

A parallelization technique applied to the computation of rotating flows in cavities

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Abstract—

In this paper, we investigate the parallel solution to the rotating internal flow problems, using the Navier-Stokes equations as proposed in [ST83] and [SPE85]. A Runge-Kutta time-stepping scheme was applied to the equations and both sequential and message-passing implementations were developed, the latter using MPI, and were tested on an SGI Origin200 distributed, global shared memory parallel computer. The results show that our approach to parallelize the sequential implementation requires little effort whilst providing good results even for medium-sized problems, on this particular computer.

Keywords— Parallel computing; Message-passing; Computational Fluid Dynamics.

I. INTRODUCTION

Speziale and Thangam [ST83], and Speziale [SPE85] have developed a formulation for the problem of rotating internal flows, i.e. determining the behaviour of “pressure-driven laminar flows in straight ducts, subjected to a steady spanwise rotation” (see [SPE85]). The governing equations are the Navier-Stokes equations and the continuity equation in a rotating framework, which can be written as follows

$$\frac{\partial \vec{v}}{\partial t} + \vec{v} \cdot \nabla \vec{v} = -\frac{1}{\rho} \nabla P + \nu \nabla^2 \vec{v} - 2\Omega \times \vec{v} \quad (1)$$

$$\nabla \cdot \vec{v} = 0 \quad (2)$$

where \vec{v} is the velocity vector, P is the modified pressure which includes both the gravitational and centrifugal force potentials, Ω is the steady spanwise rotation, ρ is the density of the fluid and ν is the kinematic viscosity of the fluid. The axial pressure gradient $\partial P / \partial z = -G$ is constant.

For a nonzero rotation rate, the velocity vector is of the form $\vec{v} = u(x, y)\vec{i} + v(x, y)\vec{j} + w(x, y)\vec{k}$, w being the axial velocity and u and v representing the secondary flow. As the rotation is around \vec{j} , it is of the form $\Omega = \Omega \vec{j}$; since the flow properties are independent of z , equations (1) and (2) may be written in component form as

$$\frac{\partial u}{\partial t} + u \frac{\partial u}{\partial x} + v \frac{\partial u}{\partial y} = -\frac{1}{\rho} \frac{\partial P}{\partial x} + \nu \nabla^2 u - 2\Omega w \quad (3)$$

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$$\frac{\partial v}{\partial t} + u \frac{\partial v}{\partial x} + v \frac{\partial v}{\partial y} = -\frac{1}{\rho} \frac{\partial P}{\partial y} + \nu \nabla^2 v \quad (4)$$

$$\frac{\partial w}{\partial t} + u \frac{\partial w}{\partial x} + v \frac{\partial w}{\partial y} = \frac{G}{\rho} + \nu \nabla^2 w + 2\Omega u \quad (5)$$

$$\frac{\partial u}{\partial x} + \frac{\partial v}{\partial y} = 0 \quad (6)$$

Due to the simple form (6) of the continuity equation, it follows that a secondary flow stream function ψ exists such that the velocity components are

$$u = -\frac{\partial \psi}{\partial y}, \quad v = \frac{\partial \psi}{\partial x} \quad (7)$$

The function ψ is the solution to the Poisson equation $\nabla^2 \psi = \partial v / \partial x - \partial u / \partial y = \zeta$, where ζ is the axial component of the velocity vector and is expressed by

$$\frac{\partial \zeta}{\partial t} + u \frac{\partial \zeta}{\partial x} + v \frac{\partial \zeta}{\partial y} = \nu \nabla^2 \zeta + 2\Omega \frac{\partial w}{\partial y} \quad (8)$$

Introducing the velocity and length scales W_0 and D , the equations to be solved numerically are written as follows:

$$\frac{\partial w}{\partial t} + u \frac{\partial w}{\partial x} + v \frac{\partial w}{\partial y} = C + \frac{1}{Re} \nabla^2 w + 2Ro u \quad (9)$$

$$\frac{\partial \zeta}{\partial t} + u \frac{\partial \zeta}{\partial x} + v \frac{\partial \zeta}{\partial y} = \frac{1}{Re} \nabla^2 \zeta + 2Ro \frac{\partial w}{\partial y} \quad (10)$$

$$\nabla^2 \psi = \zeta \quad (11)$$

$$u = -\frac{\partial \psi}{\partial y}, \quad v = \frac{\partial \psi}{\partial x} \quad (12)$$

where Re and Ro are the Reynolds and rotation numbers, C is the dimensionless pressure gradient, and the following relationships hold:

$$Re = W_0 D / \nu, \quad Ro = \Omega D / W_0, \quad C = GD / \rho W_0^2. \quad (13)$$

The initial condition for w , i.e. a non-rotating flow, satisfies the equation

$$\nabla^2 w = -\frac{G}{\rho \nu} \quad (14)$$

The reader is referred to [SPE85] for more details.

The set of equations (9)-(12) are to be computed numerically and a rectangular grid of $M \times N$ points, $M > N$ are placed over a duct with length D and height H ($H > D$), with spacings $h_x = D/(M + 1)$ and $h_y = H/(N + 1)$; therefore we may refer to the variables of interest in those equations for an specific time-step t as discrete points on the grid with indices i, j along the vertical and horizontal directions respectively.

Boundary conditions are $u = 0, v = 0, w = 0$ and $\psi = 0$ on the walls of the duct; also, by taking a Taylor's expansion of (11), boundary conditions on the axial vorticity at time-step t are obtained and written as

$$\zeta_{i,0}^t = \frac{8\psi_{i,1}^t - \psi_{i,2}^t}{2h_x^2}, \quad \zeta_{i,M+1}^t = \frac{8\psi_{i,M+1}^t - \psi_{i,M}^t}{2h_x^2} \quad (15)$$

$$\zeta_{0,j}^t = \frac{8\psi_{1,j}^t - \psi_{2,j}^t}{2h_y^2}, \quad \zeta_{N+1,j}^t = \frac{8\psi_{N+1,j}^t - \psi_{N,j}^t}{2h_y^2} \quad (16)$$

In this work, we have used a modified explicit Runge-Kutta time-stepping integration scheme, approximating time and spatial derivatives by forward and central finite-differences respectively. Other authors (see [PAF⁺97], [SOS97], [KT97]) have used different approaches for fluid dynamics simulations, mainly finite-element discretizations and implicit time-stepping integration schemes, and the parallelization is expressed via domain-decomposition.

II. DESCRIPTION OF THE EXPLICIT METHOD

The computation is divided in two main parts. Initially, we solve Equation (14) for w with an iterative method, writing the Laplacian in central finite-differences and obtaining the value of w at the (i, j) cell from

$$w_{i,j}^{k+1} = \delta \left(h_x^2 h_y^2 \frac{G}{\rho\nu} + h_y^2 (w_{i,j-1}^k + w_{i,j+1}^k) + h_x^2 (w_{i-1,j}^k + w_{i+1,j}^k) \right) \quad (17)$$

where $\delta = (2h_x^2 + 2h_y^2)^{-1}$, $i = 1, 2, \dots, N$, $j = 1, 2, \dots, M$, $k = 0, 1, \dots, k_{max}$, and we proceed with the iterations until

$$\|w^{k+1} - w^k\|_\infty < \epsilon_w \quad (18)$$

where ϵ_w is sufficiently small.

The second part is the solution of equations (9)-(12) which is made using a modified explicit Runge-Kutta time-stepping scheme [JST81], [DIC96], [BOR94]. We proceed from time-step t to $t + 1$ as follows:

$$w_{i,j}^0 = w_{i,j}^t, \zeta_{i,j}^0 = \zeta_{i,j}^t, u_{i,j}^0 = u_{i,j}^t, v_{i,j}^0 = v_{i,j}^t \quad (19)$$

$$w_{i,j}^k = w_{i,j}^0 + \alpha_k h_t \left(C + \frac{1}{Re} \nabla^2 w_{i,j}^{k-1} + 2Ro u_{i,j}^{k-1} - u_{i,j}^{k-1} \frac{\partial w_{i,j}^{k-1}}{\partial x} - v_{i,j}^{k-1} \frac{\partial w_{i,j}^{k-1}}{\partial y} \right) \quad (20)$$

$$\zeta_{i,j}^k = \zeta_{i,j}^0 + \alpha_k h_t \left(\frac{1}{Re} \nabla^2 \zeta_{i,j}^{k-1} + 2Ro \frac{\partial w_{i,j}^k}{\partial y} - u_{i,j}^{k-1} \frac{\partial \zeta_{i,j}^{k-1}}{\partial x} - v_{i,j}^{k-1} \frac{\partial \zeta_{i,j}^{k-1}}{\partial y} \right) \quad (21)$$

$$\nabla^2 \psi^k = \zeta^k \quad (22)$$

$$u_{i,j}^k = -\frac{\partial \psi_{i,j}^k}{\partial y}, \quad v_{i,j}^k = \frac{\partial \psi_{i,j}^k}{\partial x} \quad (23)$$

$$w_{i,j}^{t+1} = w_{i,j}^K, \zeta_{i,j}^{t+1} = \zeta_{i,j}^K, u_{i,j}^{t+1} = u_{i,j}^K, v_{i,j}^{t+1} = v_{i,j}^K \quad (24)$$

where h_t is the time-step length, $k = 1, 2, \dots, K$ (K is the order of the Runge-Kutta scheme) and $\alpha^{(K)} = \{\alpha_k\}_{k=1}^K$ is the set of weights for the integration, with $\alpha^{(3)} = \{1/2, 1/2, 1\}$, $\alpha^{(4)} = \{1/4, 1/3, 1/2, 1\}$ and $\alpha^{(5)} = \{1/4, 1/6, 3/8, 1/2, 1\}$ (see [DIC96]). All derivatives appearing in equations (20)-(23) are replaced by central finite-differences.

To stabilize the computation, the time-step h_t is chosen such that at each iteration it satisfies the condition

$$h_t \leq (2\nu(h_x^{-2} + h_y^{-2}) + \|u\|_\infty/h_x + \|v\|_\infty/h_y)^{-1} \quad (25)$$

Now the iterations in t proceed until

$$\|\Delta(w)\|_\infty + \|\Delta(\zeta)\|_\infty + \|\Delta(\psi)\|_\infty + \|\Delta(u)\|_\infty + \|\Delta(v)\|_\infty < \epsilon \quad (26)$$

where $\Delta(f)$ denotes $f^{t+1} - f^t$ and ϵ is sufficiently small. In case this tolerance has not been achieved, the boundary conditions on the axial vorticity ζ are updated, using (15) and (16), and another iteration is performed. The algorithm used is outlined below:

Algorithm II.1: Rotating flow algorithm

1. initialize constants, boundary conditions, etc.
- for** $k = 1, 2, \dots, k_{max}$
2. compute (17)
3. **if** $\|w^{k+1} - w^k\|_\infty < \epsilon_w$
- then break;**
- endfor**
- for** $t = 0, 1, \dots, t_{max}$
- for** $k = 1, 2, \dots, K$
4. compute (20)-(23)
- endfor**
5. **if** $\|\Delta(w)\|_\infty + \|\Delta(\zeta)\|_\infty + \|\Delta(\psi)\|_\infty + \|\Delta(u)\|_\infty + \|\Delta(v)\|_\infty < \epsilon$

- then break;
6. update boundary conditions on ζ using equations (15)-(16)
 7. correct h_t using equation (25)
- endfor
8. output the results.

III. PARALLELIZATION OF THE METHOD

The parallelization of an explicit method such as that described in the previous section requires a careful analysis of the equations in use in order to ascertain the relationships between the variables involved, since this will determine the flow of data in the code and between the many processors collaborating in the parallel computation.

The parallel algorithm developed is based on the *single-program, multiple-data, SPMD* paradigm and we consider that the number of processors available, p , is less than the number of computational cells ($M \times N$). No assumption is made with regard to as how the processors are interconnected, though both *point-to-point* communication and *reduction* operators are supposed to be available.

To partition the grid among the processors, we consider that since the domain is regular, the only major requirement to attend is that we must partition it across the largest dimension, thereby increasing the computation-communication ratio and leading to a potential good parallel performance. In our case, we divide the domain across M (as $M > N$), obtaining $m = M/p$ panels of N cells, and assign to each one of p processors $m \times N$ contiguous cells; if M is not an integer multiple of p , then one extra row of N cells is assigned to some r processors, where r is the modulus of M/p (i.e. these processors will store panels of size $(m + 1) \times N$). This partitioning leads to a *logical* interconnection of the processors as if they were on a linear array (a topology which can be easily embedded on other physical interconnections available in parallel computers, like hypercubes, 2D/3D grids and others).

While other strategies could be followed to achieve load-balance between the processors, this one makes the communication pattern regular, as each processor has to exchange at most two rows of N cells with its two neighbours (or a single row if it is at one of the ends of the linear array).

It should be stressed that though the partitioning by panels is very simple, it can be used on a variety of other problems, including those involving complex geometries, if the problem is recast using generalized coordinates and type *C*- and *O*-grids (see [FLE91, V.2, Ch. 12] and [FP96]).

Analysing the flow of data between equations (20)-(23), we note that there is a feedback mechanism in the overall Runge-Kutta scheme, as once a variable in the k -th step is produced, it is used in the computation of the next variable in sequence. This mechanism implies the need of data exchange

between the processors inside the Runge-Kutta scheme, in order to compute the finite-differences approximations to the derivatives.

Thus, once every processor has computed w^k in their assigned portion of cells using (20), they swap their left- and right-most columns of cells of w^k (say, $w_{:,1}$ and $w_{:,m}$, where the colon indicates a whole column) and also of ζ^{k-1} with their left and right neighbours (this is done in a single message of length $2N$ instead of two messages of length N to reduce the effect of message-passing latencies in the performance of the algorithm). Every processor is then able to compute ζ^k in their cells; afterwards, they exchange ζ^k in the same way in order to solve Poisson's equation for ψ^k and once this is completed, ψ^k is exchanged in order to compute the velocities u^k and v^k . Therefore in every step of the Runge-Kutta scheme there are three data-exchanges between neighbouring processors.

Also, note that the initial condition for w is the solution of Equation (14); as it is solved in the form (17), the computation can be organized such that one data exchange is suppressed, since for the first iterations (i.e. for $k = 0$ at $t = 0$), a processor will have already received columns $w_{:,m}$ and $w_{:,1}$ from its left and right neighbouring processors, which has been done in the last iteration prior to convergence using (17).

The boundary conditions on u, v, w, ψ and ζ can be computed without any communication due to their simple form. For the update of the boundary conditions on ζ – equations (15) and (16) – we use the same approach as explained in the previous paragraph, since every processor will have stored $\zeta_{:,m}^{t+1}$ and $\zeta_{:,1}^{t+1}$ of its left and right neighbours, from the last Runge-Kutta iteration.

The whole algorithm is organized by dividing the computation of equations (20)-(23) into two parts: one that refers to data stored locally in a single processor, and another which depends on the local availability of data stored in its neighbouring processors. If now we make use of *asynchronous* point-to-point communications (as present in MPI [MPI]), then we can compute any one of the variables involved using the following algorithm:

Algorithm III.1: Parallel computation of a variable f

1. asynchronously send variable f to its left and right neighbours
2. compute variable f with its local data, i.e. from columns 2 to $m - 1$
3. request the m -th column of f from processor $p - 1$ (left) and store locally into column 0 of f
4. request the 1-st column of f from processor $p + 1$ (right) and store locally into column $m + 1$ of f
5. compute columns 1 and m of variable f

It is then possible to almost completely hide the time

spent communicating between two processors, provided the amount of time spent in step 2 of the above algorithm is greater than the time needed for the two point-to-point communications between a processor and its two neighbours. A sufficiently large grid will allow this to happen; in our specific case, we are interested when the ratio M/N is large, since that will *maximize* the amount of local computation for a given p while keeping small (relative to the local computation) the communication time. This approach has been successfully applied in other parallel applications (see [CH93], [CH94], [CH95] and [CUN97]).

However, there are two penalties brought about by the parallel computation of equations (20)-(23):

1. The amount of time needed to set-up the asynchronous sends and the retrieval of data from the local communications buffer into the appropriate memory locations of the user's program;
2. The computation of *reductions*, needed to obtain the norms used in the stopping criteria of the iterations.

We can not hide these times within the local computation time and therefore they are the main causes for being unable to achieve the optimal speed-up; but we may expect that by dividing the storage of the variables among p processors, the use of processors with cache memories will provide some interesting phenomenon for large size problems and small number of processors.

The parallel algorithm can now be described as follows. Each processor stores its m columns of N cells, for each variable (w, ζ, ψ, u and v) in arrays of size $0 : (N+1), 0 : (m+1)$, where the two extra rows and columns serve to hold the boundary conditions values. Due to the simple form of some of the boundary conditions specified, one could argue that it is not needed to store them; however this would lead to a specific piece of code be written to compute the equations in the cells where the boundary conditions are involved.

We wrote the code to compute each of the equations (20)-(23) as a pair of loops scanning the columns and rows of the array holding the variable values at the cells. As an example, we will show how a sequential code to compute $\psi_{i,j}^k$ in Equation (20) was transformed into a parallel code according to Algorithm III.1. The sequential code is as follows

```
DO J = 1, M
  DO I = 1, N
    DXPSI = PSI(I, J-1) + PSI(I, J+1)
    DYPSI = PSI(I-1, J) + PSI(I+1, J)
    PSINEW(I, J) = APSI * ZETANEW(I, J) +
+               BPSI * DXPSI +
+               CPSI * DYPSI
  END DO
END DO
```

where APSI, BPSI and CPSI are constants involving h_x and h_y , derived from the central finite-differences equations. Its equivalent parallel version, using MPI, is

```
* 1. Asynchronously send PSI to its left and
```

```
*   right neighbours
CALL SNDRCV(MYID, P, PSI, NP1, MP1, 900, 1000,
+          IDSND, IDRCV)

* 2. Compute PSI with its local data
DO J = 2, MYM-1
  DO I = 1, N
    DXPSI = PSI(I, J-1) + PSI(I, J+1)
    DYPSI = PSI(I-1, J) + PSI(I+1, J)
    PSINEW(I, J) = APSI * ZETANEW(I, J) +
+               BPSI * DXPSI +
+               CPSI * DYPSI
  END DO
END DO

* 3-4. Request columns from neighbouring
*       processors
CALL GETDATA(MYID, NPROCS, IDSND, IDRCV)

* 5. Compute columns 1 and m of PSI
DO J = 1, MYM, MYM-1
  DO I = 1, N
    DXPSI = PSI(I, J-1) + PSI(I, J+1)
    DYPSI = PSI(I-1, J) + PSI(I+1, J)
    PSINEW(I, J) = APSI * ZETANEW(I, J) +
+               BPSI * DXPSI +
+               CPSI * DYPSI
  END DO
END DO
```

where MYM is m and SNDRCV and GETDATA are subroutines which call the MPI routines MPI_ISEND and MPI_IRECV, and MPI_WAIT respectively. Note that by using the MPI_ISEND and MPI_IRECV routines we have an asynchronous parallel implementation which maximizes the use of the processors. A fully asynchronous implementation, on the other hand, is not possible, for the underlying numerical method can not cope with the nonlinear instabilities that may be generated by that kind of implementation.

With this approach, once a sequential version of the code has been tested and certified to be producing the desired results, it is easy to obtain its parallel version, since the second pair of DO loops is the same as the first, apart from the indices on J. It is less error-prone, since the loop body remains unchanged; in fact, if the first pair is encapsulated in a subroutine, having the indices on J as parameters, if a modification in the body of the loops was required, then just a single part of the code would need attention. As for the performance of such code, if one uses a compiler which is capable of inlining a subroutine, then it will not be affected by this approach.

Another possible way of writing the parallel code (which we have also done) would be to provide three different parts to handle the computation, depending on the position of each processor: the first, the last, and those in the middle of the linear array. It is easy to see that this would increase three-fold the size of the code, and make it even more difficult to maintain; one could make use of subroutines which would certainly make the code more readable but, for an efficient program execution, the subroutines should be inlined, thereby increasing the object code size accordingly. As an example,

with the first approach, the ratio of source code sizes of the parallel to the sequential versions is 1.46 : 1, whereas for this latter approach it was 3.28 : 1.

The other modification required in the sequential code to produce the parallel version is in the computation of the norms. This requires a reduction operation over several values (i.e. the partial norms) stored in the processors. Due to the *SPMD* programming model used, the reduced value (i.e. the ∞ -norm of a variable) is required to be present in every processor. Therefore, a reduction, followed by a *broadcast* of the reduced value to all processors is employed, this being implemented by the MPI routine `MPI_ALLREDUCE`. It is a costly operation; note that the reduction and broadcast require sending/receiving several messages between the co-operating processors (with the associated latencies to set-up the message transfers), albeit some of those may be done in parallel.

As such, we look at Algorithm II.1 and notice that norms are required in steps 3, 5 and 7. Now we ask ourselves: can we combine the reductions in the last two steps into a single one, therefore reducing the latencies? If we consider that the computational cost of a reduction of r values followed by a broadcast is $2\lfloor \log_2 p \rfloor (\alpha + r\beta)$, where α and β are the latency and the transfer rate between two processors (directly related to each other), then it is easy to see that if we combine the five reductions needed in step 5 with the two reductions in step 7, we will be saving one latency per reduction. For the overall computation, we will have

$$2t_{\max}(\alpha \lfloor \log_2 p \rfloor + 7\beta) < 2t_{\max}(2\alpha \lfloor \log_2 p \rfloor + 7\beta) \quad (27)$$

and the savings will be greater for large p , as we shall see in §IV-C.

With the above reasoning, if the sequential code corresponding to the computation of steps 5 and 7 is written as

```
* 5. Compute norms
DO I = 1,5
  NORMS(I) = 0.0
END DO
DO J = 1,M
  DO I = 1,N
    NORMS(1) = MAX(NORMS(1), ABS(W(I,J) -
+                               WNEW(I,J)))
+   NORMS(2) = MAX(NORMS(2), ABS(ZETA(I,J) -
+                               ZETANEW(I,J)))
+   NORMS(3) = MAX(NORMS(3), ABS(PSI(I,J) -
+                               PSINEW(I,J)))
+   NORMS(4) = MAX(NORMS(4), ABS(U(I,J) -
+                               UNEW(I,J)))
+   NORMS(5) = MAX(NORMS(5), ABS(V(I,J) -
+                               VNEW(I,J)))
  END DO
END DO

NORM = NORMS(1) + NORMS(2) + NORMS(3) +
+      NORMS(4) + NORMS(5)

* 6. Update boundary conditions on ZETA
```

...

```
* 7. Time-step stabilization test
NORMS(1) = 0.0
NORMS(2) = 0.0
DO J = 1,M
  DO I = 1,N
    NORMS(1) = MAX(NORMS(1), ABS(UNEW(I,J)))
    NORMS(2) = MAX(NORMS(2), ABS(VNEW(I,J)))
  END DO
END DO
MAXHT = 2.0*NU*(INVHXSQ+INVHYSQ) +
+       INVHX*NORMS(1) + INVHY*NORMS(2)
HT = MIN(HT, 1.0/MAXHT)
```

then an equivalent parallel code, including the computation of the $\|u\|_\infty$, $\|v\|_\infty$ needed in step 7, is

```
* 5. Compute norms
DO I = 1,7
  NORMS(I) = 0.0
END DO
DO J = 1,MYM
  DO I = 1,N
    NORMS(1) = MAX(NORMS(1), ABS(W(I,J) -
+                               WNEW(I,J)))
+   NORMS(2) = MAX(NORMS(2), ABS(ZETA(I,J) -
+                               ZETANEW(I,J)))
+   NORMS(3) = MAX(NORMS(3), ABS(PSI(I,J) -
+                               PSINEW(I,J)))
+   NORMS(4) = MAX(NORMS(4), ABS(U(I,J) -
+                               UNEW(I,J)))
+   NORMS(5) = MAX(NORMS(5), ABS(V(I,J) -
+                               VNEW(I,J)))
+   NORMS(6) = MAX(NORMS(6), ABS(UNEW(I,J)))
+   NORMS(7) = MAX(NORMS(7), ABS(VNEW(I,J)))
  END DO
END DO

CALL MPI_ALLREDUCE(NORMS, REDUOUT, 7, MPI_REAL,
+                 MPI_MAX, MPI_COMM_WORLD, IERR)

NORM = REDUOUT(1) + REDUOUT(2) + REDUOUT(3) +
+      REDUOUT(4) + REDUOUT(5)

* 6. Update boundary conditions on ZETA
...

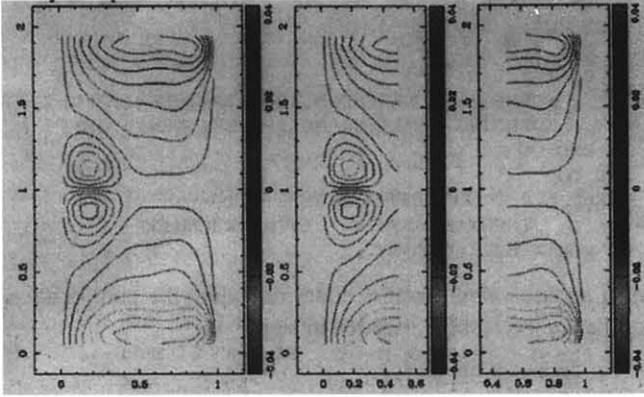
* 7. Time-step stabilization test
MAXHT = 2.0*NU*(INVHXSQ+INVHYSQ) +
+       INVHX*REDUOUT(6) + INVHY*REDUOUT(7)
HT = MIN(HT, 1.0/MAXHT)
```

where `REDUOUT` is the buffer holding the reduced `NORMS` values and which is present in all processors after the call to `MPI_ALLREDUCE`.

A. Output of the results

At the end of the overall computation, we save the values of the variables involved in files for later analysis. We consider that each processor has parallel access to the disk filesystem and each processor is thus able to open its own file, all p files being written as simultaneously as possible. In our experiments, even for the large problems, this proved to be efficient and accounted for less than 1% of the run-time. An in-house

Fig. 1. Typical results for a 64×32 mesh, $Re = 279$. From left to right: the contour streamlines for the complete domain and the contours generated by each processor.



developed visualization program (see [JUS98]) is later used, which opens the several files in sequence and exhibits the data in a variety of forms (eg. colour maps, particle traces and vector fields).

IV. EXPERIMENTS

A number of experiments were carried out on Silicon Graphics Origin200 distributed, global access memory parallel computer located at the Brazilian National Supercomputing Centre. Our software is a FORTRAN 77 code which implements Algorithm II.1 with the parallelization expressed as in Algorithm III.1. All computations were carried out in single-precision (32 bits) and with the computer in dedicated mode.

The parallel computer used is a four-processor machine in a twin-tower configuration, each tower equipped with two MIPS R10000, 180MHz processors with 1MB cache memory each, and interconnected via a CrayLink™ cable. It has an aggregate RAM memory size of 256MBytes. The machine is a “scalable, shared-memory processor (S2MP)” and it has a hierarchical memory, with increasing memory access time for data requested from farther processors. It is interconnected like a hypercube, with the use of CrayLink cables and routers. For an Origin computer with 16 and 32 processors, XpressLink™ interconnects are added to the interconnection network, making use of the spare ports on the routers, minimizing latency and increasing the bandwidth. Nonetheless, the fact remains that this machine does not have a constant latency and transfer rate between any pair of processors.

A. Typical results

With regards to the flow problem itself, typical results that were obtained are shown in Figure 1, which shows the streamlines for ψ . In that experiment, taken from [SPE85, p. 272], $Re = 279$, $Ro = 0.833$, $\Omega = 0.1 \text{ rad/s}$, $G = 6 \times 10^{-4} \text{ lb/ft}^3$

and $h_t = 10^{-4}$ (throughout the iterations). The tolerance for convergence for the initial condition on w was 10^{-5} and it was achieved in 2,595 iterations, taking 0.0008s on two processors. Convergence of the solution of the Navier-Stokes equations using the three-term Runge-Kutta scheme ($K = 3$) for a tolerance of 10^{-4} , took 406.9987s after 202,255 iterations. The figures show a similar appearance to that presented in [SPE85, p. 272].

B. Scalability

TABLE I
RUN TIME (IN SECONDS PER ITERATION) AND SPEED-UPS ON THE SGI
ORIGIN200.

$M \times N$	$p = 1$	$p = 2$	$p = 3$	$p = 4$
16×16	0.0019	0.0018	0.0022	0.0024
		1.0649	0.8836	0.7982
32×16	0.0031	0.0026	0.0026	0.0025
		1.1795	1.1997	1.2103
64×16	0.0058	0.0038	0.0034	0.0031
		1.5190	1.7272	1.9086
128×16	0.0116	0.0068	0.0053	0.0045
		1.7152	2.1810	2.5667
256×16	0.0224	0.0127	0.0094	0.0081
		1.7643	2.3875	2.7770
32×32	0.0058	0.0039	0.0035	0.0033
		1.4865	1.6426	1.7815
64×32	0.0116	0.0068	0.0053	0.0047
		1.7114	2.1770	2.4609
128×32	0.0226	0.0124	0.0093	0.0079
		1.8245	2.4448	2.8567
256×32	0.0441	0.0236	0.0166	0.0133
		1.8704	2.6489	3.3117
512×32	0.0889	0.0466	0.0321	0.0265
		1.9090	2.7719	3.3486
64×64	0.0224	0.0123	0.0092	0.0088
		1.8199	2.4278	2.5415
128×64	0.0442	0.0235	0.0168	0.0132
		1.8844	2.6384	3.3605
256×64	0.0863	0.0458	0.0317	0.0246
		1.8836	2.7232	3.5041
512×64	0.1806	0.0900	0.0624	0.0469
		2.0055	2.8940	3.8479
1024×64	42.5168	21.7635	14.4899	10.8587
		1.9536	2.9342	3.9155

The experimental results given in tables I and II shows the run-time (in seconds/iteration) for several mesh sizes. It can be seen that as the mesh sizes increase, the scalability increases as well. Also noticeable is that in two cases (1024×64 and 1024×128) a substantial increase of the run-time occurs. Using the SGI per fex performance analyser, which

TABLE II

RUN TIME (IN SECONDS PER ITERATION) AND SPEED-UPS ON THE SGI ORIGIN200 (CONTINUED).

$M \times N$	$p = 1$	$p = 2$	$p = 3$	$p = 4$
128 × 128	0.0869	0.0457	0.0317	0.0245
		1.9018	2.7394	3.5505
256 × 128	0.1816	0.0892	0.0615	0.0468
		2.0360	2.9547	3.8818
512 × 128	0.3829	0.1857	0.1232	0.0958
		2.0625	3.1093	3.9961
1024 × 128	85.0758	43.5478	28.9583	21.8141
		1.9536	2.9379	3.9000

reports among other data the number of loads and stores per floating point instruction, we see that for the 512 × 64 mesh, this value is 2.2582, whereas for the 1024 × 64 mesh it jumps to 83.3133. It appears that the increased data traffic to/from the memory is responsible for the larger run-time exhibited in the latter case.

TABLE III

DATA CACHE HIT RATES FOR SOME PROBLEM SIZES. A VALUE CLOSER TO 1 REPRESENTS A BETTER CACHE MEMORY USAGE.

	$p = 1$	$p = 2$	$p = 3$	$p = 4$
256 × 64	0.9696	0.9926	1.0	1.0
512 × 64	0.9789	1.0	1.0	1.0
1024 × 64	0.9635	1.0	1.0	1.0
256 × 128	0.9802	1.0	1.0	1.0
512 × 128	0.9753	1.0	1.0	1.0
1024 × 128	0.9635	1.0	1.0	1.0

Another important effect being shown is that of a speed-up larger than 2 for the larger problem sizes when using two processors. Theoretically, such "superlinear" effect is impossible to achieve; however, if we are using a parallel computer with a separate cache memory for each processor, what may happen if we double the number of processors in use is that we have at our disposal the double of cache memories while at the same time we are halving the amount of data being accessed locally (supposing that load balance is achieved which is our case). In this case, more data will fit in these extra cache memories when compared to using a single processor

The above reasoning was confirmed by the *data cache hit rate* reported by *perfex*; in Table III we show this rate for a few problem sizes. Note that when $p = 2$ the data cache hit rate becomes 1 but for the 256 × 64 problem size, which is small compared to the others.

C. Reducing the latency in the computation of norms

As noted in the discussion of the parallel algorithm developed, we combined the reductions needed for the computation of the norms appearing in steps 5 and 7. Table IV shows the time (in seconds) taken by two implementations of the parallel algorithm (one with combined reductions and the other with separate reductions, and the respective gains). As can be seen in that table, the combined reductions are a means of increasing the speed-up for small grids, whilst still providing a reduction in the run time with increasing p , even for a large grid.

TABLE IV

COMPARISON BETWEEN RUN TIMES FOR (A) COMBINED AND (B) SEPARATE REDUCTIONS.

	$M \times N$			
	64 × 32	128 × 32	256 × 64	512 × 64
$p = 1$	0.3170	0.6436	2.4457	5.1776
$p = 2$				
(a)	0.1852	0.3528	1.2984	2.5817
(b)	0.2244	0.3979	1.3739	2.7304
gain(%)	21.1594	12.7980	5.8145	5.7586
$p = 3$				
(a)	0.1456	0.2633	0.8981	1.7891
(b)	0.1843	0.3097	0.9773	1.9242
gain(%)	26.5539	17.6365	8.8164	7.5498
$p = 4$				
(a)	0.1288	0.2253	0.6980	1.3456
(b)	0.1694	0.2802	0.7862	1.4756
gain(%)	31.5413	24.3674	12.6447	9.6674

V. CONCLUDING REMARKS

We have presented a parallel algorithm for the solution of the rotating flow problem described by the Navier-Stokes equations, using an explicit Runge-Kutta time-stepping integration scheme.

We believe the results presented show that our approach to parallelize the computation is good and can be used for the solution of related problems. Moreover, it can be used as a framework for the parallelization of other techniques, as long as the possibility of breaking down the computation in two parts - depending on local and remote stored data - exists.

We intend to further develop and apply it to other fluid flow problems, including three-dimensional domains with complex geometries, using generalized coordinates which will allow us to use the same parallelizing technique presented here.

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