mathbf{v}(0) = \Delta x \mathbf{e}^T \mathbf{f}$ directly comes from the boundary conditions (4.30). Therefore, for assuring (4.37), it suffices to show that $(\mathbf{e}^T v_j(t))' = 0$. The matrix $A \in \mathbb{R}^{(N_x+1) \times (N_x+1)}$ is symmetric, hence $\mathbf{e}^T \mathbf{v}'(t) = \mathbf{e}^T (A \mathbf{v}(t)) = (A \mathbf{e})^T \mathbf{v}(t) = \mathbf{0}^T \mathbf{v}(t) = 0$, which yields the result. \square

The differential system (4.36) can be rewritten as a PDS of the form (1.3) with $\mathbf{y}(t) = \mathbf{v}(t) \in \mathbb{R}^{N_x+1}$, $D(\mathbf{y}(t)) = P(\mathbf{y}(t))^T \in \mathbb{R}^{(N_x+1) \times (N_x+1)}$ and

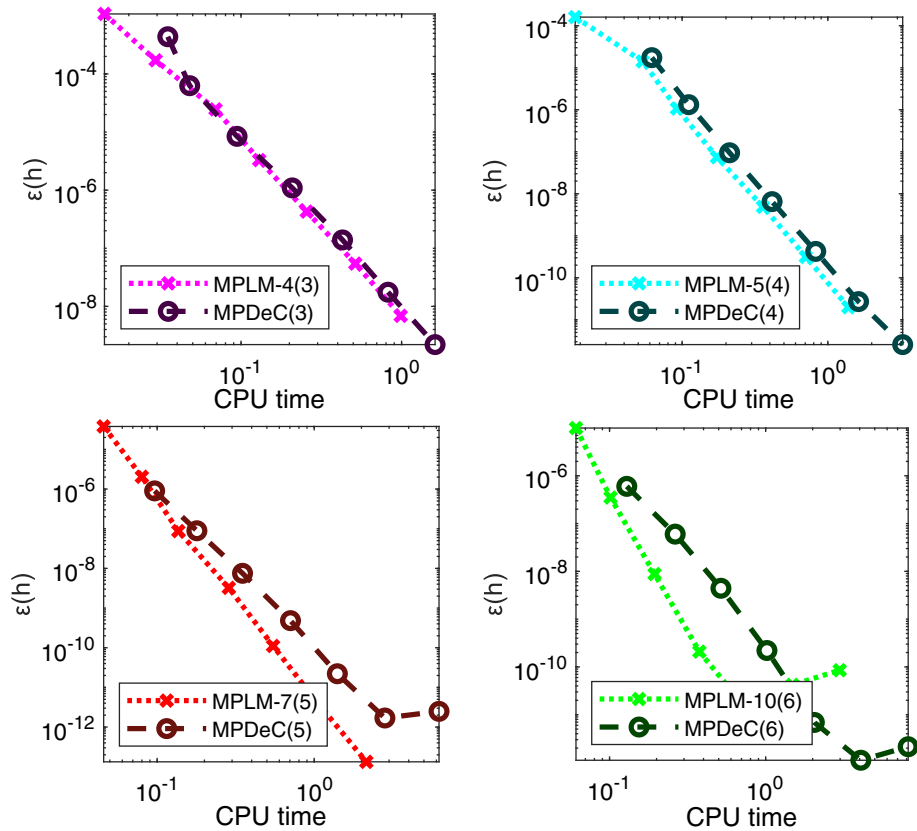


Fig. 25 Work precision diagram: mean error versus CPU time for the different methods applied to Test 4 with $y(0) = \bar{y}^0$ in (4.28). $h = 2^{-m}$, $m = 2, \dots, 8$.

$$P(y) = \frac{1}{\Delta x^2} \begin{pmatrix} 0 & y_2 \mathcal{D}_{\frac{1}{2}} & & & \\ y_1 \mathcal{D}_{\frac{1}{2}} & 0 & y_3 \mathcal{D}_{\frac{3}{2}} & & \\ & \ddots & \ddots & \ddots & \\ & & y_{N_x-2} \mathcal{D}_{N_x-\frac{3}{2}} & 0 & y_{N_x} \mathcal{D}_{N_x-\frac{1}{2}} \\ & & & y_{N_x-1} \mathcal{D}_{N_x-\frac{1}{2}} & 0 \end{pmatrix}. \quad (4.38)$$

Thus, once the equivalence of (4.36) with a fully conservative PDS is established, the property (4) proved with Theorem 6 automatically follows from (1.6).

4.5.2 Simulation Results

For our numerical experiments, we consider the PDE (4.29)–(4.30) with $L = 1$, $T = 60$ and initial condition

$$f(x) = 2 - 2 \sin^2 \left(\frac{\pi}{2} - \frac{1}{4} \right).$$

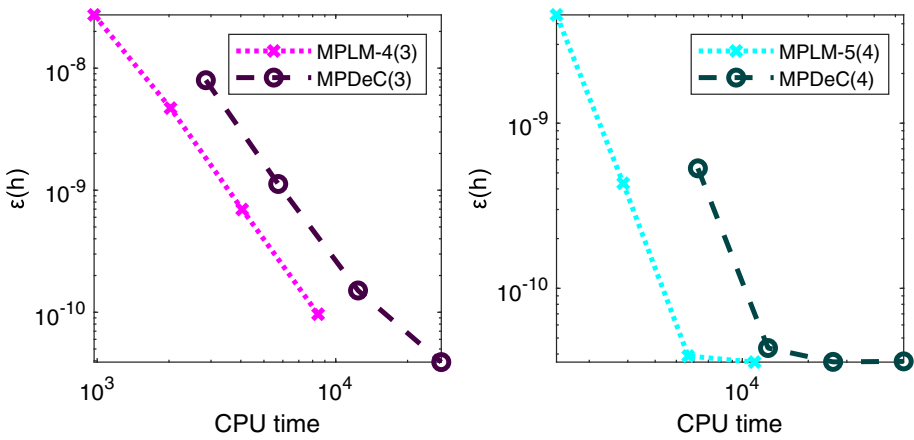


Fig. 26 Work precision diagram: mean error versus CPU time for the different methods applied to Test 5 with $\Delta x = 10^{-2}$ and $h = 2^{-m}$, $m = 9, \dots, 12$.

Table 6 Mean errors, CPU times and numerical residuals for the different methods applied to Test 5 with $\Delta x = 10^{-2}$

h	$\varepsilon(h)$	CPU time	$r(h)$	h	$\varepsilon(h)$	CPU time	$r(h)$
MPLM-4(3)				MPDeC(3)			
2^{-9}	$2.73 \cdot 10^{-8}$	$9.71 \cdot 10^2$	$1.42 \cdot 10^{-15}$	2^{-9}	$7.97 \cdot 10^{-9}$	$2.85 \cdot 10^3$	$1.47 \cdot 10^{-13}$
2^{-10}	$4.71 \cdot 10^{-9}$	$2.03 \cdot 10^3$	$4.55 \cdot 10^{-15}$	2^{-10}	$1.12 \cdot 10^{-9}$	$5.74 \cdot 10^3$	$2.84 \cdot 10^{-16}$
2^{-11}	$6.95 \cdot 10^{-10}$	$4.05 \cdot 10^3$	$3.84 \cdot 10^{-14}$	2^{-11}	$1.51 \cdot 10^{-10}$	$1.24 \cdot 10^4$	$2.85 \cdot 10^{-16}$
2^{-12}	$9.64 \cdot 10^{-11}$	$8.43 \cdot 10^3$	$6.11 \cdot 10^{-14}$	2^{-12}	$3.90 \cdot 10^{-11}$	$2.77 \cdot 10^4$	$1.56 \cdot 10^{-12}$
MPLM-5(4)				MPDeC(4)			
2^{-9}	$4.53 \cdot 10^{-9}$	$1.42 \cdot 10^3$	$1.85 \cdot 10^{-15}$	2^{-9}	$5.33 \cdot 10^{-10}$	$6.27 \cdot 10^3$	$1.42 \cdot 10^{-16}$
2^{-10}	$4.32 \cdot 10^{-10}$	$2.86 \cdot 10^3$	$3.41 \cdot 10^{-15}$	2^{-10}	$4.35 \cdot 10^{-11}$	$1.31 \cdot 10^4$	$1.42 \cdot 10^{-16}$
2^{-11}	$3.89 \cdot 10^{-11}$	$5.66 \cdot 10^3$	$3.55 \cdot 10^{-15}$	2^{-11}	$3.58 \cdot 10^{-11}$	$2.60 \cdot 10^4$	$6.11 \cdot 10^{-12}$
2^{-12}	$3.57 \cdot 10^{-11}$	$1.14 \cdot 10^4$	$2.50 \cdot 10^{-14}$	2^{-12}	$3.61 \cdot 10^{-11}$	$5.46 \cdot 10^4$	$5.30 \cdot 10^{-12}$

Moreover, a space-dependent diffusion coefficient

$$\mathfrak{D}(x) = D_0 \left(x - \frac{2}{3} \right)^2 \frac{\tan^{-1}(2x - 3)}{(2x - 3)} + 10^{-5}, \quad \text{with } D_0 = 10^{-2},$$

is introduced to simulate heterogeneous diffusion phenomena (see, for instance, [45] and references therein).

Here, the solution to the PDE is approximated by integrating the PDS (1.3)–(4.38) with modified Patankar methods. Since the stability investigation of the semi-discretization (4.36) falls outside the scope of this work, we empirically adjust the temporal step size h to be smaller than the spatial one Δx . The outcomes of the simulation by the MPLM-7(5) scheme with $h = 5 \cdot 10^{-4}$ and $\Delta x = 5 \cdot 10^{-3}$, are presented in Fig. 27.

Table 6 presents the mean errors, as defined in (4.23), along with the execution times for

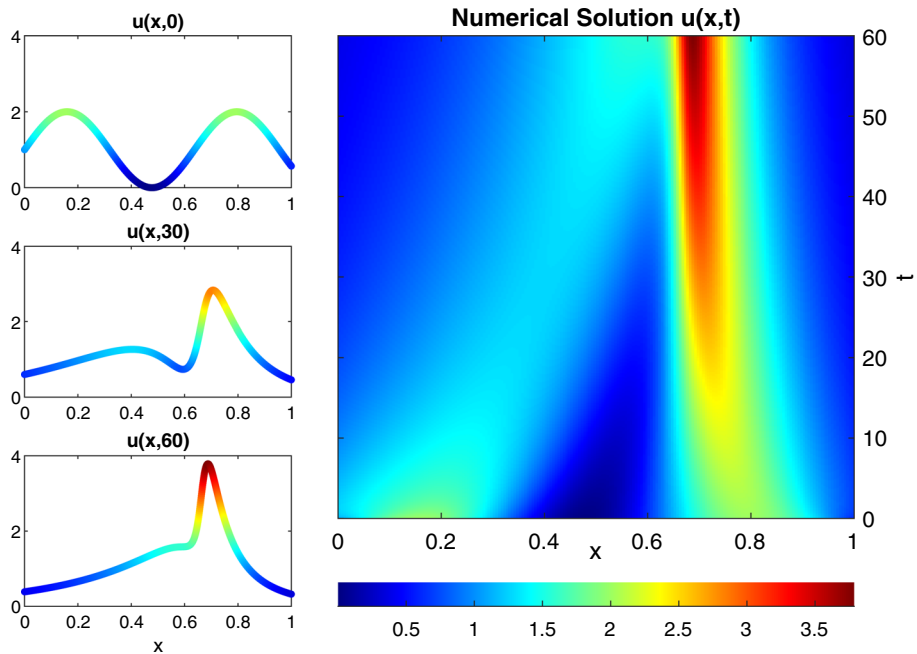


Fig. 27 Numerical solution of Test 5 by MPLM-7(5) with $N_x = 2000$ and $h = 5 \cdot 10^{-4}$.

the Julia implementations of both the MPLM and MPDeC methods. The numerical residual

$$r(h) = \Delta x \max_{0 \leq n \leq \frac{T}{h}} \left(\sum_{j=0}^{N_x} |v_j(t_n) - f(x_j)| \right),$$

on the semi-discrete conservation law (6) is there reported, as well. The experimental results reveal that the MPLM methods demonstrate superior performances (cf. Fig. 26) and exhibit a more regular behavior for the residual $r(h)$.

5 Conclusions and Perspectives

In this manuscript, we introduced accurate linearly implicit time integrators specifically designed for production-destruction differential systems. Notably, the extension of the modified Patankar technique to multistep schemes has resulted in conservative numerical methods which retain, with no restrictions on the discretization steplength, the positivity of the solution and the linear invariant of the system. We carried out a theoretical investigation of the properties of the Patankar weight denominators which ensure the consistency and convergence of the proposed schemes. Additionally, we devised an embedding technique to practically compute the PWDs and achieve arbitrarily high order of convergence. The numerical tests conducted on various problems provided experimental confirmation of the theoretical findings. The comparison with the third-order modified Patankar Runge–Kutta method presented in [26, Lemma 6, Case II with $\gamma = 0.5$] highlighted the superior performance of the proposed MPLM- $k(p)$ integrators. Furthermore, the MPLM- $k(p)$ schemes proved to be competitive

with the high order modified Patankar deferred correction discretizations in [36], especially in the case of high-dimensional systems and vanishing initial states.

Given the results of this paper, MPLM- $k(p)$ methods demonstrate considerable potential for the numerical integration of production-destruction systems. However, several aspects require deeper analysis, which we plan to address in future work. First of all, the optimization of coefficients selection for the underlying linear multistep method may be investigated, considering various combinations of k and p to ensure both positivity constraint and high order of convergence. The possibility of an extension to include negative coefficients may be considered, as well. Furthermore, since the methods of Table 1 exhibit reduced efficiency on stiff ODEs tests such as the Robertson problem [18, 27], the necessity arises of a comprehensive MPLM stability analysis. In this regard, the implementation of variable stepsize approaches and local error control strategies may reveal of high interest. However, due to the multistep nature of the MPLM methods, achieving efficient steplength adaptation appears more challenging compared to one-step modified Patankar discretizations, for which such strategies have already been successfully developed and applied [18, 26, 42].

Appendix A: Consistency Necessary Condition

In Sect. 3 we introduced, with Theorem 2, a condition on the Patankar weight denominators which leads to order p consistent MPLM schemes. Here, our objective is to prove Theorem 3 showing that (3.12) represents a necessary requirement for the consistency, as well. To do that, we consider the particular production-destruction system (1.1) with

$$P_i(\mathbf{y}) = \begin{cases} \mu y_i, & \text{if } i = j^*, \\ 0, & \text{otherwise,} \end{cases} \quad \text{and} \quad D_i(\mathbf{y}) = \begin{cases} \mu y_i, & \text{if } i = i^*, \\ 0, & \text{otherwise,} \end{cases} \quad (5.39)$$

where i^* and j^* are fixed indices in $\{1, \dots, N\}$ and μ is a given positive constant. It can be proved that

$$y_{i^*}(t) = \exp(-\mu t), \quad y_{j^*}(t) = 2 - \exp(-\mu t), \quad y_l(t) = 1, \quad \begin{matrix} 1 \leq l \leq N, \\ l \neq i^*, l \neq j^*, \end{matrix}$$

is the unique solution of the positive and fully conservative PDS (1.1)–(5.39). The following investigation outlines the behaviour of the MPLM discretizations of the class (2.9) applied to (1.1)–(5.39).

Proof of Theorem 3 The order p convergence of the underlying LM discretization and the consistency hypothesis on the MPLM- k method (2.9), imply

$$\delta_i^{LM}(h; t_n) = \mathcal{O}(h^{p+1}) \quad \text{and} \quad \delta_i(h; t_n) = \mathcal{O}(h^{p+1}), \quad n \geq k, \quad i = 1, \dots, N,$$

where the local errors $\delta_i^{LM}(h; t_n)$ and $\delta_i(h; t_n)$ are defined in (3.13) and (3.11), respectively. Subtracting the former from the latter yields

$$\begin{aligned} & \sum_{r=1}^k \beta_r \sum_{j=1}^N \left(p_{ij}(\mathbf{y}(t_{n-r})) \left(1 - \frac{y_j(t_n)}{\sigma_j(\mathbf{y}(t_{n-1}), \dots, \mathbf{y}(t_{n-k}))} \right) \right) \\ & - \sum_{r=1}^k \beta_r \sum_{j=1}^N \left(d_{ij}(\mathbf{y}(t_{n-r})) \left(1 - \frac{y_i(t_n)}{\sigma_i(\mathbf{y}(t_{n-1}), \dots, \mathbf{y}(t_{n-k}))} \right) \right) = \mathcal{O}(h^p), \end{aligned}$$

for $i = 1, \dots, N$ and $n \geq k$. Furthermore, for the particular PDS (5.39), taken $i = i^*$,

$$\mu \sum_{r=1}^k \beta_r y_{i^*}(t_{n-r}) \left(1 - \frac{y_{i^*}(t_n)}{\sigma_{i^*}(\mathbf{y}(t_{n-1}), \dots, \mathbf{y}(t_{n-k}))} \right) = \mathcal{O}(h^p).$$

Therefore, the result comes from the positivity of the system and the arbitrariness of the choice of $1 \leq i^* \leq N$. \square

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Data availability Data sharing not applicable to this article as no datasets were generated or analysed during the current study.

Declarations

Conflict of interest The authors declare that they have no Conflict of interest.

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