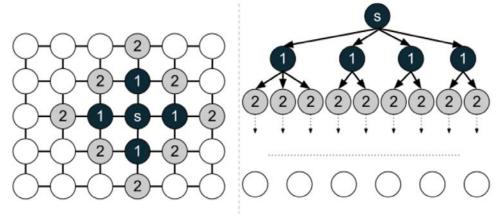
# CUDA 6 Breadth-First Search

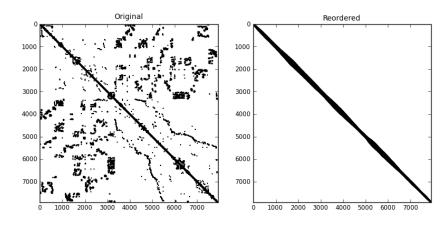
CS121 Parallel Computing Spring 2019



#### Breadth-first search

- Given a graph, explore it layer by layer.
  - □ Go wide, then go deep.
- Large number of applications.
  - Connected components, path finding, Ford-Fulkerson max flow algorithm, Cuthill-McKee ordering, bipartiteness testing, search engine crawlers, garbage collection, etc.
- Used in benchmarks such as Graph500 and Parboil to test parallel computer's memory performance.





Source: http://www.stoimen.com/blog/2012/10/08/computeralgorithms-shortest-path-in-a-graph/

Source: http://dpo.github.io/pyorder/\_images/commanche\_dual\_rcmk.png



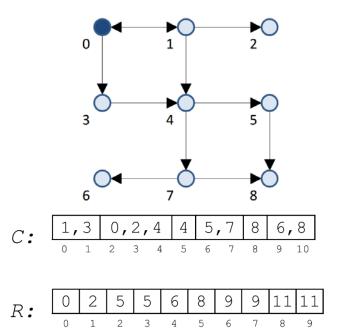
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Name	Sparsity Plot	Description	n (10 <sup>6</sup> )	<i>m</i> (10 <sup>6</sup> )	d	Avg. Search Depth
europe.osm		European road network	50.9	108.1	2.1	19314
grid5pt.5000		5-point Poisson stencil (2D grid lattice)	25.0	125.0	5.0	7500
hugebubbles-00020	X 2	Adaptive numerical simulation mesh	21.2	63.6	3.0	6151
grid7pt.300		7-point Poisson stencil (3D grid 27 lattice)		188.5	7.0	679
nlpkkt160		3D PDE-constrained optimization 8.3		221.2	26.5	142
audikw1		Automotive finite element analysis 0.9		76.7	81.3	62
cage15		Electrophoresis transition probabilities	5.2	94.0	18.2	37
kkt_power		Nonlinear optimization (KKT)	2.1	13.0	6.3	37
coPapersCiteseer		Citation network	0.4	32.1	73.9	26
wikipedia-20070206		Links between Wikipedia pages	3.6	45.0	12.6	20
kron_g500-logn20		Graph500 RMAT (A=0.57, B=0.19, C=0.19)	1.0	100.7	96.0	6
random.2Mv.128Me		G(n, M) uniform random	2.0	128.0	64.0	6
rmat.2Mv.128Me		RMAT (A=0.45, B=0.15, C=0.15)	2.0	128.0	64.0	6

- Hundreds of millions of nodes and edges.
  - □ Some graphs have billions or trillions of edges. But these don't fit into the memory of a single GPU.
- Low average degree (sparse), but high variation in degree.
  - Some nodes have a few neighbors, some nodes 100K's.
- "Small world" graphs have low diameter (~10).
- Grids and maps have high diameter (~1-10K).



## Sequential algorithm

- Assume graph is sparse, and stored in compressed sparse row format.
- Maintain a queue of unvisited nodes.
  - □ Dequeue a node, add its unvisited neighbors to the queue.
- Running time O(|V|+|E|).



Traversal from source vertex v <sub>0</sub>					
BFS Iteration	Vertex frontier	Edge frontier			
1	{0}	{1,3}			
2	{1,3}	{0,2,4,4}			
3	{2,4}	{5,7}			
4	{5,7}	{6,8,8}			
5	{6,8}	{}			
10 <b>if</b> (c	list[i] == ∞)				

```
10     if (dist[j] == ∞)
11          dist[j] := dist[i] + 1;
12          Q.Enqueue(j)
```



## Parallelizing BFS

- First BFS algorithms for GPUs focused on data parallelism.
- Initially set source distance to 0.
- Run for D rounds, where D is the diameter from s.
  - □ In round i, distance i nodes are marked.
  - Iterate through all the nodes. If a node is marked, mark its unvisited neighbors as distance i+1 nodes.
- Works well in small diameter graphs, e.g. social networks.
- Very inefficient for large diameter graphs, e.g. maps, since only a few nodes marked per round.
- O(|V||E|) running time.

```
parallel for (i in V) :
    dist[i] := ∞

dist[s] := 0
iteration := 0

do :    should add "if dist[j] == ∞" here

    done := true

parallel for (i in V) :
    if (dist[i] == iteration)
        done := false
    for (offset in R[i] .. R[i+1]-1)
        j := C[offset]
        dist[j] = iteration + 1

iteration++
while (!done)
```



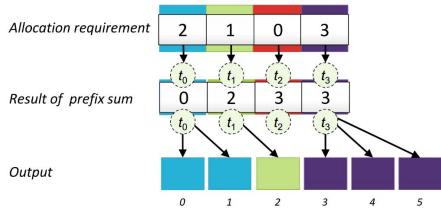
## Parallelizing BFS

- Linear, i.e. O(|V| + |E|) work parallel BFS algorithms follow the sequential algorithm.
- Two main bottlenecks
  - Maintaining explicit queue of unvisited nodes requires expensive locking operations.
  - □ If nodes have very different degrees (e.g. power law graphs), there's high load imbalance.

```
parallel for (i in V):
  dist[i] := ∞
dist[s] := 0
iteration := 0
inQ := \{ \}
inQ.LockedEnqueue(s)
while (inQ != \{\}):
  outQ := \{\}
  parallel for (i in inQ) :
     for (offset in R[i] .. R[i+1]-1)
        i := C[offset]
        if (dist[j] == \infty)
           dist[j] = iteration + 1
           outQ.LockedEnqueue(j)
  iteration++
  inQ := outQ
```

## Gathering neighbors

- We use two queues, one for nodes in current layer of BFS, other for nodes in next layer.
  - ☐ After every phase of BFS we swap the queues, to reuse memory.
  - ☐ To synchronize the layers, use a separate kernel for each layer.
- For each node in first queue, we first add all its neighbors into the second queue (gather).
  - Some of the neighbors don't belong in the next BFS layer because they've already been visited.
    - Testing each node explicitly is inefficient.
  - ☐ Also, we may add duplicates into the second queue.
  - □ We'll address both problems later.
- To add neighbors of a node into the queue, we use prefix sum instead of locking. Using prefix sum is much faster than locking.
  - If node has n<sub>i</sub> neighbors, we reserve n<sub>i</sub> queue spots for them by adding n<sub>i</sub> into the prefix sum.





## Load balanced gathering

- To load balance, we assign different numbers of threads to gather the neighbors of some node in parallel.
- If the node has moderate number of neighbors, assign a warp of threads to gather its neighbors.
  - Each thread in the warp might initially want to gather neighbors of a different node. All threads in warp write to a common location. This write is synchronized across the warp. The last to write "wins", and other threads help gather its node's neighbors.
- If node has large number of neighbors, use entire thread block for gather.
- For remaining nodes, use prefix sum based method.
  - □ There's load imbalance, but only for low degree nodes.
- "Moderate" and "large" need to be tuned.
- Eliminates most, but not all load imbalance.

## r,e

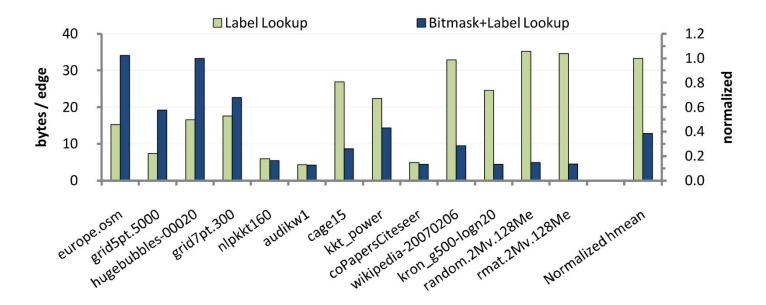
#### Visited status lookup

- A node should only be added into the frontier queue if it hasn't been visited.
  - □ Before adding a node, look up its visited status.
- To reduce memory traffic, use an integer bitmask to store status of 32 nodes.
- But then two threads might "clobber" each other by setting (different) bits in the same integer.
  - □ Can avoid using atomics, but they're slow.
  - Instead, use normal read and write ops, but treat bitmask conservatively.
    - For each node, maintain both a shared bitmask bit, and a private integer label.
    - Usually only access bitmask, saving memory traffic. Occasionally access the label.
    - If bit for a node is set, it's definitely visited.
    - If bit is unset, then not sure about node's visited status, so do another lookup on node's label.

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#### Visited status lookup

- Bitmasks are cached in texture caches.
- This is effective for low diameter graphs.
- Works less well for high diameter graphs, because each layer is processed in separate kernel, and cache flushed after each kernel launch.
- Also doesn't work well for small frontiers, since cached values aren't reused.
- Graphs on left side have high diameter; right ones are low diameter.

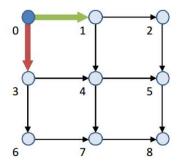


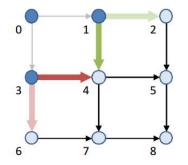


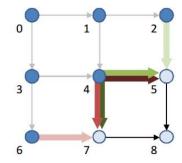
#### Duplicates in frontier

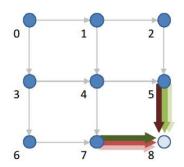
- May add same node into frontier multiple times, due to concurrent discovery.
- Problem especially severe in GPU because of SIMD and high parallelism.
- If duplicates aren't removed, the vertex frontier can grow exponentially.

BFS Iteration	Actual Vertex- frontier	Actual Edge- frontier		
1	0	1,3		
2	1,3	2,4,4,6		
3	2,4,4,6	5,5,7,5,7,7		
4	5,5,7,5,7,7	8,8,8,8,8,8		



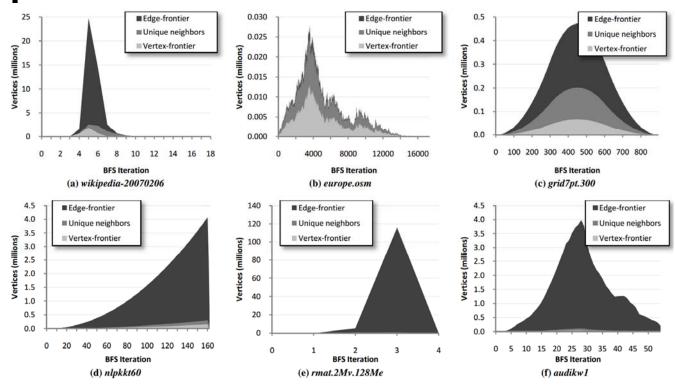






Iteration 1 Iteration 2 Iteration 3 Iteration 4

#### Duplicates in frontier



- Edge frontier: Number of nodes added to queue, allowing duplicates.
- Unique neighbors: Number of nodes added to queue, removing duplicates, but allowing visited nodes.
- Vertex frontier: Unique neighbors which haven't been visited.
- Allowing duplicates can lead to huge amount of redundant work.

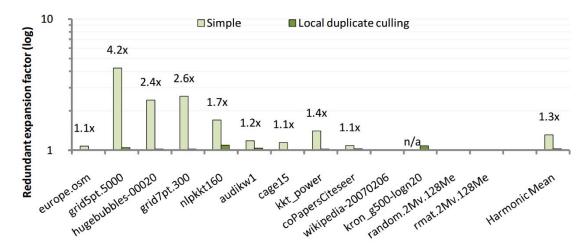


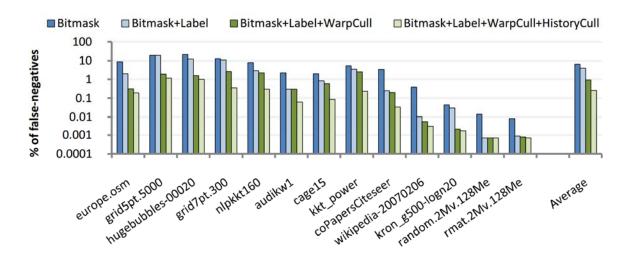
#### Duplicate culling

- Try to remove duplicates using hash table.
  - □ Won't remove all duplicates, but quite effective.
- Warp culling
  - □ Each warp allocates a hash table (with 128 entries) in shared memory.
  - □ When inserting a node, hash it into hash table.
    - If table entry empty, store the node in entry, and add node to queue.
    - If table entry filled, then if entry equals the node, don't add node to queue. Otherwise, add it.
- History culling
  - □ Same idea, but use the SM's L1 cache.

## Duplicate culling

Despite small hash table, culling surprisingly effective.



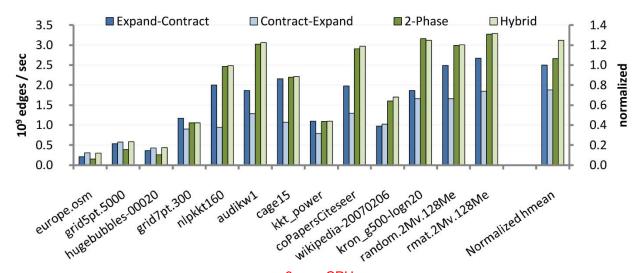


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## Putting it together

- Each kernel expands one layer of the BFS.
  - □ Input is queue containing last BFS layer (possibly with duplicate nodes).
- Threads assigned nodes from queue.
- A thread first uses warp and history culling to determine if its vertex is a duplicate.
- If not, thread gathers node's neighbors.
  - ☐ Based on neighbor list size, use a block, warp, or prefix sum gather.
    - Each thread wants to gather neighbors of a different node, and tries to "enlist" a block or warp of threads to help it.
    - Each thread writes into a variable shared by warp or block, then reads it.
    - One thread from the warp / block "wins". All other threads help it.
- Before adding a gathered node to (current layer's) queue, check if it's already visited.
- If not, the thread contributes 1 to a thread block-wide prefix sum.
- Synchronize the block and do a block-wide prefix sum to get number of enqueued nodes for block.
- First thread in block atomically adds sum to global queue index, then shares old global index with block.
- Using old global offset and prefix sum offset, each thread adds its gathered neighbor into queue.

#### Performance



8 core CPU

	CPU	CPU	NVIDIA Tesla C2050 (hybrid)				
<b>Graph Dataset</b>	Sequential <sup>†</sup>	Parallel	<b>Label Distance</b>		Label Predecessor		
1007	10 <sup>9</sup> TE/s	10 <sup>9</sup> TE/s	10 <sup>9</sup> TE/s	Speedup	10 <sup>9</sup> TE/s	Speedup	
europe.osm	0.029		0.31	11x	0.31	11x	
grid5pt.5000	0.081		0.60	7.3x	0.57	7.0x	
hugebubbles-00020	0.029		0.43	15x	0.42	15x	
grid7pt.300	0.038	0.12**	1.1	28x	0.97	26x	
nlpkkt160	0.26	0.47	2.5	9.6x	2.1	8.3x	
audikw1	0.65	1945	3.0	4.6x	2.5	4.0x	
cage15	0.13	0.23	2.2	18x	1.9	15x	
kkt_power	0.047	0.11**	1.1	23x	1.0	21x	
coPapersCiteseer	0.50		3.0	5.9x	2.5	5.0x	
wikipedia-20070206	0.065	0.19	1.6	25x	1.4	22x	
kron_g500-logn20	0.24		3.1	13x	2.5	11x	
random.2Mv.128Me	0.10	0.50***	3.0	29x	2.4	23x	
rmat.2Mv.128Me	0.15	0.70	3.3	22x	2.6	18x	

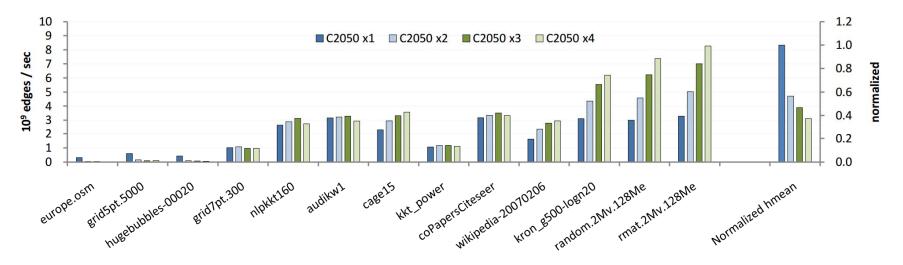
- Previous algorithm called "contract-expand", because it first takes current layer's edge frontier, contracts it (removes duplicates), then expands into next layer's edge frontier (containing duplicates).
- "Expand-contract", algorithm expands current vertex frontier, then contracts it (removes duplicates) to next layer's vertex frontier.
- 2-phase expands then contracts in two kernels.
- Hybrid combines contractexpand with 2-phase, using 2-phase for iterations with large frontiers.
- Variants differ in amount of memory traffic, latency and parallelism.
- Hybrid's performance is mostly determined by average degree (which generally increases moving down the dataset).

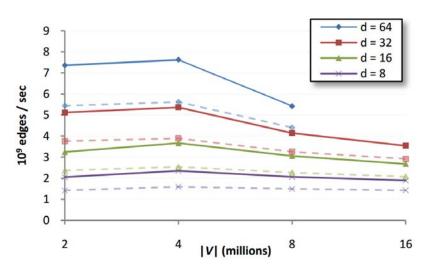
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#### Multi-GPU BFS

- Multiple GPUs can use a single logical address space.
  - □ Commuicate through PCI-e 2.0 (6.6 GB/s).
- Given p GPUs, assign n/p vertices and corresponding edges per GPU.
  - Vertices assigned in round robin order for load balancing.
  - □ Poor locality if p large.
- Each GPU expands / contracts its own vertex queue, as in the single GPU algorithm (\*).
- Then sort the new frontier into p bins, corresponding to vertices from different GPUs.
- Barrier across all GPUs.
- Run p-1 kernels, where in i'th kernel, the i'th GPU collects bin i from each other GPU.
- Then go back to step (\*) to form the next layer. Continue until all nodes visited.

#### Performance





Performance on uniform random graph. Higher average degree (d) results in better duplicate culling and higher performance.

- Only achieved speedup on graphs with small diameters and large average degrees.
  - Smaller diameter requires less synchronization.
  - Larger degree makes duplicate culling more effective.
  - Max speedups 1.5X, 2.1X and 2.5X on 2, 3, 4 GPUs.
  - Sometimes parallel algorithm performed much worse.