

```

subroutine godunov(idim,dt)
  use Hydra_commons
  use Hydra_const
  use Hydra_parameters
  use Hydra_util
  use Hydra_work_space
  use my_own_module
  implicit none

  ! Dummy arguments
  integer(kind=prec_int), intent(in) :: idim
  real(kind=prec_real),  intent(in) :: dt
  ! Local variables
  integer(kind=prec_int) :: i,j,im
  real(kind=prec_real)  :: dtdx

```

```

  ! constant
  dtdx=dt/dx

```

```

  ! Update boundary conditions
  call make_boundary(idim)
  if (idim==1)then

```

```

    ! Allocate work space for ID sweeps
    call allocate_work_space(imin,imax,ns=1)

    do j=lower+2, upper-2 ! was jmin=2,jmax=2

```

```

      ! Gather conservative variables
      do i=imin,imax
        u(i,ID)=uold(i,j,ID)
        u(i,IV)=uold(i,j,IV)
        u(i,IV)=uold(i,j,IV)
        u(i,IP)=uold(i,j,IP)
      end do
      if(nvar>4)then
        do im = 5,nvar
          do i=imin,imax
            u(i,im)=uold(i,j,im)
          end do
        end do
      end if

```

```

      ! Convert to primitive variables
      call constopr(u,q,c)

```

```

      ! Characteristic tracing
      call trace(q,dq,c,qm,qp,dtdx)

```

```

      do im = 1,nvar
        do i=1,ns-1
          qlen(i,im)=qm(i+1,im)
          qright(i,im)=qp(i+2,im)
        end do
      end do

```

```

      ! Solve Riemann problem at interfaces
      call riemann(qlen,qright,qpmv, &
        & rl,ul,pl,cl,wr,rr,pr,cr,wr,rs,vs,ps,cs,ws, &
        & rstar,vstar,ptstar,cstar,qpm,spin,spout, &
        & vshock,frac,scr,delp,pold,ind,ind2)

```

```

  ! Compute fluxes

```

# Parallelizing The Hydro Code

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# Hydro Code Explained

The hydro code is a simulation of a fluid or gas in an enclosed system. The initial conditions are set as to generate an unperturbed, homogeneous environment. Then, one can introduce points of higher or lower pressure or density to imitate an explosion or jet. The code then calculates the propagation of this perturbation using Godunov's method and outputs each frame as a single file, which can be used to create an animation.

There are five main components with different functionality to this Fortran based code:

- `main.f90`
  - calls `init_hydro` to initialize the grid with the initial conditions
  - executes the main time loop by calling
    - \* `output` to generate output data
    - \* `cmpdt` to compute the time step
    - \* `godunov` to calculate the fluid dynamics
- `module_hydro_principal.f90` contains the subroutines
  - `init_hydro`
  - `cmpdt`
  - `godunov` which calls
    - \* `make_boundary` to set the upper, lower, left and right boundaries of the grid
    - \* `allocate_work_space/deallocate_work_space` to (de)allocate space in the memory
    - \* other subroutines for calculation purposes
- `module_hydro_utils.f90` contains
  - `make_boundary`
  - other subroutines
- `module_hydro_IO.f90` contains the subroutines
  - `read_params` to read the parameters from the `input.nml` file
  - `output`
- `module_hydro_commun.f90` contains
  - `allocate_work_space/deallocate_work_space`
  - various other subroutines, mainly to define the variables used elsewhere in the code

Additionally, there are several input files to set the grid size with width  $nx$  and height  $ny$  and other initial conditions.

# Parallelization Techniques

## MPI

### Introduction

MPI stands for *message passing interface*. This refers to the fact that parallel computing occupies multiple CPUs to fasten up the execution of a code. As these CPUs work independently and with their own associated memory, they will have to communicate in some way; this is what MPI does.

The advantage of MPI over other libraries is that one has much more control over how the code is parallelized. The programmer can explicitly choose how the different cores communicate to each other and which information they share.

On the other hand, this freedom makes parallelizing a code a much more complex problem than it could be with other libraires. It requires quite a deep understanding of how the code works to know where and what to adjust.

### Parallelization Process

The main idea was to split the initial grid horizontally into equally sized parts in assigning each core a certain portion of it. Each process would then calculate the evolution of the fluid in his portion only. Finally, these different parts would be merged together to recreate the grid with its original size. This merging is where the main part of the paralellization process is happening:

For a process to be able to calculate a given pixel of its grid, it needs information of its four neighbouring pixels. Thus, to calculate the values of the lower and upper most pixels within its grid part, each process had to have access to the values the process below or above of it had calculated. This could be achieved by letting the different processes communicate to each other in the following way:

First the odd processes send their values to the above and below processes while the even ones receive from below and above, respectively. Then, this procedure is repeated with the even ones sending and the odd ones receving. Special attention has to be paid to the lower most and the upper most process, because they do only have to send and receive from one border. The main reason to implement the communication in this way, was to avoid a deadlock.

Other components of the code that needed to be adjusted were

- workspace allocation,
- time step computation and
- output formatting.

## Results

To evaluate the performance of the parallelized code we performed a strong and a weak scaling. More precisely, for the strong scaling we ran the code with 1 to 36 cores and a  $ny$  of 9000 units and repeated this ten times to get a mean time per core. Similarly, for the weak scaling we varied the number of cores from 1 to 36, each time incrementing  $ny$  by 300 units to keep the grid per core constant. The results of the respective runtimes are displayed in Figure 1.

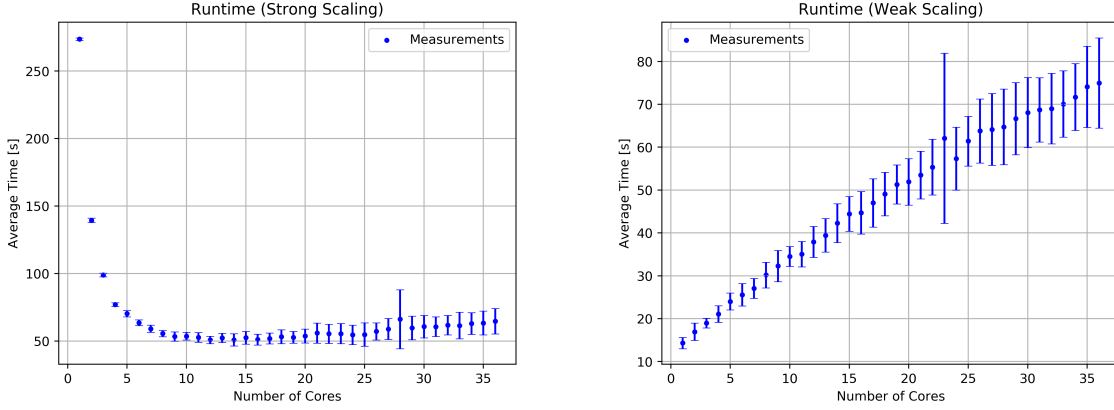


Figure 1: Respective runtimes for strong and weak scaling

To specify the percentage of the code that has actually been parallelized, we analyzed the speedup (strong scaling) and the efficiency (weak scaling).

In case of the strong scaling, the speedup  $s(N)$  is given by

$$s(N) = \frac{t_s}{t_p(N)} = \frac{1}{\alpha + \frac{1-\alpha}{N}}, \quad (1)$$

where  $t_s$  is the runtime of the serial code (i.e. one core),  $t_p(N)$  is the parallelized code as a function of the number of processes  $N$  and  $\alpha$  is the non-parallelized fraction of the code. It is displayed in Figure 2.

The actual value for  $\alpha$  can be obtained by fitting the function

$$f(N) = \frac{1}{\alpha + \frac{1-\alpha}{N}} \quad (2)$$

to the measurement points. In our case,

$$\alpha = 0.166 \pm 0.006. \quad (3)$$

This indicates that roughly 83.4% of the code is parallelized.

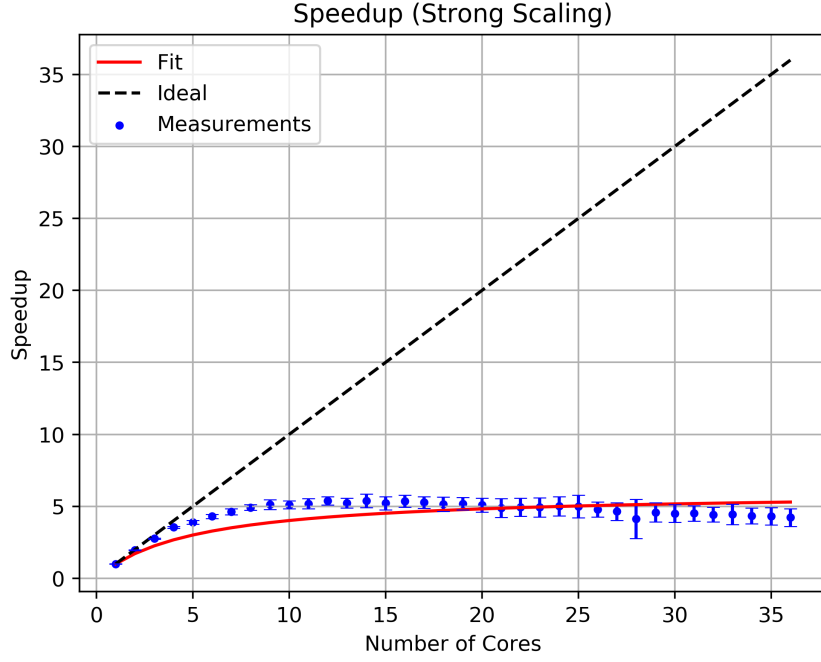


Figure 2: Speedup for strong scaling.

In case of the weak scaling, the efficiency  $\varepsilon$  is given by

$$\varepsilon(N) = \frac{t_s}{t_p(N)} = \frac{1}{\alpha \cdot N + (1 - \alpha)}.$$

It is illustrated in Figure 3. Fitting the function to the measurements yields

$$\alpha = 0.145 \pm 0.002. \quad (4)$$

Again, this indicates that around 85.5% of the code is parallelized.

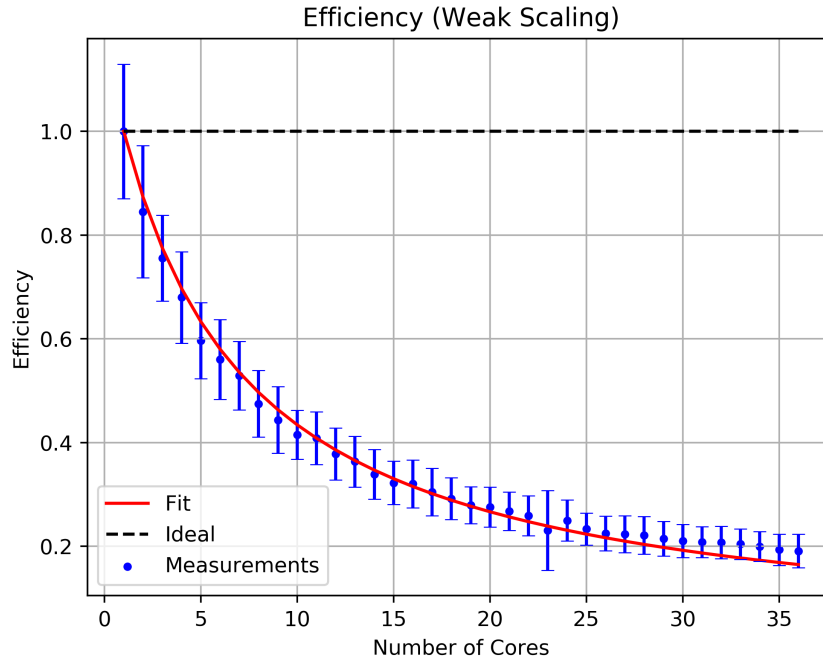


Figure 3: Efficiency for weak scaling.

## Other Techniques

### OpenMP

OpenMP stands for *open multiprocessing*. The fundamental difference between OpenMP and MPI is the parallelization process. MPI communicates between nodes, whereas with OpenMP the parallelization occurs within one node. The so-called *master thread* spawns several *slave threads* (called forking) to speed up a designated section of the code. What it may lack in flexibility (can only run on shared-memory computers), it makes up in ease of implementation and communication speed.

### MPI - OpenMP Hybrid

Implementing both of these methods can make use of the distributed memory architecture of a supercomputer, while further speeding up the runtime by running multiple threads. This combines the advantages of MPI and OpenMP.

## Discussion

As can be seen in Figure 1, the standard deviation on some of the runtimes can be quite big. This is mainly due that the whole run (going from 1 to 36 processes) was only performed 10 times. If one of these times is abnormally high or low, averaging over only 10 values will not suppress this spike well enough.

## Conclusion