Modularity Graph Clustering on the GPU

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April 10, 2012

Introduction

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- We will perform greedy clustering of large graphs.
- Clustering \approx isolating 'related' groups of vertices in a graph.
- Our primary interests are speed and parallelisation.

• Bioinformatics: group genes with similar expressions.



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- Medical imaging: image segmentation in CT/MRI.



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- Market research: consumer grouping.



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- Medical imaging: image segmentation in CT/MRI.
- Market research: consumer grouping.
- Social networks: community detection.



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$$V = \bigcup_{C \in \mathcal{C}} C$$
 as a disjoint union.

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

• Clustering modularity is defined as

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

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• We have
$$-rac{1}{2} \leq \mathsf{mod}(\mathcal{C}) \leq 1$$
.

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

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

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• Here,

$$\begin{split} \Omega &:= \sum_{e \in E} \omega(e), \\ \zeta(v) &:= \sum_{\{u,v\} \in E} \omega(\{u,v\}), \\ \text{cut}(C,C') &:= \{\{u,v\} \in E \mid u \in C \text{ and } v \in C'\}. \end{split}$$

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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).



Start with vertices in a separate cluster: $C = \{\{v\} \mid v \in V\}$.



Match clusters to increase modularity.



Merge matched clusters (sum ω and ζ).



Match clusters to increase modularity.



Merge matched clusters (sum ω and ζ).



Match clusters to increase modularity.



Return clustering with highest modularity.

0 iterations.



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Final clustering.



Parallel agglomerative clustering

Q Start with all vertices being a separate cluster: C = {{v} | v ∈ V}.
Q Find a heavy matching of clusters with edge weights

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

- Sextend matching to avoid getting stuck.
- Merge all matched clusters, summing ζ and ω .
- Go to step 2 until only a single cluster remains.
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- Merge all matched clusters, summing ζ and ω .
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We make use of parallelism in steps 2, 3, and 4.

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- Performing matching in parallel is problematic.
- Disjoint edges requirement leads to serialisation.



Suppose we match vertices simultaneously.



Vertices find an unmatched neighbour...



... but generate an invalid matching.

• To solve this we create two groups of vertices: blue and red.

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- Blue vertices propose.
- **Red** vertices respond.
- Proposals that were responded to are matched.
- Store matching in a map $\pi: V \to \mathbb{N}$:

$$\{u,v\} \in M \qquad \iff \qquad \pi(u) = \pi(v).$$

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 - its colour/matching value $\pi(v)$;
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 - its colour/matching value $\pi(v)$;
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- Both π and σ are stored in 1D arrays in global memory.



























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- Time: matching and CPU \leftrightarrow GPU data transfer, not disk I/O.
- 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

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Results (time)


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.

Results (time)

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- CUDA: road_central, |V| = 14,081,816, |E| = 16,933,413, modularity 0.996 clustering in 4.6 seconds.

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- Our algorithm is very fast for large graphs.
- CUDA: road_central, |V| = 14,081,816, |E| = 16,933,413, modularity 0.996 clustering in 4.6 seconds.
- TBB: uk-2002, |V| = 18,520,486, |E| = 261,787,258, modularity 0.974 clustering in 31 seconds.

Conclusion

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- This algorithm is suitable for both multi-core CPUs and GPUs.
- The algorithm is very fast, but quality could be improved by parallel local refinement.

Questions

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- Let $C = \{C_1, \ldots, C_k\}$ and define

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• Maximise intra-cluster edges:

$$\max_{\mathcal{C}} \left(\sum_{i=1}^{k} e_{i\,i} \right).$$

• But this leads to $C = \{V\}$ as optimal solution!

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$$a_i := \sum_{j=1}^k e_{ij}$$
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• Cut all edges and reconnect vertices randomly, then

$$e_{ij} \approx a_i a_j$$
.

• Modularity measures how much better we do than the random case:

$$\operatorname{mod}(\mathcal{C}) := \sum_{i=1}^{k} \left(e_{i\,i} - a_{i}^{2} \right).$$

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- Maximising modularity is strongly NP-complete (Brandes et al., 2008).
- Modularity maximisation fails to resolve small communities (Kumpula et al., 2007).

Star graphs



Agglomerative clustering slows down on star graphs.

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Star graphs



Merging vertices with the same neighbours is bad for clustering.

Star graphs



So we merge multiple satellites to the same centre.

• To identify star centres and satellites, we propose a centre potential.

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$$\mathsf{cp}(v) := rac{\mathsf{deg}(v)^2}{\sum\limits_{\{u,v\}\in E} \mathsf{deg}(u)}.$$

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





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• For a star graph where k satellites are connected to a clique of l vertices with 0 < l < k, we have that

$$\begin{array}{ll} \mathsf{cp}(\mathsf{satellite}) \leq \frac{1}{2} & \qquad \mathsf{and} \ \mathsf{cp}(\mathsf{satellite}) \to 0 \ \mathsf{as} \ k \to \infty, \\ \mathsf{cp}(\mathsf{centre}) \geq \frac{4}{3} & \qquad \mathsf{and} \ \mathsf{cp}(\mathsf{centre}) \to \infty \ \mathsf{as} \ k \to \infty. \end{array}$$

Colouring vertices

• To colour vertices $v \in V$, we use for a fixed $p \in [0,1]$

$$\pi(\mathbf{v}) = \begin{cases} \text{blue} & \text{with probability } p, \\ \text{red} & \text{with probability } 1 - p. \end{cases}$$

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- How to choose p? Maximise the number of matched vertices.
- For a large random graphs, the expected fraction of matched vertices can be approximated by (independent of edge density)

$$2(1-p)\left(1-e^{-\frac{p}{1-p}}\right).$$



We should choose $p \approx 0.53406$.

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• For a random graph with *n* vertices, we can approximate (independent of edge density)

$$\lim_{n\to\infty}\frac{2E(N(n))}{n}\approx 2(1-p)\left(1-e^{-\frac{p}{1-p}}\right)$$
Let $G = (\{1, \ldots, n\}, E)$ with $P(\{v, w\} \in E) = d$ for $d \in]0, 1]$. Then E(N(n)) is given by

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$$\approx n(1 - p) \left(1 - \left(1 - \frac{p d}{1 + (1 - p) (d (n - 1) - 1)} \right)^{n - 1} \right).$$

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 - ► $E' = \{\{\pi(u), \pi(v)\} \mid \{u, v\} \in E\}$ (collapse edges),

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$$E' = \{ \{ \pi(u), \pi(v) \} \mid \{ u, v \} \in E \}$$
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$$\zeta'(v') = \sum_{\pi(v)=v'} \zeta(v)$$
 (sum vertex weights),

• $\pi(u) = \pi(v)$ if and only if $\mu(u) = \mu(v)$ (compress μ to π).

►

• Implemented using the CUDA Thrust library.

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- View G as a collection of adjacency lists for each vertex.
- First, we create π and π^{-1} from μ .
- 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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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
π^{-1}												
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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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Determine different matched groups (adjacent_not_equal).



Extract boundaries for π^{-1} (copy_index_if_nonzero).

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
π^{-1}	1	4	7	8	11	13						
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π^{-1}	1	4	7	8	11	13		-				
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Perform scan to find π indices (inclusive_scan).

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