Developing Code Char ¹STFC ³EDF R&D

Introduction

Computing Large-Eddy Simulations (LES) of a full power plant reactor would require about 13 to 15 billion cells. Some algorithms designed to reach this target are presented for Code_Saturne [1, 2], which is a highly scalable multi-purpose CFD software developed by EDF-R&D since 1997 and open source since 2007 (GPL license). The code is based upon a co-located finite volume approach that can handle three-dimensional meshes built with any type of cell and with any type of grid structure. MPI is used for communication. The code is able to simulate either incompressible or weakly compressible flows, with or without heat transfer, and has a variety of turbulence models (see Fig. 1 for the toolchain).



Figure 1: Code_Saturne's toolchain

The test cases consist of LES in staggereddistributed tube bundles (see Fig. 2). The configuration is the one experimentally studied by Simonin and Barcouda [3], and the mesh is obtained by copying and shifting several elemental patterns made of a whole tube in its centre and four quarter of tubes in the corners. The elemental pattern contains about 13 million cells (2-D cross-section: 100, 040 cells; 3^{rd} direction: 128 layers) and meshes of 51 (51M), 204 (204M), 816 (816M) million and 3.2 billion (3.2B) cells are used.



Figure 2: Location of the bundle of tubes in the reactor

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Hardware - Software

		T _a and a m
	HECIOR	Jaguar
Architecture	XE6	XK6
Processor	32-Core AMD	16-Core AMD
Nodes	2,816	18,688
Cores/Nodes	32	16
Memory/node	32GiB	32GiB
Memory/core	1GiB	2GiB
Compiler	GNU 4.6.3	PGI 12.1.0
MPI	MPICH2 5.4.5	MPICH2 5.4.1

Mesh Joining - Periodicity

Meshes are generated by joining parts of existing meshes. Each new mesh is built from 4 old meshes. Periodicity is handled as mesh joining. Table 1 shows that the time to complete the overall joining (3.2B case) is similar on both machines.

	HECToR	Jaguarpf
8192	95	127
16384	146	178
32768	400	414
65536	_	1630

Table 1: Time for overall joining in seconds (3.2B case)

Partitioning - Halos

A Space-Filling algorithm based on Morton Curves (SFC) and ParMetis-4.0.2 (ParM), used as partitioners are run with the number of cores as the number of sub-domains. Partitioning and computing halos take less than 2 minutes on 65536 cores for the 3.2B case on Jaguar (see Table 2).

	Partitioning			Halos				
	HECToR		Jaguar		HECToR		Jaguar	
	SFC	ParM	SFC	ParM	SFC	ParM	SFC	ParM
8192	21.7	45.6	14.8	14.3	10.6	8.2	11.1	8.7
16384	28.2	90.7	9.7	16.1	9.5	8.3	8.9	7.8
32768	53.5	181.0	16.4	40.8	22.6	21.2	18.7	17.7
65536	-	_	57.6	94.1	-	_	81.3	63.1

Table 2: Time for partitioning in seconds (3.2B case)

Solver perfor-Figure mance comparison for the 3.2B cell mesh (HECToR and Jaguar)

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IO Performance

wo strategies to andle IOs are availole in the code, [PI-IO and a serilisation method [2]. able 3 shows that [PI-IO is faster for ie 816M on the ECToR's Lustre file system.

	MPI-IO	Ser-IO
2048	633	1203
4096	608	640
8192	859	1147
16384	732	747

Table 3: Time for IOs in seconds (816M case) on HECToR

Solver

The Navier-Stokes solver is segregated. The pressure is solved by an Algebraic Multigrid algorithm (V-cycle), with the Conjugate Gradient as a solver. The Jacobi algorithm is used for the velocity components.



Solver perfor-Figure 3: mance for various mesh sizes (HECToR)



All cases but the 816M one show that the solver exhibits better results with SFC with than ParM as a partitioner (see Fig. 3) when run on HECToR.

The 3.2B case is used for comparison on both machines, where the average CPU time per time-steps decreases as a function of the number of cores, for both partitioners. A speed-up of 1.46 (resp. 1.14) is still observed for SFC (resp. ParM) on Jaguar, going from 32768 to 65536 cores.

The time spent in the algorithms used in Code_Saturne for mesh joining, periodicity, partitioning, halo builing have proven to be modest in comparison with the IO and solver time on Cray machines with Lustre file system, especially for a 3.2 billion cell mesh.

More testing will be performed on new architecture/file systems such as IBM Blue Gene/GPFS for instance, to assess the performance of the software.

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Conclusions - Perspectives

Acknowledgements

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