Bipartite Matching

Brian A. Page bpage1nd.edu September 18, 2018

Bipartite Graphs

- Vertices of a graph G can be divided into two disjoint sets U and V.
- Every edge is of the form (u,v)
- No edges between vertices in same vertex set



• Assignment problem, transportation problem, etc.

Graph Matching

- Independent Edge Set (IES): set of edges in which no two edges share a common vertex
- **Maximal**: adding an edge not in *M* destroys matching





• Maximum : Largest possible IES (can be many)

• **Perfect**: matches all V in G

Data Sets

- Suite Sparse Matrix Collection
 - <u>https://sparse.tamu.edu/</u>
- SNAP (Stanford Large Network Dataset Collection)
 <u>http://snap.stanford.edu/data/</u>

3

• Synthetic Graphs

Edmonds–Karp

- Based on Ford-Fulkerson maximum flow method
- O(VE²) or O(V²E) time depending on implementation



Edmonds-Karp

algorithm EdmondsKarp

```
input:
            (graph[v] should be the list of edges coming out of vertex v.
    graph
             Each edge should have a capacity, flow, source and sink as parameters,
             as well as a pointer to the reverse edge.)
            (Source vertex)
    S
            (Sink vertex)
    t.
output:
            (Value of maximum flow)
    flow
            (Initialize flow to zero)
flow := 0
repeat
    (Run a bfs to find the shortest s-t path.
     We use 'pred' to store the edge taken to get to each vertex,
     so we can recover the path afterwards)
    q := queue()
    q.push(s)
    pred := array(graph.length)
    while not empty(q)
        cur := q.pull()
        for Edge e in graph[cur]
             if pred[e.t] = null and e.t ≠ s and e.cap > e.flow
                pred[e.t] := e
                q.push(e.t)
    if not (pred[t] = null)
        (We found an augmenting path.
         See how much flow we can send)
        df := \infty
        for (e := pred[t]; e ≠ null; e := pred[e.s])
            df := min(df, e.cap - e.flow)
        (And update edges by that amount)
        for (e := pred[t]; e ≠ null; e := pred[e.s])
            e.flow := e.flow + df
            e.rev.flow := e.rev.flow - df
        flow := flow + df
until pred[t] = null (i.e., until no augmenting path was found)
```

https://en.wikipedia.org/wiki/ Edmonds-Karp algorithm

> The College of Engineering at the University of Notre Dame

return flow

Hopcroft-Karp

- Based on Push-relabel (maximum flow)
- Uses BFS to partition vertices into matched and unmatched
- Swaps edges in/out of matching
- Local vs global path augmentations

```
Input: Bipartite graph G(U \cup V, E)

Output: Matching M \subseteq E

M \leftarrow \emptyset

repeat

\mathcal{P} \leftarrow \{P_1, P_2, \dots, P_k\} maximal set of vertex-disjoint shortest augmenting paths

M \leftarrow M \oplus (P_1 \cup P_2 \cup \dots \cup P_k)

until \mathcal{P} = \emptyset
```

• Runs in $O(|E|\sqrt{|V|})$

Hopcroft-Karp

```
/*
G = U \cup V \cup {NIL}
where U and V are partition of graph and NIL is a special null vertex
*/
function BFS ()
    for each u in U
        if Pair U[u] == NIL
            Dist[u] = 0
            Enqueue(Q,u)
        else
            Dist[u] = ∞
    Dist[NIL] = ∞
    while Empty(Q) == false
        u = Dequeue(Q)
        if Dist[u] < Dist[NIL]
            for each v in Adj[u]
                if Dist[ Pair V[v] ] == ∞
                    Dist[Pair V[v]] = Dist[u] + 1
                    Enqueue(Q,Pair V[v])
    return Dist[NIL] != ∞
```

```
function DFS (u)
   if u != NIL
        for each v in Adj[u]
            if Dist[ Pair V[v] ] == Dist[u] + 1
                if DFS(Pair V[v]) == true
                    Pair V[v] = u
                    Pair U[u] = v
                    return true
        Dist[u] = ∞
        return false
   return true
function Hopcroft-Karp
   for each u in U
        Pair U[u] = NIL
   for each v in V
        Pair V[v] = NIL
   matching = 0
   while BFS() == true
        for each u in U
            if Pair U[u] == NIL
                if DFS(u) == true
                    matching = matching + 1
   return matching
```

Distributed Bipartite Matching

Pros

- Performance via strong scaling
- Larger graphs
- Shorter comp. time (hopefully)

Cons

- Greatly increase complexity
- Time now dependent on system and network

Distributed Bipartite Matching

Basic Distributed Sequential Algorithm:

Given G(v,e) and process count P:

assign vertices such that $|P_i(v)| = |P_j(v)|$

distribute work

Perform BM on G via some method

if P_i interacts with vertex st. $v \notin P_i(v)$

alert P_j st v ∉P_j(v)

continue comp on P_i

return matching

Distributed Bipartite Matching

- Parallel Distributed BM possibilities:
 - Connected components
 - Cut vertex separation / sub-graph assignment
 - ???
- Questions to answer:
 - What can be parallelized?
 - How do we limit communication?
 - What other elements limit scalability and performance?

10

References

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11

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