Parallel algorithms for butterfly + clique computations

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Outline

- Problem statement + Applications
- ParButterfly framework
 - Parallel butterfly counting
 - Parallel butterfly peeling
- Implementation + evaluation of ParButterfly
- Parallel clique counting framework
 - Parallel clique counting
 - Parallel clique peeling
- Implementation + evaluation of parallel clique counting + peeling
- Conclusion + Future work

Graph processing

• Graphs are ubiquitous



https://gizmodo.com/fascinating-graphic-shows-who-owns-all-the-major-brands-1599537576



Data-driven Modeling of Transportation Systems and Traffic Data Analysis During a Major Power Outage in the Netherlands



http://bitcoinwiki.co/wp-content/uploads/ censorship-free-social-network-akasha-aimsto-tackle-internet-censorship-with-blockchaintechnology.jpg

Bipartite graphs

Bipartite graphs: Represent relationships between two groups



Measuring Long-Term Impact Based on Network Centrality: Unraveling Cinematic Citations

Representation Learning

disease phenome

Fanconi a

The human disease network

disease genome

Parallelism

• Parallelism enables us to efficiently process large graphs



Apple, Microsoft, Intel, https://www.flickr.com/photos/66016217@N00/2556707493/, HP

Bipartite graphs

• Butterflies = 4-cycles = $K_{2,2}$



Think of these as the bipartite analogue of triangles (K₃) Note: Bipartite graphs contain no triangles

Finding dense subgraphs

• Problem: Given a graph G, find dense (bipartite) subgraphs

- Applications:
 - Find communities in social networks, websites, etc.
 - Discovering protein interactions in computational biology
 - Fraud detection in finance (tampered derivatives)

- Link spam: Create many external links to a spam page, for web search ranking promotion
- Link graph: Webpages are nodes, connected by incoming / outgoing hyperlinks



Link spam detection

- Note: Web communities tend to be dense bipartite subgraphs^[1]
- Web community bipartitions: topics, page creators interested in topics



[1] Kumar, Raghavan, Rajagopalan, Tomkins (99)

Tampered derivatives

 Tampered derivatives: Backed by set of assets/loans, tampered to contain many unprofitable (lemon) asset classes^[2]



- How do we find dense subgraphs (in general)?
- Algorithms:
 - K-core
 - Triangle peeling
- How do we find dense bipartite subgraphs?

• K-core: Repeatedly find + delete min degree vertex



Formally: A k-core is an induced subgraph where every vertex has degree at least k



 Triangle peeling: Repeatedly find + delete vertex contained within the minimum # of triangles



• Problem: Bipartite graphs do not contain any triangles

 Butterfly peeling: Repeatedly find + delete vertex containing min # of butterflies^[3]

Outline (Butterflies)

- Main goal: Build a framework ParButterfly to count and peel butterflies
- New parallel algorithms for butterfly counting + peeling
- ParButterfly framework with modular settings
 - Tradeoff b/w theoretical bounds + practical speedups
- Comprehensive evaluation
 - Counting outperforms fastest seq algorithms by up to 13.6x
 - Peeling outperforms fastest seq algorithms by up to 10.7x

- Main goal: Develop efficient exact and approximate algorithms for kclique counting and peeling
- New parallel algorithms for k-clique counting + peeling
- Comprehensive evaluation
 - Counting outperforms fastest parallel algorithms by up to 9.88x
 - Peeling outperforms fastest seq algorithms by up to 11.83x
 - Compute 4-clique counts on largest publicly-available graph with > 200 billion edges

Important paradigms

• Strong theoretical bounds

- Work = total # operations = # vertices in graph
- Span = longest dependency path = longest directed path
- Running time ≤ (work / # processors) + O(span)
- Work-efficient = work matches sequential time complexity

Parallel computation graph



https://web.fe.up.pt/~jbarbosa/en/research_par.html

ParButterfly counting framework

How do we count butterflies? (per vertex)





How do we count butterflies? (per vertex)

Wedge =
$$P_2$$
 =

Wedges with the same endpoints form butterflies:



wedges w/endpoints = w = 3

butterflies on endpoints $= \binom{w}{2} = \binom{3}{2} = 3$ # butterflies on each center = w - 1 = 3 - 1 = 2

Counting framework so far



One question: How do we aggregate wedges? (will discuss wedge retrieval after)

Method 1: Semisorting (on endpoints)



• Method 1: Semisorting (on endpoints)



Method 2: Hashing (keys = endpoints)



Method 2: Hashing (keys = endpoints)



• Method 3: Histogramming (frequencies of endpoints)



Semisorting^[1], hashing^[2], and histogramming^[3] are all workefficient

w = # of wedges

O(w) expected work, O(log w) span whp

[1] Gu, Shun, Sun, and Blelloch (15)[2] Shun and Blelloch (14)[3] Dhulipala, Blelloch, and Shun (17)

Counting framework so far



One more way to count wedges: Batching (not with polylogarithmic span, but fast in practice)

Wedge aggregating (batching)

 Main idea: Process a subset of vertices in parallel, finding all wedges where those vertices are endpoints



Counting framework so far



More questions:

How do we retrieve wedges? How many wedges are there?

It depends!

 Method 1: Process wedges w/endpoints from one bipartition (Side) ^[1]

6 wedges



Is this optimal (min # wedges)? Not always.

[1] Sanei-Mehri, Sariyuce, Tirthapura (18)

(Note: Butterfly count remains the same)

 Regardless of which side we pick, butterfly count does not change – only some "useful" wedges create butterflies

6 wedges





5 wedges

2 "useful" wedges = 1 butterfly

2 "useful" wedges = 1 butterfly

Retrieve wedges

• Method 2: Degree ranking

Main idea:

Once we obtain all wedges with endpoint v, we do not have to consider wedges with endpoint v again.

• Method 2: Degree ranking

- 1. Order vertices by non-increasing degree
- 2. For each vertex v, only consider wedges with endpoint v that is formed by vertices later in the ordering than v

Retrieve wedges

• Method 2: Degree ranking



2 wedges
Retrieve wedges

• Method 2: Degree ranking



2 wedges

Retrieve wedges

• Method 2: Degree ranking



We only processed 4 wedges!

- # wedges processed using degree order = $O(\alpha m)^{[1]}$
 - α = arboricity/degeneracy (O(\sqrt{m}))
 - m = # edges
- Therefore: (using work-efficient options)

Ranking vertices = O(m) expected work, $O(\log m)$ span whp Retrieving wedges = $O(\alpha m)$ expected work, $O(\log m)$ span whp Counting wedges = $O(\alpha m)$ expected work, $O(\log m)$ span whp Computing butterfly counts = $O(\alpha m)$ expected work, $O(\log m)$ span whp

Total = $O(\alpha m)$ expected work, $O(\log m)$ span whp

[1] Chiba and Nishizeki (85)

Other rankings

• Approximate degree order

- Log degree
- Complement degeneracy order
 - Ordering given by repeatedly finding + deleting greatest degree vertex
- Approximate complement degeneracy order
 - Complement degeneracy order, but using log degree

We show these are all work-efficient

Counting framework



ParButterfly peeling framework

How do we peel butterflies?

• Goal: Iteratively remove all vertices with min butterfly count

Subgoal 1: A way to keep track of vertices with min butterfly count Subgoal 2: A way to update butterfly counts after peeling vertices

Note: We've already done subgoal 2 in counting framework

For subgoal 1, we give a work-efficient batch-parallel Fibonacci heap which supports batch insertions/decrease-keys (see paper).

Peeling framework



(with respect to peeling complexity)

ParButterfly evaluation

- m5d.24xlarge AWS EC2 instance: 48 cores (2-way hyperthreading), 384 GiB main memory
- Cilk Plus^[1] work-stealing scheduler
- Koblenz Network Collection (KONECT) bipartite graphs
- Experiments for the different modular options in our framework
- Some modifications:
 - Julienne^[2] instead of batch-parallel Fibonacci heap
 - Cannot hold all wedges in memory batch wedge retrieval

[1] Leiserson (10)

[2] Dhulipala, Blelloch, and Shun (17)

Counting:

Best aggregation method: **Batching**



Counting:

Best ranking method: Approx Complement Degeneracy / Approx Degree



Butterfly counting results

- 6.3 13.6x speedups over best seq implementations^{[1] [2]}
- 349.6 5169x speedups over best parallel implementations^[3]
 - Due to work-efficiency
- 7.1 38.5x self-relative speedups

• Up to 1.7x additional speedup using a cache-optimization^[4]

Sanei-Mehri, Sariyuce, Tirthapura (18)
ESCAPE: Pinar, Seshadhri, Vishal (17)
PGD: Ahmed, Neville, Rossi, Duffield, and Wilke (17)
Wang, Lin, Qin, Zhang, and Zhang (19)

Peeling:

Best aggregation method: Histogramming



Butterfly peeling results

• 1.3 – 30696x speedups over best seq implementations^[1]

- Depends heavily on peeling complexity
- Largest speedup due to better work-efficiency for some graphs
- Up to 10.7x self-relative speedups
 - No self-relative speedups if small # of vertices peeled

k-clique counting and peeling

How do we find k-cliques?

• Repeatedly intersect the neighborhoods of vertices

How do we find k-cliques?

• Repeatedly intersect the neighborhoods of vertices



Déjà vu: Ranking vertices

• How do we avoid double-counting k-cliques?



At each level, only store the set of vertices in the intersection of the out-neighborhood of the clique

Ranking vertices

- Degree ranking
- K-core ranking

- Arboricity ranking
 - Goodrich-Pszona
 - Barenboim-Elkin

Used in previous work Work-efficient serially

Our contribution (parallelizations of these algorithms) Space-efficient in parallel Polylogarithmic span in parallel

 $O(\alpha)$ -orientations in O(m) expected work, $O(\log^2 n)$ span whp k-clique counting in $O(m\alpha^{k-2})$ expected work, $O(k \log n + \log^2 n)$ span whp, $O(m + P\alpha)$ space • Goal: Iteratively remove all vertices with min k-clique count

Subgoal 1: A way to keep track of vertices with min k-clique count Subgoal 2: A way to update k-clique counts after peeling vertices

Note: We've already done subgoal 2 in the counting algorithm

And we've already done subgoal 1 in the butterfly peeling algorithm!

We show this algorithm is work-efficient (with respect to peeling complexity)

- k-clique peeling uses essentially the same algorithm, but must consider the undirected neighborhood of the peeled vertex
- Nash-Williams Theorem gives work-efficient bounds
 - For every subgraph S, $\propto \geq \frac{E(S)}{V(S)-1}$

k-clique evaluation

Evaluation (k-clique counting)

- 60-core GCP instance (two-way hyperthreading)
- 1.31 9.88x speedups over parallel KClist^[1]
- 2.26 79.20x speedups over serial KClist
- Up to 196.28x speedups over parallel Pivoter
 - Pivoter^[2] is faster: $k \ge 8$ on as-skitter, com-dblp; $k \ge 10$ on com-orkut
- Obtain 4-clique counts on
 - ClueWeb (74 billion edges) in < 2 hours
 - Hyperlink2014 (~100 billion edges) in < 4 hours
 - Hyperlink 2012 (~200 billion edges) in < 45 hours

[1] Danisch, Balalau, Sozio (18)[2] Jain and Seshadhri (20)

Evaluation (k-clique counting)

Comparison to KClist



Evaluation (k-clique peeling)

- 60-core GCP instance (two-way hyperthreading)
- 1.01 11.83x speedups over serial KClist
- Constrained by peeling complexity

Conclusion

Conclusion

- New parallel algorithms for butterfly counting/peeling
- Modular ParButterfly framework w/ranking + aggregation options
- Strong theoretical bounds + high parallel scalability
- Github: https://github.com/jeshi96/parbutterfly
- New parallel algorithms for k-clique counting/peeling
- Strong theoretical bounds + high parallel scalability
- Github:

https://github.com/ParAlg/gbbs/tree/master/benchmarks/CliqueCounting

Future Work

- Cycle counting (for $k \ge 6$)^[1, 2, 3]
- Dynamic/Streaming subgraph counting^[4, 5]
- Nucleus decomposition^[6]
- Objective function for butterfly peeling^[7]
- GraphIt extensions
- Hypergraph algorithms
 - [1] Bera, Pashanasangi, Seshadhri (19)
 - [2] Kowalik (03)
 - [3] Pinar, Seshadhri, Vishal (16)
 - [4] Sanei-Mehri, Zhang, Sariyuce, Tirthapura (19)
 - [5] Eppstein, Spiro (09)

[6] Sariyuce, Seshadhri, Pinar, Catalyurek (15)[7] Tsourakakis (15)

Thank you

Limitations

- Butterfly peeling is P-complete (limited speedups)
- Work-efficient butterfly counting is not the fastest in practice
 - Reducing space usage in butterfly counting
- Not easily generalized to other subgraphs

Deriving am

• # wedges = $\sum_{x \in V} \sum_{y \in N_x(x)} deg_x(y)$

 Where N_x(y) and deg_x(y) refer to neighbors / degree of y considering vertices with rank > rank(x)



 $= O(\alpha m)$

Priority queue for butterfly counts

Batch-parallel Fibonacci heap:

- *k* insertions: O(*k*) amortized expected work, O(log(*n*+*k*)) span whp
- *k* decrease-keys: O(*k*) amortized work, O(log² *n*) span whp
- delete-min: O(log n) amortized expected work, O(log n) span whp

Analysis follows directly from serial Fibonacci heap analysis, except marks are integers instead of booleans

Additionally, we use a parallel hash table to maintain buckets for butterfly peeling

 By vertex: (ρ_v = number of peeling rounds across all vertices)
O(min(max-b_v, ρ_v log m) + ∑ degree(v)²) expected work, O(ρ_v log ² m) span whp, O(n² + max-b_v) space

By edge: (ρ_e = number of peeling rounds across all edges)
O(min(max-b_e, ρ_e log m) + Σ_(u,v) Σ_{u'∈N(u)} min(degree(u), degree(u'))) expected work, O(ρ_e log² m) span whp, O(m + max-b_e) space

(Using batch-parallel Fibonacci heap and Julienne)

• By vertex: $(\rho_v = \text{number of peeling rounds across all vertices})$ $O(\rho_v \log m + \sum \text{degree}(v)^2)$ expected work, $O(\rho_v \log^2 m)$ span whp, $O(n^2)$ space

By edge: (ρ_e = number of peeling rounds across all edges)
O(ρ_e log m + Σ_(u,v) Σ_{u'∈N(u)} min(degree(u), degree(u'))) expected work, O(ρ_e log ² m) span whp, O(m) space

(Using batch-parallel Fibonacci heap)

Peeling framework bounds (Storing all wedges)

• By vertex: (ρ_v = number of peeling rounds across all vertices) O($\rho_v \log m + b$) expected work, O($\rho_v \log^2 m$) span whp, O(αm) space

By edge: (ρ_e = number of peeling rounds across all edges)
O(ρ_e log m + b) expected work, O(ρ_e log ² m) span whp, O(αm) space

(Using batch-parallel Fibonacci heap)
Peeling framework bounds (Storing all wedges)

• By vertex: (ρ_v = number of peeling rounds across all vertices) O(b) expected work, O($\rho_v \log m$) span whp, O(αm + max-b_v) space

• By edge: (ρ_e = number of peeling rounds across all edges) O(b) expected work, O($\rho_e \log m$) span whp, O(αm + max-b_e) space Edge sparsification: Keep each edge independently w/probability p

 Colorful sparsification: Assign a random color [1, ..., 1/p] to each vertex + keep each edge if the endpoints match

Scalability (Per vertex counting)



Sampling



Wedge Aggregation (Per vertex counting with cache optimization)

