Secant acceleration of sequential residual methods for solving large-scale nonlinear systems of equations∗

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Abstract
Sequential Residual Methods try to solve nonlinear systems of equations $F(x) = 0$ by iteratively updating the current approximate solution along a residual-related direction. Therefore, memory requirements are minimal and, consequently, these methods are attractive for solving large-scale nonlinear systems. However, the convergence of these algorithms may be slow in critical cases; therefore, acceleration procedures are welcome. Anderson-like accelerations are widely used in electronic structure calculations to improve a fixed point algorithm for finding Self Consistent Field (SCF) solutions of Hartree-Fock models. In this paper, it is showed how to apply this type of acceleration to Sequential Residual Methods. The performance of the resulting algorithm is illustrated by applying it to the solution of very large problems coming from the discretization of partial differential equations.

Key words: Nonlinear systems of equations, Sequential Residual Methods, acceleration, large-scale problems.

AMS subject classifications: 65H10, 65K05, 90C53.

1 Introduction
In the process of solving many real-life problems, it is necessary to handle large-scale nonlinear systems of equations. The most obvious choice for solving these systems is Newton’s method, which requires to solve a possibly large and sparse linear system of equations at each iteration. Although being very effective in many cases, Newton’s method cannot be employed for solving very large problems when the Jacobian is unavailable or when it has an unfriendly structure that makes its factorization unaffordable. On the other hand, Inexact Newton methods, that solve the Newtonian linear system approximately at each iteration, are usually effective [14, 17, 18]. Inexact-Newton methods based on linear iterative solvers as GMRES may need many matrix-vector products per iteration. Usually, matrix-vector products of the form $J(x)v$ are replaced with incremental quotients $(F(x + hv) - F(x))/h$, a procedure that does not deteriorate the overall performance of GMRES [10, 41]. However, when GMRES requires many matrix-vector products for providing a suitable approximate solution to the Newtonian linear system, the number of residual evaluations per inexact-Newton iteration may be big. Additional residual evaluations may also be necessary to decide acceptance of trial points at every iteration.

This state of facts led to the introduction of algorithms in which the number of residual evaluations used to compute trial points at each iteration is minimal, as well as the memory used to store directions and the computer effort of linear algebra calculations. DF-SANE [30] was introduced for solving large problems and was used for solving equilibrium models for the determination of industrial prices [37], multifractal analysis of spot prices [45], elastoplastic contact problems [20, 21], and PDE equations in reservoir simulations [38], among

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others. Improved versions of DF-SANE were given in [29, 36]. However, the pure form of DF-SANE may be ineffective for some large problems in which it is necessary to perform many backtracks per iteration in order to obtain sufficient descent. Therefore, on the one hand, it is necessary to investigate alternative choices of trial steps and, on the other hand, acceleration procedures are welcome.

Acceleration devices for iterative algorithms are frequent in the Numerical Analysis literature [1, 7, 8, 26, 47]. They incorporate useful information from previous iterations instead of expending evaluations at the current one. In particular, Anderson’s acceleration introduced in [1] is known to produce very good results when associated with fixed-point iterations [14, 12, 19, 26, 47], specifically those originated in Self-Consistent Field (SCF) approaches for electronic structure calculations [32, 42]. Anderson’s acceleration is closely related when associated with fixed-point iterations [4, 12, 19, 26, 47], specifically those originated in Self-Consistent Field (SCF) approaches for electronic structure calculations [32, 42]. Anderson’s acceleration is closely related when associated with fixed-point iterations [4, 12, 19, 26, 47], specifically those originated in Self-Consistent Field (SCF) approaches for electronic structure calculations [32, 42].

This work introduces a generalized and accelerated version of DF-SANE. The generalization consists of allowing non-residual (although residual-related) directions. The acceleration is based on the multipoint secant idea and Anderson’s acceleration, taking advantage of the residual-like direction steps. Global convergence results that extend the theory of DF-SANE are given.

The paper is organized as follows. Section 2 introduces the accelerated sequential residual methods. Global convergence is established in Section 3. Section 4 describes the acceleration process in detail. Implementation results that extend the theory of DF-SANE are given.

Implementation features and numerical experiments are given in Sections 5 and 6 respectively. The last section presents the conclusions.

**Notation.** The symbol $\| \cdot \|$ denotes the Euclidean norm. $\mathbb{N} = \{0, 1, 2, \ldots \}$ denotes the set of natural numbers. \( J(x) \) denotes the Jacobian matrix of \( F : \mathbb{R}^n \rightarrow \mathbb{R}^n \) computed at \( x \). For all \( x \in \mathbb{R}^n \), we denote \( g(x) = J(x)^T F(x) = \nabla_2 \|F(x)\|^2 \). If \( \{z_k\}_{k \in \mathbb{N}} \) is a sequence and \( K = \{k_1, k_2, k_3, \ldots \} \) is an infinite sequence of natural numbers such that \( k_i < k_j \) if \( i < j \), we denote

\[
\lim_{k \in K} z_k = \lim_{j \to \infty} z_{k_j}.
\]

## 2 Accelerated sequential residual methods

Given \( F : \mathbb{R}^n \rightarrow \mathbb{R}^n \), consider the problem of finding \( x \in \mathbb{R}^n \) such that

\[
F(x) = 0.
\]

A radical iterative approach for solving (1) is to employ only residuals as search directions. Given \( \sigma > 0 \), problem (1) is clearly equivalent to \( x = x - \sigma F(x) \). This trivial observation motivates the introduction of a fixed-point method given by \( x^{k+1} = x^k - \sigma_k F(x^k) \), where \( \sigma_k \) is defined at every iteration. Methods based on this approach will be called Sequential Residual Methods (SRM) in the present paper.

Popular SRM were inspired by the Barzilai-Borwein or spectral choice for the minimization of functions [3, 43, 44]. Defining

\[
s^k = x^{k+1} - x^k \quad \text{and} \quad y^k = F(x^{k+1}) - F(x^k),
\]

algorithms SANE [31] and DF-SANE [30] compute

\[
\sigma_{k+1} = \|s^k\|^2/(y^k)^T s^k,
\]

safeguarded in such a way that \( |\sigma_{k+1}| \) is bounded and bounded away from zero. This formula had been used in the context of self-scaling variable metric methods for minimization [39] as it provides a scale invariant diagonal first approximation of the Hessian. The choice of \( \sigma_{k+1} \) may be justified with the same arguments that Raydan [43] employed for the choice of the Barzilai-Borwein or spectral step in minimization problems. After the computation of \( x^{k+1} \), we consider the (generally unsolvable) problem of satisfying the secant equation [10]

\[
B_{k+1} s^k = y^k \quad \text{subject to} \quad B_{k+1} = cI. \]

This leads to the minimization of \( \|cIs^k - y^k\|^2 \), whose solution, if \( s^k \neq 0 \), is \( c = (y^k)^T s^k/\|s^k\|^2 \). Therefore, a “natural” residual-based iteration for solving problem (1) could be given by

\[
x^{k+1} = x^k - \sigma_k F(x^k),
\]

with \( \sigma_0 \) arbitrary and \( \sigma_{k+1} \) defined by a safeguarded version of (3) for all \( k \geq 0 \).
However, unlike the case of unconstrained minimization, in which \( F(x^k) \) is a gradient, the direction \( d^k = -\sigma_k F(x^k) \) may not be a descent direction for the natural merit function \( f(x) \) defined by

\[
f(x) = \frac{1}{2} \| F(x) \|^2 \text{ for all } x \in \mathbb{R}^n.
\]

In SANE [31], a test is performed in order to verify whether \( F(x^k) \) is a descent direction. If this is the case, since \( \nabla f(x) = J(x)^T F(x) \), we should have

\[
F(x^k)^T J(x^k) F(x^k) < 0.
\]

In order to avoid the employment of derivatives, SANE employs the approximation

\[
J(x^k) F(x^k) \approx \frac{F(x^k + h F(x^k)) - F(x^k)}{h},
\]

for a small \( h > 0 \). In this way, the descent test is equivalent to

\[
F(x^k)^T F(x^k + h F(x^k)) < \| F(x^k) \|^2,
\]

which requires and auxiliary functional evaluation per iteration. The necessity of an auxiliary residual evaluation per iteration in SANE motivated the introduction of DF-SANE [30]. Roughly speaking, in DF-SANE, one gets descent by starting with the trial point \( x^k - \sigma_k F(x^k) \) and proceeding to a double backtracking scheme along positive and negative directions, aiming that \( \| F(x^{k+1}) \| \) be sufficiently smaller than the maximum value of the residual norm in \( M \) consecutive past iterations, where \( M \) is given.

The description of the SRM algorithm provided in this section aims to emphasize the aspects that influence theoretical convergence properties. For this reason, acceleration steps appear only as a small detail in the description of the algorithm, although, in practice, they are essential for the algorithm robustness and efficiency. The description of the algorithm follows.

**Algorithm 2.1.** Let \( \gamma \in (0, 1) \), \( 0 < \sigma_{\min} \leq \sigma_{\max} < \infty \), \( 0 < \tau_{\min} < \tau_{\max} < 1 \), a positive integer \( M \), a sequence \( \{\eta_k\} \) such that \( \eta_k > 0 \) for all \( k \in \mathbb{N} \) and

\[
\lim_{k \to \infty} \eta_k = 0, \tag{4}
\]

and \( x_0 \in \mathbb{R}^n \) be given. Set \( k \leftarrow 0 \).

**Step 1.** If \( F(x^k) = 0 \), then terminate the execution of the algorithm.

**Step 2.** Choose \( \sigma_k \) such that \( |\sigma_k| \in [\sigma_{\min}, \sigma_{\max}] \) and \( v_k \in \mathbb{R}^n \) such that \( \|v_k\| = \|F(x^k)\| \). Compute

\[
\bar{f}_k = \max\{f(x^k), \ldots, f(x^{\max(0,k-M+1)})\}. \tag{5}
\]

**Step 2.1.** Set \( \alpha_+ \leftarrow 1 \) and \( \alpha_- \leftarrow 1 \).

**Step 2.2.** Set \( d \leftarrow -\sigma_k v_k \) and \( \alpha \leftarrow \alpha_+ \). Consider

\[
f(x^k + \alpha d) \leq \bar{f}_k + \eta_k - \gamma \alpha^2 f(x^k). \tag{6}
\]

If \([6] \) holds, then define \( d^k = d \) and \( \alpha_k = \alpha \) and go to Step 3.

**Step 2.3.** Set \( d \leftarrow \sigma_k v_k \) and \( \alpha \leftarrow \alpha_- \). If \([6] \) holds, then define \( d^k = d \) and \( \alpha_k = \alpha \) and go to Step 3.

**Step 2.4.** Choose \( \alpha^\text{new}_+ \in [\tau_{\min} \alpha_+, \tau_{\max} \alpha_+] \) and \( \alpha^\text{new}_- \in [\tau_{\min} \alpha_-, \tau_{\max} \alpha_-] \), set \( \alpha_+ \leftarrow \alpha^\text{new}_+, \alpha_- \leftarrow \alpha^\text{new}_- \), and go to Step 2.2.

**Step 3.** Compute \( x^{k+1} \) such that \( f(x^{k+1}) \leq f(x^k + \alpha_k d^k) \), set \( k \leftarrow k + 1 \), and go to Step 1.

As in DF-SANE, the sufficient decrease in \([6] \) corresponds to a nonmonotone strategy that combines the ones introduced in [24] and [33]. The main differences of Algorithm 2.1 with respect to DF-SANE are the presence of accelerations at Step 3 and the choice of the non-accelerated step in a residual-related way, but not necessarily in the residual direction. The DF-SANE method presented in [30] is the particular case of Algorithm 2.1 in which \( v^k = F(x^k) \) and \( x^{k+1} = x^k + \alpha_k d^k \).
3 Global convergence

In this section we prove global convergence properties of Algorithm 2.1. Our main purpose is to find solutions of \( F(x) = 0 \) or, at least, points at which the residual norm \( \| F(x) \| \) is as small as desired, given an arbitrary tolerance. However, this purpose could be excessively ambitious because, in the worst case, solutions of the system, or even approximate solutions, may not exist. For this reason we analyze the situations in which convergence to a (stationary) point, at which the gradient of the sum of squares vanishes, occur. The option of taking directions that are not residuals but are residual-related in the sense that their norms coincide with those of the residuals is crucial for this purpose. Roughly speaking, we will prove that, taking random residual-related directions an infinite number of times (but not at all iterations) convergence to stationary points necessarily takes place.

In Lemma 3.1 we prove that, given an arbitrary tolerance \( \varepsilon > 0 \), either Algorithm 2.1 finds an approximate solution such that \( \| F(x^k) \| \leq \varepsilon \) in a finite number of iterations or produces an infinite sequence of steps \( \{ \alpha_k \} \) that tends to zero. For this purpose we will only use continuity of \( F \). The lemma begins with a simple proof that the iteration is well defined.

**Lemma 3.1** Assume that \( F \) is continuous, \( x^k \in \mathbb{R}^n \) is an arbitrary iterate of Algorithm 2.1, and \( \| F(x^k) \| \neq 0 \). Then, the iterate \( x^{k+1} \) satisfying (6) is well defined. Moreover, if the algorithm generates an infinite sequence \( \{ x^k \} \), then there exists an infinite subset of indices \( K_1 \subset \mathbb{N} \) such that

\[
\lim_{k \in K_1} \| F(x^k) \| = 0 \tag{7}
\]

or

\[
\lim_{k \to \infty} \alpha_k = 0. \tag{8}
\]

**Proof:** The fact that \( x^{k+1} \) is always well defined follows from the continuity of \( F \) using that \( \eta_k > 0 \) and that \( \alpha_k \) can be as small as needed. Assume that the algorithm generates an infinite sequence \( \{ x^k \} \). Suppose that (7) does not hold. Then, there exists \( c_1 > 0 \) such that

\[
\| F(x^k) \| > c_1 \text{ for all } k \in \mathbb{N}. \tag{9}
\]

If (8) does not hold either, then there exists a subsequence \( K_2 \subset \mathbb{N} \) and \( c_2 > 0 \) such that

\[
\alpha_k > c_2 \text{ for all } k \in K_2.
\]

Then, by (9),

\[
\alpha_k^2 f(x^k) > (c_1 c_2)^2 \text{ for all } k \in K_2. \tag{10}
\]

Then, since \( x^{k+1} \) is well defined, for all \( k \in K_2 \),

\[
f(x^{k+1}) \leq \bar{f}_k + \eta_k - \gamma \alpha_k^2 f(x^k) \leq \bar{f}_k + \eta_k - \gamma (c_1 c_2)^2,
\]

where

\[
\bar{f}_k = \max \{ f(x^k), \ldots, f(x^{\max\{0,k-M+1\}}) \}. \tag{12}
\]

Since \( \eta_k \to 0 \), then there exists \( k_1 \in \mathbb{N} \) such that for all \( k \in K_2 \) and \( k \geq k_1 \),

\[
f(x^{k+1}) \leq \bar{f}_k - \gamma (c_1 c_2)^2 / 2. \tag{13}
\]

By (12) and (13), there exists a subsequence for which \( f(x^k) \to -\infty \), which contradicts the fact that \( f(x) \geq 0 \) for all \( x \in \mathbb{R}^n \). \( \square \)

In Lemma 3.2 we prove that we have two possibilities: Either, given an arbitrary \( \varepsilon > 0 \), we find an iterate such that \( \| F(x^k) \| \leq \varepsilon \) or, for every limit point of the sequence \( \{ x^k \} \), the gradient of \( f \) is orthogonal to every possible limit of search directions. Of course, the second possibility necessarily occurs when the system has no solutions.
Lemma 3.2 Assume that $F(x)$ admits continuous derivatives for all $x \in \mathbb{R}^n$ and $\{x^k\}$ is generated by Algorithm 2.1. Assume that $\{\|F(x^k)\|\}$ is bounded away from zero and $x_\ast \in \mathbb{R}^n$ is a limit point of $\{x^k\}$ such that $\lim_{k \in K_1} x^k = x_\ast$. Then, the set of limit points of $\{v^k\}_{k \in K_1}$ is nonempty. Moreover, for every limit point of $\{v^k\}_{k \in K_1}$, we have that

$$\langle J(x_\ast)v, F(x_\ast) \rangle = \langle v, J(x_\ast)^T F(x_\ast) \rangle = 0. \quad (14)$$

Proof: By Lemma 3.1

$$\lim_{k \to \infty} \alpha_k = 0. \quad (15)$$

By continuity, since $\lim_{k \in K_1} x^k = x_\ast$ we have that $\lim_{k \in K_1} F(x^k) = F(x_\ast)$. Therefore, the sequence $\{F(x^k)\}_{k \in K_1}$ is bounded. Since $\|v^k\| = \|F(x^k)\|$ for all $k$, we have that the sequence $\{v^k\}_{k \in K_1}$ is bounded too; therefore, it admits a limit point $v \in \mathbb{R}^n$, as we wanted to prove.

Let $K_2$ be an infinite subset of $K_1$ and $v \in \mathbb{R}^n$ be such that $\lim_{k \in K_2} v^k = v$. Since the first trial value for $\alpha_k$ at each iteration is 1, (15) implies that there exists $K_3$, an infinite subset of $K_2$, $\alpha_{k,+} > 0$, $\alpha_{k,-} > 0$, $\sigma_k \in [\sigma_{\min}, \sigma_{\max}]$, and $d^k = -\sigma_k v^k$ such that

$$\lim_{k \in K_3} \sigma_k = \sigma \in [\sigma_{\min}, \sigma_{\max}],$$

$$\lim_{k \in K_3} \alpha_{k,+} = \lim_{k \in K_2} \alpha_{k,-} = 0,$$

and, for all $k \in K_3$,

$$f(x^k + \alpha_{k,+}d^k) > f_k + \eta_k - \gamma_\alpha_{k,+}^2 f(x^k),$$

and

$$f(x^k - \alpha_{k,-}d^k) > f_k + \eta_k - \gamma_\alpha_{k,-}^2 f(x^k).$$

Therefore, by the definition (5) of $f_k$, for all $k \in K_3$,

$$f(x^k + \alpha_{k,+}d^k) > f(x^k) + \eta_k - \gamma_\alpha_{k,+}^2 f(x^k)$$

and

$$f(x^k - \alpha_{k,-}d^k) > f(x^k) + \eta_k - \gamma_\alpha_{k,-}^2 f(x^k).$$

So, since $\eta_k > 0$,

$$\frac{f(x^k + \alpha_{k,+}d^k) - f(x^k)}{\alpha_{k,+}} > -\gamma_\alpha_{k,+} f(x^k)$$

and

$$\frac{f(x^k - \alpha_{k,-}d^k) - f(x^k)}{\alpha_{k,-}} > -\gamma_\alpha_{k,-} f(x^k)$$

for all $k \in K_3$. Thus, by the Mean Value Theorem, there exist $\xi_{k,+} \in [0, \alpha_{k,+}]$ and $\xi_{k,-} \in [0, \alpha_{k,-}]$ such that

$$\langle \nabla f(x^k + \xi_{k,+}d^k), d^k \rangle > -\gamma_\alpha_{k,+} f(x^k) \quad (16)$$

and

$$\langle \nabla f(x^k - \xi_{k,-}d^k), d^k \rangle > -\gamma_\alpha_{k,-} f(x^k) \quad (17)$$

for all $k \in K_3$. Taking limits for $k \in K_3$ in both sides of (16) and (17) we get

$$\langle \nabla f(x_\ast), \sigma v \rangle = 0. \quad (18)$$

Therefore, (14) is proved. \qed

Theorem 3.1 states a sufficient condition for the annihilation of the gradient of $f(x)$ at a limit point of the sequence. For that purpose, we will assume that the absolute value of the cosine of the angle determined by $v^k$ and $J(x^k)^T v^k$ is bigger than a tolerance $\omega > 0$ infinitely many times. Note that we do not require this condition to hold for every $k \in \mathbb{N}$. The distinction between these two alternatives is not negligible. For example, if $v^k$ is chosen infinitely many times as a random vector the required angle condition holds with probability 1. Conversely, if we require the fulfillment of the angle condition for all $k \in \mathbb{N}$ and we choose random directions, the probability of fulfillment would be zero.
**Theorem 3.1** Assume that $F(x)$ admits continuous derivatives for all $x \in \mathbb{R}^n$ and $\{x^k\}$ is generated by Algorithm $2.1$. Suppose that $\sum_{k=0}^{\infty} \eta_k = \eta < \infty$, the level set $\{x \in \mathbb{R}^n \mid f(x) \leq f(x^0) + \eta\}$ is bounded, and there exists $\omega > 0$ such that

$$\langle v^k, J(x^k)^TF(x^k) \rangle \geq \omega \|v^k\| \|J(x^k)^TF(x^k)\|$$

(19)

for infinitely many indices $k \in K \subset \mathbb{N}$. Then, for any given $\varepsilon > 0$, there exists $k \in \mathbb{N}$ such that $\|J(x^k)^TF(x^k)\| \leq \varepsilon$. Moreover, there exists a limit point $x_*$ of the sequence generated by the algorithm such that $\|J(x_*)^TF(x_*)\| = 0$.

**Proof:** If the sequence generated by the algorithm stops at some $k$ such that $\|F(x^k)\| = 0$ the thesis obviously holds. By the boundedness of the level set defined by $f(x^0) + \eta$, the sequence $\{x^k\}$ is bounded. By the continuity of $F$, the sequence $\{F(x^k)\}$ is bounded as well. By the definition of $v^k$, the sequence $\{v^k\}$ is also bounded.

Therefore, there exist an infinite set $K_1 \subset K$, $x_* \in \mathbb{R}^n$, and $v \in \mathbb{R}^n$ such that $\lim_{k \in K_1} x^k = x_*$ and $\lim_{k \in K_1} v^k = v$. Therefore, by Lemma 3.2

$$\langle v, J(x_*)^TF(x_*) \rangle = 0.$$

So, by the convergence of $\{v^k\}$ and $\{x^k\}$ for $k \in K_1$ and the continuity of $F$ and $J$,

$$\lim_{k \in K_1} \langle v^k, J(x^k)^TF(x^k) \rangle = 0.$$

Therefore, by (19), since $K_1 \subset K$,

$$\lim_{k \in K_1} \|v^k\| \|J(x^k)^TF(x^k)\| = 0.$$  

(20)

If $\|F(x^k)\|$ tends to zero for some subsequence of $K_1$ the thesis obviously holds. Otherwise, there exists $c > 0$ such that $\|F(x^k)\| > c$ for all $k \in K_1$. Then, $\|v^k\| \geq c$ for all $k \in K_1$. So, by (20),

$$\lim_{k \in K_1} \|J(x^k)^TF(x^k)\| = 0.$$  

(21)

Therefore, the thesis is proved.

**Corollary 3.1** Suppose that the assumptions of Theorem 3.1 hold. Assume, moreover, that there exists $\gamma > 0$ such that for all $k \in \mathbb{N}$,

$$\|J(x^k)^TF(x^k)\| \geq \gamma \|F(x^k)\|.$$  

(22)

Then, given $\varepsilon > 0$, there exists an iterate $k$ such that $\|F(x^k)\| \leq \varepsilon$. Moreover, there exists a limit point $x_*$ of the sequence generated by the algorithm such that $\|F(x_*)\| = 0$.

**Proof:** The thesis of this corollary follows directly from (22) and Theorem 3.1.

**Corollary 3.2** Suppose that the assumptions of Theorem 3.1 hold and $J(x)$ is nonsingular with $\|J(x)^{-1}\|$ uniformly bounded for all $x \in \mathbb{R}^n$. Then, for all $\ell = 1, 2, \ldots$, there exists an iterate $k(\ell)$ such that $\|F(x^{k(\ell)})\| \leq 1/\ell$. Moreover, at every limit point $x_*$ of the sequence $\{x^{k(\ell)}\}$ we have that $\|F(x_*)\| = 0$.

**Proof:** Since $\|F(x^k)\| = \|J(x^k)^{-T}J(x^k)^TF(x^k)\| \leq \|J(x^k)^{-1}\| \|J(x^k)^TF(x^k)\|$, a sufficient condition for the fulfillment of (22) is the existence and uniform boundedness of $\|J(x^k)^{-1}\|$.

The lemma below shows that, if $\|F(x^k)\| \to 0$ for a subsequence, then it goes to zero for the whole sequence. We will need the following Assumptions A and B. Assumption A guarantees that the difference between consecutive iterates is not greater than a multiple of the residual norm. Note that such assumption holds trivially when the iteration is not accelerated and, so, $x^{k+1} = x^k + \alpha_k d^k$. Therefore, Assumption A needs to be stated only with respect to accelerated steps. Assumption B merely states that $F$ is uniformly continuous at least restricted to the set of iterates.

**Assumption A** There exists $c > 0$ such that, for all $k \in \mathbb{N}$, whenever $x^{k+1} \neq x^k + \alpha_k d^k$, we have that

$$\|x^{k+1} - x^k\| \leq c\|F(x^k)\|.$$
Assumption B  For all $\varepsilon > 0$, there exists $\delta$ such that, whenever $\|x^{k+1} - x^k\| \leq \delta$ one has that $\|F(x^{k+1}) - F(x^k)\| \leq \varepsilon$.

Lemma 3.3 below is stronger than Theorem 2 of [30] because here we do not assume the existence of a limit point.

Lemma 3.3 Suppose that $F$ is continuous, $\sum_{k=0}^{\infty} \eta_k < \infty$, Algorithm 2.1 generates the infinite sequence $\{x^k\}$, Assumptions A and B hold, and there exists an infinite subsequence defined by the indices in $K_1$ such that

$$\lim_{k \in K_1} \|F(x^k)\| = 0.$$  

Then,

$$\lim_{k \to \infty} \|F(x^k)\| = 0$$  

and

$$\lim_{k \to \infty} \|x^{k+1} - x^k\| = 0.$$  

Proof: Assume that (23) is not true. Then, there exists an infinite subsequence $\{x^k\}_{k \in K_2}$ such that

$$f(x^k) > c > 0$$  

for all $k \in K_2$.

Let us prove first that

$$\lim_{k \in K_1} \|F(x^{k+1})\| = 0.$$  

(26)

For this purpose suppose that $\varepsilon > 0$. By the hypothesis, there exists $k_1 \in K_1$ such that for all $k \in K_1, k \geq k_1$, we have that

$$\|F(x^k)\| \leq \varepsilon/2.$$  

(27)

By Assumption B there exists $\delta > 0$ such that, whenever $\|x^{k+1} - x^k\| \leq \delta$, one has that

$$\|F(x^{k+1}) - F(x^k)\| \leq \varepsilon/2.$$  

(28)

By Assumption A there exists $k_2 \geq k_1$ such that, whenever $k \geq k_2$, $k \in K_1$, we have that

$$\|x^{k+1} - x^k\| \leq \delta.$$  

(29)

Then, by (27), (28), and (29), we have that, whenever $k \geq k_2$ and $k \in K_1$,

$$\|F(x^{k+1})\| \leq \varepsilon.$$  

This proves (26).

By induction, we deduce that for all $j = 1, \ldots, M - 1$,

$$\lim_{k \in K_1} \|F(x^{k+j})\| = 0.$$  

Therefore,

$$\lim_{k \in K_1} \max \{f(x^k), \ldots, f(x^{k+M-1})\} = 0.$$  

(30)

But

$$f(x^{k+M}) \leq \max \{f(x^k), \ldots, f(x^{k+M-1})\} + \eta_{k+M-1} - \alpha_{k+M-1} f(x^{k+M-1})$$

$$\leq \max \{f(x^k), \ldots, f(x^{k+M-1})\} + \eta_{k+M-1}.$$  

Analogously,

$$f(x^{k+M+1}) \leq \max \{f(x^{k+1}), \ldots, f(x^{k+M+1})\} + \eta_{k+M} - \alpha_{k+M} f(x^{k+M})$$

$$\leq \max \{f(x^{k+1}), \ldots, f(x^{k+M})\} + \eta_{k+M}$$

$$\leq \max \{f(x^k), \ldots, f(x^{k+M-1})\} + \eta_{k+M-1} + \eta_{k+M}.$$  

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and, inductively,

\[ f(x^{k+Mj}) \leq \max\{f(x^k), \ldots, f(x^{k+M-1})\} + \eta_k + \eta_{k+M} + \cdots + \eta_{k+M+j-1} \]

for all \( j = 0, 1, \ldots, M - 1 \). Therefore,

\[ \max\{f(x^{k+M}), \ldots, f(x^{k+2M-1})\} \leq \max\{f(x^k), \ldots, f(x^{k+M-1})\} + \sum_{j=k+M-1}^{k+2M-2} \eta_j. \tag{31} \]

By induction on \( \ell = 1, 2, \ldots \), we obtain that

\[ \max\{f(x^{k+\ell M}), \ldots, f(x^{k+(\ell+1)M-1})\} \leq \max\{f(x^k), \ldots, f(x^{k+M-1})\} + \sum_{j=k+M-1}^{k+\ell M-2} \eta_j. \tag{32} \]

Therefore, for all \( \ell = 1, 2, \ldots \) we have:

\[ \max\{f(x^{k+\ell M}), \ldots, f(x^{k+(\ell+1)M-1})\} \leq \max\{f(x^k), \ldots, f(x^{k+M-1})\} + \sum_{j=k+M-1}^{\infty} \eta_j. \tag{33} \]

By \([30]\), the summability of \( \eta_j \) and \([33]\), there exists \( k_1 \in K_1 \) such that for all \( k \in K_1 \) such that \( k \geq k_1 \),

\[ \max\{f(x^k), \ldots, f(x^{k+M-1})\} + \sum_{j=k+M-1}^{\infty} \eta_j \leq c/2. \]

Therefore, by \([33]\), for all \( k \in K_1 \) such that \( k \geq k_1 \) and all \( \ell = 1, 2, \ldots \),

\[ \max\{f(x^{k+\ell M}), \ldots, f(x^{k+(\ell+1)M-1})\} \leq c/2. \tag{34} \]

Clearly, this is incompatible with the existence of a subsequence such that \([25]\) holds. Therefore, \([23]\) is proved. Then, by Assumption \([1]\) and \([23]\), \([24]\) also holds.

**Theorem 3.2** Suppose that the assumptions of Theorem 3.1 hold, \( J(x) \) is nonsingular and \( \|J(x)^{-1}\| \) is uniformly bounded for all \( x \in \mathbb{R}^n \). Then, there exists \( x_\star \in \mathbb{R}^n \) such that \( F(x_\star) = 0 \) and \( \lim_{k \to \infty} x^k = x_\star \).

**Proof:** By Corollary 3.2 there exists a subsequence \( \{F(x^k)\}_{k \in K_1} \) that converges to a point \( x_\star \) such that \( F(x_\star) = 0 \). Then, by Lemma 3.3

\[ \lim_{k \to \infty} F(x^k) = 0 \tag{35} \]

and

\[ \lim_{k \to \infty} \|x^{k+1} - x^k\| = 0. \]

Since \( J(x_\star) \) is nonsingular, by the Inverse Function Theorem, there exists \( \delta > 0 \) such that \( \|F(x)\| > 0 \) whenever \( 0 < \|x - x_\star\| \leq \delta \). Therefore, given \( \varepsilon \in (0, \delta] \) arbitrary, there exists \( c > 0 \) such that

\[ \|F(x)\| \geq c \text{ whenever } \varepsilon \leq \|x - x_\star\| \leq \delta. \]

By \([35]\), the number of iterates in the set defined by \( \varepsilon \leq \|x - x_\star\| \leq \delta \) is finite. On the other hand, if \( k \) is large enough, one has that \( \|x^{k+1} - x_\star\| \leq \delta - \varepsilon \). Since there exists a subsequence of \( \{x^k\} \) that tends to \( x_\star \), it turns out that \( \|x^k - x_\star\| \leq \varepsilon \) if \( k \) is large enough. Since \( \varepsilon \) was arbitrary this implies that \( \lim_{k \to \infty} x^k = x_\star \) as we wanted to prove. \( \square \)

We finish this section with a theorem that states that, under some assumptions, if in Algorithm 2.1 we have \( x^{k+1} = x^k + \alpha_k d^k \) for all \( k \), i.e. without accelerations, then \( \{x^k\} \) converges superlinearly to a solution. A similar result was proved in [22 Thm. 3.1] with respect to a spectral gradient methods with retards for quadratic unconstrained minimization. We will need two assumptions. In Assumption \([3]\) the sequence generated by Algorithm 2.1 is assumed to be generated by the spectral residual choice with the first step accepted at each iteration and without acceleration. In Assumption \([4]\) the Jacobian \( J(x) \) is assumed to be Lipschitz-continuous.
Assumption C Assume that, at Step 2 of Algorithm 2.1, we choose
\[ v^k = F(x^k) \]
and that there exists \( k_0 \geq 1 \) such that for all \( k \geq k_0 \),
\[ (s^{k-1})^T y^{k-1} \neq 0 \]
and
\[ \sigma_k = \frac{(s^{k-1})^T y^{k-1}}{(s^{k-1})^T s^{k-1}}. \]
where \( s^k \) and \( y^k \) are defined by (2). Assume, moreover, that, for all \( k \geq k_0 \), (6) holds with \( \alpha = 1 \) and, at Step 3,
\[ x^{k+1} = x^k + \alpha_k d^k. \]

Assumption D There exists \( L > 0 \) such that for all \( x, z \) in an open and convex set that contains the whole sequence \( \{x^k\} \) generated by Algorithm 2.1, the Jacobian is continuous and satisfies
\[ \|J(z) - J(x)\| \leq L\|z - x\|. \]

Theorem 3.3 Assume that the sequence \( \{x^k\} \), generated by Algorithm 2.1, does not terminate at Step 1 and that Assumptions C and D hold. Suppose that \( x^* \in \mathbb{R}^n \) is such that
\[ \lim_{k \to \infty} x^k = x^*, \] (36)
the Jacobian \( J(x^*) \) is nonsingular and \( s \in \mathbb{R}^n \) is such that
\[ \lim_{k \to \infty} s^k / \|s^k\| = s. \] (37)

Then,
\[ \lim_{k \to \infty} \frac{(s^k)^T y^k}{(s^k)^T s^k} = s^T J(x^*) s, \] (38)
\[ F(x^*) = 0, \] (39)
and
\[ x^k \] converges \( Q \)-superlinearly to \( x^* \). (40)

Proof: By Assumption D, there exists an open and convex set that contains the whole sequence \( \{x^k\} \) such that for all \( x, z \) in that set,
\[ F(z) = F(x) + J(x)(z - x) + O(\|z - x\|^2). \] (41)
By Assumption C, the sequence \( \{x^k\} \) is such that for all \( k \geq k_0 \),
\[ x^{k+1} = x^k - \frac{1}{\kappa_k} F(x^k), \] (42)
where
\[ \kappa_{k+1} = \frac{(s^k)^T y^k}{(s^k)^T s^k}. \] (43)
By (41),
\[ y^k = J(x^k)s^k + O(\|s^k\|^2) \] (44)
for all \( k \geq k_0 \). Therefore, by (44),
\[ \kappa_{k+1} = \frac{(s^k)^T [J(x^k)s^k + O(\|s^k\|^2)]}{(s^k)^T s^k}. \] (45)
Thus, by (36), (37), and the continuity of $J(x)$,
\[ \lim_{k \to \infty} \kappa_{k+1} = s^T J(x_*) s. \] (46)
Therefore, (38) is proved.

By (41) we have that
\[ F(x^{k+1}) = F(x^k) + J(x^k) s^k + O(\|s^k\|^2). \]
Then, by (42),
\[ -\kappa_{k+1} s^{k+1} = -\kappa_k s^k + J(x^k) s^k + O(\|s^k\|^2) \]
for all $k \in \mathbb{N}$. Therefore, for all $k \in \mathbb{N}$,
\[ s^{k+1} = \frac{\kappa_k s^k - J(x^k) s^k}{\kappa_{k+1}} - \frac{O(\|s^k\|^2)}{\kappa_{k+1}}. \]
Therefore,
\[ s^{k+1} = -\frac{1}{\kappa_{k+1}} [(J(x^k) - \kappa_k I) s^k + O(\|s^k\|^2)]. \]
Then
\[ \frac{s^{k+1}}{\|s^{k+1}\|} = -\frac{[(J(x^k) - \kappa_k I) s^k + O(\|s^k\|^2)]}{\| - [(J(x^k) - \kappa_k I) s^k + O(\|s^k\|^2)]\|}. \]
So,
\[ \frac{s^{k+1}}{\|s^{k+1}\|} = -\frac{[(J(x^k) - \kappa_k I) s^k]/\|s^k\| + O(\|s^k\|^2)/\|s^k\|]}{\| - [(J(x^k) - \kappa_k I) s^k]/\|s^k\| + O(\|s^k\|^2)/\|s^k\|\|}. \] (47)
Define
\[ \kappa_* = s^T J(x_*) s. \] (48)
Assume that
\[ \| (J(x_*) - \kappa_* I) s \| \neq 0. \] (49)
Then, by (49), (38), and (37), taking limits in both sides of (47),
\[ s = -\frac{[(J(x_*) - \kappa_* I) s] \| - [(J(x_* ) - \kappa_* I) s]\|}{\| - [(J(x_*) - \kappa_* I) s]\|}. \] (50)
Pre-multiplying both sides of (50) by $s^T$, as $s^T s = 1$, we obtain:
\[ 1 = -\frac{[s^T J(x_*) s - \kappa_*] \| - [s^T J(x_* ) - \kappa_* I] s\|}{\| - [s^T J(x_* ) - \kappa_* I] s\|}. \] (51)
Then, by (48), we get a contradiction. This implies that the assumption (49) is false. So,
\[ \|(J(x_*) - \kappa_* I) s\| = 0. \] (52)
Therefore,
\[ \lim_{k \to \infty} \frac{\kappa_k I s^k - J(x^k) s^k}{\|s^k\|} = 0. \] (53)
But (53) is the Dennis-Moré condition related with the iteration $x^{k+1} = x^k - B_k^{-1} F(x^k)$ with $B_k = \kappa_k I$. Then, we have that $F(x_*) = 0$ and the convergence is superlinear.

\[ \square \]


4 Acceleration scheme

In this section, we describe the way in which, at iteration \( k \) of Algorithm 2.1, using an acceleration scheme, we compute \( x^{k+1} \) at Step 3. The algorithmic description follows below.

Algorithm 4.1. Let the integer number \( p \geq 1 \) be given.

\textbf{Step 1.} Define \( x^{k+1}_\text{trial} = x^k + \alpha_k d^k \).

\textbf{Step 2.} Define \( k = \max\{0, k - p + 1\} \),

\[
\begin{align*}
  s^j &= x^{j+1} - x^j \text{ for } j = k, \ldots, k - 1, \\
  y^j &= F(x^{j+1}) - F(x^j) \text{ for } j = k, \ldots, k - 1, \\
  s^k &= x^{k+1}_\text{trial} - x^k, \\
  y^k &= F(x^{k+1}_\text{trial}) - F(x^k), \\
  S_k &= (s^k, \ldots, s^{k-1}, s^k), \\
  Y_k &= (y^k, \ldots, y^{k-1}, y^k),
\end{align*}
\]

and

\[
x^{k+1}_\text{accel} = x^k - S_k Y_k^\dagger F(x^k), \quad (54)
\]

where \( Y_k^\dagger \) is the Moore-Penrose pseudoinverse of \( Y_k \).

\textbf{Step 3.} Choose \( x^{k+1} \in \{x^{k+1}_\text{trial}, x^{k+1}_\text{accel}\} \) such that \( \|F(x^{k+1})\| = \min\{\|F(x^{k+1}_\text{trial})\|, \|F(x^{k+1}_\text{accel})\|\} \).

The theorem below helps to understand the behavior of Algorithm 2.1 with \( x^{k+1} \) given by Algorithm 4.1, called Algorithm 2.1-4.1 from now on. Briefly speaking, we are going to prove that, under some assumptions, the sequence generated by Algorithm 2.1-4.1 converges superlinearly to a solution of \( F(x) = 0 \). We do not mean that these assumptions are “reasonable”, in the sense that they usually, or even frequently, hold. The theorem aims to show the correlation between different properties of the method, which is probably useful to understand the algorithm and, perhaps, to seek modifications and improvements.

**Theorem 4.1** Assume that \( \{x^k\} \) is the sequence generated by Algorithm 2.1-4.1. Suppose, in addition, that

\begin{enumerate}
  \item[H1:] For all \( k \) large enough we have that \( x^{k+1} = x^{k+1}_\text{accel} \).
  \item[H2:] There exists a positive sequence \( \beta_k \to 0 \) such that for all \( k \) large enough,
    \[
    \| (S_{k+1} Y_{k+1}^\dagger - S_k Y_k^\dagger) y^k \| \leq \beta_k \| y^k \|. \quad (55)
    \]
  \item[H3:] There exists \( c > 0 \) such that
    \[
    \| S_k Y_k^\dagger F(x^{k+1}) \| \geq c \| F(x^{k+1}) \| \quad (56)
    \]
    for \( k \) large enough.
\end{enumerate}

Then, \( \|F(x^k)\| \) converges to 0 \( Q \)-superlinearly.

**Proof:** By (54) and H1, we have that, for \( k \) large enough,

\[
x^{k+1} = x^k - S_k Y_k^\dagger F(x^k). \quad (57)
\]

But, by the definition of \( S_k \) and \( Y_k \), for all \( k \in \mathbb{N} \),

\[
S_{k+1} Y_{k+1}^\dagger y^k = s^k.
\]

Therefore, by (55) in H2,

\[
\| S_k Y_k^\dagger y^k - s^k \| \leq \beta_k \| y^k \|. \quad (58)
\]
Then, by (57),
\[ \|S_k y_k^k F(x^{k+1})\| \leq \beta_k \|y_k\|. \]  
(59)
Thus, by (59) in H3,
\[ c\|F(x^{k+1})\| \leq \beta_k \|y_k\|. \]  
(60)
So,
\[ c\|F(x^{k+1})\| \leq \beta_k (\|F(x^{k+1})\| + \|F(x^k)\|). \]
Thus,
\[ (c - \beta_k)\|F(x^{k+1})\| \leq \beta_k \|F(x^k)\|. \]
Therefore,
\[ \|F(x^{k+1})\| \leq \frac{\beta_k}{c - \beta_k} \|F(x^k)\| \]
for \( k \) large enough, which means that \( F(x^k) \) tends to zero \( Q \)-superlinearly.

The rest of this section aims to shed some light on the meaning of the acceleration formula (54). Assume, for a moment, that \( A \) is an approximation of the Jacobian \( J(x^k) \) such that \( L(x) = Ax + b \) satisfies the interpolation conditions \( Ax^{k,j} + b = F(x^{k,j}) \) for \( j = 1, \ldots, p + 1 \). Then, it is natural to seek the point \( \hat{x} \) in the affine subspace generated by \( Ax^{k,1} + b, \ldots, Ax^{k,p+1} + b \) that minimizes \( \|Ax + b\|^2 \). This amounts to solve the optimization problem
\[ \text{Minimize} \quad \|Ax + b\|^2 \text{ subject to } Ax + b = \theta_1 F(x^{k,1}) + \cdots + \theta_{p+1} F(x^{k,p+1}) \quad \text{and} \quad \sum_{j=1}^{p+1} \theta_j = 1. \]  
(61)

Problem (61) is clearly equivalent to
\[ \text{Minimize} \quad \|\theta_1 F(x^{k,1}) + \cdots + \theta_{p+1} F(x^{k,p+1})\|^2 \text{ subject to } \sum_{j=1}^{p+1} \theta_j = 1. \]  
(62)
Then, by the interpolation conditions, \( A\hat{x} + b = \theta_1 F(x^{k,1}) + \cdots + \theta_{p+1} F(x^{k,p+1}) \) implies that
\[ \hat{x} = \theta_1 x^{k,1} + \cdots + \theta_{p+1} x^{k,p+1}. \]  
(63)
Therefore, solutions to (61) are obtained by solving (62) and, then, defining \( \hat{x} \) as in (63). Note that solutions to (61) do not depend on the Jacobian approximation \( A \). The DIIS acceleration method \[42\] is usually presented directly in the form (62, 63) with \( x^{k,j} = x^{k-(p+1)+j} \) for \( j = 1, \ldots, p + 1 \). The accelerated point \( x^{k+1}_{\text{acc}} \) in (54) corresponds to the minimum-norm solution to (61) with \( x^{k,j} = x^{k-p+j} \) for \( j = 1, \ldots, p \) and \( x^{k,p+1} = x^{k+1}_{\text{trial}} \). Including \( x^{k+1}_{\text{trial}} \) is important. The reason is that, if we accelerate without considering the residual \( F(x^{k+1}) \), then the accelerated point lies in the affine subspace generated by the last \( p \) iterates. Therefore, the sequence never leaves an affine subspace with dimension \( p \) which, in general, does not contain the solution. In other words, even when the acceleration provides a big improvement with respect to the basic residual method trials, residuals at these trials are necessary to define the accelerations.

The Sequential Secant Method (SSM), called Secant Method of \( n + 1 \) points in the classical book of Ortega and Rheinboldt \[2, 23, 27, 35, 40, 49\], is a classical procedure for solving nonlinear systems of equations. Given \( n + 1 \) consecutive iterates \( x^k, x^{k-1}, \ldots, x^{k-n} \), the Sequential Secant Method computes an affine function \( L : \mathbb{R}^n \to \mathbb{R}^n \) such that \( L(x^j) = F(x^j) \) for \( j = k, \ldots, k - n \); and defines \( x^{k+1} \) as the unique solution of \( L(x) = 0 \). The inconveniences of this method are obvious. On the one hand, the affine function \( L(x) \) is guaranteed to exist only if the iterates \( x^k, x^{k-1}, \ldots, x^{k-n} \) are affinely independent. On the other hand the linear system of equations \( L(x) = 0 \) may have no solutions or may have infinitely many solutions. When the residuals \( F(x^k), F(x^{k-1}), \ldots, F(x^{k-n}) \) are affinely independent, the iteration of SSM is well defined and
\[ x^{k+1} - x^k = (s^{k-1}, \ldots, s^{k-n})(y^{k-1}, \ldots, y^{k-n})^{-1} F(x^k), \]  
(64)
where \( s^j = x^{j+1} - x^j \) and \( y^j = F(x^{j+1}) - F(x^j) \) for \( j = 1, \ldots, n \). Note that (64) may be well defined even in cases in which the increments \( s^{k-1}, \ldots, s^{k-n} \) are not linearly independent. In fact, lack of linear independence of
\(s^{k-1},\ldots,s^{k-n}\) is not unusual. If some of the components of \(F(x)\) are affine functions, the corresponding linear equations tend to be satisfied after a small number of iterations and, from those iterations on, all the iterates lie in the affine subspace determined by the previous iterates, whose dimension is smaller than \(n\). Lack of linear independence of \(y^{k-1},\ldots,y^{k-n}\) also occurs, for example, when the system is linear and the Jacobian matrix is singular. Some authors developed different variations of SSM in order to overcome these inconveniences. See [35] and references therein.

5 Implementation

In this section, we discuss the implementation of Algorithm 2.1–4.1.

5.1 Stopping criterion

At Step 1 of Algorithm 2.1, given \(\varepsilon > 0\), we replace the stopping criterion \(\|F(x_k)\| = 0\) with

\[\|F(x_k)\|_2 \leq \varepsilon.\]  

(65)

5.2 Choice of the direction and the scaling factor

At Step 2 of Algorithm 2.1, we must choose \(\sigma_k\) and \(v_k\). For this purpose, we considered the residual choice \(v_k = -F(x_k)\) for all \(k\) setting, arbitrarily, \(\sigma_0 = 1\). A natural choice for \(\sigma_k\) \((k \geq 1)\) would be the spectral step given by

\[\sigma_{spg} = \frac{(x_k - x_{k-1})^T (x_k - x_{k-1})}{(x_k - x_{k-1})^T (F(x_k) - F(x_{k-1}))}\]

with safeguards that guarantee that \(|\sigma_k|\) belong to \([\sigma_{min}, \sigma_{max}]\). It turns out that defining \(\sigma_{max} > 1\), preliminary numerical experiments showed that, in the problems considered in the present work, Step 2 of Algorithm 2.1 with this choice of \(\sigma_k\) performs several backtrackings per iteration that result in a total number of functional evaluations one order of magnitude larger than the number of iterations; which is an unusual behavior of nonmonotone methods based on the Barzilai-Borwein spectral choice [6, 31, 30, 43, 44]. On the other hand, it was also observed that, at Step 3 of Algorithm 4.1, the acceleration step was always chosen. This means that the costly obtained \(x_{trial}\) was always beaten by \(x_{accel}\). These two observations suggested that a more conservative, i.e. small, scaling factor \(\sigma_k\) should be considered. A small \(\sigma_k\) could result in a trial point \(x_{trial}^{k+1} = x_k + \alpha_k d_k\) with \(\alpha_k = \pm 1\) that satisfies (6), i.e. no backtracking, and that provides the information required to compute a successful \(x_{accel}^{k+1}\) at Step 2 of Algorithm 4.1.

The conservative choice of \(\sigma_k\) employed in our implementation is as follows. We consider an algorithmic parameter

\[h_{init} > 0\]  

(66)

and we define:

\[\sigma_{min} = \sqrt{\varepsilon}, \ \sigma_{max} = 1,\]  

(67)

\[\sigma_0 = 1,\]  

(68)

\[\bar{\sigma}_k = h_{init} \frac{\|x_k - x_{k-1}\|}{\|F(x_k)\|}\]  

for all \(k \geq 1\),

(69)

and

\[\sigma_k = \bar{\sigma}_k\]  

if \(k \geq 1\) and \(\bar{\sigma}_k \in [\max\{1, \|x_k\|\} \sigma_{min}, \sigma_{max}]\).

(70)

Finally, if \(k \geq 1\) and \(\bar{\sigma}_k \notin [\max\{1, \|x_k\|\} \sigma_{min}, \sigma_{max}]\), we define

\[\tilde{\sigma}_k = h_{init} \frac{\|x_k\|}{\|F(x_k)\|}\]  

(71)

and we compute \(\sigma_k\) as the projection of \(\tilde{\sigma}_k\) onto the interval \([\max\{1, \|x_k\|\} \sigma_{min}, \sigma_{max}]\).
5.3 Procedure for reducing the step

At Step 2.4 of Algorithm 2.1, we compute \( \alpha_{\text{new}} \) as the minimizer of the the univariate quadratic \( q(\alpha) \) that interpolates \( q(0) = f(x^k) \), \( q(\alpha_k) = f(x^k - \alpha_k\sigma_k F(x^k)) \), and \( q'(0) = -\sigma_k F(x^k)^T \nabla f(x^k) = -\sigma_k F(x^k)^T J(x^k) F(x^k) \). Following [30], since we consider \( J(x^k) \) unavailable, we consider \( J(x^k) = I \). Thus,

\[
\alpha_{\text{new}}^+ = \max \left\{ \tau_{\min} \alpha_+, \min \left\{ \frac{\alpha_+^2 f(x^k)}{f(x^k - \alpha_+ \sigma_k F(x^k)) + (2\alpha_+ - 1)f(x^k)}, \tau_{\max} \alpha_+ \right\} \right\}.
\]

Analogously,

\[
\alpha_{\text{new}}^- = \max \left\{ \tau_{\min} \alpha_-, \min \left\{ \frac{\alpha_-^2 f(x^k)}{f(x^k + \alpha_- \sigma_k F(x^k)) + (2\alpha_- - 1)f(x^k)}, \tau_{\max} \alpha_- \right\} \right\}.
\]

5.4 Computing the acceleration

In Step 2 of Algorithm 4.1, computing \( x_{\text{trial}}^{k+1} \) as defined in (54) is equivalent to computing the minimum-norm least-squares solution \( \tilde{\omega} \) to the linear system \( Y_k \tilde{\omega} = F(x_{\text{trial}}^{k+1}) \) and defining \( x_{\text{trial}}^{k+1} = x_{\text{trial}}^k - S_k \tilde{\omega} \). In practice, we compute the minimum-norm least-squares solution with a complete orthogonalization of \( Y_k \). Computing this factorization from scratch at each iteration could be expensive. However, by definition, \( Y_k \) corresponds to a slight variation of \( Y_{k-1} \). In practice, when \( Y_k \) has not full numerical rank, one extra column is added to it; and if \( Y_k \) has null numerical rank, a new \( Y_k \) is computed from scratch. For completeness, the practical implementation of Algorithm 4.1 is given below.

Algorithm 5.1. Let \( 0 < h_{\text{small}} < h_{\text{large}} \) and \( p \geq 1 \) be given. Set \( r_{\text{max}} \leftarrow 0 \) and \( \ell \leftarrow 1 \).

Step 1. Define \( x_{\text{trial}}^{k+1} = x^k + \alpha_k d^k \).

Step 2.

Step 2.1. If \( k = 0 \), the set \( S_k \) and \( Y_k \) as matrices with zero columns. Otherwise, set \( S_k \leftarrow S_{k-1} \) and \( Y_k \leftarrow Y_{k-1} \).

Step 2.2. If \( S_k \) and \( Y_k \) have \( p \) columns, then remove the leftmost column of \( S_k \) and \( Y_k \).

Step 2.3. Add \( s^k = x_{\text{trial}}^{k+1} - x^k \) and \( y^k = F(x_{\text{trial}}^{k+1}) - F(x^k) \) as rightmost column of \( S_k \) and \( Y_k \), respectively; and set \( r_{\text{max}} \leftarrow \max\{r_{\text{max}}, \text{rank}(Y_k)\} \) and \( \ell \leftarrow \text{mod}(\ell, n) + 1 \).

Step 2.4. If \( \text{rank}(Y_k) < r_{\text{max}} \), then execute Steps 2.4.1–2.4.2.

Step 2.4.1. If \( S_k \) and \( Y_k \) have \( p \) columns, then remove the leftmost column of \( S_k \) and \( Y_k \).

Step 2.4.2. Add \( s_{\text{extra}} = x_{\text{extra}} - x^k \) and \( y_{\text{extra}} = F(x_{\text{extra}}) - F(x^k) \) as rightmost column of \( S_k \) and \( Y_k \), respectively, where \( x_{\text{extra}} = x^k + h_{\text{small}} \epsilon \ell \). Set \( r_{\text{max}} \leftarrow \max\{r_{\text{max}}, \text{rank}(Y_k)\} \) and \( \ell \leftarrow \text{mod}(\ell, n) + 1 \).

Step 2.5. If \( \text{rank}(Y_k) \neq 0 \), then execute Steps 2.5.1–2.5.3.

Step 2.5.1. Compute the minimum-norm least-squares solution \( \tilde{\omega} \) to the linear system \( Y_k \tilde{\omega} = F(x_{\text{trial}}^{k+1}) \) and define \( x_{\text{trial}}^{k+1} = x_{\text{trial}}^{k+1} - S_k \tilde{\omega} \).

Step 2.5.2. If Step 2.4.2 was executed, then remove the rightmost column of \( S_k \) and \( Y_k \), i.e. columns \( s_{\text{extra}} \) and \( y_{\text{extra}} \).

Step 2.5.3. If \( x_{\text{trial}}^{k+1} = x^k \), \( \|x_{\text{trial}}^{k+1}\| \leq 10 \max\{1, \|x^k\|\} \), and \( \|F(x_{\text{trial}}^{k+1})\| < \|F(x_{\text{trial}}^{k+1})\| \), then redefine \( x_{\text{trial}}^{k+1} = x_{\text{trial}}^{k+1} = x_{\text{trial}}^{k+1} = x_{\text{trial}}^{k+1}, \) substitute the rightmost column of \( S_k \) and \( Y_k \), i.e. columns \( s^k \) and \( y^k \), with \( s_{\text{accel}} = x_{\text{accel}}^{k+1} - x^k \) and \( y_{\text{accel}} = F(x_{\text{accel}}^{k+1}) - F(x^k) \), respectively, and set \( r_{\text{max}} \leftarrow \max\{r_{\text{max}}, \text{rank}(Y_k)\} \).

Step 2.6. If \( \text{rank}(Y_k) = 0 \), then execute Steps 2.6.1–2.6.3.

Step 2.6.1. Redefine matrices \( S_k \) and \( Y_k \) as matrices with zero columns.

Step 2.6.2. Execute Steps 2.6.2.1 \( p - 1 \) times.
We implemented Algorithm 2.1 and Algorithm 5.1 (the practical version of Algorithm 4.1) in Fortran 90. In numerical experiments, decomposition of $Y$ and approximating $\omega$ can be implemented with time complexity $O(np)$, the minimum-norm least-squares solution $\vec{\omega}$ is such that Step 2.6 was not executed in the previous iteration $k$. If iteration $k$ is such that Step 2.6 was not executed in the previous $p$ iterations, then matrices $S_k$ and $Y_k$ correspond to removing the leftmost column from and adding a new rightmost column to $S_{k-1}$ and $Y_{k-1}$, respectively. The leftmost column is not removed if the maximum number of columns $p$ was not reached yet. When $k = 0$, $Y_k$ is a single-column matrix for which matrices $P$, $Q$, and $R$ of the rank-revealing QR decomposition $Y_kP = QR$ can be trivially computed. For $k > 0$, the QR decomposition of $Y_k$ can be obtained with time complexity $O(np^2)$ by updating, via Givens rotations, the QR decomposition of $Y_{k-1}$. Moreover, by using the QR decomposition of $Y_k$, the minimum-norm least-squares solution $\omega$ can be computed with time complexity $O(n + p^2)$ if $Y_k$ is full-rank and with time complexity $O(n + p^3)$ in the rank-deficient case. Summing up, by assuming that $p \ll n$ and that $p$ does not depend on $n$, iterations of Algorithm 5.1 can be implemented with time complexity $O(n)$. The space complexity is $O(np + p^2)$ and it is related to the fact that we must save matrix $S_k \in \mathbb{R}^{n \times p}$ and matrices $Q \in \mathbb{R}^{n \times np}$ and $R \in \mathbb{R}^{np \times p}$ of the QR decomposition of $Y_k$. Of course, the permutation matrix $P$ can be saved in an array of size $p$. Thus, the space complexity of Algorithm 5.1 is also $O(n)$ under the same assumptions. When Step 2.6 is executed, the QR decomposition of $Y_k$ must be computed from scratch, with time complexity $O(np^2)$.

## 6 Numerical experiments

We implemented Algorithm 2.1 and Algorithm 5.1 (the practical version of Algorithm 4.1) in Fortran 90. In the numerical experiments, we considered the standard values $\gamma = 10^{-4}$, $\tau_{\min} = 0.1$, $\tau_{\max} = 0.5$, $M = 10$, $\sigma_{\min} = \sqrt{\tau}$, $\sigma_{\max} = 1/\sqrt{\tau}$, and $\eta_k = 2 - k \min\left(\frac{1}{2} \|F(x^k)\|, \sqrt{\|F(x^0)\|}\right)$, where $\epsilon \approx 10^{-16}$ is the machine precision. More crucial to the method performance are the choices of $h_{\text{init}}$, $h_{\text{small}}$, $h_{\text{large}}$, and $p$, whose values are mentioned below. All tests were conducted on a computer with a 3.4 GHz Intel Core i5 processor and 8GB 1600 MHz DDR3 RAM memory, running macOS Mojave (version 10.14.6). Code was compiled by the GFortran compiler of GCC (version 8.2.0) with the -O3 optimization directive enabled.

### 6.1 2D and 3D Bratu problem

In a first experiment, we considered 2D and 3D versions of the Bratu problem

$$-\Delta u + \theta e^u = \phi(v) \text{ in } \Omega$$

with boundary conditions $u = v$ on $\partial \Omega$. In the 2D case, $\Omega = [0, 1]^2$ and, following [28], we set $v = 10u_1u_2(1 - u_1)(1 - u_2)e^{u_1^4}$; while in the 3D case, $\Omega = [0, 1]^3$ and $v = 10u_1u_2u_3(1 - u_1)(1 - u_2)(1 - u_3)e^{u_1^4}$. In both cases, $\phi(u) = -\Delta u + \theta e^u$; so the problem has $v$ as known solution. Considering $n_p$ discretization points in each dimension and approximating $\Delta u(x) \approx \frac{u(x + he_1) + u(x - he_2) - 4u(x)}{h^2}$ and $\Delta u(x) \approx \frac{u(x + he_1) + u(x - he_2) + u(x + he_3) - 6u(x)}{h^2}$,
where $h = 1/(n_p - 1)$ and $e_i$ is the $i$th canonical vector in the corresponding space ($\mathbb{R}^2$ or $\mathbb{R}^3$), we obtain nonlinear systems of equations with $n = (n_p - 2)^2$ and $n = (n_p - 2)^3$ variables in the 2D and 3D cases, respectively. Starting from $u = 0$, fixing $\theta = -100$, and varying $n_p \in \{100, 125, \ldots, 400\}$ and $n_p \in \{10, 15, \ldots, 70\}$ in the 2D and 3D cases respectively, we run NITSOL (file nitsol.11-1-05.tar.gz downloaded from https://users.wpi.edu/~walker/NITSOL/ on October 26th, 2020) and the method proposed in the present work, which will be called Accelerated DF-SANE from now on. As stopping criterion, we considered (65) with $\varepsilon = 10^{-6} \sqrt{n}$. For NITSOL, we use all its default parameters, except the maximum number of (nonlinear) iterations, which was increased in order to avoid premature stops. By default, NITSOL corresponds to an Inexact-Newton method in which Newtonian systems are solved with GMRES (maximum Krylov subspace dimension equal to 20), approximating the Jacobian-vector products by finite-differences. For Accelerated DF-SANE, based on preliminary experiments, we set $h_{\text{small}} = 10^{-4}$, $h_{\text{large}} = 0.1$, and $h_{\text{init}} = 0.01$ in the 2D case and $h_{\text{small}} = h_{\text{large}} = 0.1$ and $h_{\text{init}} = 1$ in the 3D case. In both cases, we considered $p = 5$.

Tables 1 and 2 shows the results. In the tables, $\#it_1$ corresponds to the nonlinear iterations (outer iterations) of NITSOL; while $\#it_2$ corresponds to its linear iterations (inner or GMRES iterations). For both methods, “fcnt” stands for number of evaluations of $F$, “Time” stands for CPU time in seconds, and “SC” stands for “Stopping criterion”. Remaining columns are self-explanatory. In the case of NITSOL, stopping criterion equal to 0 means success; while 6 means “failure to reach an acceptable step through backtracking”. Accelerated DF-SANE satisfied the stopping criterion related to success in all problems. Regarding the 2D problems, it should be first noted that NITSOL failed in satisfying the stopping criterion for $n_p \geq 225$. Accelerated DF-SANE is faster than NITSOL in all problems in which both methods satisfied the stopping criterion related to success in all problems. Accelerated DF-SANE is faster than NITSOL in all problems in the table. In the larger problem in the table ($n_p = 70$), Accelerated DF-SANE is more than twenty times faster than NITSOL.

$$
\begin{array}{|c|c|c|c|c|c|c|c|}
\hline
n_p & n & \text{NITSOL (Newton-GMRES)} & & & \text{Accelerated DF-SANE} & \\
\hline
& & SC & ||F(x_i)||_2 & \#it_1 & \#it_2 & \text{fcnt} & \text{Time} & ||F(x_i)||_2 & \#it & \text{fcnt} & \text{Time} \\
\hline
150 & 21,904 & 0 & 1.5e-04 & 379 & 315,338 & 315,718 & 165.07 & 1.5e-04 & 2,946 & 6,007 & 6.59 \\
175 & 29,929 & 0 & 1.7e-04 & 401 & 331,380 & 331,782 & 265.53 & 1.7e-04 & 4,475 & 10,007 & 17.46 \\
200 & 39,294 & 0 & 2.0e-04 & 4,421 & 846,957 & 851,385 & 983.97 & 2.0e-04 & 6,775 & 14,385 & 32.24 \\
225 & 49,729 & 6 & 3.3e+01 & 3,944 & 173,310 & 177,283 & 261.03 & 2.2e-04 & 4,328 & 8,927 & 26.72 \\
250 & 61,504 & 6 & 4.1e+01 & 105 & 46,897 & 47,019 & 82.69 & 2.5e-04 & 12,661 & 26,353 & 94.47 \\
275 & 74,529 & 6 & 5.3e+01 & 1,411 & 91,194 & 95,635 & 224.74 & 2.7e-04 & 8,809 & 19,553 & 88.07 \\
300 & 88,804 & 6 & 7.2e+01 & 1,341 & 104,735 & 106,112 & 317.71 & 3.0e-04 & 15,858 & 34,194 & 190.49 \\
325 & 104,329 & 6 & 9.6e+01 & 2,430 & 194,640 & 197,126 & 765.53 & 3.2e-04 & 10,791 & 23,403 & 160.02 \\
350 & 121,104 & 6 & 1.4e+02 & 195 & 82,940 & 83,151 & 319.00 & 3.5e-04 & 10,805 & 25,915 & 176.52 \\
375 & 139,129 & 6 & 1.7e+02 & 1,127 & 121,100 & 122,243 & 649.94 & 3.7e-04 & 16,335 & 38,648 & 331.03 \\
400 & 158,404 & 6 & 2.2e+02 & 2,936 & 144,460 & 147,415 & 851.67 & 4.0e-04 & 21,106 & 55,901 & 515.65 \\
\hline
\end{array}
$$

Table 1: Performances of NITSOL and Accelerated DF-SANE in the 2D Bratu problem.

The difficulty of the Bratu problem varies with the value of $\theta$, that may be positive or negative. For the formulation given in (72), negative values of $\theta$ correspond to more difficult problem. Therefore, results in Tables 1 and 2 raise the question of how the performances of the methods compare to each other for different values of $\theta$. So, we arbitrarily considered the 3D Bratu problem with $n_p = 40$, which is affordable for both methods and varied $\theta \in [-100, 10]$. Figure 1 shows the results of applying NITSOL and Accelerated DF-SANE. The graphic shows that for $\theta \geq -20$ both methods use less than a second and NITSOL outperforms Accelerated DF-SANE. On the other hand, in the most difficult problems (i.e., $\theta < -20$), where up to one thousand seconds are required, Accelerated DF-SANE outperforms NITSOL by approximately an order of magnitude. (Note that the $y$-axis is in logarithmic scale.) Considering the number of functional evaluations as a performance metric, instead of the CPU time, results are mostly the same.

Another relevant question is how the performance of Accelerated DF-SANE varies as a function of its parameter $p$. So, considering again the 3D Bratu problem with $\theta = -100$ and $n_p = 40$, we ran Accelerated DF-SANE with $p \in \{3, 4, \ldots, 17\}$. Figure 2 shows the results. The figure on the left shows that the number of
Table 2: Performances of NITSOL and Accelerated DF-SANE in the 3D Bratu problem.

<table>
<thead>
<tr>
<th>( n_p )</th>
<th>( n )</th>
<th>( | F(x*) |_2 )</th>
<th>#it1</th>
<th>#it2</th>
<th>fcnt</th>
<th>Time</th>
<th>Accelerated DF-SANE</th>
<th>( | F(x*) |_2 )</th>
<th>#it</th>
<th>fcnt</th>
<th>Time</th>
</tr>
</thead>
<tbody>
<tr>
<td>10</td>
<td>512</td>
<td>1.7e-05</td>
<td>5</td>
<td>213</td>
<td>219</td>
<td>0.00</td>
<td>2.1e-05</td>
<td>126</td>
<td>308</td>
<td>0.00</td>
<td></td>
</tr>
<tr>
<td>15</td>
<td>2,197</td>
<td>3.7e-05</td>
<td>7</td>
<td>1,621</td>
<td>1,629</td>
<td>0.07</td>
<td>4.7e-05</td>
<td>223</td>
<td>662</td>
<td>0.05</td>
<td></td>
</tr>
<tr>
<td>20</td>
<td>5,832</td>
<td>6.1e-05</td>
<td>11</td>
<td>7,157</td>
<td>7,169</td>
<td>0.97</td>
<td>7.2e-05</td>
<td>1,221</td>
<td>4,271</td>
<td>0.92</td>
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</tr>
<tr>
<td>25</td>
<td>12,167</td>
<td>8.8e-05</td>
<td>19</td>
<td>15,893</td>
<td>15,913</td>
<td>4.62</td>
<td>1.1e-04</td>
<td>529</td>
<td>1,840</td>
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<td></td>
</tr>
<tr>
<td>30</td>
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<td>3.7e-05</td>
<td>7</td>
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<td>22,017</td>
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<td>662</td>
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<tr>
<td>35</td>
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<td>38</td>
<td>34,935</td>
<td>34,974</td>
<td>36.31</td>
<td>1.9e-04</td>
<td>1,210</td>
<td>4,530</td>
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</tr>
<tr>
<td>40</td>
<td>54,872</td>
<td>2.2e-04</td>
<td>70</td>
<td>66,115</td>
<td>66,186</td>
<td>118.56</td>
<td>2.3e-04</td>
<td>1,109</td>
<td>4,379</td>
<td>9.35</td>
<td></td>
</tr>
<tr>
<td>45</td>
<td>79,507</td>
<td>2.8e-04</td>
<td>147</td>
<td>137,626</td>
<td>137,774</td>
<td>386.09</td>
<td>2.8e-04</td>
<td>1,315</td>
<td>5,444</td>
<td>20.06</td>
<td></td>
</tr>
<tr>
<td>50</td>
<td>110,592</td>
<td>3.3e-04</td>
<td>217</td>
<td>199,042</td>
<td>199,260</td>
<td>759.61</td>
<td>3.3e-04</td>
<td>1,574</td>
<td>6,501</td>
<td>30.38</td>
<td></td>
</tr>
<tr>
<td>55</td>
<td>148,877</td>
<td>3.8e-04</td>
<td>357</td>
<td>312,023</td>
<td>312,381</td>
<td>1736.95</td>
<td>3.8e-04</td>
<td>1,706</td>
<td>7,254</td>
<td>53.10</td>
<td></td>
</tr>
<tr>
<td>60</td>
<td>195,112</td>
<td>4.4e-04</td>
<td>504</td>
<td>418,201</td>
<td>418,706</td>
<td>3074.25</td>
<td>4.4e-04</td>
<td>1,821</td>
<td>8,019</td>
<td>70.29</td>
<td></td>
</tr>
<tr>
<td>65</td>
<td>250,047</td>
<td>5.0e-04</td>
<td>988</td>
<td>691,192</td>
<td>692,181</td>
<td>6715.18</td>
<td>5.0e-04</td>
<td>2,061</td>
<td>9,379</td>
<td>109.39</td>
<td></td>
</tr>
<tr>
<td>70</td>
<td>314,432</td>
<td>5.6e-04</td>
<td>609</td>
<td>486,651</td>
<td>487,261</td>
<td>5735.90</td>
<td>5.6e-04</td>
<td>1,836</td>
<td>8,431</td>
<td>118.13</td>
<td></td>
</tr>
</tbody>
</table>

Figure 1: This figure shows the CPU time (in seconds) used by NITSOL and Accelerated DF-SANE to solve the 3D Bratu problem with \( n_p = 40 \) and \( \theta \in [-100, 10] \).

iterations of Accelerated DF-SANE is nearly constant as a function of \( p \); while the figure on the right shows, as expected, that, the larger \( p \), the larger the CPU time per iteration. Note that for the considered values of \( p \), the variation on the number of iterations is up to 20%; while the variation in CPU time is up to 300%.

Another relevant question regarding Accelerated DF-SANE is how it compares against DF-SANE, i.e. the original method without the acceleration being proposed in the present work. We were unable to run DF-SANE in the set of problems considered in Tables 1 and 2 because DF-SANE is unable to reach the stopping criterion (65) with \( \varepsilon = 10^{-6} \sqrt{n} \) within an affordable time. As an alternative, we considered the four instances of the 3D Bratu problem with \( n_p \in \{40, 70\} \) and \( \theta \in \{-100, 10\} \). In the two instances with \( \theta = 10 \), DF-SANE was able to satisfy the imposed stopping criterion within an affordable time. This result was in fact expected because for this value of \( \theta \) the Bratu problem resembles the minimization of a convex quadratic function; and Barzilai-Borwein or spectral step based methods are expected to perform especially well in this case. In the two instances with \( \theta = -100 \), we first run Accelerated DF-SANE and then we run DF-SANE using as CPU time limit the time consumed by Accelerated DF-SANE. Figure 3 shows the results. It is very clear from the four graphics that the acceleration process is very efficient in its purpose of accelerating DF-SANE. (In the cases with \( \theta = -100 \), with an excruciatingly slow progress, DF-SANE is far from reaching convergence after a couple of hours of CPU time.)
Figure 2: Performance of Accelerated DF-SANE in the 3D Bratu problem with $n_p = 40$ and $\theta = -100$ varying the number $p$ of columns in matrices $S_k$ and $Y_k$.

Figure 3: Performance of DF-SANE and Accelerated DF-SANE in four instances of the 3D Bratu problem with $\theta \in \{-100, 10\}$ and $n_p \in \{40, 70\}$. 
6.2 Modified Bratu, Driven Cavity, and Flow in a Porous Media problems

In a second set of experiments, we considered three PDE-based problems described in [41]. Fortran implementations of these problems are included as examples of usage in the NITSOL distribution. The three problems are discretization of 2D PDE-based systems of nonlinear equations. All parameters of the problems were set as suggested in [41]. We varied the value of \( n_p \) for the three problems trying to illustrate the behavior of the methods and solving as large as possible problems with both methods within an affordable time. NITSOL was run with all its default parameters, as in the previous section; while Accelerated DF-SANE was run with the same parameters already reported for the 2D Bratu problem in the previous section. Tables 3-5 show the results. Table 3 shows that NITSOL failed in satisfying the stopping criterion for \( n_p \geq 135 \) in the Driven Cavity problem. Accelerated DF-SANE was faster than NITSOL in all instances that both methods solved. In the largest instance that both methods solved, Accelerated DF-SANE is more than thirty times faster than NITSOL. Table 4 shows that NITSOL is faster than Accelerated DF-SANE in the smaller instances of the Generalized 2D Bratu problem (\( n_p \) up to 750). In the largest instance in the table (\( n_p = 1,500 \)), Accelerated DF-SANE takes around half of the time required by NITSOL. Finally, Table 5 shows that NITSOL is faster than Accelerated DF-SANE in the smaller instances of the Flow in a Porous Media problem. In the largest instance in the table (\( n_p = 1,500 \)), Accelerated DF-SANE takes around half of the time required by NITSOL.

<table>
<thead>
<tr>
<th>( n_p )</th>
<th>( n )</th>
<th>NITSOL (Newton-GMRES)</th>
<th>Accelerated DF-SANE</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>( |F(x_0)|_2 )</td>
<td>#it ( h_1 )</td>
</tr>
<tr>
<td></td>
<td></td>
<td>( |F(x_0)|_2 )</td>
<td>#it</td>
</tr>
</tbody>
</table>

Table 3: Performances of NITSOL and Accelerated DF-SANE in the “Driven Cavity Problem”.
NITSOL (Newton-GMRES)  
5.0e 8.0e  
Accelerated DF-SANE  
3.0e 1.1e  
6.9e 1.3e  
2.4e 2.0e  
1.3e 1.1e  
4.5e 1.4e  
1.0e 1.4e  
1.1e 9.9e  
1.2e 5.2e  
7.4e 9.0e  
∥8.4e 3.4e  
8.0e 5.5e  
1.9e 1.5e  
7.2e 1.3e  
5.6e 6.3e  
3.4e 1.2e  
4.1e 1.6e  
1.4e 1.1e  
6.4e 2.8e  
2.5e 6.2e  
7.0e 1.0e  

Table 4: Performances of NITSOL and Accelerated DF-SANE in “Generalized 2D Bratu Problem”.

7 Conclusions

The Sequential Secant Method, which is the most obvious multidimensional generalization of the secant method for solving nonlinear equations, seems to have been introduced by Wolfe in 1959 [49]. See also [2]. This method has been analyzed in classical books and papers [40, 46, 27]. Under suitable conditions, local convergence for solving nonlinear equations seems to have been introduced by Wolfe in 1959 [49]. See also [2]. This method, the Sequential Secant Method, which is the most obvious multidimensional generalization of the secant method, has been analyzed in classical books and papers [40, 46, 27]. Under suitable conditions, local convergence for solving nonlinear equations seems to have been introduced by Wolfe in 1959 [49]. See also [2]. This method, the Sequential Secant Method, computes

$$x^{k+1} = x^k - (s^{k-n},...,s^{k-1})(y^{k-1},...,y^{k-1})^{-1}F(x^k).$$ (73)

This iteration is well defined if the matrix $(y^{k-n},...,y^{k-1})$ is nonsingular. Moreover, the good local convergence properties need uniformly linear independence of the increments $s^{k-n},...,s^{k-1}$. The method has been updated in several ways in order to fix these drawbacks while maintaining its convergence properties. The natural limited memory version of (73) is defined by

$$x^{k+1} = x^k - (s^{k-p},...,s^{k-1})(y^{k-p},...,y^{k-1})^{-1}F(x^k),$$ (74)

where $1 \leq p \ll n$. This formula is inconvenient for solving nonlinear systems because $s^k$ necessarily belongs to the subspace generated by $s^{k-p},...,s^{k-1}$ which implies that $x^{k+1}$ is in the affine subspace determined by $x^{k-p},...,x^{k-1},x^k$, for all $j$, and so, convergence to a solution cannot occur unless the solution belongs to the same affine subspace. This is the reason why, in the present paper, we do not use the method defined by (74). Instead, we compute $x^{k+1}$ using the sequential residual approach, we define $s^k = x^{k+1} - x^k$, $y^k =$
Table 5: Performances of NITSOL and Accelerated DF-SANE in “Flow in a Porous Media Problem”.

$$F(x_{\text{trial}}^{k+1}) - F(x^k)$$

$$x_{\text{acc}}^{k+1} = x_{\text{trial}}^{k+1} - (s_{k-p+1}, \ldots, s_{k-1}, s_k)(y_{k-p+1}, \ldots, y_{k-1}, y_k)^T F(x_{\text{trial}}^{k+1})$$

and we choose $x^{k+1}$ as the best of the trials $x_{\text{trial}}^{k+1}$ and $x_{\text{acc}}^{k+1}$. In this way, we preserve the good properties of the Sequential Secant Method associated with the good global behavior of Sequential Residual approaches. The freedom on the choice of the residual step favors the employment of preconditioners when they are available.

The most popular methods for solving nonlinear systems of equations in which the application of Newton’s method is impossible or extremely expensive are based on the Inexact Newton approach with Krylov-subspace methods (as GMRES) for solving approximately the Newtonian linear systems at each iteration. These methods have a long tradition and, very likely, they deserve to be the preferred ones by practitioners in the numerical PDE community. Nevertheless, in this paper we showed that, for some very large interesting problems, an approach based in sequential-secant-like accelerations of a residual method is more efficient than a standard implementation of Newton-GMRES with its default algorithmic parameters. This indicates that those problems possess characteristics that favor the application of the secant paradigm over the inexact Newton one. Of course, the opposite situation probably occurs in many cases. This means, therefore, that efficiency for solving practical problems will be increased if practitioners have easy access to both types of methods. The codes used in this paper, that make it possible the reproducibility of all the experiments, as well as the instructions for using our algorithm may be found in [http://www.ime.usp.br/~egbirgin/sources/accelerated-df-sane/](http://www.ime.usp.br/~egbirgin/sources/accelerated-df-sane/).

Future work will include the employment of the new methods to the acceleration of KKT solvers that are necessary in the Augmented Lagrangian approach for constrained optimization [5].
References