

conjugate directions for a quadratic without performing exact searches. For non-quadratic models (See Al-Bayati, 2001).

Powell–Beale Restart Algorithm:

This method is one of the most popular conjugate gradient algorithms, which is used for minimizing nonlinear functions of many variables.

Powell developed a new procedure for restarting CG method. He checked that the new direction d_{k+1} will be sufficiently downhill if:

$$-1.2 \|g_{k+1}\|^2 > g_{k+1}^T d_{k+1} > -0.8 \|g_{k+1}\|^2 \quad \dots(1)$$

(See Powell, 1977).

Powell–Beale Restart Algorithm:

Step (1): for any starting point x_k , $k=1$, ε

Step (2): Set $d_k = -g_k$

Step (3): Compute λ_k by using the line search procedure to minimize $f(x_k + \lambda_k d_k)$

Step (4): $x_{k+1} = x_k + \lambda_k d_k$

Step (5): Check for convergence if $\|g_{k+1}\| < \varepsilon$ Stop, otherwise.

Step(6): if $k > 1$ then

$$d_{k+1} = -g_{k+1} + \frac{g_{k+1}^T y_k}{d_k^T y_k} d_k + \frac{g_{k+1}^T y_k}{d_k^T y_k} d_k$$

where $y_k = g_{k+1} - g_k$

otherwise

$$d_{k+1} = -g_{k+1} + \frac{g_{k+1}^T y_k}{d_k^T y_k} d_k$$

$k = k + 1$

Goto Step (3)

ENDIF

Step (7): Check if $k = n$

or

$$\|g_{k+1}^T g_k\| \geq 0.2 \|g_{k+1}\|^2$$

or

$$-1.2 \|g_{k+1}\|^2 > d_{k+1}^T g_{k+1} > -0.8 \|g_{k+1}\|^2$$

are satisfied then

put $k = 1$ Goto step (2)

otherwise

set $k = k + 1$

Goto step (3)

PRECONDITIONING WITH QUASI-NEWTON APPROXIMATION

Preconditioning is a mean of improving the performance of CG methods.

Effective Preconditioning requires second derivative information which is not available. However, approximate curvature information can be accumulated as it is in a Quasi-Newton method. The storage consideration is temporarily neglected so that $n \times n$ Quasi-Newton approximations may be used. With some conditions on the line search,

and for smooth nonlinear functions, the inverse Hessian approximation matrix H_k are positive definite and symmetric.

In general, a constant preconditioner H must be used throughout to obtain quadratic termination with conjugate gradient method. However, the property of quadratic termination is retained when Quasi-Newton preconditioners are used. In the BFGS updating formula for minimizing a function f (See Fletcher, 1987) we are given a symmetric and positive definite $n \times n$ matrix H_k that approximates the inverse of the Hessian of f , and a pair of n -vectors $v_k = x_{k+1} - x_k$ and $y_k = \nabla f(x_{k+1}) - \nabla f(x_k)$ satisfying the condition $v_k^T y_k > 0$. Using this we compute a new inverse Hessian approximation H_{k+1} by means of the updating formula:

$$H_{k+1} = S_k^T H_k S_k + \rho_k v_k v_k^T \quad \dots(2)$$

where

$$\rho_k = 1/y_k^T v_k \quad S_k = I - \rho_k y_k v_k^T \quad \dots(3)$$

We say that the matrix H_{k+1} is obtained by updating H_k once using the correction pair $\{v_k, y_k\}$.

Even if H_k is sparse, the new BFGS matrix H_{k+1} will generally be dense, so that storing and manipulating it is prohibitive when the number of variables is large. To circumvent this problem, the limited memory approach makes use of an alternative representation of the updating process in which Quasi-Newton matrices are not explicitly formed it follows from (2) and (3) that if an initial matrix H is update m times using the BFGS formula and the m pairs $\{v_i, y_i\}$, $i = k-1, \dots, k-m$, then the resulting matrix $H(m)$ can be written as:

$$\begin{aligned} H(m) &= (S_{k-1}^T \dots S_{k-m}^T) \bar{H} (S_{k-m} \dots S_{k-1}) + \rho_{k-m} (S_{k-1}^T \dots S_{k-m+1}^T) v_{k-m} v_{k-m}^T (S_{k-m+1} \dots S_{k-1}) \\ &+ \rho_{k-m+1} (S_{k-1}^T \dots S_{k-m+1}^T) v_{k-m+1} v_{k-m+1}^T (S_{k-m+2} \dots S_{k-1}) \\ &\vdots \\ &+ \rho_{k-1} v_{k-1} v_{k-1}^T \end{aligned} \quad \dots(4)$$

Thus instead of forming $H(m)$ we can store the scalars ρ_i and the vectors $\{v_i, y_i\}$, $i = k-1, \dots, k-m$ which determine the matrices S_i (See Gill and Murry, 1981).

The so-called BFGS method described in (See Nocedal, 1980; Gilbert and Lemarchal, 1989) updates Hessian approximation as follow. We first choose (usually diagonal) initial Hessian approximation H , and define the first m approximation through (4) as $H(1), \dots, H(m)$. At this stage the storage is full, and to construct the new Hessian approximation, we first delete the oldest correction pair from the set $\{v_i, y_i\}$, to make room for the newest one, $\{v_k, y_k\}$. The new Hessian approximation $H(m+1)$ is defined by (4), using the new set of pair $\{v_i, y_i\}$. This process is repeated during all subsequent iterations the oldest correction is removed to make space for the newest one.

Theorem: Let f be a strictly convex quadratic function and assume the line search is exact.

$$d_0 = H_0 g_0,$$

$$d_{k+1} = -H_k g_{k+1} + \frac{y_k^T H_k g_{k+1}}{y_k^T d_k} d_k, \quad k \geq 0 \quad \dots(5)$$

$$H_{k+1} = H_k - \frac{H_k y_k y_k^T H_k}{y_k^T H_k y_k} + \frac{v_k v_k^T}{v_k^T y_k} + \theta_k w_k w_k^T \quad \dots (6)$$

Where

$$w_k = (y_k^T H_k y_k)^{1/2} \left[\frac{v_k^T}{v_k^T y_k} - \frac{H_k y_k}{y_k^T H_k y_k} \right]$$

Each of H_i , $1 \leq i \leq k$ is obtained from H_{i+1} with $\theta_k \geq 0$.

The initial approximation is $H_0 = H$, a positive-definite and symmetric matrix. Then the direction (5) identical to the conjugate gradient directions preconditioned by H , that is:

$$d_{k+1} = -H_k g_{k+1} + \frac{y_k^T H_k g_{k+1}}{y_k^T d_k} d_k, \quad k \geq 0$$

and thus an algorithm with these search direction has quadratic termination. For more details (see Al-Bayati and Shareaf, 2001).

Nazareth and Nocedal (1978b) also showed that is not necessary to update the Hessian approximation at every iteration in order to have quadratic termination. The search direction (5) can also be used for general nonlinear functions. The matrices H_{k+1} have the property of hereditary positive definiteness as long as the line search guarantees that $v_k^T y_k > 0$.

NEW SAMPLING THREE-TERM PRECONDITION PB-ALGORITHM

We now present a formal description of the sampling algorithm that collects the pair $\{v_k, y_k\}$ as uniformly as possible the sampling algorithm runs parallel to the PCG method once a pair $\{v_k, y_k\}$ has been computed by the PCG method, the sampling algorithm examined the iteration index k and decides if the pair should be included in H (we denote the set of correction pairs that have been stored as H). When a new pair is accepted the algorithms checks the available space, and if the number of pairs in H is m , then a pair is chosen to leave H .

The Algorithm is started by inserting into H the first m pairs generated by the PCG process. After this, the entering and leaving pairs are chosen to keep an almost uniform distribution at any time.

Some properties of the sampling algorithm:

We now discuss some properties of the sampling algorithm. After the initialization in which the first m pairs are stored, the algorithm performs deletion and insertion operations controlled by the variable $N1$. For a given value of $N1$ the algorithm stored $m/2$ new pairs spaced by a distance 2^{N1} , and deletes the same of number of pairs. Deletion takes place in such away that the space created between two consecutive pairs is 2^{N1} . Therefore when $N1$ attains a new value, the distribution cases to be uniform and there is a transition period during which a new uniform distribution is generated: this is

achieved at the end of the second loop. It follows that the larger m is, the longer it will take to move from one uniform distribution to the next (Morales and Nocedal, 1997).

New Algorithm (Sample).

Step (1): for any starting point x_0 , ε , N , M , $k=1$, $N1=1$.

Step (2): Set $d_k = -H_k g_k$

Step (3): Set $ss_k = d_k$. Compute λ_k by using line search procedure to minimize $f(x_k + \lambda_k d_k)$

Step (4): $x_{k+1} = x_k + \lambda_k d_k$

Step (5): Check for convergence if $\|g_{k+1}\| < \varepsilon$ then stop.

Step (6): if $k < M$ then

update H by using Al-Bayati formula.

$$H_{k+1} = H_k - \frac{H_k y_k y_k^T H_k}{y_k^T H_k y_k} + sc \frac{v_k y_k^T}{v_k^T y_k} + w_k w_k^T$$

Where

$$sc = \frac{y_k^T H_k y_k}{y_k^T v_k}$$

When

$$w_k = (y_k^T H_k y_k)^{1/2} \left[\frac{v_k^T}{v_k^T y_k} - \frac{H_k y_k}{y_k^T H_k y_k} \right]$$

then compute

$$d_{kH} = -H_k g_{k+1} + \frac{g_{k+1} H_k y_k^T}{d_k^T y_k} d_k$$

Step (7): if $k = N$

or $g_k^T d_k \geq 0$ then

put $k = 1$ Go to step (2)

Else

$k = k+1$ Go To Step (3)

ENDIF

ENDIF

Step (8): If $k = 1 + (M/2 + L - 1) * 2 ** N1$

then

$k_2 = 1 + (2 * L - 1) * 2 ** (N1 - 1)$

$N1 = N1 + 1$

Where $L = 1, 2, \dots, M/2$

Step (9): If $k \leq k_2$ then

$s_1 = H_k y_k$

Else

$s_1 = -H_k y_k$

ENDIF

Step (10): Update H by

$$\mathbf{H}_{k+1} = \mathbf{H}_k + \frac{\mathbf{s}_1 \mathbf{s}_1^T}{\mathbf{y}_k^T \mathbf{s}_1} + \frac{\mathbf{y}_k^T \mathbf{s}_1}{\mathbf{y}_k^T \mathbf{v}_k} \frac{\mathbf{v}_k \mathbf{v}_k^T}{\mathbf{v}_k^T \mathbf{y}_k} + \mathbf{w}_k \mathbf{w}_k^T$$

$$\mathbf{w}_k = (\mathbf{y}_k^T \mathbf{s}_1)^{1/2} \begin{bmatrix} \mathbf{v}_k^T \\ \mathbf{v}_k^T \mathbf{y}_k - \mathbf{s}_1^T \mathbf{y}_k \end{bmatrix}$$

$$\mathbf{d}_{k+1} = -\mathbf{H}_k \mathbf{g}_{k+1} + \frac{\mathbf{g}_{k+1}^T \mathbf{H}_k \mathbf{y}_k}{\mathbf{d}_k^T \mathbf{y}_k} \mathbf{d}_k + \frac{\mathbf{g}_{k+1}^T \mathbf{y}_k}{\mathbf{d}_k^T \mathbf{y}_k} \mathbf{d}_k$$

Go To step (7).

ENDIF.

Go To step (6).

NUMERICAL RESULTS AND CONCLUSIONS

We shall discuss the computational results of the previous two algorithms (Powell–Beale preconditioned algorithm and the new sampling algorithm).

They are both written in Fortran 77 by using the computer “Pentium I” with double precision. The line search routine employed is the cubic fittings technique, fully described by (Banday, 1984). The stopping criterion is taken to be $\|\mathbf{g}_{k+1}\|_2 < 2 \times 10^{-5}$.

The comparison test involves 50 well-known test functions, by considering both the TOTAL no. of function evaluations and the total no. of iterations. The results are reported in tables (1), (2), (3) with different dimensions.

Table (1) for (low dimensions) $4 \leq n \leq 40$.

Table (2) for (medium dimensions) $42 \leq n \leq 100$.

Table (3) for (high dimensions) $200 \leq n \leq 500$.

Our numerical results indicate, in general, that the new proposed sampling algorithm is always improves the standard preconditioned PB-algorithm.

From Table (1) taking the standard preconditioned PB-algorithm as 100% NOI and NOF the new algorithm improves this standard one in about 50% NOI and 62% NOF for four selected versions namely ($m = 1, 9, n/2$ and n).

Clearly Table (2) deals with medium size test problems and the new algorithm (for all versions) has an improvement of about 56% NOI and NOF.

Finally, our numerical results in Table (3) show that there are an improvement of about 40% NOI and 45% NOF according to our calculations and our selected test functions.

In general, the new algorithm is promising in the filed of unconstrained optimization.

Table 1: Comparison of algorithms for $4 \leq n \leq 40$

n	Test function	Standard PPB NOI (NOF)	New sampling algorithm			
			m=1 NOI (NOF)	M=9 NOI (NOF)	m=n/2 NOI (NOF)	m=n NOI (NOF)
4	Wood	59(187)	13(28)	13(28)	15(36)	13(28)
	Powell	21(72)	17(41)	17(41)	30(62)	20(69)
	Wolf	11(31)	9(19)	9(19)	10(22)	10(22)
	Rosen	22(73)	16(40)	16(40)	9(25)	13(34)
	Dixon	10(32)	10(30)	10(30)	10(31)	10(29)
	Cubic	19(58)	12(31)	12(31)	15(36)	13(34)
8	Wood	82(248)	13(28)	13(28)	16(35)	13(28)
	Powell	29(101)	20(48)	32(66)	34(74)	32(66)
	Wolf	17(39)	15(31)	16(33)	17(41)	16(33)
	Rosen	43(132)	14(36)	15(39)	16(41)	15(39)
	Dixon	16(44)	15(43)	15(43)	17(50)	15(43)
	Cubic	41(168)	11(32)	11(32)	11(31)	11(32)
20	Wood	19(47)	13(28)	13(28)	14(40)	13(28)
	Powell	35(104)	27(62)	22(57)	25(61)	28(66)
	Wolf	27(58)	26(53)	28(72)	28(64)	27(55)
	Rosen	34(89)	14(36)	15(39)	14(38)	14(36)
	Dixon	25(70)	21(65)	21(65)	21(65)	22(68)
	Cubic	44(472)	11(32)	11(32)	11(32)	11(32)
40	Wood	23(55)	13(28)	14(33)	13(28)	13(28)
	Powell	38(105)	28(63)	23(60)	29(65)	28(63)
	Cubic	43(103)	11(32)	12(34)	11(32)	11(32)
Total		658(2288)	329(866)	338(850)	366(909)	358(865)

* PPB stands for preconditioned Powell-Beale algorithm.

Table 2: Comparison of algorithms for $42 \leq n \leq 100$

N	Test function	Standard PPB NOI (NOF)	New sampling algorithm			
			m=1 NOI(NO F)	m=9 NOI(NO F)	m=n/2 NOI(NO F)	m=n NOI(NO F)
42	Rosen	40(97)	14(36)	15(39)	14(38)	14(36)
0	Wood	24(60)	13(28)	14(33)	13(28)	13(28)
	Powell	47(125)	27(62)	34(70)	27(62)	27(62)
	Rosen	52(120)	14(36)	15(43)	14(36)	14(36)
	Dixon	34(98)	23(72)	24(76)	23(72)	23(72)
	Cubic	47(133)	11(32)	12(34)	11(32)	11(32)
80	Wood	23(55)	13(28)	14(30)	13(28)	13(28)
	Powell	48(141)	28(62)	34(72)	28(62)	28(62)
	Wolf	75(153)	41(83)	42(85)	42(86)	41(83)
	Rosen	52(126)	14(36)	15(43)	14(36)	14(36)
	Dixon	34(98)	22(69)	24(82)	22(69)	22(69)
82	Cubic	63(154)	11(32)	12(34)	11(32)	11(32)
100	Powell	52(162)	27(61)	33(73)	27(61)	27(61)
	Wolf	86(175)	42(85)	42(85)	42(85)	42(85)
	Rosen	64(163)	14(36)	15(39)	14(36)	14(36)
	Dixon	34(97)	22(69)	24(88)	22(69)	22(69)
	Cubic	75(199)	11(32)	12(34)	11(32)	11(32)
Total		857(2156)	347(859)	381(960)	348(864)	347(859)

Table 3: Comparison of algorithm for $200 \leq n \leq 500$

n	Test function	Standard PPB NOI (NOF)	New sampling algorithm			
			m=1 NOI(NOF)	m=9 NOI(NOF)	m=n/2 NOI(NOF)	m=n NOI(NOF)
200	Powell	52(149)	29(66)	28(73)	29(66)	29(66)
	Dixon	39(109)	22(68)	22(68)	22(68)	22(68)
	Edgar	7(19)	7(18)	7(18)	7(18)	7(18)
	Wolf	78(168)	43(87)	43(87)	43(87)	43(87)
	Rosen	37(92)	14(36)	15(39)	14(36)	14(36)
300	Powell	47(121)	26(60)	33(68)	26(60)	26(60)
	Dixon	38(107)	22(68)	22(68)	22(68)	22(68)
	Edgar	7(19)	7(18)	7(18)	7(18)	7(18)
400	Powell	49(133)	27(63)	28(76)	27(63)	27(63)
	Dixon	38(104)	22(68)	23(73)	22(68)	22(68)
500	Edgar	7(19)	7(18)	7(18)	7(18)	7(18)
Total		406(1059)	233(588)	242(624)	233(588)	233(588)

- PPB stands for preconditioned Powell-Beale algorithm.

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