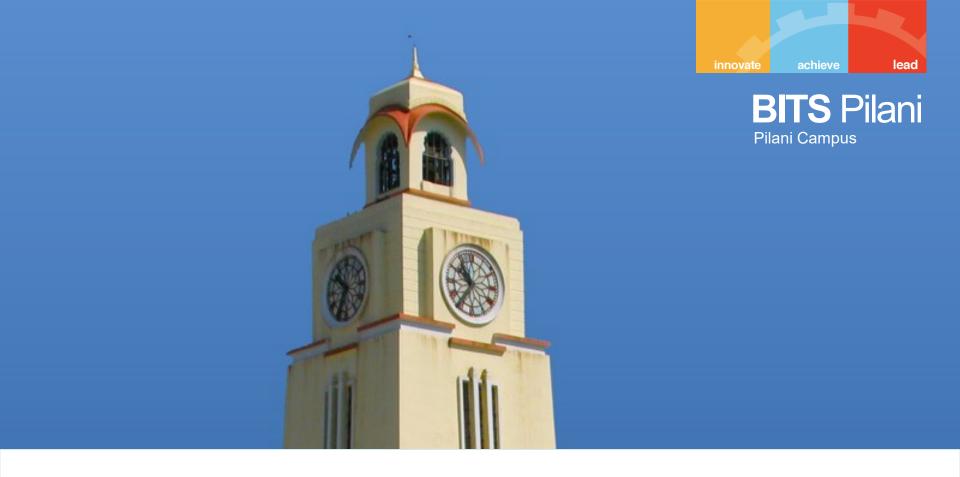




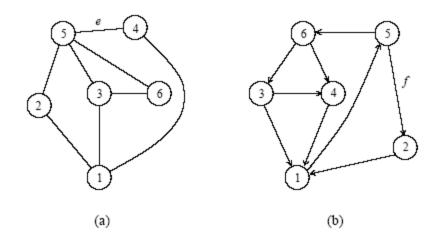
Selected Parallel Algorithms

K Hari Babu Department of Computer Science & Information Systems



Graphs-Representation

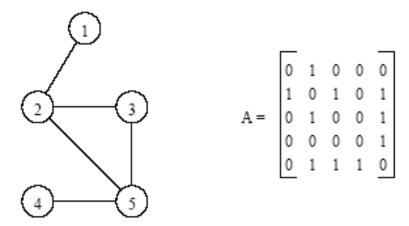
- An undirected graph G is a pair (V,E), where V is a finite set of points called vertices and E is a finite set of edges.
- An edge $e \in E$ is an unordered pair (u,v), where $u,v \in V$.
- In a directed graph, the edge e is an ordered pair (u,v). An edge (u,v) is incident from vertex u and is incident to vertex v.
- A path from a vertex v to a vertex u is a sequence $\langle v_0, v_1, v_2, ..., v_k \rangle$ of vertices where $v_0 = v$, $v_k = u$, and $(v_i, v_{i+1}) \in E$ for I = 0, 1, ..., k-1.
- The length of a path is defined as the number of edges in the path.



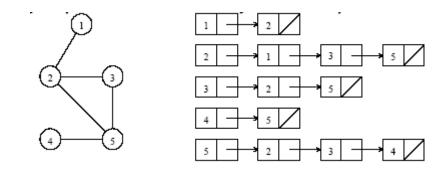
a) An undirected graph and (b) a directed graph.

- An undirected graph is connected if every pair of vertices is connected by a path.
- A forest is an acyclic graph, and a tree is a connected acyclic graph.
- A graph that has weights associated with each edge is called a weighted graph.

- Graphs can be represented by their adjacency matrix or an edge (or vertex) list.
- Adjacency matrices have a value $a_{i,j} = 1$ if nodes i and j share an edge; 0 otherwise. In case of a weighted graph, $a_{i,j} = w_{i,j}$, the weight of the edge.
- The adjacency list representation of a graph G = (V,E) consists of an array Adj[1...|V|] of lists. Each list Adj[v] is a list of all vertices adjacent to v.
- For a graph with n nodes, adjacency matrices take $\Theta(n^2)$ space and adjacency list takes $\Theta(|E|)$ space.



An undirected graph and its adjacency matrix representation.



An undirected graph and its adjacency list representation.

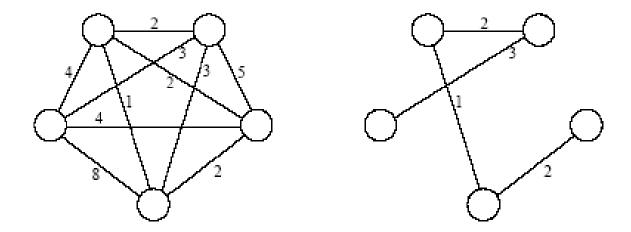


Minimum Spanning Tree – Prim's Algorithms

Minimum Spanning Tree

- A spanning tree of an undirected graph G is a subgraph of G that is a tree containing all the vertices of G.
- In a weighted graph, the weight of a subgraph is the sum of the weights of the edges in the subgraph.
- A minimum spanning tree (MST) for a weighted undirected graph is a spanning tree with minimum weight.

Minimum Spanning Tree

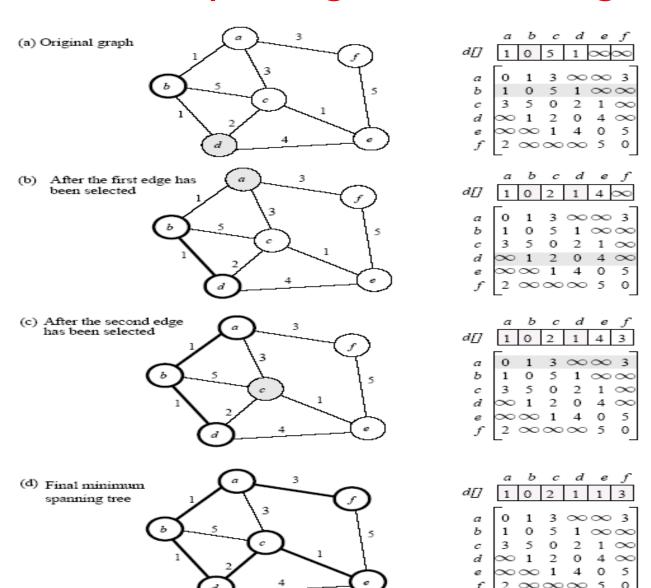


An undirected graph and its minimum spanning tree.

Minimum Spanning Tree: Prim's Algorithm

- Prim's algorithm for finding an MST is a greedy algorithm.
- Start by selecting an arbitrary vertex, include it into the current MST.
- Grow the current MST by inserting into it the vertex closest to one of the vertices already in current MST.

Minimum Spanning Tree: Prim's Algorithm



Prim's minimum spanning tree algorithm.

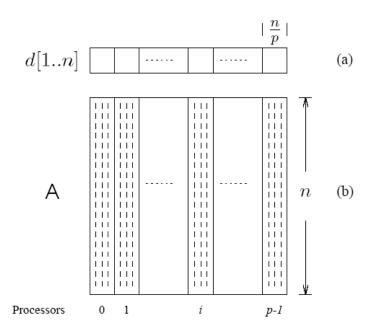
Minimum Spanning Tree: Prim's Algorithm

```
1.
          procedure PRIM\_MST(V, E, w, r)
2.
          begin
3.
               V_T := \{r\};
               d[r] := 0;
5.
               for all v \in (V - V_T) do
                    if edge (r, v) exists set d[v] := w(r, v);
6.
                    else set d[v] := \infty;
7.
8.
               while V_T \neq V do
9.
               begin
10.
                    find a vertex u such that d[u] := \min\{d[v] | v \in (V - V_T)\};
11.
                    V_T := V_T \cup \{u\};
12.
                    for all v \in (V - V_T) do
13.
                         d[v] := \min\{d[v], w(u, v)\};
14.
               endwhile
15.
          end PRIM_MST
```

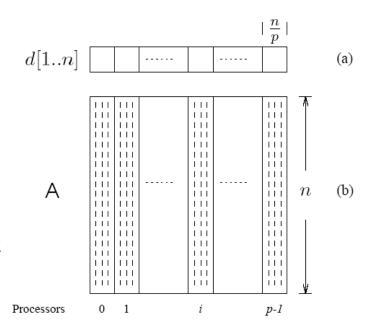
Prim's sequential minimum spanning tree algorithm.

- Prim's algorithm is iterative.
 - Each iteration adds a new vertex to the minimum spanning tree. Since the value of d[v] for a vertex v may change every time a new vertex u is added in V_T, it is hard to select more than one vertex to include in the minimum spanning tree.
 - Thus, it is not easy to perform different iterations of the while loop in parallel. However, each iteration can be performed in parallel as follows.
- Let p be the number of processes, and let n be the number of vertices in the graph. The set V is partitioned into p subsets using the 1-D block mapping.

- Each subset has n/p consecutive vertices, and the work associated with each subset is assigned to a different process.
- Let V_i be the subset of vertices assigned to process P_i for i = 0, 1, ..., p 1. Each process P_i stores the part of the array d that corresponds to V_i i.e. process P_i stores d [v] such that v ∈ Vi
- Each process P_i computes d_i[u] = min{d_i[v]|v
 ∈ (V-V_T)∩V_i} during each iteration of the while loop
- The global minimum is then obtained over all $d_i[u]$ by using the all-to-one reduction operation and is stored in process P_0 .

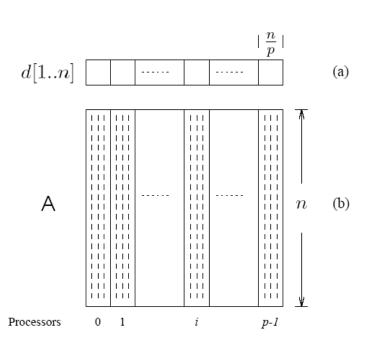


- Process P₀ now holds the new vertex u, which will be inserted into V_T. Process P₀ broadcasts u to all processes by using oneto-all broadcast
- The process P_i responsible for vertex u marks u as belonging to set V_T. Finally, each process updates the values of d[v] for its local vertices
- When a new vertex u is inserted into V_T , the values of d[v] for $vi \in (V-V_T)$ must be updated
 - The process responsible for v must know the weight of the edge (u, v). Hence, each process P_i needs to store the columns of the weighted adjacency matrix corresponding to set V_i of vertices assigned to it. This corresponds to 1-D block mapping of the matrix



- The space to store the required part of the adjacency matrix at each process is $Q(n^2/p)$
- The computation performed by a process to minimize and update the values of d[v] during each iteration is Q(n/p)
- One to all broadcast, all-to-one reduction takes Q(log p)
- The parallel run time of this formulation is given by

$$T_P = \Theta\left(\frac{n^2}{p}\right) + \Theta(n \log p).$$







Single-Source Shortest Paths - Dijkstra's Algorithm

Single-Source Shortest Paths

- For a weighted graph G = (V, E, w), the single-source shortest paths problem is to find the shortest paths from a vertex $v \in V$ to all other vertices in V.
- Dijkstra's algorithm solves the single-source shortest-paths problem on both directed and undirected graphs with nonnegative weights
- Dijkstra's algorithm is similar to Prim's algorithm. It maintains a set of nodes for which the shortest paths are known
 - The main difference is that, for each vertex u, Dijkstra's algorithm stores I[u], the minimum cost to reach vertex u from vertex s by means of vertices in V_T ; Prim's algorithm stores d [u], the cost of the minimum-cost edge connecting a vertex in V_T to u.

Single-Source Shortest Paths: Dijkstra's Algorithm

```
1.
          procedure DIJKSTRA_SINGLE_SOURCE_SP(V, E, w, s)
2.
          begin
3.
               V_T := \{s\};
              for all v \in (V - V_T) do
4.
                    if (s, v) exists set l[v] := w(s, v);
5.
6.
                    else set l[v] := \infty;
7.
               while V_T \neq V do
8.
               begin
                    find a vertex u such that l[u] := \min\{l[v] | v \in (V - V_T)\};
9.
10.
                    V_T := V_T \cup \{u\};
                    for all v \in (V - V_T) do
11.
12.
                         l[v] := \min\{l[v], l[u] + w(u, v)\};
13.
               endwhile
14.
          end DIJKSTRA_SINGLE_SOURCE_SP
```

Dijkstra's sequential single-source shortest paths algorithm.

Dijkstra's Algorithm: Parallel Formulation

- Very similar to the parallel formulation of Prim's algorithm for minimum spanning trees.
- The weighted adjacency matrix is partitioned using the 1-D block mapping.
 - Each of the p processes is assigned n/p consecutive columns of the weighted adjacency matrix, and computes n/p values of the array I.
 - During each iteration, all processes perform computation and communication similar to that performed by the parallel formulation of Prim's algorithm.
 - Each process selects, locally, the node closest to the source, followed by a global reduction to select next node.
 - The node is broadcast to all processors and the *I*-vector updated.
- The parallel performance of Dijkstra's algorithm is identical to that of Prim's algorithm.



All-Pairs Shortest Paths

All-Pairs Shortest Paths

- Instead of finding the shortest paths from a single vertex v to every other vertex, we are sometimes interested in finding the shortest paths between all pairs of vertices
- Formally, given a weighted graph G(V, E, w), the all-pairs shortest paths problem is to find the shortest paths between all pairs of vertices v_i , $v \in V$ such that i <> j.
- For a graph with n vertices, the output of an all-pairs shortest paths algorithm is an n x n matrix D = $(d_{i,j})$ such that $d_{i,j}$ is the cost of the shortest path from vertex v_i to vertex v_i .

All-Pairs Shortest Paths

- Two algorithms to solve the all-pairs shortest paths problem
- The first algorithm uses Dijkstra's single-source shortest paths algorithm, and the second uses Floyd's algorithm.
- Dijkstra's algorithm requires non-negative edge weights, whereas Floyd's algorithm works with graphs having negativeweight edges provided they contain no negative-weight cycles

Dijkstra's Algorithm

- Execute *n* instances of the single-source shortest path problem, one for each of the *n* source vertices.
- Complexity is $O(n^3)$.

Dijkstra's Algorithm: Parallel Formulation

- Two parallelization strategies
 - Execute each of the n shortest path problems on a different processor (source partitioned)
 - or use a parallel formulation of the shortest path problem to increase concurrency (source parallel).
- Source-partitioned formulation: Partition the vertices across processors
 - Works well if p<=n; No communication
 - Can at best use only n processors
- Source-parallel formulation: Parallelize SSSP for a vertex across a subset of processors
 - p processes are divided into p/n subsets
 - Each of the n-single source shortest paths problem is solved by a subset

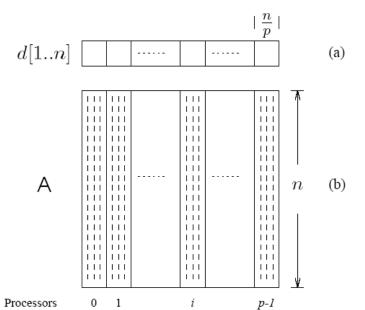
Dijkstra's Algorithm: Source Partitioned Formulation

- Use n processors, each processor P_i finds the shortest paths from vertex v_i to all other vertices by executing Dijkstra's sequential single-source shortest paths algorithm.
- It requires no interprocess communication (provided that the adjacency matrix is replicated at all processes).
- The parallel run time of this formulation is: $\Theta(n^2)$.
- While the algorithm is cost optimal, it can only use n processors.

Dijkstra's Algorithm: Source Parallel Formulation

- In this case, each of the shortest path problems is further executed in parallel. We can therefore use up to n^2 processors.
- Given p processors (p > n), each single source shortest path problem is executed by p/n processors.
- Using previous results, this takes time:
 - = time taken by a single group
 - = n iterations * (n2/p) computations

$$T_P = \Theta\left(rac{n^3}{p}
ight) + \Theta(n\log p).$$



SSSP complexity:
$$T_P = \Theta\left(\frac{n^2}{p}\right) + \Theta(n \log p)$$
.

Dijkstra's Algorithm: Parallel Formulation

- Comparing the two parallel formulations of Dijkstra's all-pairs algorithm
 - we see that the source-partitioned formulation performs no communication, can use no more than n processes, and solves the problem in time $Q(n^2)$.
 - In contrast, the source-parallel formulation uses up to n²/log n processes, has some communication overhead, and solves the problem in time Q(n log n) when n²/log n processes are used.
 - Thus, the source-parallel formulation exploits more parallelism than does the source-partitioned formulation.

Floyd's Algorithm

- For any pair of vertices v_i , $v_j \in V$, consider all paths from v_i to v_j whose intermediate vertices belong to the set $\{v_1, v_2, ..., v_k\}$. Let $p_{i,j}^{(k)}$ (of weight $d_{i,j}^{(k)}$ be the minimum-weight path among them.
- If vertex v_k is not in the shortest path from v_i to v_j , then $p_{i,j}^{(k)}$ is the same as $p_{i,j}^{(k-1)}$.
- If v_k is in $p_{i,j}^{(k)}$, then we can break $p_{i,j}^{(k)}$ into two paths one from v_i to v_k and one from v_k to v_j . Each of these paths uses vertices from $\{v_1, v_2, ..., v_{k-1}\}$.

Floyd's Algorithm

From our observations, the following recurrence relation follows:

$$d_{i,j}^{(k)} = \begin{cases} w(v_i, v_j) & \text{if } k = 0\\ \min\left\{d_{i,j}^{(k-1)}, d_{i,k}^{(k-1)} + d_{k,j}^{(k-1)}\right\} & \text{if } k \ge 1 \end{cases}$$

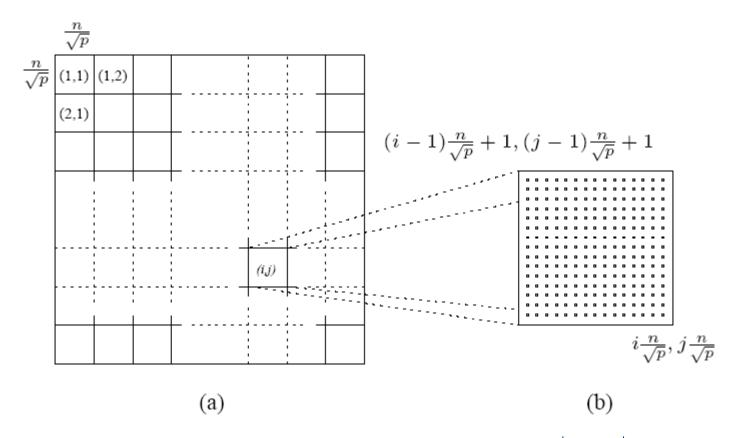
This equation must be computed for each pair of nodes and for k = 1, n. The serial complexity is $O(n^3)$.

Floyd's Algorithm

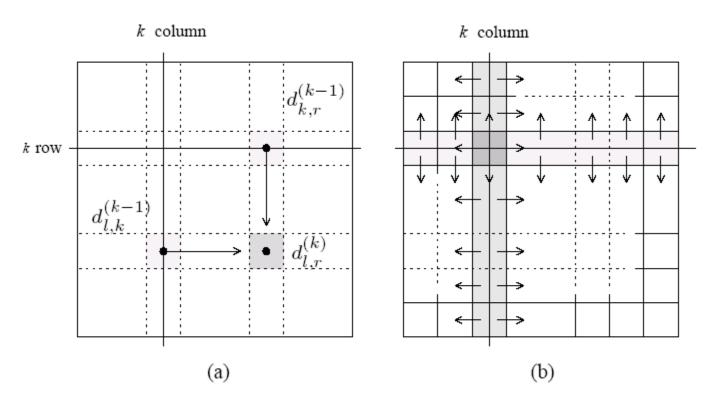
```
1.
            procedure FLOYD_ALL_PAIRS_SP(A)
2.
            begin
3.
                  D^{(0)} = A:
4.
                 for k := 1 to n do
5.
                       for i := 1 to n do
6.
                              for j := 1 to n do
                                   d_{i,j}^{(k)} := \min \left( d_{i,j}^{(k-1)}, d_{i,k}^{(k-1)} + d_{k,j}^{(k-1)} \right);
7.
           end FLOYD_ALL_PAIRS_SP
8.
```

Floyd's all-pairs shortest paths algorithm. This program computes the all-pairs shortest paths of the graph G = (V, E) with adjacency matrix A.

- Matrix $D^{(k)}$ is divided into p blocks of size $(n / \sqrt{p}) x (n / \sqrt{p})$.
- Each processor updates its part of the matrix during each iteration.
- To compute $d_{l,k}^{(k-1)}$ processor $P_{i,j}$ must get $d_{l,k}^{(k-1)}$ and $d_{k,r}^{(k-1)}$.
- In general, during the k^{th} iteration, each of the \sqrt{p} processes containing part of the k^{th} row send it to the \sqrt{p} 1 processes in the same column.
- Similarly, each of the \sqrt{p} processes containing part of the k^{th} column sends it to the \sqrt{p} 1 processes in the same row.



(a) Matrix $D^{(k)}$ distributed by 2-D block mapping into $\sqrt{p} \ x \sqrt{p}$ subblocks, and (b) the subblock of $D^{(k)}$ assigned to process $P_{i,j}$.



(a) Communication patterns used in the 2-D block mapping. When computing $d_{i,j}^{(k)}$, information must be sent to the highlighted process from two other processes along the same row and column. (b) The row and column of \sqrt{p} processes that contain the k^{th} row and column send them along process columns and rows.

```
procedure FLOYD_2DBLOCK(D^{(0)})
1.
         begin
3.
              for k := 1 to n do
4.
              begin
                   each process P_{i,j} that has a segment of the k^{th} row of D^{(k-1)};
5.
                        broadcasts it to the P_{*,j} processes;
                   each process P_{i,j} that has a segment of the k^{th} column of D^{(k-1)};
6.
                        broadcasts it to the P_{i,*} processes;
                   each process waits to receive the needed segments;
7.
                   each process P_{i,j} computes its part of the D^{(k)} matrix;
8.
              end
10.
         end FLOYD_2DBLOCK
```

Floyd's parallel formulation using the 2-D block mapping. $P_{*,j}$ denotes all the processes in the j^{th} column, and $P_{i,*}$ denotes all the processes in the i^{th} row. The matrix $D^{(0)}$ is the adjacency matrix.

- During each iteration of the algorithm, the k^{th} row and k^{th} column of processors perform a one-to-all broadcast along their rows/columns.
- The size of this broadcast is n/\sqrt{p} elements, taking time $\Theta((n \log p)/\sqrt{p})$.
- The synchronization step takes time $\Theta(\log p)$.
- The computation time is $\Theta(n^2/p)$.
- Each process runs <u>n iterations</u>. The parallel run time of the
 2-D block mapping formulation of Floyd's algorithm is

$$T_P = \Theta\left(\frac{n^3}{p}\right) + \Theta\left(\frac{n^2}{\sqrt{p}}\log p\right).$$

• The above formulation can use $O(n^2 / log^2 n)$ processors costoptimally.

$$S = \frac{\Theta(n^3)}{\Theta(n^3/p) + \Theta((n^2 \log p)/\sqrt{p})}$$

$$E = \frac{1}{1 + \Theta((\sqrt{p} \log p)/n)}$$

$$\Theta\left(\frac{\sqrt{p} \log p}{n}\right) = O(1)$$

$$\Theta(n^3/p) + \Theta((n^2 \log p)/\sqrt{p})$$

$$\Theta\left(\frac{\sqrt{p} \log p}{n}\right) = O(1)$$

• This algorithm can be further improved by relaxing the strict synchronization after each iteration.

References



• Chapter 10.1-4 of text book.



Thank You