# **Overview: Synchronous Computations**

• definition

- synchronous computation example 1: solving linear systems with Jacobi Iteration
  - solution of linear systems (problem definition)
  - fixed-point iterative linear system solvers and Jacobi iteration
  - serial and parallel code
  - partitioning and performance model
- synchronous computation example 2: solving the Heat Equation in 2D
  - problem definition and finite-difference discretization
  - serial and parallel code
  - partitioning: strip (1D) versus block (2D) partitioning; performance modelling
  - the concept of ghost layer of points (aka halo)
  - avoiding deadlocks
  - early termination
- synchronous computation example 3: Advection Equation in 2D-Assignment 1 (not covered in the lecture, similar to example 2)

Ref: Chapter 6: Wilkinson and Allen

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# Synchronous computations (definition)

- computations in which a group of processes perform local independent work BUT must periodically wait for each of other (i.e., synchronize) before proceeding
- (low-level) example: in SIMD computers the same instruction is executed on several processors on different data before proceeding with the next instruction
- in many cases, synchronization is a consequence of data exchange (e.g., to satisfy data dependency among steps)
- synchronous iteration (this lecture) is an important class of synchronous computations
  - to solve problems iteratively in such a way that several processes start together at the beginning of each iteration and the next iteration cannot begin until all processes have finished the preceding iteration
- we will illustrate synchronous iteration with two examples: Jacobi iteration and solution of the 2D Heat Equation in 2D

# Solution of Linear Systems (problem definition)

• we aim at finding  $x \in \mathbb{R}^n$  such that:

Ax = b

with  $A \in \mathbb{R}^{n \times n}$  (nonsingular matrix) and  $b \in \mathbb{R}^n$  (right-hand-side vector) given

in component-wise form, this problem reads (assuming 0-based indexing):

$$\underbrace{\begin{pmatrix} a_{0,0} & a_{0,1} & \cdots & a_{0,n-1} \\ a_{1,0} & a_{1,1} & \cdots & a_{1,n-1} \\ \vdots & \vdots & \ddots & \vdots \\ a_{n-1,0} & a_{n-1,1} & \cdots & a_{n-1,n-1} \end{pmatrix}}_{A} \begin{pmatrix} x_{0} \\ x_{1} \\ \vdots \\ x_{n-1} \end{pmatrix} = \underbrace{\begin{pmatrix} b_{0} \\ b_{1} \\ \vdots \\ b_{n-1} \end{pmatrix}}_{b}$$

- n is the number of equations/unknowns in the system
- ubiquitous problem in computational science and engineering (CSE) applications (e.g., numerical solution of <u>PDEs</u> using the <u>finite element method</u>)
- high quality parallel message-passing libraries around (e.g., <u>PETSc</u>, <u>Hypre</u>, <u>Trilinos</u>)

## **Fixed-point Iterative Linear solvers and Jacobi Iteration**

- the most basic iterative solvers are the so-called linear fixed-point methods
- in such methods, A is split as A = M N, with M being nonsingular
- starting from initial approximate solution  $x^{(0)}$ , they iterate the recurrence given by:

$$x^{(k+1)} = x^{(k)} + M^{-1} \underbrace{(b - Ax^{(k)})}_{\text{residual}}$$

till some termination criterion is fulfilled (e.g., max # of iterations reached or distance among  $x^{(k+1)}$  and  $x^{(k)}$  "sufficiently small")

- in practice one uses a cheap-to-invert approximation  $M^{-1} \approx A^{-1}$  (note that if  $M^{-1} = A^{-1}$  then  $x^{(1)}$  is already the solution)
- if they convergence, they are guaranteed to converge to *x*; however, they don't always converge (they converge if and only if ρ(*I M*<sup>-1</sup>*A*) < 1, with ρ(*B*) being the max eigenvalue of *B* in absolute value)
- Jacobi iteration (our example) choose  $M^{-1} = D^{-1}$ , with *D* being the diagonal of *A* (a quite rough approximation of  $A^{-1}$ !)

# **Sequential Jacobi Iteration**

the Jacobi recurrence in matrix form:

$$x^{(k+1)} = x^{(k)} + D^{-1}(b - Ax^{(k)})$$

can be written in component-wise form as:

$$x_i^{(k+1)} = x_i^{(k)} + \frac{1}{a_{ii}} \left( b_i - \sum_{j=0}^{n-1} a_{ij} x_j^{(k)} \right), \ i = 0, \dots, n-1$$

```
... // Init vector x
for (iter=0; iter<max_iter; iter++)
{
  for (i=0; i<n; i++) {
    sum=0.0
    for (j=0; j<n; j++) {
        sum = sum + a[i][j]*x[j];
        }
        new_x[i]=x[i]+(b[i]-sum)/a[i][i];
    }
    for (i=0; i<n; i++)
        x[i]=new_x[i];
}</pre>
```

- arrays b[] and a[][] hold b and A
- arrays x[] and new\_x[] hold  $x^{(k)}$ and  $x^{(k+1)}$
- for simplicity, we ignore early stopping condition (typically based on "sufficiently small" distance among x<sup>(k+1)</sup> and x<sup>(k)</sup>)

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# **Parallel Jacobi Iteration**

- consider a row-wise partition of A and a (naive!) static mapping of a single row per process, i.e., p = n (i.e., we have as many rows as processes)
- the vector b is partitioned/mapped accordingly to the rows of A
- HOWEVER, the vectors  $x^{(k)}$  and  $x^{(k+1)}$  are not partitioned/mapped to the processes, but replicated in all processes (why?)

message-passing parallel program (remainder: SPMD execution) at each outer loop iteration, each
 process with rank *i*, computes x<sub>i</sub><sup>(k+1)</sup>
 (entry of next iterate mapped to it)

- collective communication acts as a synchronization point
- this communication is such that all processes end up in new\_x[] with the entries of x<sup>(k+1)</sup> computed by any other processes
- let us discuss how to realize this communication step (next slides)

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## **Parallel Jacobi Iteration (communication)**

 most naive approach: p broadcasts naively implemented using point-to-point communication (not the way to go)

```
i = process_rank_id();
for (j=0; j<n; j++)
if (i!=j) send(&new_x[i],1,j);
for (j=0; j<n; j++)
if (i!=j) recv(&new_x[j],1,j);
```

```
Alternative 1 (deadlock-free?)
```

```
i = process_rank_id();
for (root=0; root<n; root++)
if (i==root)
  for (j = 0; j < n; j++)
      if (i!=j) send(&new_x[i],1,j);
else
  recv(&new_x[root], 1, root);
Alternative 2-reorder sends/recvs
```

```
(deadlock-free but still naive)
```

less naive approach: p broadcasts implemented using broadcast collective (but still not the way to go) for (root=0; root<n; root++)</p>

```
broadcast(&new_x[root], 1, root);
```

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### **Parallel Jacobi Iteration (communication)**

 smarter approach (but still not the way to go): butterfly pattern (aka recursive doubling) using point-to-point communication



*p* = 8 (thus *s* = 3)

- completes in  $s = \log_2(p)$  steps
- at each stage, we have  $\frac{p}{2}$  pairs of communication process
- at each stage, message size doubles (why?)

### **Parallel Jacobi Iteration (communication)**

smartest approach (the way to go): use MPI\_Allgather collective (it opens the door for exploiting a highly optimized algorithm available at the MPI implementation for the particular underlying high speed network at hand)



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# Partitioning and Parallel Cost Analysis of Jacobi iteration

- let us be more clever, and partition A (and b) into blocks of  $\frac{n}{p}$  rows each
- let us denote by  $\tau$  the number of Jacobi iterations
- as usual,  $t_f$  is the time/flop,  $t_s$  message start-up time,  $t_w$  per-word time
- sequential algorithm time (2 flops/inner loop + 3 flops/outer loop):

 $t_{\mathsf{seq}} = \tau \ n(2n+3)t_f$ 

parallel computation (decreases linearly with p):

 $t_{\rm comp} = \tau \, \frac{n}{p} (2n+3) t_f$ 

parallel communication

 (increases linearly with *p*):
 *t*<sub>comm</sub> =  $\tau p(t_s + \frac{n}{p}t_w) = \tau(pt_s + nt_w)$  

 parallel algorithm time:

 $t_{\text{par}} = t_{\text{comp}} + t_{\text{comm}}$ 

Assumptions:

- neglect the effect of the number of links and t<sub>h</sub>
- communication implemented inefficiently with p broadcasts
- communication cost of a broadcast equivalent to a single point-to-point communication

#### **Instantiating the Parallel Jacobi Iteration Time Model**



instantiating the model parameters as: •  $t_s = 10^5 t_f$ •  $t_w = 50 t_f$ • fixed problem size of n = 1000 (strong scaling)

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