Graph Coarsening and Clustering on the GPU

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- Clustering \simeq isolating 'related' groups of vertices in a graph.
- Relevant in: social networks, epidemiology, papers, metabolism, and ecosystems (Newman & Girvan, 2004).
- Our primary interests are speed and parallelisation.

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• Quality of a clustering is measured by its modularity mod(C), introduced in 2004 by Newman and Girvan.

• Clustering modularity is defined as

$$\mathsf{mod}(\mathcal{C}) := \frac{\sum\limits_{C \in \mathcal{C}} \sum\limits_{\{u,v\} \in E} \omega(\{u,v\})}{\sum\limits_{e \in E} \omega(e)} - \frac{\sum\limits_{C \in \mathcal{C}} \left(\sum\limits_{v \in C} \zeta(v)\right)^2}{4\left(\sum\limits_{e \in E} \omega(e)\right)^2}.$$

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• Here, vertex weights $\zeta: V \to \mathbb{R}_{\geq 0}$ are defined as

$$\zeta(\mathbf{v}) := \sum_{\{u,v\}\in E} \omega(\{u,v\}).$$

• $mod(\mathcal{C})$ can be rewritten to

$$\frac{1}{4\Omega^2} \sum_{C \in \mathcal{C}} \left[\zeta(C) \left(2\Omega - \zeta(C) \right) - 2\Omega \left(\sum_{\substack{C' \in \mathcal{C} \\ C' \neq C}} \omega(\mathsf{cut}(C, C')) \right) \right]$$

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• Here, $\Omega := \sum_{e \in E} \omega(e)$ and

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 To calculate modularity, we only need to keep track of summed vertex weights of clusters and summed edge weights between clusters.

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$$\frac{1}{2\Omega^2}\left(2\Omega\omega(\operatorname{cut}(\mathcal{C},\mathcal{C}'))-\zeta(\mathcal{C})\zeta(\mathcal{C}')\right).$$

• This suggests a greedy agglomerative strategy (e.g. Zhu et al., 2008).

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Solution Merge all matched clusters, summing ζ and ω (graph coarsening).

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We make use of parallelism in steps 2 and 3.











Final clustering.



Star graphs



Agglomerative clustering slows down on star graphs.

Fagginger Auer, Bisseling (UU)

GPU Graph Coarsening & Clustering

Star graphs



Merging vertices with the same neighbours is bad for clustering.

Star graphs



So we merge multiple satellites to the same centre.

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• We use $cp(\cdot)$ to identify satellites and match these to centres.

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$$\zeta'(v') = \sum_{\pi(v)=v'} \zeta(v)$$
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• $\pi(u) = \pi(v)$ if and only if $\mu(u) = \mu(v)$ (compress μ to π).

►

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- Then, we create the new adjacency lists and weights for G'.
- Use μ , π , π^{-1} , and a bookkeeping array ρ in global GPU memory.

ρ	1	2	3	4	5	6	7	8	9	10	11	12
μ	9	2	3	22	9	9	22	2	3	3	2	4
π^{-1}												
π												

Initialise ρ sequentially and store μ .

ρ	1	2	3	4	5	6	7	8	9	10	11	12
μ	9	2	3	22	9	9	22	2	3	3	2	4
π^{-1}									-			
π												

Sort by increasing μ -value (sort_by_key).

ρ	2	8	11	3	9	10	12	1	5	6	4	7
μ	2	2	2	3	3	3	4	9	9	9	22	22
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Determine different matched groups (adjacent_not_equal).

ρ	2	8	11	3	9	10	12	1	5	6	4	7
μ	1	0	0	1	0	0	1	1	0	0	1	0
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Extract boundaries for π^{-1} (copy_index_if_nonzero).

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Perform scan to find π indices (inclusive_scan).

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Extract π as $\pi(\rho(i)) = \mu(i)$ (scatter).

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π	4	1	2	5	4	4	5	1	2	2	1	3

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- Sort the neighbour list by index.
- Compress the neighbour list by replacing subsequences $(j', \omega_1, j', \omega_2, \dots, j', \omega_l)$ with $(j', \omega_1 + \omega_2 + \dots + \omega_l)$.

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- Calculate centre potentials and match satellites in parallel.
- Coarsen in parallel as described previously.
- This gives us a fine-grained shared-memory parallel clustering algorithm.
- We do not perform local improvement (Kernighan-Lin): changing cluster weights makes parallelising this very hard.
• Created an implementation on the GPU using CUDA and on the CPU using TBB.

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- Test set: 10th DIMACS challenge.
- Test hardware: dual quad-core Xeon E5620 and an NVIDIA Tesla C2050 (thanks: the Little Green Machine project).

Results (scaling)



Clustering time scaling



Results (quality)

	V	<i>E</i>	CUDA	TBB	Ovelgönne et al. (2010)
karate	34	78	0.363	0.383	0.412
jazz	198	2,742	0.314	0.369	0.444
email	1,133	5,451	0.440	0.473	0.572
PGP	10,680	24,316	0.809	0.841	0.880

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• Lower quality, because we do not use local refinement.

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- TBB: uk-2002, |V| = 18,520,486, |E| = 261,787,258, modularity 0.974 clustering in 31 seconds.
- Comparison to state of the art: DIMACS challenge.

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- We propose the centre potential to deal with star-like graphs.
- The algorithm is very fast, but quality could be improved by parallel local refinement.

Questions

 \exists any questions?

• Performing matching in parallel is problematic.

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- Disjoint edges requirement leads to serialisation.



Suppose we match vertices simultaneously.



Vertices find an unmatched neighbour...



... but generate an invalid matching.

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- Proposals that were responded to are matched.

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 - and its proposal/response value $\sigma(v)$.
- Both π and σ are stored in 1D arrays in global memory.
























