



### cuRipples: Influence Maximization on Multi-GPU Systems

ICS 2020

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PNNL is operated by Battelle for the U.S. Department of Energy



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Introduction

![](_page_1_Picture_2.jpeg)

### **The Team**

![](_page_1_Picture_4.jpeg)

### Marco Minutoli (PNNL)

![](_page_1_Picture_6.jpeg)

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![](_page_1_Picture_8.jpeg)

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![](_page_1_Picture_10.jpeg)

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![](_page_1_Picture_12.jpeg)

![](_page_1_Picture_13.jpeg)

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![](_page_2_Picture_0.jpeg)

## The Influence Maximization Problem

Algorithms

- The objective of the Influence Maximization Problem (Inf-Max) is to identify a small set of individuals in a social network, which when activated, will very likely result in the activation of the maximum number of vertices
  - Problem statement coming from the social sciences
    - ✓ Pedro M. Domingos and Matthew Richardson. "Mining the network value of customers". In: Proceedings of the seventh ACM SIGKDD international conference on Knowledge discovery and data mining, San Francisco, CA, USA, August 26-29, 2001. ACM, 2001, pp. 57-66
  - How does (word-of-mouth) information propagate?
  - Who are the key individuals that optimize the information diffusion?

![](_page_3_Figure_0.jpeg)

Arun Sathanur

### Sam Silva

Madelyn Dunning 5

![](_page_4_Picture_0.jpeg)

## Influence Maximization: Problem Definition

- Given: A graph G(V, E), a diffusion model (how a vertex gets activated based on the state of its neighbors), and a budget  $\mathbf{k}$ , the influence maximization problem is stated as follows:
- Find a set of  $\mathbf{k}$  vertices called the seed set  $\mathbf{S}_{i}$ , that when initially activated result in maximal activations on the network amongst all possible sets of kvertices
- Two diffusion models studied in our work:
  - Linear Threshold: A vertex can get activated if a fraction of neighboring vertices that are active is greater than a threshold  $\lambda_{v}$
  - Independent Cascade: One shot chance for an activated vertex to activate its neighbor

![](_page_5_Picture_0.jpeg)

## The Greedy Hill Climbing Algorithm

- Uses the sub-modularity property of the Influence Function
- Approximation Factor: (1-1/e) ε
  - 1. Generate a set of **n** random samples **SG** 
    - Different instantiations of G are computed based on the edge probabilities
- 2. Repeat until k most influential nodes are chosen:
  - Compute the influence of all remaining nodes across different samples w.r.t. the current seed set S
  - 2. Pick the best influential node, and add to S

Kempe, David, Jon Kleinberg, and Éva Tardos. "Maximizing the spread of influence through a social network." *Proceedings of the ninth ACM SIGKDD international conference on Knowledge discovery and data mining*. 2003.

![](_page_5_Picture_10.jpeg)

![](_page_6_Picture_0.jpeg)

## The IMM algorithm

### "Who is influencing me?" instead of "Who am I influencing?"

- Decide the number of experiments
- Generating the reverse reachability information with graph explorations
- Greedy seed selection
- EstimateTheta builds on Sample and SeedSelect

```
Input : G, k, \epsilon
Output: S
begin
        \langle \mathbb{R}, \theta \rangle \leftarrow \texttt{EstimateTheta}(\mathsf{G}, \mathsf{k}, \epsilon)
       \mathbb{R} \leftarrow \texttt{Sample}(\mathsf{G}, \, \theta - |\mathbb{R}|, \, \mathbb{R})
       \mathsf{S} \leftarrow \texttt{SelectSeeds}(\mathsf{G},\,\mathsf{k},\,\mathbb{R})
        return S
end
```

Tang, Youze, Yanchen Shi, and Xiaokui Xiao. "Influence maximization in near-linear time: A martingale approach." Proceedings of the 2015 ACM SIGMOD International Conference on Management of Data. 2015.

![](_page_7_Picture_0.jpeg)

## **Influence Maximization: Challenges**

- Approximation Factor: (1 1/e) ε
- Algorithms are computationally expensive (high order polynomial for the greedy hill climbing approach of Kempe et al.)
- Alternative methods (random reverse reachable paths) are memory and compute intensive (HPC)
- Nonlinear growth in work relative to approximation factor (ɛ) and number of seeds required (k)

![](_page_7_Figure_6.jpeg)

![](_page_7_Figure_7.jpeg)

![](_page_8_Picture_0.jpeg)

### **DGX-1** Volta

- 8 x V100 GPUs, NVLINK2
  - 5120 CUDA Cores/640 Tensor Cores/4096-bit memory bus/16 **GB HBM2**
- GPUs not fully interconnected (some at 2 hops)
  - Not all GPUs with peer-to-peer atomic memory operations
- V100 has 6 peer-to-peer links, some connections are faster

![](_page_8_Figure_7.jpeg)

Architectures

![](_page_9_Picture_0.jpeg)

### Newell@PNNL

- 2 Power9 CPUs with a total of 128 logical cores per system
- 4 NVIDIA V100 GPUs with NVLINK2 (16GB per GPU)
  - 3 links (GPU-GPU, CPU-GPU)
- X-Bus allows atomic memory operations for GPUs connected to different sockets
- 1TB of system memory per node
- EDR Infiniband internal network

![](_page_9_Figure_8.jpeg)

### **Summit: Unprecedented Parallelism**

![](_page_10_Figure_1.jpeg)

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![](_page_11_Picture_0.jpeg)

- The engine instantiates a thread pool
  - Usually 1 thread per core on the CPUs of system
- Each GPU has a dedicated CPU thread offloading work with the possibility to over-subscribe
  - More than 1-thread pushing work to the same device (Hyper-Q)
- The engine builds a representation of the topology of GPUs
  - To structure reductions between GPUs
  - Topology built query the CUDA runtime
- CPU and GPU workers steal from the same "task queue"

![](_page_12_Picture_0.jpeg)

### Sampling

- Two different strategies for IC and LT models
- For the LT model
  - Each GPU thread performs a randomized BFS, but is limited to visit 8 vertices at most
  - When the limit is exceeded the tasks is invalidated and replayed on the CPU
- For the IC model
  - Parallel BFS derived from the nvgraph.
- Each worker has Parallel Random Number Generator
  - Sequences split with the leap-frog scheme
  - GPU threads do round-robin among them

![](_page_12_Figure_11.jpeg)

![](_page_12_Figure_12.jpeg)

![](_page_12_Figure_13.jpeg)

200

### IC model

RRR Size (#)

18

![](_page_13_Picture_0.jpeg)

## **Seed Selection**

- Greedily select the covering the greatest number of RRR sets
  - Build a histogram of the vertices occurrences for those not yet selected as seeds
- The histogram can be updated or rebuilt from scratch
  - Partition the RRR in covered and uncovered
  - Rebuild works on the uncovered
  - Update works on the newly covered.
- CPU and GPU histograms are then reduced
  - GPUs use a local histogram later reduced using a tree reduction

**Experimental Results: Strong scaling on Summit** 

![](_page_14_Figure_1.jpeg)

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### Results

![](_page_15_Picture_0.jpeg)

Sample 📕 SeedSelection 📕 Total

![](_page_15_Figure_1.jpeg)

Sample SeedSelection Total

![](_page_15_Figure_2.jpeg)

![](_page_15_Figure_3.jpeg)

![](_page_15_Figure_4.jpeg)

(b) web-BerkStan

Sample SeedSelection Total

Configuration

(e) soc-LiveJournal1

![](_page_15_Figure_5.jpeg)

![](_page_15_Figure_6.jpeg)

![](_page_15_Figure_7.jpeg)

Figure 2: DGX-1v IC Model ( $\epsilon = 0.5, k = 100$ ). The configuration reports the number of CPU workers(C) and GPU workers(G)

32C-8G

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![](_page_15_Figure_11.jpeg)

![](_page_15_Picture_13.jpeg)

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### **Summary of our contributions**

 Scalable implementations (shared and distributed memory systems) https://github.com/pnnl/ripples

![](_page_16_Figure_3.jpeg)

CuRipples achieves a speedup of 790x over a state-of-the-art serial implementation, while also significantly improving the quality. The input network is com-Orkut.

Cores (C), GPUs (G), Nodes (N).

System	Time (s)	Speedup	Scale
<b>com-Orkut</b> ( $\epsilon$ =0.5, $k$ =100)			
IMM <sub>seq</sub>	28024.56	$1.00 \times$	1C
<b>IMM</b> <sub>opt</sub>	9027.50	$3.10 \times$	1C
<b>IMM</b> <sub>mt</sub>	1319.21	$21.24 \times$	20C (1N)
CuRipples <sub>dgx-1v</sub>	35.47	790.09  imes	80C+8G (1N)
CuRipplesnewell	43.72	$641.00 \times$	128C+4G (1N)
<b>com-Orkut</b> ( <i>ϵ</i> =0.13, <i>k</i> =200)			
<b>IMM</b> <sub>edison</sub>	294.51	$95.16 \times$	3,072C (64N)
<b>IMM</b> <sub>edison</sub>	47.77	$586.61 \times$	49,152C (1024N)
<b>CuRipples</b> <sub>summit</sub>	36.30	772.03  imes	2,688C+384G (64N)
soc-LiveJournal1 ( <i>ϵ</i> =0.5, <i>k</i> =100)			
IMMseq	16434.81	$1.00 \times$	1C
<b>IMM</b> <sub>opt</sub>	3954.59	$4.16 \times$	1C
<b>IMM</b> <sub>mt</sub>	1026.21	$16.02 \times$	20C
CuRipples <sub>dgx-1v</sub>	70.23	$234.01 \times$	80C+8G (1N)
CuRipplesnewell	65.26	$251.84 \times$	128C+4G (1N)
<b>soc-LiveJournal1</b> ( <i>ϵ</i> =0.13, <i>k</i> =200)			
<b>IMM</b> <sub>edison</sub>	190.94	$86.07 \times$	3,072C (64N)
<b>IMM</b> <sub>edison</sub>	55.12	$298.16 \times$	49,152C (1024N)
<b>CuRipples</b> <sub>summit</sub>	106.43	$154.42 \times$	2,688C+384G (64N)

TABLE II: Comparative evaluation of cuRipples relative to previous implementations of IMM-both serial (IMMseq) [2] and parallel (IMM<sub>opt/mt/edison</sub>) [3]. Abbreviations used: No.

![](_page_17_Picture_0.jpeg)

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# Thank you

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https://github.com/pnnl/ripples

![](_page_18_Picture_10.jpeg)