

## impVRDFON: Implementation details of VRDFON

Implementation details of the VRDFON solver, called `impVRDFON`, is provided. The VRDFON package is publicly available at [17].

In Section 1, we first describe several heuristic enhancements added to the VRDFON algorithm and discuss under what conditions the complexity results for an implemented version of VRDFON can be guaranteed.

In Section 2, we give extensive numerical results:

- Subsection 2.1 defines performance measures.
- Subsection 2.2 provides details on the solvers compared.
- Subsection 2.3 makes a testing and tuning for VRDFON.
- Subsection 2.4 compares VRDFON with 20 state-of-the-art DFO solvers.

## 1 Heuristic enhancements

In this section we propose many practical improvements that make VRDFON a very competitive solver. Two of the most important of these are **surrogate quadratic models in adaptively determined subspaces** and finding and updating step sizes in an implemented version of MLS. An implemented version of DS uses MLS with different search directions and heuristically reconstructs step sizes in each unsuccessful iteration, resulting in small extrapolation step sizes. VRDFON calls DS repeatedly until an  $\varepsilon = \sqrt{\omega}$ -approximate stationary point is found.

### 1.1 Random approximate coordinate directions

The use of random directions is preferable to the use of deterministic directions (see [3]). On the other hand, it is known that coordinate directions are useful to estimate the gradient, since at least one of these directions has a good angle with the gradient (see [8]). Inspired by these points, we construct an effective version of scaled random directions, which we call **random approximate coordinate directions**, that combine the advantages of both scaled random and coordinate directions. The random coordinate direction  $p$  multiplies a standard random direction by a scaling vector having one component equals to  $1/\|p\|$  and the others equal to  $\gamma_{rd}/\|p\|$  with  $0 < \gamma_{rd} < 1$ , where is a small tuning parameter.

## 1.2 Subspace information

The points  $Z_i$  with the best inexact function values are stored as columns of a matrix  $Z$ , their inexact function values  $\tilde{f}(Z_i)$  in the vector  $F$ , and their step sizes  $\alpha_i$  in the vector  $S$ . We adjust the matrix  $Z$  if its entries are contaminated by NaN or  $\pm\infty$  by inserting a large positive tuning parameter  $\gamma_Z > 0$ .

We denote the points stored in  $Z$  as **sample points**, whose maximum number is defined by  $m_{\max} := \min \left\{ \overline{m}, \frac{1}{2}n(n+3) \right\}$ , where  $\overline{m}$  is a tuning parameter. The **number of sample points** is denoted by  $m \in [2, m_{\max}]$  and the **subspace size**  $m^\circ$  is defined as the largest integer satisfying  $\frac{1}{2}m^\circ(m^\circ + 3) \leq m$ .

If  $m$  is greater than  $m_{\max}$ , we update  $Z$ ,  $F$ ,  $S$  by replacing the worst point, its inexact function value, and its step size with the current best point, its inexact function value, and its step size, respectively. Otherwise, we append the current best point to  $Z$ , its inexact function value to  $F$ , and its step size to  $S$ .

## 1.3 Random subspace directions

In [19], it was shown by extensive numerical results that after using a derivative-free line search algorithm with coordinate directions, the use of random subspace directions by such an algorithm is very useful. Inspired by this, after using random coordinate directions by VRDFON, random subspace directions are used by such an algorithm in the hope of reducing the influence of noise.

We write  $A_{II} := (A_{ij})_{i \in I, j \in I}$  for the submatrix of  $A$  with row and column indices of  $I$  by  $A_{II}$ ,  $A_{:k}$  for the  $k$ th column of a matrix  $A$ , and  $b$  for the index of the best point.

Random subspace directions are constructed based on the information of sample points with good function values. As in [19], we generate an  $(m-1) \times 1$  standard random vector  $\alpha^{\text{rs}}$  and then scale  $\alpha^{\text{rs}}$  as  $\alpha^{\text{rs}} := \alpha^{\text{rs}} / \|\alpha^{\text{rs}}\|$ . Then we

compute the **random subspace direction** by  $p := \sum_{i=1, i \neq b}^m \alpha_i^{\text{rs}} (Z_{:i} - Z_{:b})$ .

## 1.4 Reduced quadratic models

It is well-known [15] that model-based algorithms need at least

$$N := n + \frac{1}{2}n(n+1) = \frac{1}{2}n(n+3)$$

sample points and  $\mathcal{O}(N^3)$  operations for estimating the gradient vector and Hessian matrix of an objective quadratic model. For medium or large scale problems, this is prohibitively expensive. To overcome this problem, we construct quadratic models in adaptively determined subspaces called **reduced quadratic models**, one of which is a fully quadratic model when  $m = N$ .

For all  $i = 1, \dots, m$ , let  $z_i$  and  $\tilde{f}_i := \tilde{f}(z_i)$  be the sample points and their inexact function values stored in  $Z$  and  $F$ , respectively, and define  $\mathbf{s}_i := z_i - Z_{:,b}$ . Before defining the model errors, we need to know whether or not the number of sample points  $m$  is large enough to build a full or reduced quadratic model. Therefore, we compute all the subspace sizes that are admissible to construct quadratic models. The subspace size defined in Subsection 1.2 is calculated as follows

$$m^\circ := \left\lfloor \frac{1}{2}(-3 + \sqrt{9 + 8m}) \right\rfloor. \quad (1)$$

It is clear that for  $m < N$  no fully quadratic model can be constructed; reduced quadratic models are constructed instead. When the dimension is larger than  $m_{\max}$ , the  $n \times m_{\max}$  matrix  $Z$  and the  $n \times 1$  vector  $z^m$  are reduced to the  $m_{\max} \times m^\circ$  matrix  $Z^\circ$  and the  $m^\circ \times 1$  vector  $z^\circ$ , respectively. In other words, the restriction of the entries of  $Z$  and  $z^m$  is done by choosing a random subset of size  $m^\circ$ .

Some components of each best point stored in  $Z$  may be ignored. To overcome this shortcoming, reduced quadratic models are constructed several times before  $m$  exceeds  $m_{\max}$ , each time choosing  $Z^\circ$  ( $z^\circ$ ) from a random subset of the entries of  $Z$  ( $z_{\text{best}}$ ). The different subsets are not drawn separately, so entries between two consecutive samples may overlap.

We denote by  $g(x)$  the gradient of  $f$  at  $x$ , by  $\tilde{g}(x)$  the approximation of  $g(x)$ , by  $B(x)$  the Hessian matrix of  $f$  at  $x$  and by  $\tilde{B}(x)$  its approximation. We write  $\tilde{g}^\circ$  and  $\tilde{B}^\circ$  for the estimated gradient vector and the symmetric Hessian matrix in a subspace, respectively. Let  $M := \frac{1}{2}m^\circ(m^\circ + 3)$  and

$\mathbf{K} := \min(2M, m - 1)$ . To evaluate the inexact  $\tilde{g}^o$  and  $\tilde{B}^o$ , we define the model errors by

$$\varepsilon_i := \frac{\tilde{f}_i - F_b - (\tilde{g}^o)^T \mathbf{s}_i - \frac{1}{2} \mathbf{s}_i^T \tilde{B}^o \mathbf{s}_i}{\mathbf{sc}_i} \quad \text{for all } i = 1, \dots, \mathbf{K}, \quad (2)$$

where  $\mathbf{sc}$  is an appropriate scaling vector. It is shown in DEUFLHARD & HEINDL [9] that numerical methods can generally perform much better when they preserve affine invariance. The following choice ensures the affine invariance of the fitting method.

The numerator of (2) is  $\mathcal{O}(\|\mathbf{s}_i\|^3)$  if  $m = N$ ; otherwise it is  $\mathcal{O}(\|\mathbf{s}_i\|^2)$ . To obtain  $\varepsilon_i$  of a uniform magnitude, we choose  $\mathbf{sc}$  as follows:

- (1) We form the matrix  $\mathbf{S} := (\mathbf{s}_1 \ \cdots \ \mathbf{s}_{\mathbf{K}})^T$ , where  $\mathbf{s}_i := z_i^o - Z_{:,b}^o$  for all  $i = 1, \dots, \mathbf{K}$ .
- (2) We compute the matrix

$$H^o := \left( \sum_l \mathbf{s}_l \mathbf{s}_l^T \right)^{-1} \quad (3)$$

by constructing a reduced QR factorization  $\mathbf{S} = \mathbf{Q}\mathbf{R}$ , where  $\mathbf{Q} \in \mathbb{R}^{\mathbf{K} \times m^o}$  is an orthogonal matrix and  $\mathbf{R} \in \mathbb{R}^{m^o \times m^o}$  is a square upper triangular matrix.

- (3) We compute the scaling vector  $\mathbf{sc}$

$$\mathbf{sc}_i := (\mathbf{s}_i^T H^o \mathbf{s}_i)^{e/2} \quad \text{for } i = 1, \dots, \mathbf{K}, \quad (4)$$

with  $\mathbf{s}_i^T H^o \mathbf{s}_i = \|\mathbf{R}^{-T} \mathbf{s}_i\|^2$  for  $i = 1, \dots, \mathbf{K}$ . In (4),  $e = 3$  if  $m = N$  holds (full quadratic model) and  $e = 2$  otherwise (reduced quadratic model).

We calculate  $H^o$  and  $\mathbf{sc}$  in the same way as SNOBFIT [15] by performing a reduced QR factorization, except that they are performed in adaptively determined subspaces.

We have  $\varepsilon = Ay - a$ , where, for  $i = 1, \dots, \mathbf{K}$ ,

$$a_i := \frac{F_b - \tilde{f}_i}{\mathbf{sc}_i} \quad \text{and} \quad A_{ij} := \begin{cases} \frac{\mathbf{s}_i^j}{\mathbf{sc}_i} & \text{if } j \in \{1, \dots, m^o\}, \\ \frac{(\mathbf{s}_i^{j-m^o})^2}{2\mathbf{sc}_i} & \text{if } j \in \{m^o + 1, \dots, 2m^o\}, \\ \frac{\mathbf{s}_i^{j'} \mathbf{s}_i^{j''}}{\mathbf{sc}_i} & \text{if } j \in \{2m^o + 1, \dots, M\}. \end{cases} \quad (5)$$

Here  $j'$  and  $j''$  are the remainders of the division of  $j - 2m^\circ$  and  $j - 2m^\circ + 1$  by  $m^\circ$ , respectively, and  $\mathbf{s}_i^j$  is the  $j$ th component of the vector  $\mathbf{s}_i$ . To find the entries of the inexact  $\tilde{g}^\circ$  and  $\tilde{B}^\circ$  we solve the linear least squares problem

$$\min_{y \in \mathbb{R}^M} \|Ay - a\|_2^2. \quad (6)$$

In finite precision arithmetic, each of the vectors  $a$ ,  $\mathbf{sc}$ ,  $y$ , and hence  $\tilde{g}^\circ$ ,  $\tilde{B}^\circ$  can have entries with value NaN or  $\pm\infty$ . We replace the components of the vectors  $a$ ,  $\mathbf{sc}$  and  $y$  with value NaN or  $\pm\infty$  by a large positive tuning parameter  $\gamma_v > 0$ .

We construct the vector  $a$  and the matrix  $A$  by (5), adjust  $a$ , and find all multipliers by solving (6). Then we adjust  $y$  and define  $\tilde{g}^\circ$  by the first  $m^\circ$  components of  $y$ , the diagonal entries of  $\tilde{B}^\circ$  by the next  $m^\circ$  components of  $y$ , and the off-diagonal entries of  $\tilde{B}^\circ$  symmetrically by the remaining entries of  $y$ .

In summary, we construct reduced quadratic models. Two advantages of such models are the use of limited sample points and the ability to construct them multiple times by increasing the size of the subspace from 2 to  $m_{\max}$ . To obtain a robust model, we adjust the matrix  $Z$  and the vector  $y$  whenever they are contaminated by NaN or  $\pm\infty$ . Second, we compute  $\mathbf{sc}$  and adjust it. Third, we calculate  $y$  and adjust it. Finally, we estimate  $\tilde{g}^\circ$  and  $\tilde{B}^\circ$ .

## 1.5 Perturbed random directions

In addition to scaled random directions and random subspace directions, the use of other directions, such as **perturbed random directions** (see below) and **improved trust region directions** (explained later in Subsection 1.6) can improve the efficiency of the method in the presence of noise.

**perturbed random directions** are used if  $\tilde{B}^\circ$  is not computable, which implies that at least one entry of  $\tilde{B}^\circ$  is contaminated by NaN or  $\pm\infty$ . Otherwise, reduced quadratic models are constructed in Subsection 1.4 and improved trust region directions are computed.

A perturbed random direction  $\tilde{p} := p^\circ - \alpha^\circ \tilde{g}^\circ$  is a perturbation of a standard random direction  $p^\circ$  by the steepest descent direction  $-\tilde{g}^\circ$  with the approximate gradient. Both  $\tilde{g}^\circ$  and  $p^\circ$  are restricted to a subspace which is

a random subset  $\mathcal{J}$  of  $\{1, \dots, n\}$  whose size is  $m^\circ$ . For all  $i \notin \mathcal{J}$ ,  $\tilde{p}_i = 0$ . To be numerically appropriate, both are scaled by the heuristic step size  $\alpha^\circ := (1 + \kappa(\tilde{g}^\circ)^T p^\circ) / \|\tilde{g}^\circ\|^2$  and the decreasing sequence  $\kappa := 1/(1 + n_f)^{\gamma_\kappa}$  with the tuning parameter  $0 < \gamma_\kappa < 1$  and the number of function evaluations  $n_f$ . It is easy to show that  $\tilde{p}^T \tilde{g} = (\kappa p^\circ - \alpha^\circ \tilde{g}^\circ)^T \tilde{g}^\circ < 0$ ; hence the perturbed random directions are descent directions.

## 1.6 An improved trust region direction

The goal of trust region methods is to restrict steps within a trust region to increase the accuracy of surrogate models. Therefore, trust region directions can be very useful, even in the presence of noise.

We now solve the trust region subproblem in a subspace

$$\begin{aligned} \min \quad & \zeta^T \tilde{g}^\circ + \frac{1}{2} \zeta^T \tilde{B}^\circ \zeta \\ \text{s.t.} \quad & \|\zeta - z^\circ\|_\infty \leq d, \end{aligned}$$

whose solution is denoted by  $\zeta_{\text{best}}$ . Here  $d$  denotes the **trust region radius**. The trust region direction is  $p_{\text{tr}}^\circ := \zeta_{\text{best}} - z^\circ$  in a subspace which is a random subset of  $\{1, \dots, n\}$  whose size is  $m^\circ$ . The idea is to construct the **improved trust region direction** by scaling  $p_{\text{tr}}^\circ$  with the positive tuning parameter  $\gamma_p$  and perturbing it by  $p_{\text{mean}} := z_{\text{mean}} - Z_{:,b}$ , where  $z_{\text{mean}}$  is the mean of  $Z$ . This perturbation transforms  $p_{\text{tr}}^\circ$  into a full subspace direction enriched by  $p_{\text{mean}}$ .

## 1.7 An implemented version of MLS

In this section, we discuss the implementation of MLS. MLS is enriched by heuristic techniques for finding and updating step sizes.

The efficiency of line search methods depends on how their step sizes are updated. The line search algorithms discussed in [19] and [22] find their step sizes in a way that does not seem to be effective, especially for large scale problems, because step sizes are updated independently; they depend only on the corresponding step size generated by the previous executions. To address this shortcoming, we generate and update step sizes in a new way.

Let  $0 \leq \underline{\alpha}_{\text{init}} < \overline{\alpha}_{\text{init}} \leq \infty$  be the lower and upper bounds of an initial interval, which are the tuning parameters. We start with the initial interval

$$[\underline{\alpha}, \overline{\alpha}] = [\underline{\alpha}_{\text{init}}, \overline{\alpha}_{\text{init}}].$$

Then  $\underline{\alpha}$  or  $\overline{\alpha}$  is updated. We describe this below.

• **The initial interval.** When the trial point  $z = z_{\text{best}} \pm \alpha p$  and its inexact function value  $\tilde{f}(z)$  are computed in lines 1 and 4 of **EP** in [18], here the gain  $\tilde{f}(z_{\text{best}}) - \tilde{f}(z)$  is stored in the vector  $\mathbf{dF}$  and its step size  $\alpha$  in the vector  $\mathbf{a}$ . Namely, when an extrapolation with many sufficient gains along  $p$  is found, much information is stored in the vectors  $\mathbf{dF}$  and  $\mathbf{a}$ . Then, an index set of the stored points with the decreasing inexact function values is found by

$$\text{ind} := \{i \mid \mathbf{dF}_i < 0 \text{ for } i = 1, \dots, n\}.$$

If  $\text{ind}$  is non-empty, the lower bound of the interval  $[\underline{\alpha}, \overline{\alpha}]$  can be found by  $\underline{\alpha} := \max(\mathbf{a}_{\text{ind}})$ ; if  $\underline{\alpha} = \underline{\alpha}_{\text{init}}$  is given as a positive tuning parameter, then  $\underline{\alpha}$  is updated by  $\underline{\alpha} := \min(\underline{\alpha}, \max(\mathbf{a}_{\text{ind}}))$ . Next, an index set of positive gains or the corresponding step sizes strictly larger than  $\overline{\alpha}$  is found by

$$\overline{\text{ind}} := \{i \mid \mathbf{dF}_i \geq 0 \text{ or } \mathbf{a}_i > \overline{\alpha} \text{ for } i = 1, \dots, n\}.$$

If  $\overline{\text{ind}}$  is nonempty, an upper bound on the initial interval is found by  $\overline{\alpha} := \min(\mathbf{a}_{\overline{\text{ind}}})$ ; if  $\overline{\alpha} = \overline{\alpha}_{\text{init}}$  is given as a positive tuning parameter, the upper bound of the initial interval is updated by  $\overline{\alpha} := \max(\overline{\alpha}, \min(\mathbf{a}_{\overline{\text{ind}}}))$ .

• **The initial extrapolation step size.** If extrapolation step sizes are too small, extrapolations may be performed slowly with many function evaluations. To overcome this problem, we only change  $\alpha = \delta$ , in line 1 in **MLS**, to  $\alpha := \max\{\sqrt{\underline{\alpha}\overline{\alpha}}, \delta\}$  whenever the interval  $[\underline{\alpha}, \overline{\alpha}] \subseteq (0, \infty)$  is found.

• **Reducing extrapolation step sizes.** Since the source of the inefficiency of the line search algorithm is generating step sizes that are too small when the interval has already been found, we change  $\alpha = \alpha/\gamma_e$ , in line 10 in **MLS** in [18], to

$$\alpha := \max \left\{ \overline{\alpha}_{\min}, \min \left\{ \sqrt{\underline{\alpha}\overline{\alpha}}, \frac{\alpha}{\gamma_e} \right\} \right\}$$

with the goal of slowly reducing step sizes. Here  $\overline{\alpha}_{\min} \in (0, 1)$  is a minimum threshold for the step size which is a tuning parameter.

• **Updating the interval.** After finding the new step size  $\alpha$ , we update either the lower or the upper bound of the interval. This can be done in the

following two cases:

(i) After extrapolation has been performed with  $\gamma$ -reduction of  $\tilde{f}$  along one of  $\pm p$ .

(ii) After no  $\gamma$ -reduction of  $\tilde{f}$  along  $\pm p$  has been found.

To do this, we insert the following statement between lines 10 and 11 of MLS [18]:

<b>if</b> $\alpha > \underline{\alpha}$ <b>then</b> , set $\bar{\alpha} = \alpha$ ;	$\triangleright$ upper bound is updated
<b>else</b> , set $\underline{\alpha} = \alpha$ ;	$\triangleright$ lower bound is updated
<b>end if</b>	

• **Decrease of  $f$  in flat regions.** If the  $r$ th iteration of MLS is unsuccessful (no  $\gamma$ -reduction of  $\tilde{f}$  along  $\pm p$  is found), the best point, its inexact function value, and its step size are updated when  $\tilde{f}(z_{\text{best}} + \alpha p) < \tilde{f}(z_{\text{best}})$ . In particular, this can happen in very flat regions of the feasible domain that do not contain a nearby stationary point. Therefore, we insert the following statement between lines 9 and 10 in MLS [18]:

**if**  $\tilde{f}(z) < \tilde{f}(z_{\text{best}})$ , set  $z_{\text{best}} := z$  and  $\tilde{f}(z_{\text{best}}) := \tilde{f}_{\text{penult}}$ ; **end if**

• **Updating the best point.** If the  $r$ th iteration of MLS is successful ( $\gamma$ -reductions of  $\tilde{f}$  are found along one of  $\pm p$ ). A point with lowest inexact function value is chosen as the new best point. As discussed in Subsection 1.2, the matrices  $X$ ,  $F$ , and  $S$  are then updated. In fact, lines 7 of EP [18] should be changes to

find  $i' := \min_{i \geq 0} \{ \tilde{f}(z_{\text{best}} + \alpha \gamma_e^i p) \mid \tilde{f}(z_{\text{best}} + \alpha \gamma_e^i p) < \tilde{f}(z_{\text{best}}) \};$   
compute  $\tilde{\alpha} = \gamma_e^{i'} \alpha$ ,  $z_{\text{best}} := z_{\text{best}} + \tilde{\alpha} p$  and choose  $\tilde{f}(z_{\text{best}}) := \bar{F}_{i'}$ ;

here  $\bar{F}$  is the list of inexact function values at the trial points evaluated by extrapolation.

## 1.8 The implementation of DS

This section discusses the implementation of DS that extended to include different search directions and heuristic techniques for updating step sizes and reconstructing the lower and upper bounds of the interval. It also describes how many worst case function evaluations are used in each successful and unsuccessful iteration of DS.

Denote by  $d_t$  the trust region radius in the iteration  $t$  and let  $C$  be the number



of random approximate coordinate directions. Let  $\gamma_{d_1} > 1$  and  $\gamma_{d_2} \in (0, 1)$  be the parameters for updating  $d_t$  and let  $0 < d_{\min} < d_{\max} < \infty$  be the parameters for controlling  $d_t$ . As described in Section 4 of [18], **DS** has  $R$  calls to **MLS** with scaled random directions. An implemented version of **DS** retains this procedure and additionally has  $C$  calls to **MLS** with random approximate coordinate directions with the goal of restoring and updating the  $m$  sample points and their inexact function values. Then, **DS** constructs a quadratic model or a linear model to produce an improved trust region direction or a perturbed random direction, and makes  $T_0$  calls to **MLS** along these directions as long as **MLS** finds decreases in  $\tilde{f}$ . If **DS** cannot find any decrease in  $\tilde{f}$  for all  $T_0$  calls to **MLS**, the lower bound  $\underline{\alpha}$  of the current interval  $[\underline{\alpha}, \bar{\alpha}]$  becomes too small, so the length of this interval becomes large. Therefore, we need to restart the interval and heuristically reconstruct the lower and upper bounds of this interval using the information from the  $m$  sample points. Let  $\gamma_a > 0$  be the parameter for adjusting heuristic step size. The structure of **DS** is shown below.

## 1.9 An implemented version of VRDFON

As mentioned earlier, **VRDFON** uses the **DS** and **MLS** procedures, while using all practical enhancements to achieve an  $\varepsilon = \sqrt{\omega}$ -approximate stationary point.

If step sizes are too small, line search methods may end up in strongly non-convex regions before a minimizer is found. To overcome this drawback, line 7 of **VRDFON** is changed as

**if**  $\sim\text{successDS}$  **then**  $\delta := \delta/Q$ ; **else**,  $\delta := \max(\delta, \sqrt{\underline{\alpha}\bar{\alpha}})$ ; **end if**

Here  $\delta := \max(\delta, \sqrt{\underline{\alpha}\bar{\alpha}})$  is used if the interval  $[\underline{\alpha}, \bar{\alpha}] \subseteq (0, \infty)$  has already been found.

## 1.10 Complexity results for the implemented version of VRDFON

To obtain the complexity results for **VRDFON**, random scaled directions should be used; this is not a problem to guarantee the complexity results when other directions are used. In fact, the order of the complexity bounds of **VRDFON** does not change for all cases after applying the heuristic improvements, although the constant factors may become larger, as in the deterministic

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**Algorithm 1** An implemented version of DS
 

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1: for  $t = 1, \dots, T_0$  do
2:   perform MLS with either  $R$  scaled random directions,  $C$  random
3:   approximate coordinate directions, or both;
4:   if  $m \geq 3$  then
5:     while true do
6:       perform MLS with random subspace directions;
7:       if no decrease in  $\tilde{f}$  by MLS, break; end
8:     end while
9:   end if
10:  if  $m \geq 2$  then ▷ a reduced quadratic or a linear model
11:    if both estimations are computable then
12:      construct a reduced quadratic model;
13:      generate the trust region radius by  $d_t := \|z_{\text{mean}} - Z_{:b}\|$ ;
14:      restrict the trust region radius by

$$d_t := \max(d_{\min}, \min(d_{\max}, \gamma_{d_1} d_t));$$

15:      minimize the model to get an improved trust region direction;
16:      while true do
17:        run MLS with improved trust region directions;
18:        if no decrease in  $\tilde{f}$  by MLS, break; end
19:        update  $d_t := (\gamma_{d_2} + \text{rand})d_t$ ;
20:      end while ▷ rand  $\in (0, 1]$  is a random value
21:    else
22:      while true do
23:        perform MLS with perturbed random directions;
24:        if no decrease in  $\tilde{f}$  by MLS, break; end
25:      end while
26:    end if
27:  end if
28:  if no decrease in  $\tilde{f}$  by MLS in all  $T_0$  calls then
29:    for  $i = 1, \dots, n$  do
30:      compute  $\mathbf{dz}_i := Z_{:i} - Z_{:b}$ ; ▷ the difference of the old points
31:      find  $\mathcal{I}_i := \{j \mid \mathbf{dz}_j \neq 0 \text{ and } (Z_{:b})_j \neq 0\}$ ;
32:      compute  $\beta_i^t := \min\{|(Z_{:b})_j / \mathbf{dz}_j| \mid j \in \mathcal{I}_i\}$ ;
33:    end for
34:    generate two random values  $\mu_1$  and  $\mu_2$  satisfying  $0 < \mu_1 < \mu_2 < 1$ ;
35:    reconstruct  $[\underline{\alpha}, \bar{\alpha}] := [\gamma_a \mu_1 \beta_{\min}^t, \gamma_a \mu_2 \beta_{\min}^t]$  with  $\beta_{\min}^t := \min_{i=1, \dots, n} \beta_i^t$ ;
36:  end if
37: end for

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cases (as mentioned in [18] the complexity of deterministic DFO methods is in the worst case by a factor  $n$  worse than that of randomized methods). However, the robustness and efficiency of **VRDFON** are numerically increased. If  $R \sim n$ , then the factors of our bounds in the nonconvex, convex, and strongly convex regions will be large by a factor  $n$ , as in the deterministic case. We are shown numerically in Section 2.4 that the efficiency and robustness of an improved version of **VRDFON** with  $R \sim 1$  are reduced, although our factors are better by a factor  $n$  than in the deterministic case (see BANDEIRA et al. [3] and KIMIAEI & NEUMAIER [19]). When  $R \sim n$  is satisfied, the efficiency and robustness of the implemented version of **VRDFON** are actually increased.

The variable **comBound** takes values of 0, 1, 2 and is a tuning parameter for considering three cases for the complexity results of the implemented version of **VRDFON**. Accordingly, we discuss the conditions under which complexity results can be found for **VRDFON**:

CASE 1 (**comBound** = 0). In this case, random approximate coordinate directions are used and random scaled directions are ignored. Other proposed directions and heuristic techniques are also used. In this case, [18, Proposition 3.1] may not be valid and no complexity result is found for **VRDFON**.

CASE 2 (**comBound** = 1). Both random scaled directions and random approximate coordinate directions are used. Other proposed directions and heuristic techniques are also used. In this case, the order of complexity bounds does not change for all cases after heuristic improvements are used, but only their factors become larger.

CASE 3 (**comBound** = 2). In this case, random scaled directions are used and random approximate coordinate directions are ignored. Other proposed directions and heuristic techniques are also used. The order of complexity bounds does not change for all cases after applying the heuristic improvements, only their factors become larger.

In CASE 2 all theorems for complexity of **VRDFON** in [18] remain valid with the minor changes. The number  $R$  of scaled random steps in each **MLS** must be replaced by the number  $T = \mathcal{O}(C + R)$  of all trial steps. In [18, Proposition 4.5],  $\mathcal{O}(RT_0 \log \omega^{-1})$  is changed to  $\mathcal{O}((R + C)T_0 \log \omega^{-1})$ , which does not change the result given in Corollary 4.6 in [18]. But in CASE 3 the results are unchanged.

## 2 Numerical results

In this section, we provide numerical experiments on a number of test problems of dimensions  $1 < n \leq 300$ . We used all 192 noiseless problems of small dimensions  $1 < n \leq 30$  and all 171 noiseless problems of medium dimensions  $30 < n \leq 300$ . With the different noise levels  $\omega = 10^{-3}, 10^{-2}, 0.1, 0.9$  for small scale problems and  $\omega = 10^{-4}, 10^{-3}, 10^{-2}, 0.1$  for medium scale problems, these produced  $4 \times 192$  small noisy problems and  $4 \times 171$  medium noisy problems. The results were averaged over five runs. We compare **VRDFON** with the 20 state-of-the-art DFO solvers on small and medium scale test problems.

### 2.1 Performance measures

Two important tools for figuring out which solver is **robust** and **efficient** are the data profile of MOREÉ & WILD [23] and the performance profile of DOLAN & MOREÉ [10], respectively.  $\mathcal{S}$  denotes the list of compared solvers and  $\mathcal{P}$  denotes the list of problems. The fraction of problems that the solver  $s$  can solve with  $\kappa$  groups of  $n_p + 1$  function evaluations is the data profile of the solver  $s$ , i.e.,

$$\delta_s(\kappa) := \frac{1}{|\mathcal{P}|} \left| \left\{ p \in \mathcal{P} \mid cr_{p,s} := \frac{c_{p,s}}{n_p + 1} \leq \kappa \right\} \right|. \quad (7)$$

Here  $n_p$  is the dimension of the problem  $p$ ,  $c_{p,s}$  is the **cost measure** of the solver  $s$  to solve the problem  $p$  and  $cr_{p,s}$  is the **cost ratio** of the solver  $s$  to solve the problem  $p$ . The fraction of problems that the performance ratio  $pr_{p,s}$  is at most  $\tau$  is the performance profile of the solver  $s$ , i.e.,

$$\rho_s(\tau) := \frac{1}{|\mathcal{P}|} \left| \left\{ p \in \mathcal{P} \mid pr_{p,s} := \frac{c_{p,s}}{\min(c_{p,\bar{s}} \mid \bar{s} \in S)} \leq \tau \right\} \right|. \quad (8)$$

Note that  $\rho_s(1)$  is the fraction of problems that the solver  $s$  wins compared to the other solvers, while  $\rho_s(\tau)$  ( $\delta_s(\kappa)$ ) is the fraction of problems for sufficiently large  $\tau$  ( $\kappa$ ) that the solver  $s$  can solve. **The data and performance profiles are based on the problem scales, but not on the noise levels. The other two plots are based on the noise levels. These four plots are used to identify the behaviour of the compared solvers with respect to problem scales and noise levels.**

**Efficiency and robustness.** The **efficiency**  $e_{p,s}$  of the solver  $s$  to solve the problem  $p$  is the inverse of the performance ratio  $pr_{p,s}$ . Efficiency measures the ability of a solver  $s \in \mathcal{S}$  relative to an ideal solver. The number of function evaluations is taken as a suitable cost measure, and the efficiency relative to this measure is called the **nf** efficiency. The **robustness** of a solver counts the number of problems it solves.

## 2.2 Codes compared

The VRDFON package is publicly available at [17]. We compare it with:

- VRBBO – an efficient random algorithm by KIMIAEI & NEUMAIER [19]. It can be downloaded from

<https://www.mat.univie.ac.at/~neum/software/VRBBO/>.

- NOMAD (version 3.9.1) , obtained from

<https://www.gerad.ca/nomad>

is a Mesh Adaptive Direct Search algorithm (MADS) [1]. NOMAD1 uses the following option set

```
opts = nomadset('max_eval',nfmax,'max_iterations',2*nfmax,
               'min_mesh_size','1e-008','initial_mesh_size','10')
```

while NOMAD2 uses the following option set

```
opts = nomadset('max_eval',nfmax,'max_iterations',2*nfmax,
               'min_mesh_size','1e-008','initial_mesh_size','10','model_search','0').
```

- SNOBFIT, obtained from

<http://www.mat.univie.ac.at/~neum/software/snobfit>,

is a combination of a branching strategy to enhance the chance of finding a global minimum with a sequential quadratic programming method based on fitted quadratic models to have good local properties by HUYER & NEUMAIER [15].

- **GRID** – a grid algorithm for bound constrained optimization by ELSTER & NEUMAIER [11], available at

<http://www.mat.univie.ac.at/~neum/software/GRID>.

This solver was originally written in Fortran with auxiliary routines that are no longer available. We reimplemented **GRID** in Matlab. The trust region subproblem is solved with `minq8` [16].

- **UOBYQA** and **NEWUOA**, obtained from

<https://www.pdfdocs.net/docs.html>,

are model-based solvers by POWELL [25, 26].

- **BFO** – a trainable stochastic derivative free solver for mixed integer bound-constrained optimization by PORCELLI & TOINT [24], available at

<https://github.com/m01marpor/BFO>.

- **DSPFD** – a direct search Matlab code for derivative-free optimization by GRATTON et al. [12], available at

[pages.discovery.wisc.edu/%7Ecroyer/codes/dspfd\\_sources.zip](http://pages.discovery.wisc.edu/%7Ecroyer/codes/dspfd_sources.zip).

- **MCS** – a deterministic global optimization by a multilevel coordinate search by HUYER & NEUMAIER [14], downloaded from

<https://www.mat.univie.ac.at/~neum/software/mcs/>.

It used the following parameters:

```
iinit = 1; nfMCS = nfmax; smax = 5n + 10; stop = 3n; local = 50;
gamma = eps; hess = ones(n, n); prt = 0.
```

- **BCDFO** – a deterministic model-based trust region algorithm for derivative-free bound constrained minimization by GRATTON et al. [13], obtained from ANKE TROELTZSCH (personal communication).

- **FMINUNC** – a deterministic quasi Newton or trust region algorithm, available at the Matlab Optimization Toolbox at

<https://ch.mathworks.com/help/optim/ug/fminunc.html>.

The following options set was used:

```
opts = optimoptions(@fminunc,'Algorithm','quasi-newton'  
'Display','Iter','MaxIter',Inf,'MaxFunEvals',limits.nfmax  
'TolX',0,'TolFun',0,'ObjectiveLimit',-1e-50);
```

- SDBOX – a derivative-free algorithm for bound constrained optimization problems discussed in [22], downloaded from

<http://www.iasi.cnr.it/~liuzzi/DFL/index.php/list3>.

- CMAES, obtained from

<http://cma.gforge.inria.fr/count-cmaes-m.php?Down=cmaes.m>,

is the stochastic covariance matrix adaptation evolution strategy by AUGER & HANSEN [2]. We used CMAES with the tuning parameters

```
oCMAES.MaxFunEvals = nfmax, oCMAES.DispFinal = 0,  
oCMAES.LogModulo = 0, oCMAES.SaveVariables = 0,  
oCMAES.DispModulo = 0, oCMAES.MaxIter = nfmax,  
oCMAES.Restarts = 7.
```

- LMMAES by LOSHCHILOV et al. [21], fMAES by BEYER [5], BiPopMAES by BEYER & SENDHOFF [6], obtained from

<https://homepages.fhv.at/hgb/downloads.html>,

are three various covariance matrix adaptation evolution strategies.

- subUOBYQA, subNEWUOA and subNMSMAX are UOBYQA, NEWUOA and NMSMAX, respectively, in the random subspace generated by the columns of  $S := 2 * \text{rand}(n, m) - 1$  (in Matlab) to handle problems in medium and high dimensions. Here  $n$  is the dimension of the problem and  $m$  is the subspace dimension. Using these solvers, we recursively minimize the problem  $\tilde{f}(x + \Delta Sz)$  with respect to  $z \in \mathbb{R}^n$ . Let  $\Delta$  be the trust region radius, which is initially one, and let  $0 < \zeta < 1$  be a parameter for the decrease of the function value. If  $\tilde{f}(x) > \tilde{f}(x + \Delta Sz) - \zeta \Delta^2$  holds, the trial point  $x + \Delta Sz$  can be accepted as a new point and  $\Delta$  is expanded to  $\Delta = \max(\Delta, \|Sz\|)$ ; otherwise  $\Delta$  is reduced to  $\Delta = \frac{1}{2} \min(\Delta, \|Sz\|)$ . For each call to these solvers, the maximum number of function evaluations is  $\max(\lceil n/m \rceil, 10m)$ . According to our findings, the best value for the subspace dimension  $m$  is 10. For

$m < 10$  the efficiency and robustness of these subspace solvers decrease when they are used to solve problems in medium and high dimensions, while for  $10 < m \leq 30$  the number of sample points to construct the quadratic models increases, making each call to the original solver (UOBYQA, NEWUOA and NMSMAX) costly.

Unfortunately, software for FDLM by BERAHAS et al. [4] and STRRS by CHEN [7] was not available to us.

To obtain the improved trust region directions discussed in Section 1.6, we solve the trust region subproblem by minq8 [16], available at

<https://www.mat.univie.ac.at/~neum/software/minq/>.

minq8 requires the tuning parameters `minqmax` = 10000 and `minqeps` =  $10^{-8}$ .

## 2.3 Tuning of VRDFON

In this section, we perform a testing and tuning for five important tuning parameters are `comBound` (kind of the complexity results), `model` (model-based?),  $R$  (number of scaled random directions in MLS),  $C$  (number of random approximate coordinate directions), and  $T_0$  (number of times MLS is called by DS). Other tuning parameters were chosen fixed as they did not change the efficiency and robustness of our solver. Accordingly, for a testing and tuning for small scale problems ( $1 < n \leq 30$ ), we consider the 30 versions of VRDFON given in Table 1. Then we chose the four best versions of VRDFON and ran them for medium scale problems ( $31 < n \leq 300$ ) and large scale problems ( $300 < n \leq 5000$ ) to select the best version of VRDFON to compare with the other solvers. Finding the optimal tuning parameters for VRDFON and the other goal, such as the development of an effective mixed integer solver, is still a work in progress [20].

We do not intend to estimate the scaling vector  $s$  in [18]; it is assumed that  $s_i = 1$  for  $i = 1, 2, \dots, n$ .

Table 1 contains the values of the tuning parameters of VRDFON, leading to the best numerical performance for VRDFON shown in Figures 2–4.



$C$	$T_0$	$R$	comBound	model	complexity	chosen as solver
$n$	$n$	—	0	0	no	no
$n$	$n$	—	0	1	no	no
$n$	10	—	0	0	no	no
$n$	10	—	0	1	no	no
$n$	5	—	0	0	no	no
$n$	5	—	0	1	no	no
$n$	2	—	0	0	no	no
$n$	2	—	0	1	no	VRDFON1
10	$n$	—	0	0	no	no
10	$n$	—	0	1	no	no
5	$n$	—	0	0	no	VRDFON2
5	$n$	—	0	1	no	no
2	$n$	—	0	0	no	no
2	$n$	—	0	1	no	no
$\lceil n/2 \rceil$	$n$	$\lceil n/2 \rceil$	1	0	yes	no
$\lceil n/2 \rceil$	$n$	$\lceil n/2 \rceil$	1	1	yes	VRDFON3
—	$n$	$n$	2	0	yes	no
—	$n$	$n$	2	1	yes	no
—	10	$n$	2	0	yes	no
—	10	$n$	2	1	yes	no
—	5	$n$	2	0	yes	no
—	5	$n$	2	1	yes	VRDFON
—	2	$n$	2	0	yes	no
—	2	$n$	2	1	yes	no
—	$n$	10	2	0	yes	no
—	$n$	10	2	1	yes	no
—	$n$	5	2	0	yes	no
—	$n$	5	2	1	yes	no
—	$n$	2	2	0	yes	no
—	$n$	2	2	1	yes	no

Figure 1: 30 variants of tuning of VRDFON

Table 1: The values of tuning parameters of VRDFON

$\bar{m} = 230$	$m_{\max} = \min(0.5n(n+3), \bar{m})$	$\delta_{\min} = 10^{-50}$	$\delta_{\max} = 1$	$\gamma_Z = 100$	$\gamma_{\text{rd}} = 10^{-30}$
$\gamma_{d_1} = 2$	$Q = 1.5$	$d_{\min} = 10^{-4}$	$\gamma_v = 100$	$\gamma = 10^{-6}$	$\gamma_e = 3$
$d_{\max} = 10^3$	$\underline{\alpha}_{\text{init}} = 0.01$	$\bar{\alpha}_{\text{init}} = 0.99$	$\gamma_\kappa = 0.85$	$\gamma_{d_2} = 0.5$	$\gamma_p = 0.25$
$\gamma_a := 10^{-5}$	$\bar{\alpha}_{\min} = 10^{-3} * \text{rand}$	$R = n$	$T_0 = 5$	<b>model</b> = 1	<b>comBound</b> = 2

For the noise levels  $\omega \in \{10^{-4}, 10^{-3}, 10^{-1}, 0.9\}$  and small scale problems ( $1 < n \leq 30$ ), Figure 2 shows in its subfigures the cumulative (over all noise levels used) performance and data profiles in terms of the number of function evaluations and the other two plots show (the **nf** efficiency versus the noise level  $\omega$  and the number of solved problems versus the noise level  $\omega$ ). The result of this comparison is that VRDFON3 is slightly more robust than the others, since all the proposed directions are used. For this version, the complexity results are valid since random scaled directions were tried. VRDFON1 and VRDFON2 are slightly more efficient than VRDFON3 and VRDFON, except for high noise. In fact, using scaled random directions guarantees the existence of complexity results and increases the robustness of our solver under low to high noise and the efficiency of our solver only under high noise. Moreover, since VRDFON1, VRDFON3, and VRDFON are model-based, it is confirmed that VRDFON is effective when it is model-based.

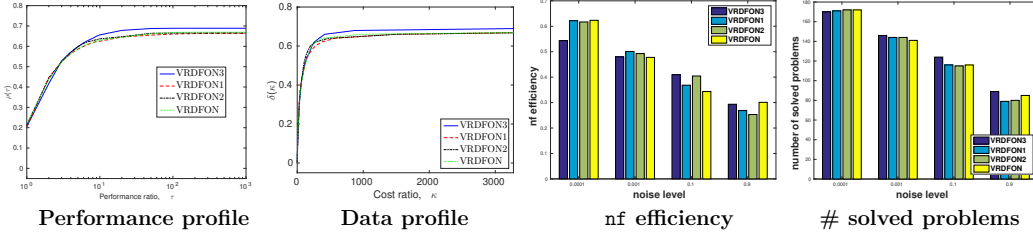


Figure 2: For the noise levels  $\omega \in \{10^{-4}, 10^{-3}, 10^{-1}, 0.9\}$  and small dimensions  $1 < n \leq 30$ . Data profile  $\delta(\kappa)$  in dependence of a bound  $\kappa$  on the cost ratio while performance profile  $\rho(\tau)$  in dependence of a bound  $\tau$  on the performance ratio. Problems solved by no solver are ignored. Here ‘# solved problems’ counts the number of solved problems.

For the noise levels  $\omega \in \{10^{-4}, 10^{-3}, 10^{-2}, 10^{-1}\}$  and medium scale problems ( $30 < n \leq 300$ ), Figure 3 shows in its subfigures the cumulative (over all noise levels used) performance and data profiles in terms of the number of function evaluations and the other two plots show (the **nf** efficiency versus the noise level  $\omega$  and the number of solved problems versus the noise level

$\omega$ ). We conclude from these subfigures that VRDFON1 and VRDFON are slightly more efficient and robust than VRDFON2 and VRDFON3.

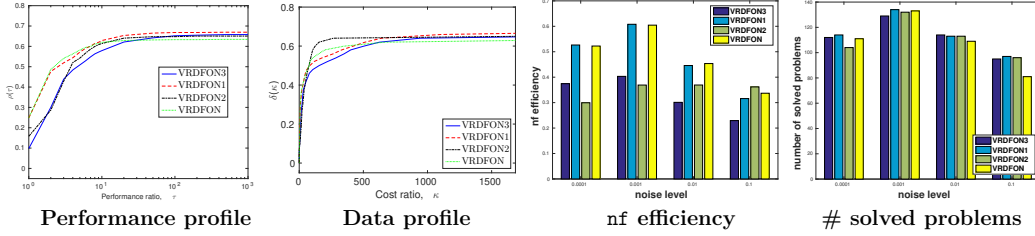


Figure 3: For the noise levels  $\omega \in \{10^{-4}, 10^{-3}, 10^{-2}, 10^{-1}\}$  and medium dimensions  $30 < n \leq 300$ . Other details as in Figure 2.

For the noise levels  $\omega \in \{10^{-5}, 10^{-4}, 10^{-3}\}$  and large scale problems ( $300 < n \leq 5000$ ), Figure 4 shows in its subfigures, the cumulative (over all noise levels used) performance and data profiles in terms of the number of function evaluations and the other two plots show (the nf efficiency versus the noise level  $\omega$  and the number of solved problems versus the noise level  $\omega$ ). We conclude from these subfigures that VRDFON1 and VRDFON are more efficient and robust than VRDFON2 and VRDFON3.

As a result, we choose VRDFON as default version to compare with the other solvers for problems in low to high dimensions. Note that our complexity results hold for this version.

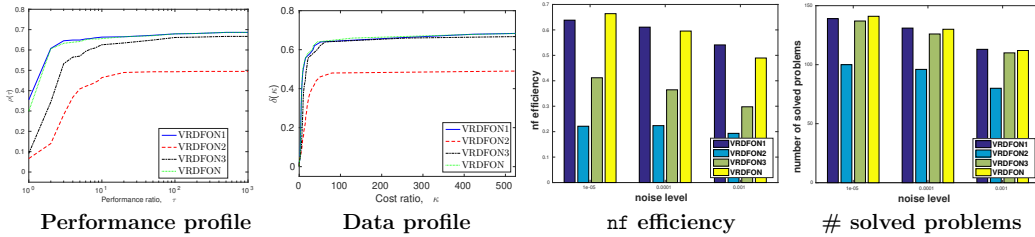


Figure 4: For the noise levels  $\omega \in \{10^{-5}, 10^{-4}, 10^{-3}\}$  and large dimensions  $300 < n \leq 5000$ . Other details as in Figure 2.

## 2.4 Comparison between VRDFON and other DFO solvers

This section compares the default version of VRDFON with the other solvers for problems in low and medium dimensions. In each figure, we display only the best five solvers, but the best four solvers in high dimensions.

### 2.4.1 Small scale: $1 < n \leq 30$

For the noise levels  $\omega \in \{10^{-4}, 10^{-3}, 10^{-1}, 0.9\}$  and small scale problems ( $1 < n \leq 30$ ), this section contains a comparison between VRDFON and the well-known model-based solvers (NOMAD1, UOBYQA, NEWUOA, BCDFO, GRID, and SNOBFIT), the well-known direct search solvers (NOMAD2, NMSMAX, BFO, MCS, DSPFD), the well-known line search solvers (VRBBO, SDBOX, FMINUNC), and the well-known matrix adaptation evolution solvers (CMAES, BiPopMAES, fMAES, LMAES). Moreover, a comparison is made between VRDFON and the five most efficient and robust solvers with both standard and shifted initial points.

To compare our algorithm with the well-known model-based and direct search solvers, Figure 5 shows the cumulative (over all noise levels used) performance and data profiles in its subfigures in terms of the number of function evaluations and the other two plots show (the **nf** efficiency versus the noise level  $\omega$  and the number of solved problems versus the noise level  $\omega$ ). From the subfigures of Figure 5,

- NEWUOA is more efficient than the well-known model-based solvers and VRDFON, while UOBYQA is more robust than others and VRDFON, the third robust solver, is more efficient than SNOBFIT and GRID and less efficient than others. Moreover, NOMAD1 is the second robust solver compared to the well-known model-based solvers;
- VRDFON is the second efficient solver and the first robust solver compared to the well-known direct search solvers, while NMSMAX is the first efficient solver and NOMAD2 is the second robust solver.

To compare our algorithm with the well-known line search and matrix adaptation evolution solvers, Figure 6 shows in its subfigures, the cumulative (over all noise levels used) performance and data profiles in terms of the number of function evaluations and the other two plots show in terms of the noise levels (the **nf** efficiency versus the noise level  $\omega$  and the number of solved problems versus the noise level  $\omega$ ). From the subfigures of Figure 6, we conclude that VRDFON is more robust and efficient than the known line search solvers, while VRDFON is more robust and efficient than the well-known matrix adaptation evolution solvers at low noise and fMAES is more robust and efficient than others at high noise. At low to high noise, VRDFON is slightly more efficient than the well-known matrix adaptation evolution solvers, while fMAES is more robust than others.

Since our solver is model-based and line search-based, we now provides two comparisons between VRDFON and the best model-based (NOMAD1, UOBYQA,

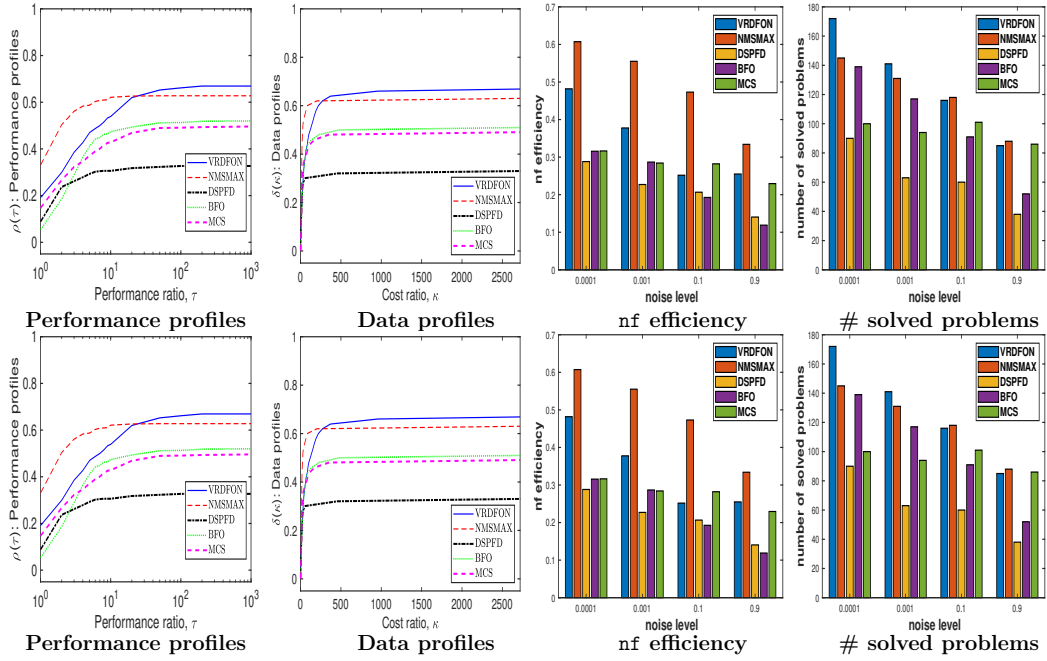


Figure 5: Comparison between VRDFON and model-based solvers (first row) and direct search solvers (second row) for the noise levels  $\omega \in \{10^{-4}, 10^{-3}, 10^{-1}, 0.9\}$  and small dimensions  $1 < n \leq 30$ . Other details as in Figure 2.

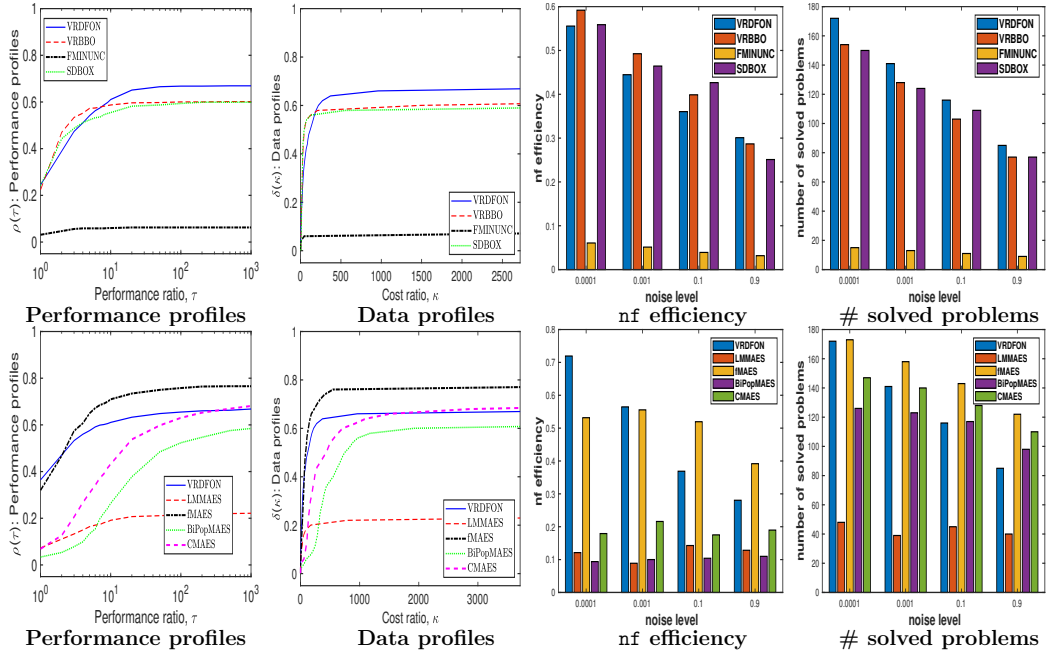


Figure 6: Comparison between VRDFON and line search solvers (first row) and matrix adaptation evolution solvers (second row). Other details as in Figure 2.

NEWUOA, and BCDFO) and line search solvers (SDBOX and VRBBO) with the shifted and initial starting points for small scale problems.

If the default starting points are used, we conclude from Figure 7 that VRDFON is comparable to NEWUOA in terms of efficiency and is more robust than others. It is also more robust than others. In fact, the efficiency and robustness of VRDFON are higher when the standard initial points are used than when the shifted initial points are used, see Figure 8. This is reason why we used the shifted points for all test problems in our comparisons.

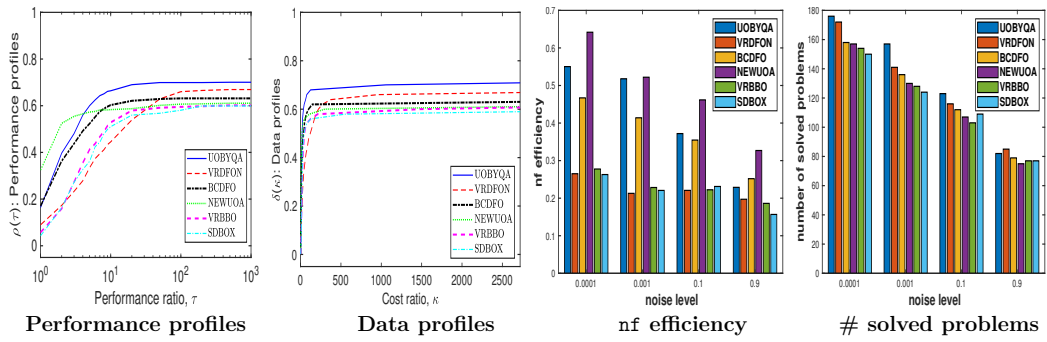


Figure 7: For the noise levels  $\omega \in \{10^{-4}, 10^{-3}, 10^{-1}, 0.9\}$  and small dimensions  $1 < n \leq 30$ . The standard initial points are used. Other details as in Figure 2.

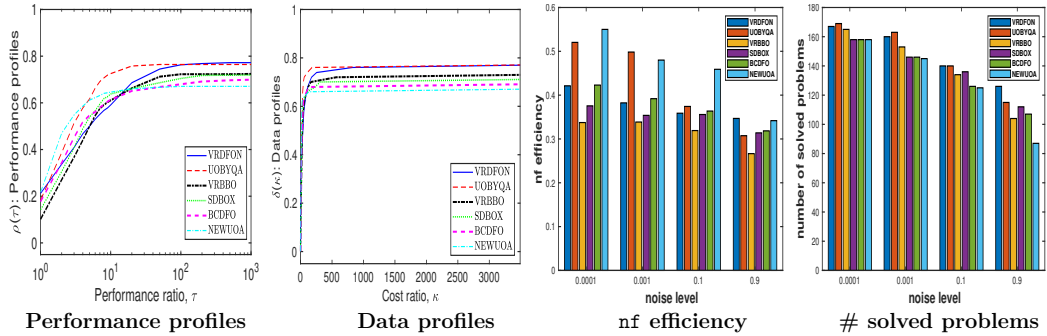


Figure 8: For the noise levels  $\omega \in \{10^{-4}, 10^{-3}, 10^{-1}, 0.9\}$  and small dimensions  $1 < n \leq 30$ . The shifted initial points are used. Other details as in Figure 2.

#### 2.4.2 Medium scale: $30 < n \leq 300$

For the noise levels  $\omega \in \{10^{-4}, 10^{-3}, 10^{-2}, 10^{-1}\}$  and medium scale problems ( $30 < n \leq 300$ ), this section contains a comparison between VRDFON and the

well-known direct search solvers, line search solvers, and matrix adaptation evolution solvers.

Figures 9 and 10 show in their subfigures, the cumulative (over all noise levels used) performance and data profiles in terms of the number of function evaluations and the other two plots show (the **nf** efficiency versus the noise level  $\omega$  and the number of solved problems versus the noise level  $\omega$ ). From the subfigures of Figure 9, we conclude that **VRDFON** is more robust and efficient than the well-known direct search solvers. From the subfigures of Figure 10, we can also conclude that **VRDFON** is more robust and efficient than the well-known line search solvers and matrix adaptation evolution solvers at low and medium noise, while **LMMAES** is more robust than **VRDFON** and the other matrix adaptation evolution solvers at high noise.

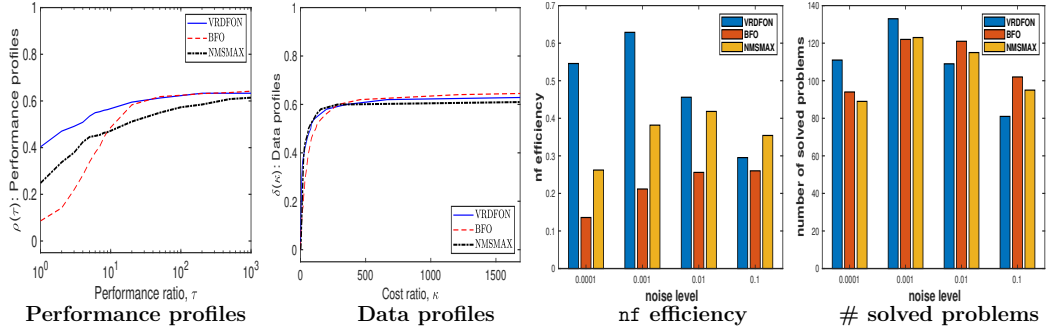


Figure 9: Comparison between **VRDFON** and direct search solvers for the noise levels  $\omega \in \{10^{-4}, 10^{-3}, 10^{-2}, 10^{-1}\}$  and medium dimensions  $30 < n \leq 300$ . Other details as in Figure 2.

Since the model-based solvers cannot handle problems in high dimensions, as described earlier, we denote **UOBYQA** in the random subspace by **subUOBYQA** and **NEWUOA** in the random subspace by **subNEWUOA**. Moreover, we denote **NMSMAX** in the random subspace by **subNMSMAX**.

The subfigures of Figure 11 show that **VRDFON** are much more efficient than others at low to high noise, but **VRDFON** are more robust than others at low noise, while **subNEWUOA**, **subUOBYQA**, and **NMSMAX** are slightly more robust than **VRDFON** only at high noise.



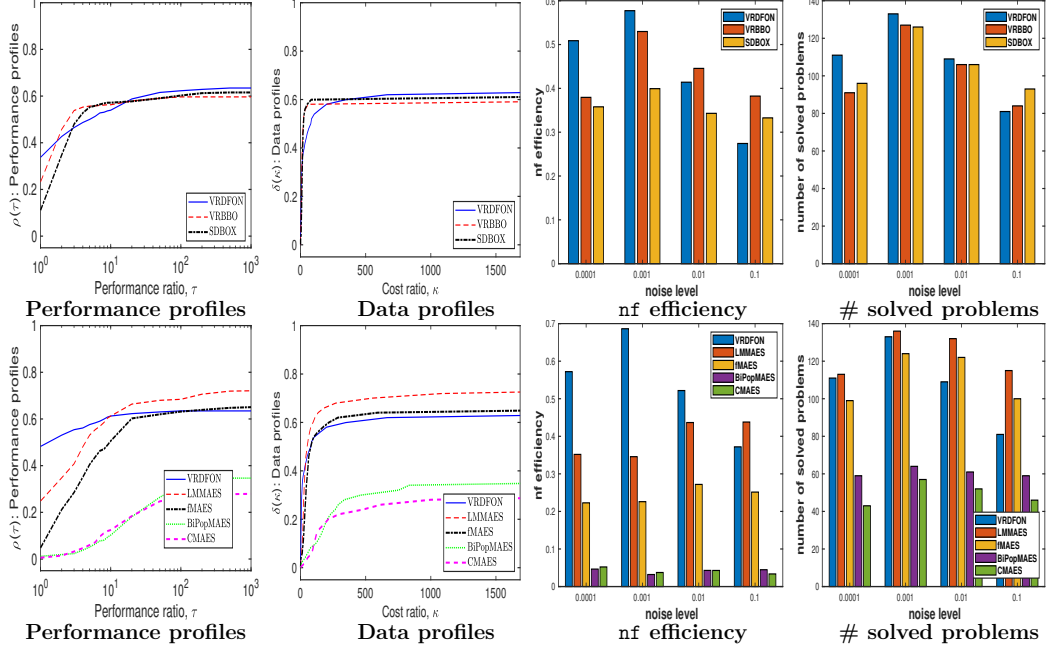


Figure 10: Comparison between VRDFON and line search solvers (first row) and matrix adaptation evolution solvers (second row) for the noise levels  $\omega \in \{10^{-4}, 10^{-3}, 10^{-2}, 10^{-1}\}$  and medium dimensions  $30 < n \leq 300$ . Other details as in Figure 5.

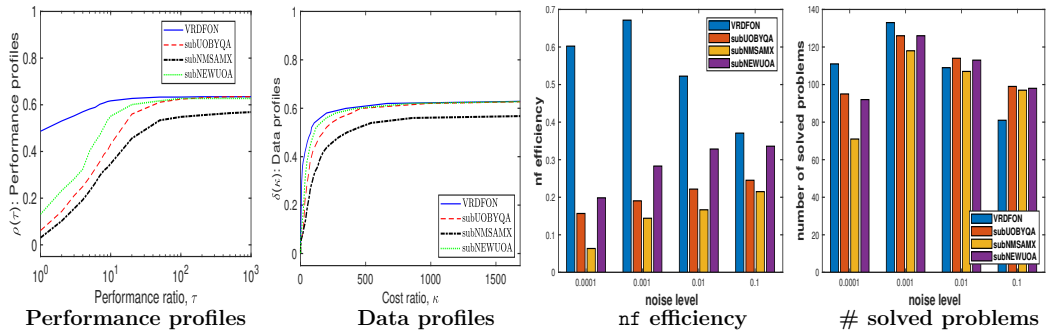


Figure 11: Comparison between VRDFON and model-based solvers in a random subspace for the noise levels  $\omega \in \{10^{-4}, 10^{-3}, 10^{-2}, 10^{-1}\}$  and medium dimensions  $30 < n \leq 300$ . Other details as in Figure 2.

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