RELAXED MODULUS-BASED SYNCHRONOUS MULTISPLITTING MULTI-PARAMETERS TOR (TWO-PARAMETERS OVER-RELAXATION) METHODS FOR LINEAR COMPLEMENTARITY PROBLEMS

Li-Tao Zhang^{1,2,3,†}, Ying-Chao Zhao⁴, Yi-Fan Zhang¹ and Sheng-Kun Li^5

Abstract In 2013, Bai and Zhang [Numerical Linear Algebra with Applications, 20(2013), 425–439] constructed modulus-based synchronous multisplitting methods by an equivalent reformulation of the linear complementarity problems into a system of fixed-point equations and studied the convergence of them. In 2014, Zhang and Li [Computers and Mathematics with Application, 67(2014), 1954–1959] analyzed and obtained the weaker convergence results for linear complementarity problems. In 2008, Zhang et.al. [International Journal of Computer Mathematics, 85(2), 2008, 211–224] presented global relaxed non-stationary multisplitting multi-parameter method by introducing some relaxed parameters. In this paper, we generalize Bai and Zhang's methods and study relaxed modulus-based synchronous multisplitting multi-parameters TOR (two-parameters over-relaxation, abbreviated as TOR) methods for linear complementarity problems. Furthermore, the convergence results of our new method in this paper are given when the system matrix is an H_+ -matrix.

Keywords Modulus-based method, linear complementarity, successive relaxation, H_+ -matrix.

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[†]The corresponding author. Email: litaozhang@163.com(L. Zhang)

¹School of Mathematics, Zhengzhou University of Aeronautics, Zhengzhou, Henan, 450015, China

²College of Mathematics and Information Science, Henan Normal University, Xinxiang, Henan, 453007, China

³Henan province Synergy Innovation Center of Aviation economic development, Zhengzhou, Henan, 450015, China

⁴Department of Sport and Public Art, Zhengzhou University of Aeronautics, Zhengzhou, Henan, 450015, China

⁵College of Applied Mathematics, Chengdu University of Information Technology, Chengdu, Sichuan, 610225, China

1. Introduction

Consider the linear complementarity problems, abbreviated as LCP(q, A), for finding a pair of real vectors r and $z \in \mathbb{R}^n$ such that

$$r := Az + q \ge 0, z \ge 0 \text{ and } z^T (Az + q) = 0,$$
 (1.1)

where $A = (a_{ij}) \in \mathbb{R}^{n \times n}$ is a given large, sparse and real matrix and $q = (q_1, q_2, ..., q_n)^T \in \mathbb{R}^n$ is a given real vector. Here, z^T and \geq denote the transpose of the vector z and the componentwise partial ordering between two vectors, respectively.

Many problems in scientific computing and engineering applications may lead to solutions of LCPs of the form (1.1). For example, the linear complementarity problems may arise from application problems such as the convex quadratic programming, the Nash equilibrium point of the bimatrix game, the free boundary problems of fluid dynamics etc. (e.g. see [12, 18, 23] and the references therein). Some solvers for LCP(q, A) with a special matrix A were proposed [2-8, 13-15, 17, 20, 31]. Recently, many people have focused the solvers of LCP(q, A) with an algebra equation [7,8,17,20,31,35]. Bai introduced modulus-based matrix splitting methods [7], and Bai and Zhang proposed modulus-based matrix multisplitting methods for linear complementarity problems [1,8]. Zhang and Ren [35] extended the condition of a compatible H-splitting to that of an H-splitting. Li [21] extended the modulusbased matrix splitting iteration method to more general cases. Zhang et al. [37–39] gave the weaker convergence results when the system matrix is an H_+ -matrix. Dai et al. [16] provided a comparison theorem between preconditioned two-step modulusbased Gauss-Seidel (PTMGS) iteration method and two-step modulus-based Gauss-Seidel (TMGS) iteration method, which shows that PTMGS method improves the convergence rate of original TMGS method for linear complementarity problem. Mezzadri et al. [22] analyzed the solution of horizontal linear complementarity problems arising from finite-difference discretizations of differential problems. Ren et al. [26, 27] considered the preconditioned general two-step modulus-based matrix splitting iteration method and the general two-sweep modulus-based matrix splitting iteration method for linear complementarity problems of H_+ -matrices. Wen et al. [30] studied the relaxation modulus-based matrix splitting iteration method for solving linear complementarity problems of positive definite matrices. Wu et al. [32] provided the preconditioned general modulus-based matrix splitting iteration method for linear complementarity problems of *H*-matrices. Zhang et al. [34] studied the modified modulus-based multigrid method for linear complementarity problems arising from free boundary problems.

The rest of this paper is organized as follows: In section 2, we give some notations and lemmas. In section 3, we propose relaxed modulus-based synchronous multisplitting multi-parameters TOR (two-parameters over-relaxation, abbreviated as TOR) method for solving LCP(q, A). For more information about the TOR method, readers can refer to the literature [21]. In section 4, we give the convergence analysis for the proposed method.

2. Notations and Lemmas

A matrix $A = (a_{ij})$ is called an *M*-matrix if $a_{ij} \leq 0$ for $i \neq j$ and $A^{-1} \geq 0$. The comparison matrix $\langle A \rangle = (\alpha_{ij})$ of matrix $A = (a_{ij})$ is defined by: $\alpha_{ij} = |a_{ij}|$,

if $i = j; \alpha_{ij} = -|a_{ij}|$, if $i \neq j$. A matrix A is called an H-matrix if $\langle A \rangle$ is an M-matrix and is called an H_+ -matrix if it is an H-matrix with positive diagonal entries [2, 11, 30]. Let $\rho(A)$ denote the spectral radius of A. A representation A = M - N is called a *splitting* of A when M is nonsingular. Let A and B be M-matrices. If $A \leq B$, then $A^{-1} \geq B^{-1}$. Let A be an H-matrix, and A = D - B, D = diag(A), then $\rho(|D|^{-1}|B|) < 1$. Moreover, D is nonsingular. Finally, we define by $R^n_+ = \{x | x \geq 0, x \in R^n\}$.

Lemma 2.1.([19]) Let A be an H-matrix. Then A is nonsingular, and $|A^{-1}| \leq \langle A \rangle^{-1}$.

Lemma 2.2.([29]) Let $H^{(1)}, H^{(2)}, ..., H^{(l)}$...be a sequence of nonnegative matrices in $\mathbb{R}^{n \times n}$. If there exist a real number $0 \le \theta < 1$, and a vector $\nu > 0$ in \mathbb{R}^n , such that

 $H^{(l)}\nu \leq \theta\nu, \ l=1,2,\ldots$

then $\rho(K_l) \leq \theta^l < 1$, where $K_l = H^{(l)} H^{(l-1)} ... H^{(1)}$, and therefore $\lim_{l \to \infty} K_l = 0$.

Lemma 2.3. ([34]) Let $A = (a_{ij}) \in Z^{n \times n}$ which has all positive diagonal entries. A is an *M*-matrix if and only if $\rho(B) < 1$, where $B = D^{-1}C$, D = diag(A), A = D - C.

Lemma 2.4. ([4, 6, 9]) $A \in \mathbb{R}^{n \times n}$ be an H_+ -matrix. Then, the LCP(q, A) has a unique solution for any $q \in \mathbb{R}^n$.

Lemma 2.5. ([7]) Let A = M - N be a splitting of the matrix $A \in \mathbb{R}^{n \times n}, \Omega$ be a positive diagonal matrix, and γ a positive constant. Then, for the LCP(q, A)the following statements hold true:

(i) if (z,r) is a solution of the LCP(q,A), then $x = \frac{1}{2}\gamma(z - \Omega^{-1}r)$ satisfies the implicit fixed-point equation

$$(\Omega + M)x = Nx + (\Omega - A)|x| - \gamma q; \qquad (2.1)$$

(ii) if x satisfies the implicit fixed-point equation (2), then

$$z = \gamma^{-1}(|\mathbf{x}| + x) \text{ and } \mathbf{r} = \gamma^{-1}\Omega(|\mathbf{x}| - \mathbf{x})$$
 (2.2)

is a solution of the LCP(q, A).

3. RMSMMTOR methods

To suit computational requirements of the modern high-speed multiprocessor environments, in this paper, we further present synchronous parallel multisplitting multi-parameters TOR for the modulus-based splitting iteration methods in [7] by making use of multiple splittings of the system matrix A [10, 26]. This class of modulus-based synchronous multisplitting (MSM) iteration methods is advantageous over the synchronous multisplitting iteration methods discussed by Machida et al. [25] and by Bai [2], as at each iteration step, it only needs to solve subsystems of linear equations rather than linear complementarity subproblems (see also [6,9]).

At first, we introduce the concept of multisplitting method and the detailed process of parallel iterative method.

 $\{M_k, N_k, E_k\}_{k=1}^l$ is a multisplitting of A if 1) $A = M_k - N_k$ is a splitting for k = 1, 2, ..., l; 2) $E_k \ge 0$ is a nonnegative diagonal matrix, called weighting matrix; 3) $\sum_{k=1}^{l} E_k = I$, where I is the identity matrix.

Given a positive diagonal matrix Ω and a positive constant γ , from Lemma 2.5, we know that if x satisfies either of the implicit fixed-point equations

$$(\Omega + M_k)x = N_k x + (\Omega - A)|x| - \gamma q, k = 1, 2, ..., l,$$
(3.1)

then

$$z = \gamma^{-1}(|\mathbf{x}| + x) \text{ and } \mathbf{r} = \gamma^{-1}\Omega(|\mathbf{x}| - \mathbf{x})$$
 (3.2)

is a solution of the LCP(q, A).

Let

$$A = D - L_k - F_k - U_k, \ k = 1, 2, ..., l,$$

where D = diag(A), L_k and F_k are strictly lower triangular, and U_k are such that $A = D - L_k - F_k - U_k$, then $(D - L_k - F_k, U_k, E_k)$ is a multisplitting of A. With the equivalent reformulations (3.1), (3.2) and TOR method of the LCP(q, A), we can establish the following relaxed modulus-based synchronous multisplitting multiparameters TOR method (RMSMMTOR), which is similar to Method 3.1 in [1].

Method 3.1. (The RMSMMTOR method for LCP(q, A))

Let $(M_k, N_k, E_k)(k = 1, 2, ...l)$ be a multisplitting of the system matrix $A \in \mathbb{R}^{n \times n}$. Given an initial vector $x^{(0)} \in \mathbb{R}^n$ for m = 0, 1, ... until the iteration sequence $\{z^{(m)}\}_{m=0}^{\infty} \subset \mathbb{R}^n_+$ is convergent, compute $z^{(m+1)} \in \mathbb{R}^n_+$ by

$$z^{(m+1)} = \frac{1}{\gamma} (|x^{(m+1)}| + x^{(m+1)})$$

and $x^{(m+1)} \in \mathbb{R}^n$ according to

$$x^{(m+1)} = \omega \sum_{k=1}^{l} E_k x^{(m,k)} + (1-\omega) x^{(m)}$$

where $x^{(m,k)}, k = 1, 2, ..., l$, are obtained by solving the linear systems

$$[\alpha_k \Omega + D - (\beta_k L_k + \gamma_k F_k)] x^{(m,k)}$$

=[(1 - \alpha_k)D + (\alpha_k - \beta_k)L_k + (\alpha_k - \gamma_k)F_k + \alpha_k U_k] x^{(m)} + \alpha_k [(\Omega - A)|x^{(m)}| - \gamma q],
k = 1, 2, ..., l, (3.3)

respectively.

Remark 3.1. In Method 3.1, when $\alpha_k = \alpha, \beta_k = \beta, \gamma_k = \gamma, \omega = 1$, the RMSMM-TOR method reduces to the MSMTOR method; when $\alpha_k = \alpha, \beta_k = \beta, \gamma_k = \gamma$, the RMSMMTOR method reduces to the RMSMTOR method; When $\gamma_k = 0, \omega = 1$, the RMSMMTOR method reduces to the MSMMAOR method; When $\gamma_k = 0$, the RMSMMTOR method reduces to the RMSMMAOR method; When $\alpha_k = \alpha, \beta_k = \beta, \gamma_k = 0, \omega = 1$, the RMSMMTOR method reduces to the MSMAOR method; When $\alpha_k = \alpha, \beta_k = \beta, \gamma_k = 0$, the RMSMMTOR method reduces to the RMSMMAOR method; When $\alpha_k = \alpha, \beta_k = \beta, \gamma_k = 0$, the RMSMMTOR method reduces to the RMSMAOR method; When $\alpha_k = \alpha, \beta_k = \beta, \gamma_k = 0$, the RMSMMTOR method reduces to the RMSMAOR method.

4. Convergence analysis

Based relaxed modulus-based synchronous multisplitting multi-parameter TOR method, we will present a weaker convergence results of the multisplitting methods for the linear complementarity problem for H_+ -matrix, which is as follows:

Theorem 4.1. Let $A \in \mathbb{R}^{n \times n}$ be an H_+ -matrix, with D = diag(A) and B = D - A, and let $(M_k, N_k, E_k)(k = 1, 2, ..., l)$ and $(D - L_k - F_k, U_k, E_k)(k = 1, 2, ..., l)$ be a multisplitting and a triangular multisplitting of the matrix A, respectively. Assume that $\gamma > 0$ and the positive diagonal matrix Ω satisfies $\Omega \ge D$. If $A = D - L_k - F_k - U_k(k = 1, 2, ..., l)$ satisfies $\langle A \rangle = D - |L_k| - |F_k| - |U_k|(k = 1, 2, ..., l)$, then the iteration sequence $\{z^{(m)}\}_{m=0}^{\infty}$ generated by the RMSMMTOR iteration method converges to the unique solution z_* of LCP(q, A) for any initial vector $z^{(0)} \in \mathbb{R}^n_+$, provided the relaxation parameters α_k and β_k, ω satisfy

$$0 < \beta_k, \ \gamma_k \le \alpha_k \le 1, \ 0 < \omega < \frac{2}{1+\rho'}$$

or $0 < \beta_k, \ \gamma_k < \frac{1}{\rho(D^{-1}|B|)}, \ 1 < \alpha_k < \frac{1}{\rho(D^{-1}|B|)}, \ 0 < \omega < \frac{2}{1+\rho'},$ (4.1)

where $\rho = \rho(J), J = D^{-1}|B|, \rho' = \max_{1 \le k \le l} \{1 - 2\alpha_k + 2\alpha_k\rho_\epsilon, 2\delta_k\rho_\epsilon - 1, 2\alpha_k\rho_\epsilon - 1\}, \delta_k = \max\{\beta_k, \gamma_k\}$. Moreover, β_k, γ_k should be greater than or less than α_k at once. **Proof.** From Lemma 2.3 and (3.3), for the RMSMMTOR method, it holds that

$$(\alpha_k \Omega + D - (\beta_k L_k + \gamma_k F_k))x_*$$

=[(1 - \alpha_k)D + (\alpha_k - \beta_k)L_k + (\alpha_k - \gamma_k)F_k + \alpha_kU_k]x_* + \alpha_k[(\Omega - A)|x_*| - \gamma q],
k = 1, 2, ..., l, (4.2)

By subtracting (4.2) from (3.3), we have

$$\begin{aligned} x^{(m,k)} - x_* &= (\alpha_k \Omega + D - (\beta_k L_k + \gamma_k F_k))^{-1} \\ &\times [(1 - \alpha_k) D + (\alpha_k - \beta_k) L_k + (\alpha_k - \gamma_k) F_k + \alpha_k U_k] (x^{(m)} - x_*) \\ &+ (\alpha_k \Omega + D - (\beta_k L_k + \gamma_k F_k))^{-1} \alpha_k (\Omega - A) (|x^{(m)}| - |x_*|), \end{aligned}$$

$$k = 1, 2, ..., l, \end{aligned}$$

which immediately results in the error about the RMSMMTOR method as follows:

$$x^{(m+1)} - x_* = \omega \sum_{k=1}^{l} E_k (\alpha_k \Omega + D - (\beta_k L_k + \gamma_k F_k))^{-1} \\ \times [(1 - \alpha_k)D + (\alpha_k - \beta_k)L_k + (\alpha_k - \gamma_k)F_k + \alpha_k U_k](x^{(m)} - x_*) \\ + \omega \sum_{k=1}^{l} E_k (\alpha_k \Omega + D - (\beta_k L_k + \gamma_k F_k))^{-1} \\ \times \alpha_k (\Omega - A)(|x^{(m)}| - |x_*|) + (1 - \omega)(x^{(m)} - x_*).$$

The above error relationship is the base for proving the convergence of RMSMM-TOR method. By taking absolute values on both sides of the above equality, making use of Lemma 2.1 and estimate $||x^{(m)}| - |x_*|| \le |x^{(m)} - x_*|$, defining $\epsilon^{(m)} = x^{(m)} - x_*$ and arranging similar terms together, we can obtain

$$|\epsilon^{(m)}| = |x^{(m+1)} - x_*| \le \mathcal{T}_{RMSMMTOR} |x^{(m)} - x_*|, \tag{4.3}$$

where

$$\mathcal{T}_{RMSMMTOR} = \omega \sum_{k=1}^{l} E_k H_{RMSMMTOR} + |1 - \omega|I$$

$$= \omega \sum_{k=1}^{l} E_k (\alpha_k \Omega + D - (\beta_k |L_k| + \gamma_k |F_k|))^{-1} \qquad (4.4)$$

$$\times [|1 - \alpha_k|D + |\alpha_k - \beta_k||L_k| + |\alpha_k - \gamma_k||F_k|$$

$$+ \alpha_k |U_k| + \alpha_k |\Omega - A|] + |1 - \omega|I.$$

Case 1: Let $0 < \beta_k, \gamma_k \leq \alpha_k \leq 1, 0 < \omega < \frac{2}{1+\rho'}$. For this subcase, the matrix M_k, N_k read as

$$M_{k} = \alpha_{k}\Omega + D - (\beta_{k}|L_{k}| + \gamma_{k}|F_{k}|),$$

$$N_{k} = (1 - \alpha_{k})D + (\alpha_{k} - \beta_{k})|L_{k}| + (\alpha_{k} - \gamma_{k})|F_{k}| + \alpha_{k}|U_{k}| + \alpha_{k}|\Omega - A|.$$
(4.5)

By (4.5) and $|\Omega - A| = (\Omega - D) + |B|, |B| = |L_k| + |F_k| + |U_k|, k = 1, 2, ..., l$, we have $N_k = M_k - 2\alpha_k D + 2\alpha_k |B|$. So

$$H_{RMSMMTOR} = M_k^{-1} N_k = M_k^{-1} (M_k - 2\alpha_k D + 2\alpha_k |B|) = I_k - 2\alpha_k M_k^{-1} (D - |B|).$$

 \mathbf{So}

$$|H_{RMSMMTOR}| \le M_k^{-1} [M_k - 2\alpha_k (D - |B|)]$$

$$\le I - 2\alpha_k M_k^{-1} D(I - D^{-1} |B|).$$

Let e denote the vector $e = (1, 1, ..., 1)^T \in \mathbb{R}^n$. Since J is nonnegative, the matrix $J + \epsilon e e^T$ has only positive entries and irreducible for any $\epsilon > 0$. By the Perron-Frobenius theorem for any $\epsilon > 0$, there is a vector $x_{\epsilon} > 0$ such that

$$(J + \epsilon e e^T) x_{\epsilon} = \rho_{\epsilon} x_{\epsilon},$$

where $\rho_{\epsilon} = \rho(J + \epsilon e e^T) = \rho(J_{\epsilon})$. Moreover, if $\epsilon > 0$ is small enough, we have $\rho_{\epsilon} < 1$ by continuity of the spectral radius. Because of $0 < \alpha_k \le 1$, we also have $1 - 2\alpha_k + 2\alpha_k\rho < 1$, and $1 - 2\alpha_k + 2\alpha_k\rho_{\epsilon} < 1$. So

$$\begin{split} |H_{RMSMMTOR}| &\leq I - 2\alpha_k M_k^{-1} D[I - (D^{-1}|B| + \epsilon e e^T)] \\ &= I - 2\alpha_k M_k^{-1} D[I - J_\epsilon]. \end{split}$$

Multiplying x_{ϵ} in two sides of the above inequality, and $M_k^{-1} \ge D^{-1}$, we can obtain

$$|H_{RMSMMTOR}|x_{\epsilon} \leq x_{\epsilon} - 2\alpha_k M_k^{-1} D[1 - \rho(J_{\epsilon})] x_{\epsilon}$$
$$\leq x_{\epsilon} - 2\alpha_k D^{-1} D[1 - \rho(J_{\epsilon})] x_{\epsilon}$$
$$= (1 - 2\alpha_k + 2\alpha_k \rho(J_{\epsilon})) x_{\epsilon}.$$

By (4.1), we have

$$\begin{aligned} |\mathcal{T}_{RMSMMTOR}|x_{\epsilon} &\leq \omega \Sigma_{k=1}^{l} E_{k} (1 - 2\alpha_{k} + 2\alpha_{k}\rho(J_{\epsilon}))x_{\epsilon} + |1 - \omega|x_{\epsilon} \\ &\leq \omega (1 - 2\alpha_{k} + 2\alpha_{k}\rho_{\epsilon})x_{\epsilon} + |1 - \omega|x_{\epsilon} \\ &= (\omega\rho_{1} + |1 - \omega|)x_{\epsilon} \\ &= \theta_{1} x_{\epsilon} (\epsilon \to 0), \end{aligned}$$

where $\theta_1 = \omega \rho_1 + |1 - \omega| < 1$.

So, the iteration sequence $\{z^{(m)}\}_{m=0}^{\infty}$ generated by the RMSMMTOR iteration method converges to the unique solution z_* of LCP(q, A) for any initial vector $z^{(0)} \in \mathbb{R}^n_+$, provided the relaxation parameters α_k and β_k, ω satisfy $0 < \beta_k, \gamma_k \leq \alpha_k \leq 1, 0 < \omega < \frac{2}{1+\rho'}$.

Case 2: $0 < \beta_k, \gamma_k < \frac{1}{\rho(D^{-1}|B|)}, 1 < \alpha_k < \frac{1}{\rho(D^{-1}|B|)}, 0 < \omega < \frac{2}{1+\rho'}.$ **Subcase 1:** $\alpha_k \ge \beta_k$ and $\alpha_k \ge \gamma_k$. For this subcase, the matrix M_k, N_k read as

$$M_{k} = \alpha_{k}\Omega + D - (\beta_{k}|L_{k}| + \gamma_{k}|F_{k}|),$$

$$N_{k} = (\alpha_{k} - 1)D + (\alpha_{k} - \beta_{k})|L_{k}| + (\alpha_{k} - \gamma_{k})|F_{k}| + \alpha_{k}|U_{k}| + \alpha_{k}|\Omega - A|$$
(4.6)

$$= M_{k} - 2D + 2\alpha_{k}|B|.$$

 So

$$|H_{RMSMMTOR}| \le M_k^{-1} [M_k - 2(D - \alpha_k |B|)]$$

$$\le I - 2M_k^{-1} D(I - \alpha_k D^{-1} |B|).$$

Similar to the Case 1, let e denote the vector $e = (1, 1, ..., 1)^T \in \mathbb{R}^n$, and $x_{\epsilon} > 0$ such that $J_{\epsilon}x_{\epsilon} = (J + \epsilon e e^T)x_{\epsilon} = \rho(J_{\epsilon})x_{\epsilon}$. Moreover, if $\epsilon > 0$ is small enough, we have $\rho_{\epsilon} < 1$ by continuity of the spectral radius. Because of $1 < \alpha_k < \frac{1}{\rho(D^{-1}|B|)}$, we also have

$$2\alpha_k\rho - 1 < 1$$
 and $2\alpha_k\rho_{\epsilon} - 1 < 1$.

 So

$$|H_{RMSMMTOR}| \leq I - 2M_k^{-1}D[I - \alpha_k(D^{-1}|B| + \epsilon ee^T)]$$
$$= I - 2M_k^{-1}D[I - \alpha_k J_\epsilon].$$

Multiplying x_{ϵ} in two sides of the above inequality, and $M_k^{-1} \ge D^{-1}$, we can obtain

$$|H_{RMSMMAOR}|x_{\epsilon} \leq x_{\epsilon} - 2M_{k}^{-1}D[1 - \alpha_{k}\rho(J_{\epsilon})]x_{\epsilon}$$
$$\leq x_{\epsilon} - 2(1 - \alpha_{k}\rho(J_{\epsilon}))]x_{\epsilon}$$
$$= (2\alpha_{k}\rho(J_{\epsilon}) - 1)x_{\epsilon}.$$

By (4.1), we have

$$\mathcal{T}_{RMSMMTOR}|x_{\epsilon} \leq \omega \Sigma_{k=1}^{l} E_{k}(2\alpha_{k}\rho(J_{\epsilon}) - 1)x_{\epsilon} + |1 - \omega|x_{\epsilon}$$
$$\leq \omega(2\alpha_{k}\rho_{\epsilon} - 1)x_{\epsilon} + |1 - \omega|x_{\epsilon}$$
$$= (\omega\rho_{2} + |1 - \omega|)x_{\epsilon}$$
$$= \theta_{2}x_{\epsilon}(\epsilon \to 0),$$

where $\theta_2 = \omega \rho_2 + |1 - \omega| < 1$.

So, the iteration sequence $\{z^{(m)}\}_{m=0}^{\infty}$ generated by the RMSMMTOR iteration method converges to the unique solution z_* of LCP(q, A) for any initial vector $z^{(0)} \in R^n_+$, provided the relaxation parameters $\alpha_k \ge \beta_k$ and $\alpha_k \ge \gamma_k$.

Subcase 2: $\alpha_k \leq \beta_k$ and $\alpha_k \leq \gamma_k$. For this subcase, the matrix M_k, N_k read as

$$M_{k} = \alpha_{k}\Omega + D - (\beta_{k}|L_{k}| + \gamma_{k}|F_{k}|),$$

$$N_{k} = (\alpha_{k} - 1)D + (\beta_{k} - \alpha_{k})|L_{k}| + (\gamma_{k} - \alpha_{k})|F_{k}| + \alpha_{k}|U_{k}| + \alpha_{k}|\Omega - A|$$

$$= M_{k} - 2D + 2\beta_{k}|L_{k}| + 2\gamma_{k}|F_{k}| + 2\alpha_{k}|U_{k}|$$

$$\leq M_{k} - 2D + 2\delta_{k}|B|.$$
(4.7)

where $\delta_k = \max\{\beta_k, \gamma_k\}$. So

$$|H_{RMSMMTOR}| \le M_k^{-1} [M_k - 2(D - \delta_k |B|)]$$

$$\le I - 2M_k^{-1} D(I - \delta_k D^{-1} |B|).$$

Similar to the Case 1, let e denote the vector $e = (1, 1, ..., 1)^T \in \mathbb{R}^n$, and $x_{\epsilon} > 0$ such that $J_{\epsilon}x_{\epsilon} = (J + \epsilon e e^T)x_{\epsilon} = \rho(J_{\epsilon})x_{\epsilon}$. Moreover, if $\epsilon > 0$ is small enough, we have $\rho_{\epsilon} < 1$ by continuity of the spectral radius. Because of $0 < \beta_k, \gamma_k < \frac{1}{\rho(D^{-1}|B|)}$, we also have

$$2\delta_k \rho - 1 < 1$$
 and $2\delta_k \rho_\epsilon - 1 < 1$.

So

$$|H_{RMSMMTOR}| \leq I - 2M_k^{-1}D[I - \delta_k(D^{-1}|B| + \epsilon ee^T)]$$
$$= I - 2M_k^{-1}D[I - \delta_k J_{\epsilon}].$$

Multiplying x_{ϵ} in two sides of the above inequality, and $M_k^{-1} \ge D^{-1}$, we can obtain

$$|H_{RMSMMTOR}|x_{\epsilon} \le x_{\epsilon} - 2(1 - \delta_k \rho(J_{\epsilon}))]x_{\epsilon} = (2\delta_k \rho(J_{\epsilon}) - 1)x_{\epsilon}$$

By (4.1), we have

$$\begin{aligned} |\mathcal{T}_{RMSMMTOR}|x_{\epsilon} &\leq \omega \Sigma_{k=1}^{l} E_{k}(2\delta_{k}\rho(J_{\epsilon}) - 1)x_{\epsilon} + |1 - \omega|x_{\epsilon} \\ &\leq \omega(2\delta_{k}\rho_{\epsilon} - 1)x_{\epsilon} + |1 - \omega|x_{\epsilon} \\ &= (\omega\rho_{3} + |1 - \omega|)x_{\epsilon} \\ &= \theta_{3}x_{\epsilon}(\epsilon \to 0), \end{aligned}$$

where $\theta_{3} = \omega \rho_{3} + |1 - \omega| < 1$.

So, the iteration sequence $\{z^{(m)}\}_{m=0}^{\infty}$ generated by the RMSMMTOR iteration method converges to the unique solution z_* of LCP(q, A) for any initial vector $z^{(0)} \in \mathbb{R}^n_+$, provided the relaxation parameters $\alpha_k \leq \beta_k$ and $\alpha_k \leq \gamma_k$.

Remark 4.1. Obviously, we can find that the conditions of Theorem 4.4 in this paper are weaker than those of Theorem 2.3 in [41]. Moreover, the parameters can be adjusted suitably so that the convergence property of method can be substantially

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improved. That is to say, we have more choices for the splitting A = B - C which makes the multisplitting iteration methods converge. Therefore, our convergence theories extend the scope of multisplitting iteration methods in applications.

Remark 4.2. In this paper, RMSMMTOR method is also the generalization of MSMAOR method in [1] and MSMMAOR method in [38]. The readers can refer to the following three theorems.

Theorem 4.2. ([1]) Let $A \in \mathbb{R}^{n \times n}$ be an H_+ -matrix, with D = diag(A) and B = D - A, and let $(M_k, N_k, E_k)(k = 1, 2, ..., l)$ and $(D - L_k, U_k, E_k)(k = 1, 2, ..., l)$ be a multisplitting and a triangular multisplitting of the matrix A, respectively. Assume that $\gamma > 0$ and the positive diagonal matrix Ω satisfies $\Omega \ge D$. If $A = D - L_k - U_k(k = 1, 2, ..., l)$ satisfies $\langle A \rangle = D - |L_k| - |U_k|(k = 1, 2, ..., l)$, then the iteration sequence $\{z^{(m)}\}_{m=0}^{\infty}$ generated by the MSMAOR iteration method converges to the unique solution z_* of LCP(q, A) for any initial vector $z^{(0)} \in \mathbb{R}^n_+$, provided the relaxation parameters α and β satisfy

$$0 < \beta \le \alpha < \frac{1}{\rho(D^{-1}|B|)}.$$

Theorem 4.3. ([38]) Let $A \in \mathbb{R}^{n \times n}$ be an H_+ -matrix, with D = diag(A) and B = D - A, and let $(M_k, N_k, E_k)(k = 1, 2, ..., l)$ and $(D - L_k, U_k, E_k)(k = 1, 2, ..., l)$ be a multisplitting and a triangular multisplitting of the matrix A, respectively. Assume that $\gamma > 0$ and the positive diagonal matrix Ω satisfies $\Omega \ge D$. If $A = D - L_k - U_k(k = 1, 2, ..., l)$ satisfies $\langle A \rangle = D - |L_k| - |U_k|(k = 1, 2, ..., l)$, then the iteration sequence $\{z^{(m)}\}_{m=0}^{\infty}$ generated by the MSMMAOR iteration method converges to the unique solution z_* of LCP(q, A) for any initial vector $z^{(0)} \in \mathbb{R}^n_+$, provided the relaxation parameters α_k and β_k satisfy

$$0 < \beta_k \le \alpha_k \le 1 \text{ or } 0 < \beta_k < \frac{1}{\rho(D^{-1}|B|)}, 1 < \alpha_k < \frac{1}{\rho(D^{-1}|B|)}.$$

Theorem 4.4. ([38]) Let A be an H-matrix, and for k = 1, 2, ..., l, L_k and F_k be strictly lower triangular matrices. Define the matrix $U_k, k = 1, 2, ..., l$, such that $A = D - L_k - F_k - U_k$. Assume that we have $\langle A \rangle = |D| - |L_k| - |F_k| - |U_k| = |D| - |B|$. If

$$0 \le \beta_k \le \gamma_k, 0 \le \alpha_k \le \gamma_k, 0 < \gamma_k < \frac{2}{1+\rho}, 0 < \omega < \frac{2}{1+\rho_{\gamma_k}},$$

then GRNMMTOR method converges for any initial vector $x^{(0)}$, where $\rho = \rho(J), J = |D|^{-1}|B|, \rho_{\gamma_k} = \max_{1 \le k \le \alpha} \{|1 - \gamma_k| + \gamma_k \rho_\epsilon\}, q(m,k) \ge 1, m = 0, 1, ..., k = 1, 2, ..., l.$

Remark 4.3. From Table 1, we obviously see that the MSMMAOR method in [1] uses the same parameters α, β in different processors, but the RMSMMTOR method in this paper uses different parameters $\alpha_k, \beta_k, \gamma_k (k = 1, 2, ..., l)$ in different processors. In RMSMMTOR method, we may choose proper E_k to balance the load of each processor and avoid synchronization.

5. Numerical experiments

In this section, numerical examples are used to illustrate the feasibility and effectiveness of the relaxed modulus-based synchronous multisplitting multi-parameter

Method	$lpha_k,eta_k,\omega$	Description	Ref
MSMJ	$\alpha_k = 1, \beta_k = 0, \omega = 1$	Modulus-based synchronous	[1]
		multisplitting Jacobi method	
MSMGS	$\alpha_k = \beta_k = 1, \omega = 1$	Modulus-based synchronous	[1]
		multisplitting Gauss-Seidel method	
MSMSOR	$0 < \alpha(\alpha_k) = \beta(\beta_k) < \frac{1}{\rho(D^{-1} B)}, \omega = 1$	Modulus-based synchronous	[1]
		multisplitting SOR method	
MSMAOR	$0 < \beta(\beta_k) \le \alpha(\alpha_k) < \frac{1}{\rho(D^{-1} B)}, \omega = 1$	Modulus-based synchronous	[1]
	P = 1 - 1	multisplitting AOR method	
MSMAOR	$0 < \beta(\beta_k) \le \alpha(\alpha_k) < \frac{1}{\alpha(D^{-1} B)}$	Modulus-based synchronous	[1]
		multisplitting AOR method	
MSMMAOR	$\omega = 1, 0 < \beta_k \le \alpha_k \le 1$ or	Modulus-based synchronous	[34]
	$0 < \beta_k < \frac{1}{a(D^{-1} B)}, 1 < \alpha_k < \frac{1}{a(D^{-1} B)}$	multisplitting multi-parameters	
		AOR method	
RMSMMTOR	$0 < \beta_k, \gamma_k \leq \alpha_k \leq 1, 0 < \omega < \frac{2}{1+\alpha'}$ or	Relaxed modulus-based	this paper
	$0 < \beta_k, \gamma_k < \frac{1}{\rho(D^{-1} B)}, 1 < \alpha_k < \frac{1}{\rho(D^{-1} B)}$	synchronous multisplitting	
	$0 < \omega < \frac{2}{1+a'}$	multi-parameter TOR method	
	where $\rho' = \max_{1 \le k \le l} \{1 - 2\alpha_k + 2\alpha_k \rho_\epsilon,$		
	$2\delta_k\rho_\epsilon - 1, 2\alpha_k\rho_\epsilon - 1\}, \delta_k = \max\{\beta_k, \gamma_k\}$		

Table 1. The relaxed modulus-based synchronous multisplitting multi-parameter method and corresponding convergence results.

AOR methods (RMSMMAOR) (F = U) in terms of iteration count (denoted by 'IT') and computing time (denoted by 'CPU'), and norm of absolute residual vectors (denoted by 'RES'). Here, 'RES' is defined as

$$\operatorname{RES}(z^{(k)}) := \|\min(Az^{(k)} + q, z^{(k)})\|_2$$

where $z^{(k)}$ is the kth approximate solution to the LCP(q, A) and the minimum is taken componentwise in [10].

All initial vectors are chosen to be

$$x^{(0)} = (1, 0, 1, 0, \cdots, 1, 0, \cdots)^T \in \mathbb{R}^n.$$

In the table, α, β denote the iteration parameters in the relaxed modulus-based synchronous multisplitting multi-parameter AOR methods (RMSMMAOR) and the modulus-based synchronous multisplitting multi-parameter methods (MSMAOR). In addition, we take $\Omega = \frac{1}{2\alpha}D$ in [11] for RMSMMAOR and MSMAOR methods. Let *m* be a prescribed positive integer and $n = m^2$. Consider the LCP(*q*, *A*), in

which $A \in \mathbb{R}^{n \times n}$ is given by $A = \hat{A} + \mu I$ and $q \in \mathbb{R}^n$ is given by $q = -Mz^*$ where

$$\hat{A} = \operatorname{tridiag}(-rI, S, -tI) = \begin{pmatrix} S & -tI & 0 & \cdots & 0 & 0 \\ -rI & S & -tI & \cdots & 0 & 0 \\ 0 & -rI & S & \cdots & 0 & 0 \\ \vdots & \vdots & \ddots & \vdots & \vdots \\ 0 & 0 & \cdots & \cdots & S & -tI \\ 0 & 0 & \cdots & \cdots & -rI & S \end{pmatrix} \in R^{n \times n}$$
(5.1)

is a block-tridiagonal matrix,

$$S = \operatorname{tridiag}(-1, 4, -1) = \begin{pmatrix} 4 & -1 & 0 & \cdots & 0 & 0 \\ -1 & 4 & -1 & \cdots & 0 & 0 \\ 0 & -1 & 4 & \cdots & 0 & 0 \\ \vdots & \vdots & \ddots & \vdots & \vdots \\ 0 & 0 & \cdots & \cdots & 4 & -1 \\ 0 & 0 & \cdots & \cdots & -1 & 4 \end{pmatrix} \in R^{n \times n}$$
(5.2)

is a tridianonal matrix, and

$$z^* = (1, 2, 1, 2, \cdots, 1, 2, \cdots)^T \in \mathbb{R}^n$$

is the unique solution of the LCP(q, A), one can see [11] for more details.

For symmetric case, we take r = t = 1, which is considered in [11]. In this case, the system matrix $A \in \mathbb{R}^{n \times n}$ is symmetric positive definite for $\mu \ge 0$. Obviously, the LCP(q, A) has a unique solution.

In Table 2, the iteration steps, the CPU times, and the residual norms of RMSMMAOR and MSMAOR methods for the symmetric case are listed for different parameters and different problem sizes of m. When both RMSMMAOR and MSMAOR methods are applied to solve the LCP(q, A), the iteration parameters α, β about MSMAOR method satisfy Theorem 4.1 in [33] and Theorem 4.2 in this paper, but the iteration parameters α, β about RMSMMAOR method only satisfy Theorem 4.2 in this paper and don't satisfy Theorem 4.1 in [33].

lase.							
		m	20	30	40	50	60
$\mu = 0.5$	RMSMMAOR	IT	22	22	22	22	22
	$\alpha = 1$	CPU	0.1560	0.7800	2.4336	5.9280	12.4957
	$\beta = 1.2$	Error	7.2225×10^{-6}	7.2598×10^{-6}	7.2970×10^{-6}	7.3390×10^{-6}	7.3707×10^{-6}
$\mu = 0.5$	MSMAOR	IT	30	30	30	31	31
	$\alpha = 1$	CPU	0.2184	1.0764	3.2916	8.3773	17.6905
	$\beta = 0.7$	Error	9.7188×10^{-6}	9.8399×10^{-6}	9.9531×10^{-6}	7.3792×10^{-6}	7.4496×10^{-6}
$\mu = 1.5$	RMSMMAOR	IT	19	19	19	19	19
	$\alpha = 1$	CPU	0.1716	0.6552	2.0748	5.0856	10.7797
	$\beta = 1.2$	Error	6.6884×10^{-6}	6.8943×10^{-6}	7.0943×10^{-6}	7.2888×10^{-6}	7.4782×10^{-6}
$\mu = 1.5$	MSMAOR	IT	23	24	24	24	24
	$\alpha = 1$	CPU	0.1716	0.8424	2.6520	6.4584	13.6657
	$\beta = 0.7$	Error	9.5945×10^{-6}	6.6969×10^{-6}	7.0677×10^{-6}	7.4200×10^{-6}	7.7563×10^{-6}
$\mu = 2.5$	RMSMMAOR	IT	17	17	17	17	17
	$\alpha = 1$	CPU	0.1404	0.6084	1.8720	4.5552	9.6565
	$\beta = 1.2$	Error	7.6793×10^{-6}	8.2513×10^{-6}	8.7861×10^{-6}	9.2902×10^{-6}	9.7683×10^{-6}
$\mu = 2.5$	MSMAOR	\mathbf{IT}	20	20	20	21	21
	$\alpha = 1$	CPU	0.1404	0.7020	2.2932	5.6472	11.9341
	$\beta = 0.7$	Error	8.3861×10^{-6}	9.4078×10^{-6}	6.1592×10^{-6}	6.6458×10^{-6}	7.0992×10^{-6}

Table 2. IT, CPU and Error for RMSMMAOR and MSMAOR with different parameters in symmetric

From Table 2, for RMSMMAOR and MSMAOR methods with $\alpha = 1, \beta = 1.2$ and $\alpha = 1, \beta = 0.7$, fixing the value of μ , it is easy to see that the iteration steps unchange as the increasing of the problem size m, however, CPU times increase as the increasing of the problem size m. Moreover, for RMSMMAOR and MSMAOR methods, fixing the value of m, it is also easy to see that the iteration steps and CPU times decrease as the increasing of the problem size μ . In our numerical experiments, we find that the iteration steps and CPU times of RMSMMAOR are less than that of MSMAOR under certain conditions.

6. Conclusions

In this paper, we establish relaxed modulus-based synchronous multisplitting multiparameters TOR methods and analyze its convergence properties in detail when the system matrix is either a positive-definite matrix or an H+-matrix. Numerical experiments show that the RMSMMTOR methods are feasible under certain conditions. In future work, we can consider extending RMSMMTOR method for nonlinear complementarity problems and horizontal linear complementarity problems.

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